All Microarray Data Following Oxythioquinox Exposure

	1 DMSO - Average	1 DMSO - Stdev	1 15 - Average	1 15 - Stdev	1 60 - Average	1 60 - Stdev	1 120 - Average	1 120 - Stdev	2 dmso - Average	2 dmso - Stdev	2 15 - Average	2 15 - Stdev	2 60 - Average	2 60 - Stdev	2 120 - Average	2 120 - Stdev	3 dmso - Average	3 dmso - Stdev	3 15 - Average
S78187_at	2090.15	547.23	1196.5	1382.39	2294.3	439.96	2039.05	517.25	2420.9	558.05	1981.55	641.98	2580.55	1240.76	2380.2	632.44	2167.45	14.78	2313
X69636_at	506.9	74.39	654.15	755.83	1552.5	984.72	559.4	195.59	837.85	218.57	693.25	733.48	694.4	70.43	784.8	678.82	1108.1	491.58	882.4
X04688_at	206.05	248.83	130.6	150.9	54	26.87	173.3	4.38	350.8	448.59	403.55	329.3	336.75	352.78	339.3	422.28	201.35	215.74	27.25
U16306_at	299.6	359.07	21.75	25.1	372.7	251.31	251.45	168.79	393.65	251.8	229.75	21.85	66.3	13.29	193.25	169.49	240.15	251.09	306.75
M36542_s_at	1463.35	56.21	346.5	400.36	1124.4	383.53	2161.15	1222.09	1002.35	62.44	1683.75	1803.19	1254.2	257.1	1195.7	487.48	1384	285.25	2161.3
HG2171-HT2	226.7	53.17	239.2	276.34	409.25	57.49	333.15	217.44	167.2	14.28	380.8	289.77	405.5	95.88	430	64.77	207.45	20.01	241.9
M35531_at	1589.6	311.55	510	589.3	2053.55	1064.13	1094.75	84.78	1553.7	115.68	1225.1	1075.23	1713.85	42.5	1264.7	461.32	1554.75	281.5	2145.05
X97671_at	189.9	216.09	34.45	39.81	400	491.16	586.7	655.91	265.05	91.85	380.05	229.17	40.95	17.89	372	465.7	60.35	58.05	278.15
AC002477_s_	1228.55	25.24	853.55	986.2	973.1	338	808.35	90.16	1090.15	204.71	535.7	97.3	857.25	814.52	949.65	151.53	834.6	367.98	1281.65
L76528_s_at	423.2	152.45	160.55	185.47	539.55	130.89	321.1	101.82	180.9	55.01	272.15	135.27	197.05	109.96	757.65	412.74	176.25	82.38	206.5
U33267_at	212.35	11.24	127.85	147.71	230.95	135.69	223.25	228.89	617.55	463.51	234.9	57.7	358.25	28.07	450.3	131.8	303.15	2.19	382.55
AFFX-BioB-3_	1402.6	169.85	776.45	897.11	4483.75	4081.49	3829.15	3538.15	2487.65	1137.24	3876.3	3827.99	1024.25	989.31	1929	202.23	1123.8	65.2	2245.7
M64673_at	5334.45	1604.92	3212.55	3711.67	4789.05	2494.6	4929.95	534.22	4933.4	1636.1	5488.6	1926.44	4499.35	165.11	4864.8	1407.99	5524.2	1363.02	5400.4
U02031_at	3552.3	1521.27	1541.75	1781.27	1562.35	951.55	2133.55	1469.72	2908.35	410.76	2708.8	1402.48	3588.2	114.27	3192.85	462.24	4103.6	199.97	3238.3
U81607_at	579.75	469.45	613.9	709.23	337.95	410.48	268.6	287.51	161.5	90.93	270.55	292.53	484.2	181.59	113.85	14.5	259.1	33.94	369.65
AFFX-LysX-5	146.05	130.89	174.6	201.67	318.55	48.01	397.45	387.99	112.35	82.66	220.55	251.66	139.75	180.24	501.35	167.09	109.7	130.67	464.9
M33317_f_at	360.4	171.26	335.7	387.92	328.45	249.54	602.1	564.55	833.05	288.43	431	59.26	369.9	9.05	221.5	17.39	225.65	56.92	533.85
D10704_at	648.55	623.46	326.4	377.17	1541.25	1080.81	1318.05	413.73	1328.3	369.68	1853.8	618.44	794.35	418.96	1089.4	642.48	1293.85	301.44	1569.35
D86978_at	1283.4	174.66	468	540.65	825.5	473.34	865.35	160.58	1460.8	568.94	1760.85	564.34	1155.45	261.42	1269.55	72.9	1325.2	421.86	1313.2

Z37986_at	5422.95	939.82	2637.7	3047.49	4500.8	536.41	5879.2	906.65	8293.35	1607.04	5058.15	440.32	6776.85	534.22	8471.4	2653.06	7842.5	2119.06	6098.55
U51903_at	500.65	5.02	230.75	266.65	326.2	418.47	216.7	126.57	460.55	472.7	559	404.75	318.15	1.06	177.45	165.11	229.65	283.76	195.15
D50914_at	1890.05	483.73	1156	1335.58	1509.05	1152.94	3322.85	479.21	2246.6	1204.91	2695.2	422.71	2025.5	509.97	3141.55	452.19	2476	75.24	2752.35
M13207_at	2369.85	333.12	1251.75	1446.25	1045.45	782.41	2745.2	259.23	1057.9	434.87	1792.65	1101.18	1314.75	424.19	997.05	1165.52	1550.9	642.05	1029.25
U25956_at	916.3	356.81	86.95	100.48	268.8	96.17	366.95	71.49	669.6	436	779.1	13.29	438.35	111.93	579.35	193.68	293.5	66.33	268.05
X12953_at	337.6	197.57	91.3	105.5	291.4	58.41	385.7	164.9	432.55	52.54	379.8	386.79	610.45	488.54	644.3	197.42	343.85	297.76	419.4
M81650_rna1	54.8	28.99	153.15	176.99	113.8	126.71	158.25	43.35	146.9	7.92	312.1	363.17	104.05	29.63	140.75	52.4	109.05	82.24	91.9
X52005_at	599.1	470.65	295.65	341.6	465.95	526.58	420.7	193.32	273.5	267.43	275.9	159.81	734.45	656.97	1043.55	127.21	391.35	22.98	699.75
X86816_at	126.45	89.17	38.25	44.19	56.35	24.54	144.95	107.41	188.85	188.44	57.5	26.3	90.1	24.04	68.95	4.31	72.05	18.6	79.7
U88666_at	436.8	8.06	436.2	504.03	617.45	149.69	1209.35	993.7	388.8	101.12	702.1	100.83	801.5	34.51	467.25	5.73	398.25	367.2	307.2
D89289_at	488.95	182.22	425.2	491.3	638.95	5.87	628.95	444.98	236.25	134.42	384.95	96.38	397.1	377.31	411.95	463.51	488.1	192.47	734
Z78285_f_at	17.1	5.8	12	13.86	73.2	68.31	68.95	76.16	39.35	26.94	56.85	65.97	32.55	16.05	49.4	7.5	24.45	3.89	48.2
U54804_at	86.9	45.96	406.95	470.16	116	45.82	188.65	188.87	579.15	379.93	282.1	146.51	482.3	200.39	327.7	196.72	196.25	190.99	565.25
AFFX-HUMIS	658.35	153.37	288.5	333.33	997	242.54	242.3	258.94	1380.7	563	832.95	39.53	465.95	364.51	997.85	594.04	762.15	197.78	799
U76992_at	411.4	193.75	466.3	538.82	521.45	312.05	983.4	663.12	565.45	146.02	725.25	399.44	495.1	112.01	406.3	433.03	478.55	32.74	290.15
HG3494-HT3	8743.1	1638.51	6641.6	7673.52	8285.45	3201.28	9224.95	4406.76	9727.1	2539.65	9590.95	2943.76	7860.25	265.8	6982.3	3585.17	8015.45	994.97	8403.2
D82070_at	1592.1	281.71	862.9	997.02	1341.5	413.94	1701.35	796.7	1687.75	612.43	1605.3	967.75	1818.3	604.15	1466.85	578.91	1251.1	363.17	1816.65
U56998_at	5268.4	593.4	3497.65	4041.04	2913	197.42	1904.4	861.82	3587.15	1819.6	4111	1638.22	2408.05	1892.43	3548.3	1284.67	4019.3	2139.71	4491.65
D49488_at	381.65	144.32	37.15	42.92	451.25	388.7	233.6	120.21	158.35	102.18	613.5	360.34	112.9	75.24	394.5	345.35	245.05	16.48	104.4
D11094_at	1694.3	484.65	1148.4	1326.82	2326.95	162.56	2187.6	106.91	2614.9	238.01	1544	975.52	2309.6	297.97	1797.85	536.2	2163.95	75.73	1739

U31814_at	1009.25	154.08	727.35	840.4	1311.2	193.18	1080.3	241.12	1302.15	96.38	1747.95	402.98	1324.15	163.7	1331.55	735.46	1299.25	106	1121.55
M17219_at	253.4	225.14	175.55	202.87	438.7	155	366.75	6.15	588.6	147.93	389.45	455.16	320.5	243.81	324.3	306.32	348.35	22.13	348.15
M25393_at	222.95	255.34	87.5	101.12	51.9	34.37	421.75	496.18	453.9	430.49	153.2	169.85	158.65	138.52	52.75	45.33	47.45	7	193.3
Z37166_at	5791.75	682.15	4242	4901.1	5032.8	1145.65	5808.45	2527.69	5323.8	1998	5983.85	2307.22	4858.5	244.52	4430.15	772.66	6144.35	2382.31	5893.6
U40434_at	933.85	35.43	253.85	293.24	506.5	324.84	576.3	645.31	637.3	470.65	442.9	238.72	585.25	30.33	974.65	636.18	159.05	42.78	460.35
U23143_at	5565.9	1035.2	4367.95	5046.69	4985.65	1435.07	6449.2	2858.97	3719	84.99	6217.65	646.79	3810.1	1204.63	3342.3	524.39	5365.8	2843.7	5891.85
J00129_at	1007	464.14	539.6	623.39	582.4	526.51	1004.75	404.39	722.85	84.07	684.2	415.21	964	132.65	1004.45	581.31	747.45	201.03	925.3
Y07604_at	5049.25	368.19	1449.9	1675.14	3224.3	2021.9	5040.95	1237.65	2155	1701.58	5068.9	523.26	3192.4	771.03	4322.3	1862.66	3463.75	794.86	3696
L32179_at	219.15	87.47	349.8	404.18	266.75	98.22	739.8	909.76	715.55	753.71	328.65	227.48	455.1	339.98	462.85	387.28	110.9	87.96	606.15
D31846_at	6826.8	342.81	3855.7	4454.77	7599.45	2682.41	7576.25	362.11	6311.1	956.15	9076.2	3024.3	9063.1	522.98	6287.05	1199.89	6979.5	2702.28	6637.05
D25217_at	524.8	89.66	195.65	226.06	110.15	19.73	170.85	164.26	507.05	366.63	591.35	704.35	712.85	284.75	251.1	140.15	549.05	195.8	767.75
X05153_rna1	169.75	92.98	93.75	108.26	670.1	537.12	215.8	76.79	152.65	10.68	485.3	552.67	216.95	234.69	363.85	272.73	378.15	314.59	172.8
J04823_rna1	18746.5	1877.37	10274.4	11870.77	22196.95	3133.69	25899.05	7119.08	19378.3	4451.09	18174.7	804.26	17749	8753.56	21538.2	2773.84	16285.05	1469.44	17611.3
U79295_at	234.1	262.05	226.55	261.7	133.15	63.99	287.85	280.51	364.2	296.56	97.8	94.89	119.05	42.21	249.9	255.12	115.15	36.27	547.9
X96849_at	1241.5	147.08	592.95	685.12	1288.65	238.51	758.05	896.4	1331.25	531.82	399.15	84.78	1000.1	21.64	1302.15	315.02	1126	1195.72	740.4
U85658_at	1298.9	227.12	851.7	984.01	590.5	221.04	896.75	295.5	1318.85	35	1097.95	193.11	959.75	37.83	1096.35	47.59	1050.3	131.38	712.25
M59499_at	186.3	107.9	102.2	118.09	86.15	68.66	471.2	610.52	271	313.81	95.85	10.54	185.3	166.59	257.45	333.54	86.65	67.53	79.6

X51441_at	7394.65	1497.86	4433.1	5121.86	6872.65	908.14	6972.6	1334.03	6752.3	1177.62	4830.45	1327.03	3667.8	2269.81	3536.35	669.84	2219.3	636.82	5488.35
M31241_s_at	578.4	141.56	90.85	105.01	594.9	592.7	158.8	37.19	300.45	156.62	223.75	69.08	217.9	135.76	154.6	149.06	258.6	268.28	312.55
X73113_at	1956.8	279.87	881.25	1018.16	1485.4	726.06	1946.05	239.92	2305.05	225.92	2447.3	931.12	2608.65	883.67	2355.6	188.09	1196.35	234.12	1582.1
Z38026_at	914.2	532.73	638.35	737.58	441.3	167.3	554.55	213.05	1141.3	429.5	1309.15	831.2	487.85	163.84	899.75	565.19	1808.75	1680.3	677.2
D64154_at	4626	1305.32	1723.65	1991.42	5336.35	166.38	6835	297.83	5865.6	392.87	5125.35	1316.42	4445.55	872.5	5425.55	705.9	5176.15	672.81	5002.9
U05861_at	751.15	686.67	553.5	639.51	1901.4	671.75	754.1	175.22	606.3	381.84	814.95	690.49	872.2	71.13	1582.75	398.45	978.05	163.55	1338.45
X59350_at	1607.2	886.43	1091.15	1260.7	2314.2	526.94	2099.1	1383.24	1733.6	861.11	1523	1479.55	1157.2	348.18	1311.95	313.74	1859.9	290.06	1798.4
HG1747-HT1	921	368.26	849.75	981.82	900	189.93	1282.9	641.91	665.25	578.2	1292.8	742.04	953.9	139.72	695.6	72.41	907.3	108.61	731.25
X62515_s_at	1462.35	807.3	788.75	911.25	906.75	235.96	654.3	101.54	1189.75	956.93	1051.1	477.86	827.5	384.24	303.25	69.93	684.6	586.9	1230
X76223_s_at	1354.2	224.58	907	1047.93	997.65	366.63	1237.25	490.8	905.65	429.57	1001.55	663.34	2244.9	709.94	702.9	469.38	1809.8	75.52	958.85
HG3415-HT3	3357	758.16	1625.9	1878.5	3195.55	515.83	3093.55	302.43	3184.15	357.16	2849.2	583.93	2654.05	283.05	2583.35	361.69	2769.45	1921	4132.2
M58583_at	402.8	83.16	387.7	447.88	568.45	683.7	198.8	33.66	571.2	211.71	462.8	532.88	579.4	81.32	147.35	130.46	296.25	270.19	248.4
X53065_f_at	1389.5	375.47	574.95	664.33	1101.1	542.49	1084.9	183.42	824.15	48.72	894.3	444.63	905.55	167.23	639	560.74	861.2	141.56	858.05
X60484_at	179.2	87.54	57.35	66.26	166.55	124.8	191.9	200.54	352.55	310.35	136.5	99.98	124.05	36.7	79.15	44.34	196.25	222.24	302.65
U66711_rna1	7816.3	1878.36	2916.5	3369.65	6087.45	2121.67	6175	293.73	5101.65	2126.34	4654.9	56.85	7969	248.05	5279.45	50.7	8073.25	981.68	8627
D50911_at	1699.85	124.66	1137.4	1314.09	1921.5	90.37	1687.75	703.5	2794.3	1088.38	2097	32.24	2413.8	311.98	2067.05	554.16	2161.2	45.25	2310.8
U11870_rna1	726.95	177.27	66.2	76.51	803.45	441.02	288.9	34.08	1032.95	546.38	755.55	260.85	1491.45	203.72	281.8	136.9	541.5	377.17	1169.65
M25809_at	1057.25	38.54	970.7	1121.47	711.75	180.52	687.15	621.05	598.85	448.8	1515.75	169.07	1028.75	986.77	805.85	575.66	731.25	37.12	259.8
M55210_at	1071	18.81	918.85	1061.58	990.8	132.94	800.75	75.87	1466.3	603.44	1393.85	168.93	530.7	480.69	189.4	33.38	556.9	438.55	1404.55
J03827_at	22956.85	7912.6	11284.8	13038.2	18501	7239.93	19489.75	14848.18	17694.65	7040.73	16654.2	8195.37	16475.9	2772.28	17936.45	5832.71	17998.4	3327.22	18585

AB000462_at	1254.95	182.5	1063.4	1228.67	3981.25	4617.62	1854	1355.95	1806.8	745.15	2111.1	1696.63	5914.65	4633.6	2767.85	1701.94	1573.15	750.88	2639.8
M90657_at	209.9	178.76	321.75	371.73	853.35	190.71	493.05	419.95	717.85	816.07	999.75	39.39	235.5	165.32	675.8	195.02	604	270.54	681.6
L07956_at	542.9	196.01	445.85	515.13	1067.2	363.74	440.6	409.7	485.1	340.26	473.35	481.19	495.1	487.9	381.25	292.39	753.4	236.74	686.7
Z47087_at	1148.55	162.42	660.95	763.6	1540.25	368.9	1275.7	314.95	1583.6	163.2	1391.2	462.73	1199.5	47.38	1408.6	114.83	1111.45	29.2	798.85
L26081_at	70.45	30.33	28.35	32.74	86.65	53.39	92.3	58.55	221.05	195.23	120.7	32.81	261.9	295.71	82.55	8.56	61.6	20.65	140.05
K02402_at	58.3	26.73	14.9	17.25	32.45	12.52	176.2	128.69	106.5	60.25	66.15	33.02	149.85	94.96	91.85	40.94	93.85	53.67	58.5
HG33-HT33_	46326.95	22308.44	19068.65	22031.54	56969.8	52.89	58467.3	31961.79	59380.7	5064.44	48679.15	10715.71	37246.6	4247.31	38849.1	12031.28	46578.65	13487.42	43848.6
U19147_s_at	176.35	108.97	71.95	83.09	377	177.77	293.5	166.45	77.55	58.19	236.95	266.51	144.8	134.35	94.05	101.61	192.3	196.72	31.65
X82207_at	1642.05	12.94	1073.15	1239.91	1446.3	424.12	1575.55	73.89	839.95	318.98	1017.65	336.23	1430.2	289.35	1948.7	750.38	1653.75	683.56	1604.5
X80199_at	3028.5	160.94	2048	2366.26	2741.35	354.05	2541	180.17	3125.5	371.23	3706.8	516.9	3551.5	358.79	2943.5	129.12	3815.9	806.95	3532.7
U53786_at	764.35	605.5	965	1114.97	1373.9	478.15	948.5	540.8	780.3	657.18	780.25	186.46	1608.55	626.28	1050.05	430.98	934.3	497.66	1515.5
S57132_s_at	3647.9	1304.61	1446.45	1671.25	3561.8	1994.75	2471.75	289.42	2116.6	24.61	2799.55	657.68	3180.05	818.48	2894.85	1098.07	3448.65	166.95	3402.15
X13967_at	2688	249.33	1611.85	1862.31	2041.35	114.62	2475.35	271.46	2654.95	283.9	2416.8	635.69	1775.25	906.16	1778.75	127.77	1676.8	200.96	1641.05
M63262_at	598.95	750.45	138.05	159.45	161.75	93.41	252.6	115.26	740.05	251.38	981.65	14.78	713.95	701.8	408.3	259.79	331.95	289.84	330.6
X80062_at	208.6	95.74	295.7	341.67	122.2	151.32	204.3	258.52	543.1	338.99	543.3	141.42	154	102.11	283.55	305.26	135.4	64.21	154.95
U25771_at	332.85	131.73	520.25	601.11	550	375.05	240.75	248.12	507.55	680.87	103.85	30.19	408.5	277.89	178.9	108.05	358.85	334.96	129
U15641_s_at	1356.95	1590.64	294.35	340.05	1149.35	1292.66	339.5	198.84	643.75	372.57	1125	780.65	390.5	108.05	1181.2	967.6	950.3	906.09	629.65
S78774_at	157.55	28.64	52.85	61.02	279.05	225.5	64.05	32.17	118	41.58	106.75	52.4	114.75	50.98	110.7	0.42	133.3	52.33	150.8
L40586_at	454.5	250.46	142.05	164.12	395.55	93.27	633.85	459.27	393.4	33.8	195.3	118.09	717.55	450.36	463.75	568.58	303.05	215.6	216.7

D83703_at	1093.4	595.38	413.2	477.44	1490.6	444.63	1629.35	360.7	816.05	555.86	1538.2	76.79	1752.05	109.96	1232.25	160.58	810.95	59.75	977.4
U28727_at	196.1	61.66	67.8	78.35	236.35	33.45	282.85	226.34	215.4	88.95	216.95	98.78	103.3	49.21	117.45	8.84	181.1	39.74	188.45
U20230_at	310.45	24.11	126.95	146.72	190.8	148.92	321.75	150.68	618.7	734.83	211.5	26.73	248.1	188.37	163.75	34.86	119.5	31.82	220.85
HG620-HT62	569.2	28.14	352.55	407.36	859.85	174.87	1122.7	274.64	502.4	465.13	750.7	278.74	1346.85	402.84	757.55	608.18	470.95	101.47	1051.05
U49957_s_at	356.75	266.93	266.25	307.66	325.9	370.67	106.9	50.06	196.55	146.16	89.35	51.55	252.75	244.87	313.4	363.45	503.75	130.32	63
X52425_at	2344.2	417.48	1066.05	1231.71	1796.35	640.71	907.95	683.7	2417.8	13.58	2300.2	543.06	2794.45	708.73	2307.15	706.47	2874.7	554.51	2631.15
S80343_at	2109.6	44.83	1007.3	1163.76	2006.55	40.38	1832.3	97.72	2146.2	632.72	1398.85	767.71	2414.3	80.04	2105.75	341.32	1764.1	332.06	1673.3
D86960_at	291.15	279.38	241.9	279.45	700	308.16	221.55	187.03	441.15	346.69	192.25	98.92	92.65	19.59	1382.75	575.09	377.85	206.83	244.3
U27326_s_at	295.9	129.26	819.4	946.67	410.45	1.2	225.5	53.88	1250.65	366.78	742.35	59.04	977.6	184.27	511.8	235.61	166.25	91.57	255.15
U52077_s_at	408.7	220.33	364.3	420.87	310.45	364.37	718.5	787.43	641.45	287.72	339.75	355.32	449.8	327.39	305.4	9.33	515.3	205.63	264.5
X99802_at	306.65	114.9	203.6	235.18	348.55	63.71	375.35	236.67	312.2	35.92	384.4	185.4	288.65	79.55	333.15	71.21	266.2	8.77	301.35
X57959_at	17280.25	3874.31	9273.25	10714.15	13209.2	5894.44	12388	10468.29	16966.25	3390.22	14586.7	6894.86	11116.35	1196.78	14221.9	2675.55	15470.7	3923.88	15699.05
U38846_at	5317.2	1204.34	3284.5	3794.9	5817.3	329.37	5620	2731.27	6384.75	683.84	6148	1250.87	5693.6	808.65	5938.75	174.44	6898.65	416.56	6512.65
J04029_s_at	3411.25	177.98	2678.4	3094.58	2956.65	366.35	2837.55	568.02	4954.35	2018.72	3176.85	1158.88	2586.95	77.57	2643.95	680.87	3691.45	119.85	2452.6
HG3995-HT4	8271	444.2	3109.95	3593.16	9462.25	1136.25	8891.2	577.85	9114.15	539.73	10067.9	4565.93	10288.95	80.54	9666.35	549.63	7028.95	353.77	8045.35
X93920_at	3894.3	896.19	1683.6	1945.25	3197.4	794.36	2183.2	395.84	3016.35	101.75	1873.1	374.48	2558.85	539.59	2739.7	344.5	3747.4	1072.82	3497.3
X63131_s_at	122.6	37.76	101.75	117.59	454	477.01	391.4	379.72	221.75	34.44	305.95	99.49	626.1	245.22	369.25	83.79	244.15	155.63	314.25
M92269_f_at	630.8	475.46	92.8	107.2	548.2	498.09	1154.25	1275.27	622.7	118.51	997.1	367.27	866.8	746.85	533.5	286.8	405.2	128.83	877.15
Z50053_at	523.55	102.46	141.65	163.7	824.85	571.41	374.7	108.33	485.5	427.38	253.3	254.98	820.1	273.93	935.65	754.98	438.85	477.93	451.85
AB003177_at	3926.9	1486.2	2604.65	3009.38	4055.2	1148.62	3574.7	2560.29	4215.8	902.27	3560.25	328.03	3803.35	49	3878.6	134.07	3537.55	204.85	3838.7

X63563_at	540.3	424.97	484.05	559.25	593.75	274.99	446.45	288.71	794.4	52.89	930.8	109.18	760.9	823.64	1262.6	557.06	606.9	87.68	716.4
M59216_s_at	684.35	19.73	162.9	188.23	445.5	431.34	470.5	97.44	399.4	97.86	284.55	169.92	356.95	351.5	308.05	390.82	378.3	389.9	562.6
X07743_at	100.65	28.5	45.35	52.4	266.25	84.08	820.75	617.09	174.95	128.2	311.55	300.03	448.1	473.48	528.8	196.72	296.95	218.57	70.85
D87735_at	26703.7	13561.04	12340.6	14258.1	33185	6077.58	37437.1	17604.7	28495.6	5125.11	26391.9	7102.75	26733.7	10242.87	27198.15	3055.9	28222.3	9152.51	28144.45
S66896_at	191.8	169.28	308.35	356.31	186.1	197.85	724.55	700.67	417.1	181.59	184.85	89.17	241.5	149.77	727.5	42	237.65	275.42	135.3
AFFX-HUMTI	1489.65	597.43	434.35	501.83	2029.75	1308.64	1460.15	225.5	1004.85	525.17	1010.85	49.99	1497.9	664.68	1672.75	91.71	1291.8	372.36	1384.95
L22214_at	287.25	31.61	196.2	226.7	317.8	142.55	1083.3	860.27	408.65	126.5	287.65	40.94	294.95	36.56	176.05	81.39	596.3	562.29	442.25
X68688_rna1	782.9	494.69	373.95	432.11	1210.1	564.84	2535.3	1046.94	1626.9	952.33	1859.35	1966.11	2107.85	1006.14	1563.2	269.27	988.2	106.21	439.4
U14187_at	2241.2	480.41	929.3	1073.67	3269.75	737.44	2589.8	2175.63	2012.05	290.55	876.5	697.49	2394.25	274.71	2630.4	354.68	2604.5	628.34	3335.4
X55330_at	50.5	22.63	35.3	40.73	673.7	255.55	492.85	615.54	306.7	103.94	505.2	415.5	75.25	30.05	151.3	156.69	41.3	29.27	195.7
U60805_at	116.35	121.83	60	69.3	386	466.27	157.1	21.78	219.15	100.06	258.5	47.38	99.5	35.07	112.05	38.54	245.8	268.13	270.25
M85276_at	197.35	110.24	89.75	103.73	302.65	125.79	569.05	331.56	558.5	434.73	937.3	311.41	227.6	105.36	167.95	49.43	222.05	100.76	181.45
U79255_at	911.9	912.87	797.65	921.57	418.9	67.18	764.35	127.21	426.5	70.29	1518.2	1579.25	762.1	44.12	430.05	87.19	792.3	535	532.15
X66401_cds1	941.25	271.18	823.5	951.48	1238.8	253.99	1888.8	387.49	857.65	81.11	757.95	655.42	722.25	278.81	702.5	379.15	909.45	484.58	1313.2
X97230_f_at	637.75	320.53	678.75	784.25	1014	4.95	617.85	612.99	1216.3	925.18	1421.15	655.28	1425.2	895.2	845.35	212.34	957.75	99.49	1398.9
HG2705-HT2	1492.85	360.41	838.75	969.09	1817.1	576.15	1038	983.73	1796.2	78.91	1854.85	1012.93	1264.15	317.7	591.45	440.32	1393.95	448.52	1138.7
U61500_at	403.95	123.39	281.25	324.92	100.3	86.13	176.85	154.93	456	175.93	221.9	245.08	322.25	180.81	300.4	327.25	174.15	66.82	378.75
M60854_at	67272.35	53476.16	24926.45	28799.54	108373.8	4867.72	97571.5	25951.81	102063.66	13179.83	75536.85	5357.39	73907.2	33717.25	76368.45	14727.83	71635.5	40958.45	80521.9
U15174_at	2006.1	496.67	655.05	756.82	2427.6	428.22	2269.4	808.93	1396.7	28.43	1098.45	116.6	1277.25	676.77	1754.75	519.79	1190	442.65	1901.25

L28010_at	1646.8	3.11	1516.1	1751.64	815.5	409.41	1211.15	696.29	1998.2	1175.07	1739.35	900.08	1647.05	153.51	1726.65	162.99	1705.4	351.86	1687.95
D31628_s_at	189.75	5.3	510.65	589.94	498.75	137.82	423.6	377.03	190.25	84.22	390.05	84.36	411	317.21	666.3	672.03	163.8	22.06	239.7
U09410_at	293.55	310.91	398.2	460.04	820.65	155.21	826.6	626.21	829.75	147.29	484.3	274.22	472.15	211.21	119.2	67.18	735.75	433.95	841.25
X51755_cds5	385.45	71.77	200.9	232.07	189.95	70.78	1034.05	885.79	321.3	113	320.95	24.68	302.25	27.65	304.75	30.19	339.2	105.5	323.75
X59303_s_at	3318.95	853.83	654	755.61	3004.9	494.83	1982.85	792.74	2460.15	1235.24	1235.95	1000.77	2084.75	1673.51	2867.1	495.26	3205.15	494.48	2498.6
X83705_s_at	99.4	3.25	36.45	42.07	774.65	186.61	80.2	41.44	718	859.84	345.6	302.78	141.75	79.27	278.5	324	346.1	307.87	804.9
L37362_at	524.45	62.3	602.55	696.15	522.35	229.74	291	234.62	151.25	151.82	662.85	647.5	240.05	23.55	252.15	70.36	239.55	134.14	866.15
U38964_s_at	1074.4	347.19	794.6	918.11	1043.65	325.06	1146.1	491.72	819.45	174.58	1399.25	386.29	576.75	474.96	718.85	281.5	594.2	35.64	1148.5
V01515_cds1	130.2	122.61	58.75	67.81	199.75	217.72	404.05	550.2	124.3	1.13	436.55	475.11	134.45	3.61	95.4	113	55	39.32	101.65
U40343_at	596.4	360.91	476	549.99	1041.1	329.37	673.05	254.77	1516.65	976.59	1475	342.95	759.6	639.93	1268	117.24	946.1	47.23	547.65
HG987-HT98	2394.3	217.22	1369.9	1582.79	3384.95	759.36	3028.1	604.01	1823.6	1263.18	2141.3	339.69	1267.15	719.34	1545.4	237.31	676.05	385.59	915
U79251_at	269.6	76.37	66.7	77.07	1077.5	1041.14	713.05	885.23	879.8	327.53	315.35	154.79	249.65	52.54	429.65	381.63	780.05	102.88	99.95
X95592_at	337.05	62.44	163.9	189.36	469.35	294.65	178.1	162.78	523.25	7.57	251.2	300.1	399.65	111.93	228.15	166.81	351.55	397.32	92.05
U67171_at	4530.8	969.73	2056.2	2375.74	4566.8	795.49	5388.65	2490.64	4800.05	1601.38	4284.95	1181.36	5391.6	244.23	4767.6	839.48	4797.35	832.62	4597.4
X63417_at	171.05	28.21	363.2	419.6	414.3	78.49	106.15	12.09	392.3	428.93	109.7	8.49	412.2	412.24	175.25	36.13	375.95	412.88	134.45
K01383_at	1316.75	522.34	773.55	893.71	1648.65	1449.07	539.5	277.33	1104.9	83.16	1787.45	475.1	1699.25	834.74	542.9	476.45	720.9	184.13	530.5
M31303_rna1	2172.4	159.52	1799.1	2078.61	1803.6	114.27	2061.5	903.26	2400.8	693.53	2850.8	525.24	3489.05	162.56	4068.65	1668.84	3231.75	474.54	3271.95
M57567_at	7207.25	59.61	3909.2	4516.57	5940.85	929.63	6511.2	1161.35	6784.45	1182.21	5713.45	2110.08	5598.55	1135.54	6009.8	1569.78	6410.4	284.26	6798.1

U73704_at	106.55	1.91	32.1	37.05	288	239.43	229.65	131.17	78.05	22.13	712.45	602.24	211.3	66.04	453.05	537.9	107.15	110.1	482.85
U50708_at	201.3	185.54	27.65	31.89	488.65	236.95	364.8	450	135.4	108.75	226.9	264.46	138.25	93.55	331.25	68.09	82.4	4.81	458.5
U92457_s_at	862.25	132.02	385.1	444.91	1196.7	524.25	683.3	248.76	599.15	257.03	788.1	87.82	1112.8	290.48	1392	60.39	1287.15	837.57	763.2
HG2383-HT4	1935.85	527.29	1502.4	1735.81	1623.7	806.67	1158.6	1050.76	2305.3	1222.02	1282.5	1291.74	1953.35	1074.59	1535.7	619.71	1731.15	1560.09	1744.05
X04327_at	674.1	318.76	498.95	576.5	234.15	256.04	610.55	809.43	267.05	85.35	502.5	198.84	96.65	6.15	259.5	229.67	490.8	20.36	385.8
D13168_at	145.9	152.88	319.75	369.46	559.1	248.19	164.8	102.67	426	110.31	385.4	158.96	402.9	194.17	52.65	42.5	227.55	143.75	570.65
M76766_at	554.9	180.45	32.35	37.41	268.25	44.05	134.55	46.6	776.15	141.49	190.5	194.88	355.3	375.9	674.45	220.97	516.5	125.44	640.2
U05012_s_at	58.35	5.59	96.85	111.93	498.75	599.56	207.9	78.49	358.1	463.15	323.6	390.32	161.4	126.85	201.05	192.55	79.55	45.75	109.55
M97016_s_at	183.05	44.62	94.05	108.68	168.3	17.54	220.4	198.27	104.5	10.75	186.7	26.45	166.15	8.7	150.15	75.17	155.4	78.21	140.65
HG1612-HT1	662.8	462.45	602.1	695.65	431.45	87.61	639.7	512.65	1585.1	198.56	980.9	191.06	741.3	464.43	1568	146.8	485.75	260.29	847.2
X93036_at	11988.9	4679.49	4771.8	5513.17	15576.45	804.48	14801	1620.97	14918.3	1592.83	14373.8	1658.02	16697.1	8432.67	15965.1	2882.03	10593.3	3364.7	19184.35
U20760_at	738.65	191.13	602.9	696.64	1337.85	460.54	942.6	510.53	869.2	58.69	725.55	129.9	1124.5	492.71	812.65	815.51	831.3	302.78	1112.15
D13666_s_at	257.85	6.01	193.05	223.09	644.95	861.47	93.65	100.2	348.35	424.19	322.95	231.15	107.9	69.3	338.35	397.32	76	65.34	255.35
D56495_at	904.1	116.96	648.6	749.39	1207.5	317.07	751.65	340.33	775.8	276.34	1301.1	386.08	1066.3	289.21	817.45	306.96	1020.05	565.47	1325.2
U15173_at	532.95	226.34	178.1	205.77	458.3	239.99	763.9	535.99	460	63.22	273.45	36.84	390.75	266.37	890.7	232.21	304.35	36.42	425.8
HG2868-HT3	1041.3	145.24	361.6	417.76	1135.25	566.04	954.5	629.47	753.65	576.08	334.8	226.42	713.2	359.07	735.9	18.67	1010.35	117.73	616.3
M31774_s_at	54.8	16.55	178.3	206.05	277.75	354.33	96.25	77.43	103.65	4.45	176.5	225.14	134.45	41.93	779.9	1029.69	90.2	14.42	119.95
U32576_rna1	453.75	78.28	250.9	289.91	534.55	195.8	607.85	63.71	394.95	25.53	1071.45	1052.1	659.95	151.67	449	43.7	459	37.34	442.25
D42123_at	7896.65	1187.16	4008.45	4631.34	7937.6	1147.78	8562.15	1796.26	8051.85	497.45	8617.6	430.2	8377.8	128.55	6280.55	2008.11	8462.2	952.61	8808.15

U06155_at	300.65	122.82	603.5	697.21	1196.85	846.9	1040.9	959.54	839.85	94.96	847	108.33	857.75	160.73	130.85	16.48	587.2	229.53	552.9
U71374_at	300.7	324.14	381.05	440.32	694.65	139.37	814.7	1057.55	942.85	271.18	521.3	210.58	1132.45	279.24	480.9	524.53	714.45	92.56	745.8
M11313_s_at	283.15	186.89	252.1	291.33	642.85	728.81	164.45	104.72	134.8	0.85	314	289.21	468.25	536.48	130.3	69.86	227.45	110.24	156.45
AFFX-DapX-I	321	213.69	256.45	296.35	123.9	148.21	143	159.24	228.75	190.71	809.7	267.43	114.3	118.37	194.6	74.67	392.95	142.91	438.15
M21574_at	270.55	207.54	256.5	296.42	374.5	191.77	411.15	168.36	879.55	91.85	908.25	186.61	874.1	153.3	895.6	338	809.55	366.21	900.2
L32832_s_at	358.15	38.96	201.6	232.92	697.35	797.69	373.8	119.08	384.6	133.78	311.45	185.62	566.15	121.27	622.95	68.66	410.1	368.26	548.2
U13666_at	682	11.03	264.9	306.04	295.25	83.65	899.25	555.15	771.7	665.1	786.6	298.96	549.5	537.83	767.05	192.12	702.8	250.74	651.4
S70609_at	675.65	583.01	441.9	510.53	292	181.16	413.9	204.92	262	136.9	614.85	77.57	316.15	2.62	797.25	698.69	160.8	58.97	176.7
HG1155-HT4	1624.75	685.12	1158.1	1337.99	2631.75	765.58	2136	293.31	1101.7	302.36	2492.45	932.89	2396	602.17	2093.3	352	2008.2	246.36	1976
M81695_s_at	1320	453.82	879.7	1016.4	1241.35	82.94	1674.55	96.94	1087.7	250.88	1814.15	1016.75	1862	204.64	1021.75	622.47	698.6	78.77	727.4
U18271_cds1	495.7	478.71	60.3	69.72	111.65	5.44	411.15	511.03	294	271.1	302.5	208.74	164.2	165.46	331	101.4	95.35	46.46	584.2
M26004_s_at	177.45	12.09	118.65	137.11	367.45	157.33	86.75	30.33	239.5	185.4	169.75	167.65	171.5	148.78	324.85	304.13	51.1	20.36	294.85
M36653_s_at	1850.15	1249.67	343.95	397.32	2319.65	683.14	641.65	123.81	1012.75	493.21	1551.35	48.44	2112.85	1875.04	857.65	405.81	876.65	77.29	688
X53296_s_at	601.45	301.02	444.2	513.22	818.55	909.41	182.45	66.54	1013.9	15.27	532.45	125.37	604.9	280.3	578.2	275.49	629	515.76	638
L04483_s_at	90280.95	38533	36304.75	41945.79	124858.1	29355.26	98903	12571.65	126233.1	12900.89	100383.66	13360.85	103860.6	54830.62	92592.45	42804.49	90310.45	33964.53	80461.9
D13900_at	7837.1	1450.84	3507.6	4052.57	8548.75	1684.97	7118.05	501.13	6107.5	1026.72	7211.3	2031.38	6500.6	785.88	7523.6	940.31	6590.3	1773.14	5797.35

U09937_rna1	6962.35	1048.99	2761.95	3191.1	4468.65	2541.55	7932.5	3756.43	8468.7	69.01	4896.9	482.53	4396.3	464.43	6612.9	4098.39	5449.4	1666.65	4402.55
M97388_at	1094.1	353.55	578.55	668.43	1750.9	687.17	1059	246.36	945.05	115.05	1471.95	73.04	628.2	45.25	1046.15	132.3	1068.25	304.69	1157.95
X64624_s_at	405.6	501.9	11.35	13.08	387.7	14.71	131.8	84.43	355.4	181.59	390.65	100.2	417.5	50.35	138.35	84.92	325.05	144.6	424.25
U32989_at	82.2	65.48	167.15	193.11	125.05	5.3	51.35	28.21	190.9	222.88	42.2	4.38	76.1	17.11	224.2	247.35	73.25	2.19	226.9
L37033_at	10754.9	3357.91	2434.7	2813.01	8368.55	453.75	7250.6	681.09	6697.55	2305.66	6587.65	1535.34	8059.9	256.82	6807.05	1991.85	7469.3	1372.64	7299.1
D26598_at	6185.75	2981.94	2605.35	3010.22	7619.9	234.19	8030.6	1704.13	7669.95	680.31	7621.85	213.76	7882.05	3102.43	7716.8	884.31	5994.7	987.55	6765.95
U15637_s_at	2243	1362.88	594.15	686.53	1828.9	244.09	1872.8	1206.47	1898.1	238.58	2206.5	349.59	1187.7	564.98	1552.6	798.61	1230.9	125.16	2086.9
S75313_at	146.5	180.45	15.35	17.75	638.45	531.96	191.5	218.78	246.45	8.41	270.2	362.89	149.4	128.55	150.1	150.47	42.6	18.53	67.95
U77664_at	612.9	125.87	1057.75	1222.09	766	238.86	661.3	77.36	403.65	104.58	980.35	327.04	801.75	335.38	1091.4	420.73	767.7	119.78	665.2
HG3484-HT3	218.45	272.59	322.25	372.29	241.1	60.25	58	3.96	506.4	171.26	454.45	328.88	398.35	14.78	44.8	18.81	363.75	387.28	426.55
U65416_rna1	3374.25	621.33	1267.6	1464.56	4387.65	2316.27	2666.55	811.26	3405.35	1516.39	4068.3	2294	3175.15	1127.34	2586	608.68	3139.5	421.01	3559.35
X91868_at	220.15	163.27	118.6	137.04	859.35	1022.97	444	55.58	446.15	558.97	595.15	337.08	427.45	147.57	233.1	154.71	775.25	513.15	500.55
M24485_s_at	39192.1	14848.54	23181.8	26783.79	57710.4	4654.18	60752.6	8857.64	50788.55	5408.73	42366.5	8519.22	44790.95	13687.68	48692.4	5998.53	42154.15	2782.68	51314.8
M57731_s_at	734.8	154.57	1078.45	1245.99	1124.65	295.5	854.65	342.73	769.35	167.23	413.4	205.91	303.1	182.86	648.05	829.93	665.25	140.64	363.9
M63896_at	340.35	444.28	77	88.95	283.1	114.83	349	417.33	402.4	424.12	476.3	278.46	47.15	35.14	201.35	134.85	274.95	272.73	208.7
S78825_at	1022.75	956.08	199	229.95	1185.25	1266.36	302.5	49.92	682	622.68	483.35	230.87	429.85	24.54	210.85	0.07	862.55	682.29	913.55
X64559_at	159.3	41.01	52.2	60.25	150.65	103.17	155.35	121.13	143.8	24.04	124.85	12.66	128.1	5.37	137.55	14.78	265.6	204.78	291.3

L05425_at	980.95	362.53	1026.55	1186.03	1162.5	254.28	1095.85	128.34	1882.8	180.88	1038.4	47.38	1806.35	237.8	1577.15	430.84	1429.45	960.75	1320.4
Z29064_at	544.55	29.2	387	447.17	400.55	406.94	589.05	19.02	247.1	79.76	527.95	262.55	279.2	171.26	497.55	478.78	569	266.01	425.6
X15822_at	9073.75	5084.31	4410.7	5095.98	11478.05	918.04	13605.3	6244.04	12164.9	7007.57	9079.55	3277.79	9339.9	3236.57	11675.7	782.34	8520.2	503.88	10631.75
U37690_at	11560.45	288.99	6597.4	7622.47	10872.8	84.99	13401.1	2039.58	14663.8	2771.01	13043.1	2701.29	16132.75	2110.36	17179.3	6216.32	11324.35	5233.51	9548.95
S50223_at	37.9	6.65	30.1	34.79	99.3	98.99	76.55	81.25	72.35	42.07	99.75	84.22	32.6	6.65	66.55	15.49	78.4	31.96	42.55
X15341_at	20878.3	13436.3	8443.9	9755.95	26282.55	2988.44	26386.7	6799.82	19831.2	2180.29	20747.15	3172.15	17003.6	6909.71	16898.35	3460.51	16290.55	6520.44	19038.6
X94232_at	546.5	284.96	379.5	438.41	693.8	103.52	918.55	784.25	980.75	440.03	599.85	271.32	486.4	410.55	232.55	39.24	813.25	176.56	626.95
X57398_at	4598.7	708.95	2788.55	3221.79	3219.25	630.81	4137.8	1312.67	4165.65	2796.68	3579.1	959.69	4840.95	504.8	4430.1	504.31	5947	1744.15	4547.4
S34389_at	3126.15	228.61	1621.55	1873.48	2914.4	753.21	3049.75	611.86	2310.85	482.88	3516.65	1134.55	2523.8	761.27	2858.8	621.69	2616.5	351.86	3330.6
D10537_s_at	1500.8	198.56	945	1091.77	1582.85	147.71	1639.85	436.36	1707.1	1.98	1031.85	715.38	1084.5	78.21	920	824.49	1596.7	207.04	2038.25
X83863_at	36.65	6.29	195.5	225.85	112.2	38.75	77.15	71.77	52.55	34.58	175.35	62.58	49.75	22.13	70.25	23.41	37.75	5.87	42.55
X72879_at	130.35	28.07	217.45	251.24	662.75	536.91	329.95	0.64	513.95	237.38	389.25	27.93	114.6	137.32	548.45	262.97	375.6	427.66	153.3
M17262_at	752.7	593.55	582.85	673.38	1433.4	659.02	1225.2	293.17	1829.8	124.59	1207.55	231.58	1217.5	75.66	1348.3	3.54	1546.25	234.69	1847.8
Y00264_at	2025.7	182.43	876.8	1013	1885.25	317.7	1761.15	32.17	2603.4	886.57	2428.8	29.13	2319.95	32.17	1597.45	341.89	2508.6	199.12	2393.5
X02612_at	147.3	9.33	279.4	322.86	141.5	16.12	688.3	707.53	137.4	82.87	426.45	473.97	550.05	529.84	204.1	85.14	353.2	274.64	384.1
X96969_at	1365.9	268.28	854.3	986.98	1505.3	494.83	1211.05	365.22	1270.8	351.29	1353.9	113.28	1524.95	255.62	1226.1	495.96	1441.1	199.4	1016.75
Z80787_at	343.35	90.72	374.7	432.89	462.05	262.97	431.15	323.5	568.45	265.52	444.3	130.53	196.6	175.36	382.9	67.32	250.7	213.97	677.85
U06088_at	1010.25	779.02	637.8	736.95	1856.95	1136.39	842.9	913.02	1219.05	436.36	1352.6	1506.7	916.15	459.27	771.85	783.4	964.65	963.15	714.75
M86917_at	1075.25	100.76	455.9	526.79	685.1	29.42	1013.55	988.89	465.25	452.76	1109.9	313.39	673.05	714.39	738.35	69.65	860.65	140.36	1035
U90878_at	12703.9	68.45	6131.6	7084.36	10796.85	1715.37	12448.5	6087.91	10813.95	2639.13	8318.55	2927.78	8574.55	4085.73	9062.4	2690.4	10228.4	2311.81	9965.95

D26135_at	97.85	27.93	58.6	67.74	118.25	39.95	106.9	32.95	90.4	16.26	149.65	7.85	167.2	82.02	69.25	20.15	90.35	56.92	177.55
D10522_at	2426.45	1081.66	687.5	794.36	2337.55	147.57	1942.95	572.54	2109.5	370.1	1858.75	438.19	1620.45	432.68	1259.75	555.01	1790.25	121.41	1797.35
U10693_at	198	130.96	38.25	44.19	633.05	728.39	226.2	70.99	272.65	207.82	328.45	221.25	153.25	150.54	322.65	151.67	167.45	171.33	182.05
D87467_at	398.95	266.23	252.15	291.4	422.75	148	362.05	56.21	518.2	122.47	678.65	292.53	725.7	144.82	429	211.99	532.95	12.37	453.2
J05582_s_at	364.9	63.5	166.8	192.76	1029.35	863.17	431.6	347.61	594.85	311.9	1137.35	801.93	1451.15	211.21	1159.95	698.98	859.9	754.06	478.55
M19045_f_at	3098.15	909.27	1220.1	1409.69	2772.9	655.35	2370.5	1267.84	2333.95	64.98	2553.35	758.8	3096.65	120.42	2473.7	98.99	1805.15	865.57	2169.55
Y07867_at	220.6	210.15	163.8	189.22	704.35	10.11	283.9	31.4	353.25	66.11	372.55	232.57	345	418.18	85.4	5.52	602.25	96.38	611.05
Z71460_at	2066.15	106.84	1112.2	1284.95	2200.55	367.48	2292.2	1396.82	2463.15	423.49	2044.9	196.29	1877.4	90.93	1746.8	589.16	2104.6	679.25	1927.3
U78095_at	15248.6	5042.24	10516.55	12150.57	16347.45	4327.85	16447.6	7612.99	15800.4	485.78	15295.8	5378.96	14789.5	772.44	15221.9	1296.98	14019.9	1609.66	14046.5
D86974_at	7461.45	6462.89	3707.35	4283.44	2582.25	782.41	2248.65	1781.41	5338.55	1689.35	3740.9	169.85	6451.9	1571.19	4670.2	10.32	5877.45	4626.53	3267.85
U92027_at	300.75	290.83	358.95	414.72	359.1	28.85	463.15	184.63	502.65	449.93	654	340.83	478.8	400.93	260.4	19.8	350.1	364.87	556.9
X66397_at	789.6	364.16	281.2	324.84	666.8	487.2	513	173.67	836.8	78.63	915.95	327.6	791.45	4.31	863.8	204.78	869	471.07	792.15
U28249_at	657.5	141.7	110.8	127.99	1081.9	207.32	1229.35	302.01	862.05	12.09	452.7	356.66	646.65	610.59	611	10.61	351.6	330.5	571.25
M98343_at	2850	649.12	1201.7	1388.47	3235.85	176.28	2589.75	263.96	3290	616.74	1738.6	1263.03	3712.7	176.78	3002	550.98	2569.9	561.44	3539.5
X15331_s_at	2362.85	782.56	921.25	1064.41	2893.15	780.72	2521.15	342.59	2863.05	339.91	1490.05	1326.46	1801.35	332.55	1753.85	307.38	2719.55	20.01	2396.3
M30257_s_at	445.35	13.65	68.2	78.77	286.9	47.94	186.25	153.09	60.85	5.02	172.5	95.04	460.55	389.69	484.65	289.42	448.1	193.46	319.6
U71300_at	264.45	107.55	78.3	90.51	588.45	295.36	886.15	885.65	436.2	113.28	220.4	192.9	719.15	199.05	593.5	38.33	467.9	163.06	277.9
M86757_s_at	277.65	113.07	79.9	92.35	581.35	500.7	212.15	99.07	1069.55	1023.4	228.85	146.3	472.25	397.75	204.65	209.37	113.8	13.44	189.95
S82472_at	39.1	19.52	12.25	14.21	96.5	74.25	65.7	23.05	252.95	230.87	213.7	246.36	90.5	43.27	50.2	8.91	63.3	48.51	66.4
X89430_at	148.35	138.1	32.7	37.76	366.25	336.51	631.95	798.54	64.9	1.84	529.3	695.37	89.7	41.86	517.55	502.12	88.25	1.63	84.85

U28811_at	1711.25	186.18	1639.6	1894.34	1296.65	1089.58	813.75	817.49	1348.3	772.16	2239.85	1060.02	1852.25	440.03	1498.85	347.26	1895.35	530.4	1217.7
U73799_at	2131.4	1390.88	604.75	698.69	1920.5	1478.84	1294.9	255.69	1263.3	339.69	2529.6	1926.16	1863.8	1116.24	2212.15	384.88	3165.6	715.88	2742.85
X99140_at	1284.55	513.01	630.75	728.81	805.4	523.4	858.7	388.48	1607.75	153.51	1787.6	1588.3	851.55	760.92	1164.15	1126.07	1735.5	474.47	745.65
J05032_at	310.2	380.42	212.6	245.65	667.05	60.88	432.45	226.77	732.95	263.4	716.3	187.67	948.45	41.65	580.45	217.29	478.7	163.06	630.95
U25029_at	19.7	13.58	20.95	24.25	48.4	12.02	45.5	42.99	51.9	12.16	27.55	21.43	25.5	17.54	29.35	13.08	59.15	6.72	194.1
U38175_at	1553.15	702.23	977.85	1129.74	1148.35	888.48	987.35	783.97	834.05	580.04	1085.3	211	681.35	301.72	1876.15	785.24	1610.45	247.7	869.95
M37721_at	1339.95	519.09	958.3	1107.19	1359	73.54	886.65	290.27	1332	101.96	1529.95	113.21	1463.6	12.59	1127.9	356.81	1229.05	614.12	1107.9
M12783_at	301.55	95.95	125.7	145.24	295.05	82.1	258.45	154.93	227.75	0.92	201.1	69.58	212.05	20.58	265.55	33.87	125.3	48.22	177.45
D79984_s_at	1556.4	218.5	699.15	807.73	904.15	987.47	1250.6	146.8	1610.4	523.54	1248.4	315.09	1431.75	211.07	1268.2	253.99	1809.05	84.5	1934.3
U19796_at	6138.55	1128.19	2675.1	3090.76	6357.8	247.06	6720.85	1546.23	5029.4	264.46	7202.9	711.21	5882.65	472.14	5389.45	1224.5	4578.5	1203.5	4160.65
U30998_at	100.9	119.78	86	99.42	58.55	2.76	224.15	297.9	71.85	20.29	371.2	401.35	223.65	219.7	292.8	369.39	429.75	290.83	274.75
U34880_at	1830.7	130.67	1263.9	1460.32	2706.9	640.78	2842.9	252.86	2555.85	56.21	2810.8	208.17	2439.75	127.35	2221.85	1144.17	1852.9	204.21	2174.45
X54637_at	1155.8	641.91	471.3	544.47	372.1	10.32	1019.3	271.81	1086.5	1122.6	654.35	526.3	844.7	650.4	933.2	387.21	1515.65	586.55	976.95
U90065_s_at	1163.5	1104.64	615.65	711.28	1363.8	707.67	612.7	156.84	2027.05	435.79	1870.75	1842.37	967.15	373.42	335.15	135.55	1549	730.02	1591.7
D64158_at	951.25	78.42	636.2	735.11	1068.5	132.51	924.65	233.7	1188.25	211.07	1071.3	2.97	1296.5	104.65	915.25	171.76	1101.85	223.8	971.75
L11695_at	36.3	4.53	128.2	148.07	256	79.05	335.85	419.1	267.8	265.17	466.65	334.67	60.8	5.8	67.45	30.76	169.3	111.44	370.7
L14813_at	583.25	476.94	297.1	343.23	1063.2	274.92	995.8	570.92	565.15	130.32	646.25	230.59	1064.3	811.62	792.4	98.43	682.15	224.22	1109.2
U90913_at	9033.25	3749.72	3821.9	4415.74	10080.55	2835.14	11515.15	1828.08	5803.95	2498.7	6412.7	703.85	5826.25	2048.84	5913.6	1131.65	6954.6	2617.85	6309.7
U37012_at	2192.3	651.67	642.35	742.11	2427.8	254.42	2509.85	422.21	1804.9	159.66	2731.15	633.36	2029.4	783.62	2347.2	330.36	2801.1	330.08	2551.2

U41740_at	535.05	39.1	424.3	490.17	318.9	217.22	338.6	211.71	578	89.94	310	266.72	444.6	9.19	699.6	88.25	189.7	11.6	121.5
D00003_s_at	15.85	5.59	10.95	12.66	371.45	464.92	324.85	243.32	162.2	203.36	87.65	27.22	37.95	22.7	39.95	17.75	17.65	0.07	17.3
AB000896_at	2176.9	57.7	1078.1	1245.64	3790	2651.37	3873.25	3400.26	3544.65	2517.51	5626.2	4606.38	2697.1	284.26	3992.75	2827.79	2154	93.06	3606.6
L11353_at	917.85	686.81	905.75	1046.45	640.85	129.33	680	303.49	1122.7	71.28	868.8	92.63	980.8	1144.1	1149.4	596.52	978.9	394.42	1043.65
M91438_at	92.3	3.25	293.5	339.13	392.25	362.82	143.6	37.62	156.5	24.61	207.05	151.11	386.4	419.88	421.2	368.26	367.75	42.36	139.85
U12387_s_at	634.5	0.99	264.35	305.4	491.15	162.56	682.8	20.36	478.6	70.85	526.5	299.11	585.4	81.74	573.2	525.24	453.85	292.67	359.55
U41515_at	1738.75	114.76	659.15	761.62	1878.8	83.01	1801.8	97.58	2106.35	193.82	1488.4	63.07	1056.3	175.79	1704.9	3.82	1491.3	344.36	987.2
HG2259-HT2	8970.4	2506.55	2572.2	2971.83	9258.55	1647.63	8351.2	1174.22	7891.25	1929.2	4836.65	74.03	7645.7	3916.1	8394.45	190.57	11319.4	9091.27	5209.1
M18255_cds/	134.75	86.34	45.3	52.33	485.45	55.23	830.05	1048.57	252.5	69.3	171.55	223.8	203.15	151.53	243.6	35.78	213.75	174.02	190.6
D21205_at	155.15	59.47	92.1	106.35	192.2	16.55	268.05	134.56	309	40.02	310.1	102.53	331.25	76.3	628.55	351.22	222.4	91.92	284.5
U71207_at	503.1	288.78	167.4	193.46	171.95	37.26	216.6	199.4	447.7	204.64	137.45	134.85	290.3	257.67	633.45	719.91	445.3	441.23	636.65
U10687_s_at	334.7	313.11	307.75	355.6	244.2	71.98	152.75	124.24	218.6	61.52	795	957.85	175.45	49	140.45	94.68	100.55	41.08	278.3
S80437_s_at	6984.5	4519.54	2133.85	2465.47	2418.75	790.33	3418.2	756.75	4377.65	26.94	2768.55	696.01	5284.15	2159.86	5427.15	3646.2	5855.05	2611.84	3948.3
X99374_s_at	108.05	43.91	628.65	726.27	67.05	9.4	469.15	617.66	104.9	13.72	186	190.07	75	42.14	70.75	13.51	87.15	14.21	138.5
U12775_at	705	68.59	264.4	305.47	981.25	793.59	396.75	346.98	632.45	638.87	1062	125.44	195.65	23.41	735.9	368.97	455.7	458.21	1006.2
L41680_at	105.1	60.53	58.45	67.53	116.25	15.63	262.55	297.9	123.5	50.35	163.3	15.13	110.95	64.56	97.05	10.82	73.15	26.8	82.95
D38550_at	546.55	182.93	463.3	535.28	953	231.37	628.25	300.31	748.6	245.37	290.35	266.79	831.25	108.97	992.3	91.22	388.65	485.85	682.85
M29874_s_at	517.95	78.28	203.6	235.18	603.35	585.98	514.85	31.47	713.75	46.32	848.8	464.71	819.9	28.57	486.05	402.56	670.55	177.13	482.6
Y09836_at	1829.85	1792.59	1435.35	1658.38	992.95	1099.06	1486.05	939.82	1141.6	337.15	998.4	1029.69	1131.2	1079.19	1242.05	941.94	541.15	449.79	847

U63541_at	698.95	537.61	683.25	789.34	1034.55	1017.46	809.75	159.03	218.9	146.8	1159.6	403.19	635.2	577.28	1233.5	678.54	1177.2	60.25	398.7
K03189_f_at	4470.9	572.76	2277.75	2631.64	3788.4	64.63	2632.6	1664.67	2696.15	763.04	3082.95	1505.5	4342.65	32.88	3683.8	1252.14	3128.5	1964.63	3032.6
U58091_at	187.75	233.56	26.75	30.9	100.8	21.78	48	41.86	207.45	230.16	220.7	194.88	244.25	207.39	325.5	378.58	251.8	43.7	115.2
U23028_at	1062.15	90.72	823.75	951.7	624	270.68	1065.95	243.88	1080.05	429.14	1051.35	514.56	1024.95	130.18	1015.25	180.1	853.65	34.58	1052.75
HG3227-HT3	1940.9	824.77	1213.65	1402.26	2251.85	391.24	3101.5	1012.15	2324.7	826.18	2783.75	1243.87	2870.05	785.24	1589.4	408.42	2170.65	400.29	3173.15
D30036_at	1539.7	419.03	692.8	800.44	1440.6	385.37	2023.85	169.07	1478.65	577.07	1456.5	351.86	1750.85	406.8	1241.65	795.71	1315.85	109.67	1916.45
X54667_s_at	982.35	727.4	785.15	907.15	1627.1	565.26	2131.85	2023.24	1603.5	853.05	1190.45	1362.24	1257.5	373.21	1575.4	1319.89	1328.15	257.32	1377.3
U07856_at	27.35	16.48	26.25	30.33	128.15	29.2	576.5	455.94	123.2	58.83	159.05	135.41	133.85	113.77	28	6.79	38.45	1.48	72.7
M88461_s_at	253.2	77.36	301.25	348.11	249.55	135.98	179.8	68.73	181	45.68	271.1	176.92	176.65	149.13	243.2	60.53	173.2	85.7	74.7
X99350_rna1	583.25	655.28	120.05	138.66	149.45	84.5	296	248.9	88.85	4.03	249.05	122.12	202.8	199.83	238.55	45.04	54.6	6.79	109.95
U31116_at	135.7	126.71	188.95	218.28	34.75	23.55	116.75	60.88	131.25	115.19	75.3	49.21	247.75	2.9	41.3	8.34	193.35	229.6	369.85
U71364_at	471.8	337.86	318.7	368.26	1393.2	252.15	692.6	281.43	675.6	615.89	654.95	559.53	553.2	352.56	453.8	249.61	1214.55	3.61	685.9
X60992_at	753.1	692.4	421.7	487.2	1048.85	1197.63	395.2	208.03	425.75	82.94	525.95	170.91	511.45	492.78	541.7	355.96	1165	84.85	328.6
Y08837_at	59.05	19.87	409.6	473.2	390.5	480.55	119.25	136.97	91.4	37.34	236.7	102.25	121.55	20.58	301.35	365.08	42.7	8.34	367.45
U63336_at	572.3	287.65	587.4	678.68	1038.15	427.87	480.35	289.42	581.65	217.44	1466	60.1	712.35	503.25	695.75	643.68	471	485.5	1216.6
M64572_at	1380.1	1092.62	702.4	811.48	1426	562.15	366.45	13.65	545.6	542.21	817.2	659.87	1124.25	778.74	1029.2	94.75	1471.45	578.77	1094.95
X85545_at	669.05	42.78	167	192.9	396.6	187.38	379.8	40.16	803.2	31.25	344.25	256.04	456.55	354.9	452.05	450.07	232	196.43	784.5
X12433_at	1032.55	420.09	677.75	783.12	1122.7	637.95	996.3	3.11	1361.2	109.46	1517.05	698.55	1043.4	120.07	1204.1	342.52	1369	394	1293.1
S40719_s_at	2147.3	810.34	1235.8	1427.79	2784.2	1281.7	2923.25	1609.73	2293.6	493.56	2746.25	1556.27	1363.8	56.29	2042.15	279.52	1993.6	102.39	2508

U06233_at	239.3	200.39	183.95	212.49	81.9	33.23	65.85	30.76	126.65	60.6	336.75	261.56	83.6	62.08	79.6	63.5	56.75	20.86	308.45
U51336_at	6349.45	341.6	4307.95	4977.25	5736.95	368.76	6183.45	554.16	5245.2	635.55	6445.55	212.49	6962.5	452.12	7053.15	405.1	6954.5	66.89	5882.05
AC002045_x	3248.55	2180.51	2206.9	2549.83	1869.45	548.64	1381.25	704.63	2219	632.86	2283.1	1293.58	2286.2	436.57	1733.55	164.26	1545.65	285.6	3041.15
U51477_at	2359.55	166.52	1407.6	1626.35	1716.7	467.11	2340.6	53.17	2457.2	625.08	1819.6	731.29	1364.2	692.4	2372.05	703.22	2482.45	643.11	2052.05
M89957_at	705.25	500.84	231.75	267.78	420.45	153.94	331.65	189.86	479.35	151.25	717.1	591.71	412.1	79.48	809.15	651.6	276.3	60.81	424.45
D28589_at	1707.7	1045.53	441.6	510.25	1648.1	66.33	1441.6	1266.15	1582.95	482.6	1049.05	953.25	1781.9	10.47	1676.4	703.29	1489.95	313.32	1673.3
X74295_at	2327.55	1016.47	1011.55	1168.78	2675.95	1284.6	2798.15	1265.93	1638.2	270.26	2433.7	536.55	1932.45	877.73	1548.3	883.46	1280.2	697.91	1470.45
X52009_s_at	449.65	384.03	62.75	72.48	124.9	111.86	219.55	195.37	352.95	422.21	116.3	61.09	78.1	20.36	89.35	80.54	233	245.79	267.75
M82809_at	607.35	51.97	418.55	483.59	913	257.67	1153.2	909.34	955.3	382.97	831.75	581.45	877.7	238.86	1081.6	141	756.35	37.97	828.85
U16997_at	561.15	27.65	719.95	831.77	1183.1	1259.22	1107.1	94.33	821.85	272.59	989.9	911.88	289.05	74.6	1176.55	172.46	741.45	144.6	815.45
D90279_s_at	138.9	94.47	58.9	68.02	104.95	17.47	182.3	147.08	145.05	87.75	357.2	84.99	301.65	216.45	151	37.76	156.2	15.56	152.75
AB000450_a	1069	496.81	1089.8	1259.07	4133.95	3582.56	3779.2	3842.98	2490.45	708.03	2382.9	1762.39	3718.15	1093.54	3341.9	2315.49	1439.55	268.77	2165.45
U02493_at	5908.75	298.33	2371.25	2739.69	4696.4	957	4474.9	1720.39	5502.35	902.48	4238.2	1355.81	5569.9	949.22	5513.75	1161	6849.9	293.87	5781.2
D90070_s_at	38.85	11.1	473.25	546.81	398.4	195.73	172.15	162.85	155.85	183.78	376.4	197.42	28.25	8.56	44.6	13.15	179.5	19.94	130.75
U03858_at	649	325.98	74.1	85.56	858.6	137.46	301.7	78.06	763.7	165.04	803.6	774.28	986	322.86	443.3	567.95	580.35	24.68	830.65
M13981_at	2960.95	2204.12	243.85	281.78	547.9	663.97	562.35	149.13	691.6	631.45	613.85	346.13	1343.2	166.88	274.9	293.45	1056	158.39	952.85
M26682_at	634.1	74.53	325.85	376.53	391.8	58.69	443.75	319.97	481.8	72.27	377.7	6.65	299.35	124.66	292.8	109.32	323.9	20.65	608.35

M69023_at	1496.9	114.69	823.35	951.27	716.05	779.3	1422.4	368.4	1164.9	539.52	892.85	452.05	1208.35	454.03	1368.65	117.87	1376.6	103.8	1147.45
X17094_at	2080.35	457.29	1880.85	2173.15	1059.3	745.29	1387.45	944.91	1512.9	1008.33	1682.45	365.08	962.95	860.34	1915.65	342.45	2023.85	302.01	1047.85
L16782_at	144	68.02	34.35	39.67	109.4	64.77	144.2	173.24	516.05	260.85	612.45	481.89	171.3	41.86	111.8	13.15	250.35	222.81	50.6
U35340_at	294	14.42	105.7	122.19	476.25	171.05	740.2	781.21	505.1	360.34	269.5	97.02	412.35	294.93	311.6	44.83	255.25	145.31	479.6
U17579_rna1	422.7	193.61	117.35	135.55	898.65	959.47	633.4	483.38	449.2	254.42	1028	164.61	291.95	116.74	422.65	347.4	498	257.53	839.8
D14660_at	151.65	132.16	79.4	91.78	308.2	337.01	186.2	111.44	123.7	97.86	351.85	148.56	139.95	147.43	191.95	159.88	292.15	339.91	88.05
M92843_s_at	1835.7	862.67	1961.15	2265.92	2386.2	279.45	1689.2	446.18	2166.7	22.49	3340.4	673.31	2777.5	1186.38	1852.85	944.48	2118.2	41.01	2888.2
L09708_at	277.6	157.54	360.35	416.27	231.15	48.01	258.5	113.28	519.7	538.25	199.95	44.76	663.2	533.3	229.2	193.46	244.55	10.39	557.45
M90696_at	297.85	183.21	156.75	181.09	262.95	135.98	549.4	745.15	234.15	229.46	500.3	229.67	230.35	264.95	274.4	285.67	263	125.02	167.85
X80923_at	190.6	234.62	578.65	668.57	390.65	175.01	639.5	574.45	282.1	304.34	572.85	81.81	472.05	195.52	560.05	62.58	442.35	97.93	298.35
M94633_at	307.95	355.18	65.9	76.08	223.9	47.23	318.1	27.15	516.65	682.29	194.45	169.63	351.8	248.19	131.65	77.15	234.85	159.74	447
M81780_cds3	353	51.34	163	188.37	329.2	251.59	230.3	122.61	640	378.73	324.25	46.03	748.55	792.88	1001.5	1036.05	200.4	85.28	1136.55
HG4272-HT4	1431.5	183.56	957.95	1106.83	1703.6	481.54	1275	50.77	1765.65	865.57	2237.1	249.47	1404.85	149.69	1730.1	479.84	1610.35	596.59	1381.85
X69433_at	1429.15	206.69	418.85	483.87	1289.6	359.78	1235.25	585.56	1389.7	151.75	1455.3	261.06	960.15	217.58	1752.1	504.87	1969.95	1104.01	1629.9
M20778_s_at	655.5	230.23	846.1	977.5	674.65	498.3	1084.95	703.92	617.9	571.06	602.85	12.23	457.8	68.31	515.35	458.13	775.95	755.69	1073.1
U19261_at	1852.85	877.45	266.4	307.73	1986.1	310.14	1607.8	635.69	1886.65	122.68	1921.15	100.48	1817.35	346.27	1822.15	587.82	1759.4	358.93	2190.15
U31973_s_at	45.25	3.75	27.3	31.54	269.6	363.31	80.55	89.31	22.9	0.85	32.4	3.82	40.15	25.95	129.25	115.61	21.05	2.33	344.35

M81882_at	263.5	203.36	34	39.32	325.95	357.02	277.6	70.85	151.65	160.3	168.65	73.89	117.5	122.19	436.1	435.86	357.1	294.44	168.2
X17644_s_at	2054.55	363.24	1258.3	1453.81	1950.55	223.23	1045.25	1044.18	1394.7	610.94	1562.8	251.73	1722.7	289.77	1747.5	324.14	2052.15	878.3	1582.9
D17427_at	811.2	300.24	103.75	119.85	854.95	602.1	799.8	727.19	449.6	215.1	377.05	26.38	617.75	163.27	298.25	171.9	554	158.82	250.05
L04569_at	198	40.59	68.15	78.7	171.5	94.19	300.5	183.56	197.05	56.92	378.75	43.35	167.2	32.67	263.75	76.01	130.4	4.67	188.65
U57452_at	29.5	22.49	8.7	10.04	93.45	86.9	456	630.88	108.35	79.55	53.05	19.3	45.15	10.82	126.9	145.24	23.35	10.25	51
U54617_at	234.2	84.71	62.6	72.27	351.95	469.59	294.95	211.92	127.3	129.97	379.7	55.3	223.45	273.72	48.85	24.4	161	112.15	53.45
X00734_at	1126.35	337.93	172.6	199.4	481.8	282.28	702.95	478.07	442	1.56	787.45	723.3	482.2	220.33	425.05	29.06	1234.4	387.35	681.2
X99586_s_at	424.8	209.16	167.55	193.54	65.85	32.31	182	112.57	687.1	337.01	504.75	144.18	464.15	8.98	365.85	198.2	247.35	128.91	726.65
M35999_at	107.8	1.27	70.85	81.81	58.15	14.78	139.25	51.12	203.9	117.38	72.9	9.76	79.8	62.65	362.7	173.95	52.1	16.12	106.15
M57464_s_at	493.4	44.41	346.5	400.36	426.3	135.62	924.7	1198.69	602.9	55.86	795.65	674.37	446.4	405.03	289.25	167.23	394.55	88.32	411.35
U90914_at	1007.65	166.52	433.8	501.2	635.9	356.24	343.15	206.55	674.3	3.96	376.9	245.22	745.7	91.64	129.75	3.18	659.3	552.53	1096.9
L08666_at	12942.95	6193.48	6921.4	7996.81	17885.65	4592.87	17890.05	1763.59	16787.65	1423.76	13833.7	2841.58	11503.35	232.14	13358.9	1982.02	12039.95	6558.77	17423.3
X64177_f_at	23536	7849.31	9708.35	11216.76	20315.9	1239.13	18233.4	1679.1	24109.35	2012.21	16468.45	4937.09	23050.85	350.37	20679.8	5679.91	21871.1	3312.09	16691.1
U92459_at	42.95	6.15	48.75	56.36	124.45	102.88	62.3	76.51	236.3	281	61.55	27.65	330.75	267.07	274.1	353.69	174.25	68.38	155.5
S78271_s_at	1345.95	116.04	780.05	901.21	938.6	772.44	788.45	109.39	843.65	615.54	1343.45	570.71	1206.95	425.32	1449.8	324.42	1595.75	1040.08	976.15
M10950_cds2	27.5	6.51	105.9	122.33	94.9	89.94	160.7	198.27	24.45	21.14	104.75	119.01	18.4	7.64	144.25	176.42	15.75	0.92	33.35
Z35309_at	87.95	21.57	158.5	183.14	105	42.57	132.6	81.88	81.1	0.28	265.95	206.4	249.75	126.64	347.25	203.58	115.85	24.25	494.4
Y00451_s_at	508.05	356.03	931.6	1076.36	547.9	128.41	835.05	136.97	1110.35	748.9	648.4	360.77	880.5	90.23	573.15	3.32	811.2	311.27	553.8
M73720_at	637.7	348.18	280.75	324.35	988.7	334.89	1103.8	981.46	463.65	251.38	1000.5	112.15	422.5	494.69	526	225	610.8	126.43	629.05

X93510_at	4503.6	2039.86	3226.2	3727.44	5979.1	382.12	6255.05	750.45	4764.45	882.4	5452.3	953.46	4723.1	2153.85	4351.8	297.13	4455.5	1616.02	4203.8
D25539_at	388	18.1	85.1	98.29	360.85	441.59	586.9	408	574.55	361.26	470.4	96.31	141.9	6.08	315.25	338.07	855.1	730.3	545.5
HG3432-HT3	1961.3	92.07	1042.15	1204.13	1373	587.61	1597.65	608.18	1871.9	94.47	1474.4	1555.92	1187	973.97	1729	288.78	1255.45	24.4	1295.65
M20203_s_at	226.1	124.73	152.6	176.35	447.85	261.13	1003.3	970.57	267.25	93.41	227.65	19.02	505.9	476.73	463.45	398.88	431.3	403.19	242.2
D87440_at	261.9	182.15	146.1	168.86	607.95	35.43	280.65	195.52	528.55	555.72	549.2	284.96	462.55	478.92	96.3	12.87	381.65	140.22	484.9
Z47038_s_at	3742.8	250.32	1629.2	1882.32	3067.45	288.99	2919.85	290.69	2602.7	341.11	3489.15	998.51	3634	775.84	2446.3	357.23	2500.45	103.87	3036.3
D49817_at	802.5	404.18	204.15	235.82	474.95	33.3	1045.65	275.28	549.3	304.76	550.05	305.12	543.9	23.19	383.45	86.48	981.85	713.12	367.4
M90656_at	1419.85	502.4	1062.4	1227.54	1953.85	555.43	2038.75	76.72	2396.35	698.41	2568	1405.59	2235.05	360.98	2721.45	1268.62	1803.7	1009.61	2386.1
M30773_at	110.35	115.75	147.4	170.27	273.1	17.54	177	137.89	57.1	9.76	305.45	143.33	210.9	15.84	241.7	235.47	189	211.99	184.25
X15217_at	28.7	10.18	159.35	184.06	90.95	60.17	66.5	29.56	53.55	14.21	35.65	11.81	190.5	200.96	80.05	70.36	58.4	28	68.85
AFFX-CreX-3	5019.5	906.93	2935.15	3391.21	19995.05	18420.34	22118.25	24370.51	8957.3	5530.85	19241.15	22101.68	7754.2	4035.6	7488.8	726.76	6040.35	51.12	9309.35
U80226_s_at	161.4	45.25	360.45	416.42	527.65	412.6	277.95	6.29	709.35	196.65	290.05	73.89	595.4	185.4	181.25	192.55	106.2	35.5	509.25
HG1153-HT1	23817.65	9449.28	13834.7	15984.29	34485.9	345.78	36823.95	10465.53	30772.2	5717.81	23784.85	10126.97	24816.15	4943.88	30505.5	3836.2	25961.4	5617.12	26553.2
U05875_at	2150.75	809.85	1859.85	2148.83	4520.4	1839.89	3609.35	474.82	4612.5	536.13	4279.55	727.97	4007.3	312.68	2402.6	374.2	4090.5	1322.71	4624
D13305_at	1066	435.29	413.05	477.23	720.25	10.11	442.25	140.36	1315.35	867.41	1526.2	746.7	1173.4	156.27	1018.4	586.05	970.2	147.5	985.7
L10123_at	254.9	170.98	395.55	457	531.75	444.56	521.7	353.55	247.25	6.01	842.8	488.75	745.05	261.42	210.55	197.92	514.55	83.09	475.95
M68520_at	1218.95	265.8	764.7	883.46	1073.3	189.08	583.65	284.61	1535	98.15	1601.65	494.9	1270.1	165.89	986.9	10.04	1201.6	187.38	846.7
U81802_at	1679.35	119.57	1030.6	1190.77	1242.6	1285.94	1653.75	329.16	1153.9	688.44	1772.85	371.73	1550.65	492.22	1971.7	43.84	2500.35	281.78	1689.5
M24748_cds2	64.35	13.65	77.95	90.01	185.95	172.18	178.1	147.79	199.3	109.46	120.75	29.2	258.8	255.83	75.5	17.11	128.6	93.48	129
X02419_rna1	4826.45	755.68	4227.2	4883.99	3346.65	121.27	4870.05	1457.98	6786.15	552.46	5120.4	81.32	3833.15	240.49	4656.6	434.87	4811.95	2125.78	5386.95
U03911_at	205	164.76	47.65	55.08	125.6	64.91	56.75	6.29	196.75	106.7	66	5.52	263.95	339.62	117	87.68	63.15	40.38	106.15

X04143_at	83.95	8.13	101.2	116.96	331.6	47.23	150.7	22.49	177.65	122.12	245.4	224.86	119.15	56.5	175.75	125.51	133.2	40.73	130.45
U43083_at	780.65	148.14	528.45	610.59	1222.4	421.86	773.75	296.21	824.2	181.3	686.7	256.82	773.9	286.38	621.5	277.04	1123.65	484.16	803.1
M57732_at	200.85	125.94	130	150.19	274.05	34.72	187.1	16.12	393.75	184.91	259.6	27.72	290.2	130.81	239.8	4.81	183.45	19.73	192.5
D87683_at	121.85	54.09	233.3	269.55	449.15	134.7	99.2	25.17	161.55	2.05	171.15	114.34	121.95	1.34	71.55	51.41	277.65	185.62	292.65
M80359_at	415.7	338.42	772.7	892.79	398.7	300.1	197.95	45.33	1015.15	432.68	374.35	228.18	561.35	461.95	490.7	253	699.95	250.39	621.65
L19297_at	194.1	131.24	71.05	82.1	211.05	32.74	467.75	176.85	301.35	95.53	204	48.08	556.6	465.42	465.25	358.57	408.15	300.59	237.95
HG1111-HT1	1396.4	301.23	873	1008.62	2098.8	1019.65	1558.85	518.38	1517.45	550.48	1311.55	717.93	1308.85	229.6	684.25	752.29	1417.85	372.29	1757.7
U62800_at	235.9	69.72	86.2	99.56	579.6	542.78	170.45	7.28	359	181.73	1626.1	1114.68	190.15	56.36	423.55	438.34	146.85	51.12	264.7
M95610_at	1573.9	317.07	746.15	862.03	1636.25	869.95	1624.55	722.88	1569.2	176.21	1802.7	23.62	1336.4	971.99	1084	544.76	1966.25	285.6	1705.6
D87011_at	171.15	103.87	82.2	95.04	114.9	51.48	154.75	109.39	110.2	25.6	167.1	93.34	100.55	33.59	131.6	90.93	111.5	38.33	138.25
U70321_at	3421.1	115.97	1215.55	1404.38	2817.8	1341.1	3395.7	1775.83	2682.35	578.34	2397.8	1419.59	3259.15	399.73	2774.7	21.07	3021.4	1007.77	3296.25
U43672_at	448.25	89.87	23.95	27.65	170.15	65.27	86.3	55.86	78.7	54.02	302.3	164.9	288.1	232.78	157.55	109.53	77.95	3.32	195.15
U43843_at	376.85	142.48	201.2	232.5	163.45	80.12	490.9	517.6	163	85.7	211.8	16.83	234.15	23.26	365.65	351.64	198.45	88.88	317.35
Z49208_at	19.4	2.69	17.45	20.15	44.85	30.48	384.8	523.97	31.75	17.75	31.7	22.63	35.25	8.84	69.8	48.51	25.3	4.81	55.35
U53468_at	67.5	7.64	25.7	29.7	38.3	17.96	133.25	110.1	81.55	20.01	60.95	22.56	362.4	41.44	90.8	41.3	86.6	6.22	56.85
M32373_at	179.35	2.47	121.85	140.78	979.45	793.59	515.6	490.02	717.45	659.09	1088.25	447.81	324.85	28.35	217.45	137.67	188.9	1.98	435.05
X68090_s_at	158.1	79.48	100.5	116.11	149.5	110.59	77.6	56.71	673.85	134	136.55	78.14	219.75	38.54	122.85	73.04	103.2	7.92	128.55
X66533_at	236.85	309.92	13.25	15.34	61.55	53.39	729.2	596.8	461.05	579.19	61.8	58.27	195.8	175.08	68.8	26.3	22.4	0.99	37.9
U26403_at	487.6	377.74	152.8	176.49	343.75	329.58	948.25	428.15	169.8	31.11	1238.5	1150.04	627.4	109.46	622.45	522.76	466.4	465.7	486.7
L06633_at	207.05	181.23	33.55	38.82	469.4	357.37	79.3	61.8	55.35	25.24	92	74.1	323.95	192.12	67.35	22.7	270.4	330.93	110.55

U89355_at	163.65	111.37	341.1	394.14	467	302.64	152.9	54.73	474.4	81.46	364.8	312.26	80.55	24.82	42.05	14.07	173.9	136.47	447.55
L04953_at	741	10.75	1090	1259.36	372.15	79.41	531.25	132.3	914.05	171.33	667.95	53.67	1123.25	334.67	1035.25	106.84	728.6	712.62	1163.55
J05401_at	90	5.52	144.3	166.74	170.85	104.02	412.15	429.85	705.3	300.94	631.95	278.95	163.35	10.25	265.45	183.07	200.25	192.4	564.15
Y10517_at	92.55	32.17	18.9	21.78	283.05	291.12	34.1	25.46	60.3	1.84	129.3	129.68	198	223.16	37.45	16.33	103	33.38	76.3
U59423_at	433.55	211.92	195.65	226.06	243.1	138.73	169.25	41.37	430.85	523.19	1170.55	775.91	916.05	204.85	481.05	200.04	308.8	257.81	869.05
X53800_s_at	812.8	20.93	1118.25	1291.95	733.5	1.56	279	200.68	573.05	465.77	330.65	218.28	243	10.18	630	203.93	451.7	122.47	765.25
M26901_s_at	781.05	406.8	888.8	1026.86	957.05	268.63	620.05	182.93	1113.15	313.6	470.15	216.45	917.85	278.39	963.75	924.68	626.9	283.13	1278.2
U67988_at	67.7	55.44	12.8	14.85	715.05	833.89	247.5	16.55	215.9	263.75	420.75	449.79	329.75	332.55	75.1	59.68	357.15	411.18	116.8
U52522_at	3601.4	872.15	3186.15	3681.27	4377.2	730.87	4066.8	1994.61	4523.8	717.43	4465.5	601.46	4562.5	1452.26	3074.6	1157.82	4752.45	1707.45	4670.55
M32598_at	570.65	125.65	453.5	523.97	1608.25	680.17	1118.3	944.98	426.45	139.51	541.5	435.58	909.8	928.86	975.3	886.43	920.65	813.95	954.35
U01120_at	474.45	192.55	44.85	51.83	1199.75	241.19	453.35	118.86	675.95	736.73	343.85	214.75	483.9	220.19	49.8	24.32	396.75	144.6	439.45
J03592_at	22126.05	13567.19	11329.35	13089.75	33166.3	5255.78	30692.6	5794.6	26136.8	614.05	23931.5	1273.64	19489.9	4142.09	20862.05	2404.66	24588.75	10912.85	31037.9
M61826_s_at	364.75	109.25	32.75	37.83	219.1	238.15	501.15	589.51	487.05	495.75	579.35	148.56	194.25	101.33	72.8	6.65	282.7	73.4	173.1
X56807_at	157.95	117.03	85.65	98.92	232.45	82.94	154.2	83.3	181.9	185.69	467.35	350.23	262.7	299.39	293	297.55	325.2	79.62	299.4
M21142_cds2	18999.15	4558.51	8170.7	9440.3	13376.45	6924.77	11804.3	9444.68	15365.8	4109	13672.9	6984.38	15619.3	2119.76	14548.5	2904.09	15898.45	3816.75	12694.85
HG1102-HT1	4882.7	2175.2	2192.15	2532.79	2803.7	1006.78	2137.8	542.63	2093.3	106.91	2020.6	191.48	2580.85	1384.3	2262.1	1194.73	2994.05	145.45	2793.1
L05568_at	566.1	455.52	350.25	404.68	172.4	74.67	203.45	189.15	76.9	36.35	266.5	236.46	487.25	378.51	126.35	6.15	306.15	140.22	224.3
HG4679-HT5	71.25	12.66	62.8	72.55	361.15	429.57	137.5	80.47	66.8	39.88	87.8	47.23	302.15	311.62	239.25	25.53	361.95	48.01	46.2
M95809_at	763.05	77	1201.3	1387.91	801.85	81.67	770.45	977.43	1480.85	206.4	1383.4	103.38	998.2	108.33	1274.25	101.19	1130.65	40.23	1421.15

X01060_at	2912.1	918.67	969.5	1120.2	2423.05	948.3	2057.85	14.64	3060.45	295.08	993.7	978.92	3230.5	1698.61	2577.9	2119.2	2646.8	2346.18	1728.3
U31201_cds1	1157.5	510.67	721.4	833.54	665.8	355.53	907.25	885.23	777.05	526.72	970.05	268.63	189.75	0.07	785.25	387.57	991	369.25	736.3
U73514_at	1993.4	1828.01	819.7	947.1	2758.7	153.16	3224.9	1038.88	3306.35	202.87	1450.35	784.82	1232.35	1214.74	1635.55	309.64	1469.85	17.75	951.75
U57341_at	1251.55	290.69	275.75	318.55	1395.35	778.88	2051.85	1371.43	839.65	175.29	391.2	449.01	1252.7	296.84	547.95	323.22	389.15	471.43	1146
X04654_s_at	6644.15	142.91	3688.9	4262.02	4920.5	804.97	5635.05	1028.2	5792.55	1987.04	4830.25	8.7	5168.5	2195.71	5360.3	833.4	6955.5	1467.1	5026.8
U48730_at	671.9	239.14	97.7	112.85	279.35	55.23	129.05	24.96	166.45	30.33	419.65	280.37	197.5	61.24	101	42.71	285.95	49.71	408.85
AFFX-M2783	645.1	214.39	243.8	281.71	562.25	21.99	570.35	22.84	568.1	196.15	887.55	25.24	993.85	275.7	1721.35	1505.36	709.8	34.37	608
U49020_cds2	262.1	38.04	496.05	573.11	502.7	189.08	471.6	437.42	421.45	19.87	446.85	180.38	317.6	150.76	486.05	30.33	382.3	71.98	208.4
HG2264-HT2	615.9	354.54	220.35	254.63	1536.7	562.86	835.95	532.24	869.45	258.59	1585.75	368.19	710.6	434.59	1231.25	6.43	1075.5	949.79	1530.75
S71043_rna1	1809	496.95	476.8	550.84	1195.15	1041.64	1113.55	937.98	1142.35	118.3	1087.35	450.64	1337.1	193.04	1085.95	737.44	1702.75	331.14	1448.9
M34516_r_at	297.8	249.75	332.7	384.38	742.3	382.83	224.65	153.09	345	199.55	579.75	427.02	611.15	125.51	692.95	718.77	460.6	326.82	902.5
K03494_s_at	117.75	14.92	103.75	119.85	95.8	6.93	179.7	117.66	88.1	4.24	257.5	189.36	158.35	42.92	114.05	2.76	104.05	41.93	203.5
X98263_at	1470.7	213.26	1095.55	1265.79	1455.6	641.2	1838.7	591	1198.55	52.26	1115.35	760.63	1516.8	203.79	1259.55	333.4	1689.95	509.75	1546
D11086_at	2769.7	861.54	1125.5	1300.37	2952	1387.34	3013.5	1253.7	1910.85	1081.1	3465.15	1457.13	1907.6	631.16	2092.55	1438.61	526.85	325.34	2844.3
M81883_at	223.95	246.43	188.3	217.51	148.95	33.16	92.75	88.18	88.2	1.98	243.15	112.5	183.65	147.71	284.45	253.92	106.9	24.04	67.65
S78693_f_at	329.55	224.51	248.5	287.09	524.05	131.03	244.05	129.75	641.7	254.7	324.6	319.33	498.55	675.22	318.1	101.12	171.95	61.45	168.75

D26361_at	372.95	383.46	369.65	427.02	566.75	321.1	887.4	462.59	656.65	100.48	749.6	286.24	776.35	47.16	651.25	3.32	630.2	357.09	308.7
X89399_s_at	237.75	106.84	107.75	124.52	164.65	181.8	77.5	35.07	216.4	145.95	374.45	401.99	194.65	65.97	271.75	327.32	142.15	18.46	167.35
U79751_at	164.45	155.49	171.45	198.06	144.8	58.12	53.65	52.68	157.3	113	29.45	17.32	329.9	428.51	218.5	239.85	117	69.58	46.35
Y08302_at	3251.75	787.65	991.1	1145.09	1309.05	1189	2156.7	492.85	1568.95	531.67	2522	1362.59	1752.3	467.96	1270.35	475.81	2083.25	409.34	2362.5
U19247_rna1	736.55	158.32	787.4	909.76	950.45	60.03	716.95	6.01	1260.2	271.67	778	341.25	850.9	427.66	615.2	588.6	566.5	397.54	686.5
X81198_at	1835.8	499.5	1392.6	1608.95	1505.05	295.78	1383.15	497.73	1813.55	557.84	1031.6	1089.65	2361.55	1002.04	1647.4	108.75	2210.4	726.2	1786
M65199_at	533.45	41.79	247	285.39	466.05	93.41	557.75	133.57	555.8	112.29	615	197.57	657.45	232.14	493.7	36.91	480.25	19.45	605.6
HG1103-HT1	1251.95	187.45	452.8	523.12	746.1	620.13	723.5	712.2	608.95	194.38	532.6	665.39	653.2	63.92	784.65	162.42	1282.3	19.52	1132.45
U83303_cds2	524.4	0	301.4	348.18	158.05	57.91	451.6	499.36	310.25	321.38	224.85	170.91	199.1	14.42	349.95	52.82	101.95	43.91	378.95
M11591_at	483.9	175.08	131.55	151.96	1095.2	464	604.85	434.52	628.7	516.05	625.35	124.1	686.65	622.89	275.25	263.11	220	89.38	241.85
U10550_at	421.25	347.97	50.5	58.41	690.1	703.43	571.1	539.66	309.9	96.31	156.05	60.46	403.3	34.22	66.8	22.49	464.85	13.08	183.9
U58032_at	165.55	21.57	48.05	55.51	684.05	775.63	417.8	291.89	587.65	617.8	152.6	19.37	109.35	50.28	557.25	711.42	691.1	82.73	565.75
L40402_at	161.25	74.88	41.1	47.52	575.6	682.92	323.95	315.58	181.75	134	259.85	245.58	504.6	460.33	337.6	285.81	164.85	70.07	321
M27093_s_at	34.9	1.84	223.3	257.95	134.4	56.71	79.35	76.72	216.45	183.07	238.8	31.54	69.6	27.01	93.6	58.69	55.1	18.81	265.05
X73874_at	317.35	160.3	326.75	377.52	184.7	112.71	469.9	363.74	311.9	30.12	593.3	199.4	134.65	124.95	267.85	102.04	293.1	151.04	314.85
U21943_at	358.4	68.87	13.15	15.2	227.15	174.58	541.65	723.3	483.75	131.03	713.2	603.02	401.25	20.44	126.45	148.42	481.6	256.82	562.25
AFFX-BioDn-	3133	215.1	1251.9	1446.46	5981.85	3403.66	6243.25	5219.51	4379	2028.83	4918.3	4989.06	2617.8	1923.05	3652.7	884.31	3416.15	386.15	3001.5
U18271_cds3	498.95	129.05	389.5	450	555.8	303.91	720	738.64	325.9	96.59	526.8	227.97	609.15	45.33	293.6	294.02	437.75	34.86	426.5
U44755_at	3538.3	190.92	1903.75	2199.6	4127.35	770.39	3713.9	511.38	3010.95	516.12	3441.6	1473.89	3463.35	377.81	3224.3	469.09	3488.1	190.92	3623.4

M77349_at	5747.7	1771.87	2287	2642.32	9298.65	2891.15	6770.6	743.31	4985.65	275.7	4228.75	1229.31	4661.65	24.4	4883.15	1028.49	5324.3	1110.3	6300.1
AFFX-BioB-5	1369.65	150.26	1026.5	1185.96	4555.15	3709.41	4242.35	4137.21	1826.35	893.57	2925.65	2943.05	5020.95	3261.81	2189.3	840.04	1871.35	182.93	2093.75
L35269_at	355.1	304.9	599.05	692.19	1011.55	528.99	330.9	250.03	868.1	154.29	242.5	99.7	727.55	207.25	881.55	679.6	717.65	283.9	782.95
M20137_at	732.6	346.06	501.9	579.83	449.25	337.08	350.25	327.18	309.4	147.22	779.1	130.25	707.05	191.56	755.4	591	947.2	260.64	767.05
X03663_at	548.95	323.08	206.2	238.29	549.45	478.5	1352.7	486.91	945.85	165.82	688.8	162.21	690.85	134.28	883.35	18.88	1606.7	574.03	844
Z30425_at	624.5	97.16	142.95	165.11	253.6	100.55	464.7	113	297.45	70.78	482.65	409.63	206.1	82.02	222.4	34.51	594.6	6.22	167
U32674_s_at	2167.65	339.06	469.1	541.93	1269.35	120.84	382.15	21.43	508.9	525.95	1673.65	1304.12	1457.1	965.91	1670.85	251.52	1005.75	204.28	487.45
Y00486_rna1	9270.45	2073.73	4845.1	5597.88	11483.25	3430.81	10392.95	242.61	8902.55	1260.7	10561.6	3683.46	9475.7	1785.16	10815.35	446.96	6952.7	1188.08	7703.8
D38047_at	17301	9171.32	9415.7	10878.7	17473.3	5496.2	21784.9	6181.39	15495.5	550.13	17488.9	4890.92	15176.4	6873.22	14110	3555.19	11756.95	2560.5	14360.15
U80456_at	119.75	12.8	48.2	55.72	130.3	93.2	404.1	461.18	140.9	18.24	304.8	204.92	441.75	496.74	289.25	303.56	97.8	35.92	132.15
X65550_at	385.45	381.2	30.45	35.14	84.25	44.34	128.3	47.66	310.9	234.05	222.3	167.87	394.2	366.14	187.7	28.14	93.85	10.68	260.05
D50915_at	66.8	30.55	35.3	40.73	185.6	108.89	156.35	131.59	482.6	437.42	317.8	134.49	187.05	175.01	366.2	406.16	102.15	26.94	262.6
X59710_at	138.65	163.13	20.95	24.25	216.65	60.46	264.85	247.28	320.65	305.82	221	100.13	56.35	29.2	30.9	17.96	212.95	72.2	94.8
U41815_at	168.6	94.05	49.65	57.35	921.6	604.72	258.05	255.9	320.45	6.86	172.35	119.71	256.15	149.55	113.75	40.52	236.6	20.51	179.1
U00946_at	528.75	166.67	129.95	150.12	767	229.81	206.25	128.91	173.85	81.95	758.55	181.23	402.05	379.93	604.4	701.73	252.2	194.6	155.25
L40397_at	6893.35	2407.34	3851.95	4450.46	5691.9	39.46	7173.95	4588.49	4295.75	2288.41	4846.65	1534.92	3920.65	1848.02	6357.2	1060.38	6605.35	1678.46	7534.1
X52896_s_at	336.2	39.32	189.55	218.99	532.1	338.14	529	297.55	416.4	225.99	422.75	371.58	402.05	11.67	722.8	215.38	296.95	55.65	604.8
U09196_at	3591.05	704.49	1658.9	1916.68	3839.2	1036.19	4064.85	1548.49	2460.05	661.22	4299.75	861.04	4114.65	924.12	3243.55	954.66	3576.9	613.49	3687.7

M84711_at	51070.35	33303.11	25597.2	29574.46	60811.95	3665.43	61417.85	32602.93	71494.5	1115.25	56677.75	20334.34	51252.25	4597.82	55420.9	7569.86	46644.6	13519.6	57535.2
X75962_at	1138.45	164.12	387.6	447.88	1450.7	1051.61	523.4	235.47	1446.3	498.79	1354	183.99	624.95	516.12	891.5	113.84	897	221.75	1262.8
L36870_at	487.55	192.26	141.3	163.2	171.85	7.57	504.5	151.6	692.4	409.41	377.6	273.93	332.3	75.66	562.8	588.74	471.45	325.76	547.5
M29550_at	761.65	82.38	475.1	548.86	541.4	485.64	540.55	154.93	515.4	67.03	675.75	717.22	1020.5	63.64	398.25	372.15	588.75	608.89	480.25
U69140_s_at	329	127.14	426.05	492.22	497.15	310.21	336.5	26.73	746.35	644.53	563.9	157.68	582.05	30.48	559.65	178.4	337.85	25.39	303.9
X84908_at	62.8	5.94	101	116.67	112.7	64.21	181.4	46.53	267.15	339.2	186.2	185.69	317.1	402.63	556.45	717.64	196.45	124.24	170.35
D79999_at	2160.55	247.42	2269.35	2622.02	2799.6	763.11	2548.75	583.01	2643.6	729.88	2340.8	284.54	4124.4	337.43	2906.05	521.07	2675.85	476.52	1614.65
M21539_at	1563.4	524.53	1037.75	1199.04	1280.05	432.25	883.7	223.02	1773.7	34.22	1242.65	129.61	1197.75	136.12	1673.15	835.16	792.65	212.34	1334.1
U62739_at	5356.7	142.98	2834.8	3275.32	5003.15	726.27	5293.15	890.74	5062.95	146.72	6348.3	1053.87	4468.4	302.5	4384.15	896.26	4898.55	200.61	4564.05
AFFX-HSAC(	143882.09	39464.62	50325.7	58145.25	114825.25	48506.46	104905.45	7183.71	83759.59	78854.42	95052	13133.38	106503.55	39252.56	92987.8	3001.67	117640.95	16324.34	93759.66
L19183_at	650.9	356.66	823.3	951.2	866.3	102.67	1000.45	32.88	700.9	159.81	594.55	668.43	1684.7	969.58	1461.1	821.94	1272.75	1226.76	486.95
S37730_s_at	19751.55	1103.58	11551.55	13346.43	16941.35	5345.94	19556.3	7490.1	10313.6	4339.23	11042.5	2942.98	11068.7	4116.49	11117.7	4775.38	14912.1	3076.9	17138.15
M23197_at	229.25	149.13	274.2	316.78	487.05	189.01	284	188.51	386.6	51.48	796.5	604.58	267.2	34.37	585.95	379.65	218.2	211.14	424.4
U63312_at	80.2	16.4	49.8	57.56	314.15	326.47	83.35	33.3	89.3	5.8	109.45	101.89	52.1	7.78	52.9	8.34	46.15	4.03	91.15
L10678_at	5308.25	1041.92	2473.45	2857.77	3927.4	1498.08	5142.9	2147.48	5363.45	1148.41	4133.55	2507.47	4134.85	172.18	3759.65	461.67	5277.7	812.61	6279.4
M22490_at	1894.65	298.19	1325.7	1531.73	2177.3	156.55	2612.8	257.81	2260.65	667.44	2811.45	927.65	2219.55	817.63	2648.75	1389.39	1655.55	1906.71	2132.1
U21049_at	485	215.81	319.75	369.46	86.5	41.3	660.6	662.28	348.35	364.8	933.65	596.44	245.15	122.82	943.1	1198.12	216.8	31.4	682.7
U83117_at	421.6	138.17	198.1	228.82	224	37.76	129.2	45.11	300.45	181.94	366.55	232.43	480.8	129.97	89	39.74	188.15	144.89	97
X81003_at	2364.7	561.3	1252.45	1447.09	2349.65	103.59	2253.05	423.2	2171	804.69	2548.85	279.94	2111.8	201.53	1720.25	791.89	2299.8	1408.13	2484.35

L29339_at	254.8	50.91	93.25	107.69	432.8	50.91	256	110.31	172.1	12.87	248.9	28.14	206.9	155.85	263.1	15.13	157.2	53.32	152.25
M20543_at	520.8	142.98	200.75	232	202.45	106.42	552.15	125.51	163.45	98.36	417.3	229.1	1321.65	920.44	785.65	150.83	284.8	235.47	862.5
X72727_at	1606.05	56.64	1310.8	1514.48	1074.3	507.84	1030.9	444.49	1843.6	1316.07	1405.65	366.21	1287.9	389.9	1798.35	216.02	2128	272.24	1634.3
D28124_at	8275.6	1137.45	4578.95	5290.36	9386.55	399.59	8960.9	3259.48	9963.05	1535.62	7773.95	1250.94	8887.05	482.32	9640.05	738.29	8087.95	2054.78	10811.8
U01337_at	3957.05	216.02	2291.35	2647.34	3770.6	317.63	3446.25	84.92	3468.75	452.76	3388.2	234.19	2939.2	800.02	2257	1512.64	3838.15	623.03	3064
X92110_at	608	183.28	448.6	518.31	191.25	145.17	1091.45	763.04	745.75	117.45	552.95	242.18	231.15	59.47	215.45	121.27	495.05	76.86	570.9
U45255_s_at	213.4	9.19	144	166.31	258.85	11.81	226.3	51.05	224.8	24.89	244.35	33.73	232.6	15.7	198.9	42.14	164.6	5.37	317
L27624_s_at	249.75	254.77	207.85	240.2	496.35	550.62	214.8	116.11	1030.8	51.76	1291	607.12	1377.9	747.13	1216.55	547.51	2719.75	354.76	3052.85
HG1604-HT1	96.5	36.2	11.7	13.58	155.8	161.93	188.45	3.32	204.6	106.07	135.2	95.74	56.4	34.08	75.65	16.48	172	129.54	145.85
X71348_at	172.8	57.13	40.4	46.67	163.35	15.77	220.4	206.19	153.8	18.38	144.55	33.45	128.45	54.8	164.45	7.99	139.25	33.02	150.25
U76388_at	3153.85	643.4	1632.55	1886.21	3236.65	620.34	2400.95	320.39	1906.75	627.27	2804.2	693.39	3550.1	1149.61	4000.15	13.36	2963.45	591.49	2882
U77735_at	137.95	83.23	448.5	518.17	80.15	32.17	160.7	26.16	100.95	62.58	185.2	76.93	131.05	27.93	187.25	81.25	127	52.33	162.85
U08015_at	114.2	61.24	314.2	363.03	288.45	233.84	439.5	500.91	162.9	43.56	149.7	13.86	393.75	6.29	429.45	370.45	201.65	110.1	194.75
L35592_at	112.75	12.52	53.45	61.73	488.3	575.16	194.3	72.12	268.85	118.02	556.5	542.92	173.3	103.8	171	65.9	126.95	50.56	172.7
HG2238-HT2	2822.55	21.57	1065.45	1231	2897.8	327.39	2859.15	876.32	2753.9	16.83	3006.95	292.81	3328.6	1067.17	2971.2	1259.07	2798.25	1223.22	3121
U03494_at	401.05	38.82	785.1	907.08	665.95	593.19	208.95	8.7	419.7	160.8	199.05	153.65	773.9	617.59	225.75	71.06	282.1	93.2	986.15
D90042_at	269.7	269.12	258.45	298.61	95.25	75.02	93.25	32.31	75.75	17.61	183.4	50.2	458.25	237.52	501.45	364.8	399.25	134.42	241.8
U19345_at	639.65	111.51	364.25	420.8	605.75	132.58	848.9	416.34	900.95	414.72	609.75	271.88	696.05	335.95	757.85	84.07	362.7	238.86	865.3
X55019_s_at	1362.45	157.9	524.05	605.5	2213.75	1681.29	911.5	331.35	1930.9	651.25	2282.15	1214.31	1804.55	891.59	1711.1	155.99	1604.35	59.18	1579.45
D83657_at	117.9	53.46	49.1	56.71	133.45	97.09	108.35	38.96	127.6	43.84	563.55	586.26	139.35	58.48	151.4	42.28	79.3	4.24	101.55
U94320_at	296.15	287.58	7.9	9.19	118.95	110.66	257.75	293.8	397.05	523.75	276.25	44.34	312.2	243.24	321.7	114.27	388.1	196.43	211.4

X65663_at	23.6	11.88	263.95	304.98	242.6	143.26	106.55	47.87	157.3	52.33	115.65	99.35	61.6	33.09	109.6	16.12	45.9	14	89.25
U43885_at	139.05	154.08	173.8	200.82	70.3	9.62	63.85	0.35	258.85	260.43	41.85	1.63	357.55	99.07	173.2	198.56	245.4	194.88	95.8
L14778_s_at	952.6	435.58	532.05	614.69	1394.9	308.44	954.3	323.01	1665.9	1012.86	702.1	415.92	1426.6	121.34	1233.8	588.74	1263.9	3.25	1490.3
D79989_at	796.15	401.99	636.5	735.39	1497.95	451.63	1021.3	640.36	891	91.08	978.05	19.02	228	95.46	549.4	251.31	447.6	464.99	1274.8
X77166_at	184.25	8.13	487.25	562.93	232.65	72.62	163	108.05	140.4	13.01	201.35	128.34	150.1	0.71	142.3	70.71	296.9	279.31	122.05
U52112_rna1	2852.95	1056.91	1031.75	1192.11	2677.7	470.65	1994.55	326.61	2254.2	100.69	2496.6	620.42	3081.1	333.9	2109.85	910.54	2544.95	294.23	2509.4
U46194_at	462.6	185.54	356.85	412.31	301.65	246.99	637.25	760.21	583.35	680.73	707.15	406.23	312.05	335.66	209.35	27.22	423.65	14.5	450.5
Z48510_at	220.85	142.76	16.7	19.23	542.05	257.46	212	126.85	194.55	56.36	153.15	182.22	147.95	87.89	76.2	41.58	127.2	115.26	226.25
U66497_at	200.1	171.26	184.6	213.26	155.4	133.78	207.6	19.94	338.05	100.48	131.5	118.23	107.55	108.26	246.6	235.04	403.1	418.04	293.4
M92439_at	741.8	3.54	583.1	673.73	958.2	154.15	350.4	344.36	1195.1	268.84	750.95	148.85	793.15	361.97	365.1	248.76	1246.7	669.49	1047.7
L41351_at	3810.05	73.47	1377.35	1591.34	3597.3	13.86	5241.75	413.3	4034.5	1277.18	4814.4	517.18	4348.85	178.12	4017.75	1274.98	2976.3	1095.45	4317.25
S62028_s_at	538.1	350.72	94.95	109.67	400.15	107.98	253.4	128.41	214.75	27.79	285.45	211.78	335.7	39.74	703.95	18.74	190.8	152.31	593
U41163_s_at	1074.95	867.27	296.3	342.38	1255.1	213.97	780.2	763.68	322.8	1.84	355.1	148.63	1651.8	115.12	1518.95	336.51	1457.65	834.6	1306.85
AF001620_at	45.15	35.85	148.4	171.4	429.8	558.33	68.2	27.15	76.1	2.26	58.6	46.67	64.2	10.75	218.1	244.8	196.35	214.32	70.7

V00536_rna1	199.4	100.41	255.1	294.72	411.2	98.71	351.7	395.27	593.7	559.89	219.8	36.63	685.95	358.01	350.5	344.5	425.7	90.51	681.15
J05096_rna1_	86.9	39.03	16.2	18.67	142.05	147.43	254.6	101.68	96.1	89.8	456.9	198.84	294.2	365.29	279	95.32	265.6	132.09	470.35
D26561_cds1	219.95	61.02	239.9	277.19	263.25	107.13	75.5	25.31	290.15	149.27	117.55	118.58	183.3	227.12	46.55	6.29	71.95	15.91	310.35
X81836_s_at	244.5	216.66	70.9	81.88	137.55	168.22	177.95	57.49	677.8	456.79	384.1	94.47	480.05	170.2	88.3	31.96	400.65	13.93	302.6
L05188_f_at	919	281.15	394.95	456.3	1611.25	1088.45	1347.15	379.08	889.45	154.64	974.5	422.71	665.6	512.79	1388.1	440.95	757.3	92.07	751.65
L13042_at	438.85	43.49	473.65	547.23	595	628.34	736.75	627.27	469.4	464.71	846.65	466.76	228.1	66.33	188	84.99	427.15	418.82	698.1
U94836_at	1533.6	501.06	767.8	887.14	653.3	575.44	1033.35	102.18	1547.4	363.88	1513.35	269.9	1140.15	178.26	1539.55	30.33	1634.75	426.03	1546.9
Y10209_at	543.3	19.8	302.85	349.95	1048.35	844.36	322.45	284.33	437.95	89.31	1265.15	870.8	287.65	226.49	304.9	75.8	429.65	317.56	191.85
M32053_at	834.15	169.63	248.85	287.58	712.25	672.95	392.85	171.76	165.45	42.5	574.15	299.04	806.5	661	483.35	471.43	762.9	238.58	843.5
M65292_s_at	357.75	23.26	308.75	356.74	142.2	114.69	156.35	102.74	404.7	46.53	87.9	33.09	219.55	241.34	450.15	268.77	215.65	127.63	490.15
HG36-HT410	194.2	88.95	226.95	262.27	492.1	291.75	301.6	188.94	553.85	165.25	405.6	311.41	504.2	7.5	606.95	384.03	574	70.14	620.4
U32986_s_at	5214.05	28.5	3158.05	3648.74	3871.3	191.2	5880.2	389.76	5270.3	1086.26	4108.6	1079.19	5178.9	1311.54	5211.8	176.78	5493.95	822.01	4610.65
X79200_at	6947.4	204.35	2225.5	2571.32	5771.25	713.97	5034.7	1811.89	4467.6	512.23	5079.45	450.92	6739.45	1536.61	4710.2	1279.86	5981.15	2780.84	5786.1
D38496_s_at	760	410.12	579.35	669.42	1427.95	669.14	803.2	606.98	677.5	23.62	1447.8	388.06	515.8	419.31	1245.3	213.26	1158.5	1152.87	614.35
X12451_at	2444.6	745.01	1676.7	1937.19	2261.7	895.9	2232	16.83	2167.5	2036.47	1711.55	141.21	1874.4	88.39	2451.65	262.69	3324.25	744.51	2809.8
M23294_at	2393.55	363.67	1391.4	1607.54	1647.35	314.31	1365	1384.66	3733.05	388.84	1537.75	1217.57	2716.5	44.97	1937.5	70.99	1701	47.52	2478.45

D00408_at	132.35	71.77	8.05	9.26	248.45	338.49	64.7	57.28	59.3	9.62	56.1	50.35	231.05	123.11	36.25	34.01	19.7	14.99	113.75
D88460_at	79.5	22.91	56.3	65.05	172.3	168.86	108.05	92.7	147.2	72.12	249.2	254.42	132.75	2.05	421.6	231.65	157.85	82.24	227.5
L27071_at	44.3	1.56	19.65	22.7	38.6	6.22	147.9	159.24	407.1	480.41	472.5	365.86	257.65	287.44	174.45	227.33	134.25	97.65	409.2
X86401_s_at	139.75	36.13	42.65	49.29	243.75	102.46	352.05	39.81	185.35	159.88	174.6	97.86	153.7	67.03	89.65	24.54	120.55	9.69	179.75
HG2994-HT4	1485.3	496.81	862.9	997.02	816.4	129.97	1499.6	492.29	915.6	58.55	2184.8	1362.88	1548.8	209.59	1687.7	692.12	1476.65	1107.4	1044.25
D29833_at	291.55	165.53	86.8	100.27	360.65	197.21	282.5	148.35	219.85	89.02	157.05	69.37	151.3	2.55	123.7	16.97	271.65	192.26	881.95
M68516_rna1	97.95	11.24	19.55	22.56	115.95	52.4	146.6	55.01	148.25	52.54	81.4	4.53	73.15	11.67	201.5	159.66	63.55	5.59	286.9
D89501_at	313.8	291.75	264.35	305.4	167.95	48.58	201.85	113.35	211.1	110.03	120.35	6.86	427.1	105.22	539.5	705.98	223.6	222.88	221.45
HG4582-HT4	110.2	71.56	278.05	321.24	637.5	235.61	142.5	139.87	239.4	146.37	233.3	106.77	261.95	276.69	426.35	272.17	210.6	134.77	271.1
U24153_at	434	413.52	246.45	284.75	997.2	915	737.7	943.85	724.95	869.1	370	229.1	647.85	739.42	1390.5	964.21	257.9	99.56	608.1
U34844_at	57.5	4.95	59.75	69.08	507.85	567.74	98.45	76.44	314.25	192.26	804.05	643.4	290.8	18.38	186.9	132.79	107.15	36.56	78.2
M82827_s_at	447.45	355.18	105.1	121.48	83.65	13.65	605.85	712.55	374.25	400.01	358.45	235.96	355.7	90.37	140.15	66.11	98.25	24.68	416.5
U48231_at	237.6	73.26	309.35	357.44	603.4	478.15	893.35	716.94	353.5	258.66	416.1	421.01	410.15	381.2	602.1	144.25	432.35	64.84	324.95
M28212_at	495.15	45.47	306.15	353.77	615.85	295.22	294.5	211.57	721.4	137.46	1077.4	464.43	686.75	299.6	618.25	126.22	387.25	190.99	660.2
L48546_at	2473.15	45.89	1605.65	1855.09	2142.3	1325.54	2682.7	2261.47	2596.25	551.47	3325.25	1601.38	2301.35	1483.02	2754.75	647.78	4570.55	162.42	3381.45
D78132_s_at	2528.8	149.48	1172.4	1354.53	1862.75	23.12	1924.05	512.72	3098.25	658.53	1981.5	654.07	2279.05	521.07	2469.8	260.36	1968.85	514.42	2795
M96132_at	91	52.18	29.4	33.94	345.5	416.63	148.55	171.19	383.45	479.21	187.05	124.8	57.3	1.7	111.15	76.16	53.35	0.21	109.7
Y00815_at	2855.45	1490.23	2243.55	2592.18	1778.7	788	1978.45	1389.96	2212.3	1917.96	2418.4	998.43	2455.8	1325.97	2801.35	1081.1	2864.55	664.89	1869.05

M81757_at	100567.9	91651.5	37381.6	43189.94	158135.09	13467.56	133701.45	37080.61	120950.66	12930.23	118888.2	1932.95	98575.6	40006.83	98411.2	22752.86	90395.25	42794.88	104922.7
U20980_at	172.4	44.12	114.25	132.02	122.5	25.6	206.45	46.6	192.2	77.5	238.25	63.57	192.6	21.35	145.95	76.01	145.2	70.71	148.85
Z14093_at	333.55	56.21	212.2	245.22	512	100.27	545.4	221.61	361.25	125.37	302.25	63.99	302.6	74.81	317	2.69	374.3	332.76	364.3
X59739_at	193.85	121.41	225.2	260.22	36.4	11.74	32.65	17.89	173.45	109.96	71.8	77.22	304.25	49.57	32.35	20.29	219.1	236.17	45.8
Y10207_at	125.25	9.55	538.3	621.97	200.05	131.45	382.05	218.43	188.95	44.9	172.5	41.3	470.7	314.38	679.6	90.93	158.9	1.84	211.4
U27768_at	214.1	250.74	28.6	33.09	82.5	46.81	408.4	543.2	334	419.31	87.95	25.39	427.6	218.07	56.55	19.3	112.45	70.78	90.75
X57522_at	1921.45	328.03	879.1	1015.69	2597.8	394.42	2439.95	734.61	1250.6	105.36	1945.6	282.7	2318.6	30.55	2256	633.14	2062	165.6	2148.05
U58130_at	336.15	64.13	277.05	320.11	437.95	66.96	852.9	945.26	120.75	59.33	293.6	249.89	458.7	305.89	252.75	90.3	344.6	222.6	543.15
U90918_at	365.8	303.91	206.7	238.86	671.3	162.07	128	56.29	184.35	59.47	339.15	188.59	140.15	28.35	180.35	48.15	203.2	103.24	550.3
AFFX-HUMR	15411.3	11547.2	7055.75	8152.02	40812.85	20122.91	18105.55	21548.44	33260	15197.7	15823.35	11892.62	22074.95	9186.66	51232.7	41372.96	16435.05	6290.21	18266
U47025_s_at	8236.3	145.81	5250.8	6066.69	9483.15	1928.07	9208.4	431.62	13419.7	2773.84	15726.05	699.68	20306.1	3252.83	18851.75	3515.52	14906.95	1286.72	13531.55
HG1205-HT1	1948.45	390.96	673.15	777.75	1468.45	842.66	1561.5	772.3	941.7	878.23	1405.55	421.51	2210.2	327.96	1398.5	405.03	1653.3	332.2	760.5
HG2380-HT2	312.1	121.91	76.55	88.46	582.55	383.89	188.9	127.56	390.25	353.62	311.1	221.04	338.55	318.13	318.5	227.12	113.8	7.07	234
M15881_at	404.35	261.56	863.85	998.08	408.6	265.02	1025.45	868.11	142.05	9.4	820.9	864.65	255	11.88	342.25	113.21	553	325.55	347.8
M14123_xpt1	32.45	9.26	11.6	13.44	404.75	510.6	33.55	25.24	66.65	56.36	184.9	29.27	159.85	185.33	223.2	253.85	37.75	2.05	71.8
Z78289_at	448.65	483.31	187.65	216.87	255.35	187.17	431.3	221.89	320.1	240.56	391.75	407.65	276.1	148.78	219.25	56.5	225.85	8.7	227.2
U92015_at	793.95	34.44	594.2	686.46	774.1	731.43	687.7	717.15	864.45	110.8	582.95	562.08	761.4	88.81	774.1	223.73	251.05	135.98	1040.2

U24186_at	153.05	164.69	531.3	613.91	711.45	544.4	167.6	106.07	250.75	245.44	315.3	135.62	101.85	56.64	252.75	206.55	189.1	128.55	142.75
HG662-HT66	9039.25	528.7	4115.75	4755.22	11100.5	1237.3	8595.5	4941.26	10920.8	1474.46	10289.15	4762.44	9864.75	1808.99	9449.45	2088.16	9047.1	1283.54	9424.85
HG3313-HT3	96.6	49.36	165.35	190.99	194.7	167.3	77.25	13.22	197.45	115.05	89.65	36.13	143.2	37.34	98.1	49.92	481.4	541.79	118.55
M90356_f_at	596	190.64	703.25	812.54	805.6	319.33	754.15	27.79	910.1	371.23	519.8	161.36	695.2	237.87	779	320.46	614	600.05	709.65
U26712_at	28.15	4.03	292.55	338.07	31.2	20.36	590.6	441.8	42.1	41.44	340	439.11	284.15	198.63	226.4	201.81	142.05	115.75	171.3
D83735_at	8557.05	420.23	3976.05	4593.86	9569.8	959.26	10685.95	2487.96	8066.45	900.64	8400.8	13.44	6371.95	1653.99	8206.3	692.82	6792.3	2360.89	6641.35
U00928_at	384.4	245.37	112.35	129.75	309.95	287.86	154.05	37.41	279.55	268.35	146.9	15.56	285.2	277.75	481.6	589.87	90	10.18	656.5
M28585_f_at	294.8	92.91	439.75	508.06	358.65	406.09	116.95	124.52	213	260.5	62.4	29.27	530.6	209.16	312.1	348.89	205.95	56.36	159.5
U41518_at	523.15	20.58	210.85	243.6	366.1	360.62	478.55	196.93	453.35	169.07	1237.65	160.87	584.1	466.97	1082.85	917.75	185.3	135.62	942.25
X69910_at	3688	1833.67	2573.3	2973.1	3174.15	1320.1	2336	805.39	2509.3	972.55	1988.15	169.07	2459.8	807.37	2137.85	369.6	1924.15	80.4	2113.8
HG2887-HT3	5278.6	3224.97	2028.25	2343.42	1587.25	1442.57	2502.7	1946.81	1196.5	558.76	3932.45	1868.95	3364.25	280.37	1927.6	1558.75	3358.5	2186.66	2263.6
U33202_s_at	251.95	279.24	354.6	409.7	278.65	264.53	93	83.86	105.2	58.41	125.2	38.04	518.9	176.07	317.9	329.94	332.85	125.94	307.75
U59831_rna1	1965.45	0.5	1631.35	1884.79	1945.75	840.11	1403.7	16.69	2273.5	333.33	1864.55	449.51	1544.6	85.7	2314.3	230.66	2660.85	699.12	2851.15
U59914_at	106.15	76.01	30.7	35.5	338.1	376.75	268.2	306.88	39.4	18.24	56.7	4.53	211.35	264.39	82.3	84.99	120.45	77.99	132.95
AF012270_at	60.7	5.37	27.15	31.32	128.85	109.39	129.15	114.62	82.5	17.39	119.65	66.54	86.8	11.46	133.35	77.43	79.55	49	75.9
D79992_at	938.6	181.3	266.3	307.73	1314.65	628.12	1120.1	977.65	1026.85	149.27	943.25	346.69	1318.85	144.46	1604.35	316.01	769.75	147.57	1651.8
M31166_at	162	172.11	428.45	495.05	125.25	35.99	306.95	236.1	372.8	76.51	202.25	185.47	153.85	87.47	198.95	174.16	145.4	90.09	141.1
D26308_at	4002.15	2990	1395.6	1612.49	4994.1	1414.5	4609.15	708.31	3128.7	236.17	4040.2	730.72	3102.3	679.25	3692.75	286.17	3564.6	63.36	4186.45
D13897_rna2	571.6	549.42	291.85	337.22	1443.55	1097.36	1437.4	664.82	1312.7	870.45	1271.1	22.63	1297.3	250.32	831.15	773.65	981.2	1068.86	594.2
M14113_at	77.5	50.06	21.2	24.47	418.3	538.53	111.35	41.93	271.7	150.19	234.9	169.14	77.7	40.16	305.35	263.4	130.55	14.21	183.35

X78992_at	6788.3	2526.21	304.85	352.21	2889.75	3363.49	2148.25	1488.53	4504.5	4344.32	2322.45	2466.46	4796.95	2388.54	4591.05	2796.54	4823.35	1102.59	3577.6
J04102_at	172.5	48.22	803.85	928.78	441.55	83.51	446.55	281.5	151.75	38.82	787	95.74	137.05	10.82	558.7	564.7	649.2	608.11	557.15
Z83805_at	334.25	173.03	114.65	132.44	153.35	46.32	115.65	7.99	260.8	25.17	742.5	138.45	311.35	118.58	204.9	89.94	166.2	43.42	225.25
HG3893-HT4	204	110.59	392.5	453.54	259.85	55.79	403.2	210.86	667.3	55.58	415.05	127.35	776.9	736.24	249.55	24.68	473.15	333.97	496.15
M95549_at	378.85	6.29	52.9	61.09	319.45	66.26	1549.1	2049.76	441.05	3.46	283.1	101.96	527.5	195.44	784.25	341.46	223.8	26.02	369.3
U72935_cds3	211	211.42	174.3	201.38	164.35	68.66	160.2	20.22	120.05	50.98	387.5	242.11	314.35	185.62	516.7	237.31	275.95	67.95	333.95
M29335_s_at	38.1	19.94	217.35	251.09	100.35	76.44	1073.95	1358.99	85.65	94.26	101.05	41.22	91.75	12.37	260.65	262.41	47.45	1.63	55.2
U28251_cds2	91.4	48.93	95.7	110.59	162.55	36.42	91.35	3.32	117.25	54.24	79.7	2.83	123.2	63.5	87.6	42.99	156.85	2.76	115.55
AC000066_at	46.6	3.11	22.6	26.16	131.7	65.34	503.65	625.86	343.15	273.3	252.1	49.21	118.05	67.39	106.8	43.13	246.5	134.35	833.1
M13928_s_at	642	531.6	1104.05	1275.55	314.8	132.65	422.4	199.97	348.15	11.95	608.05	88.46	717.95	326.19	901.7	408	304.4	250.88	467.8
L31860_at	62.95	6.72	40.6	46.95	137.1	27.72	160.25	14.21	62.5	2.97	75.05	49.29	328.7	35.78	70.25	66.4	41.6	18.1	69.75
HG3104-HT3	953.3	184.98	317.65	367.06	961.05	112.92	1119.85	304.98	1013.2	21.07	1047.8	143.83	1120.85	182.5	936.85	1.91	826.1	30.55	984.6
D86965_at	3537.5	1273.78	1523.65	1760.34	3456.1	188.51	3309.25	866.14	2608.55	16.19	4099.95	751.3	3660.1	345.63	3860.8	283.27	3243.5	1011.02	2605.05
M60450_s_at	228.1	67.32	92.25	106.56	563.25	528.28	424.2	498.23	590.25	9.83	542.85	292.81	490	342.95	378.9	341.96	108.3	38.47	162.5
X52056_at	161.6	36.91	35.9	41.44	144.5	43.98	127.05	93.69	101.1	4.1	120.7	40.02	110	18.24	142.25	79.97	156.9	46.39	147.3
L02867_at	2158.05	984.65	853.9	986.56	1246.1	83.58	1094.4	408.71	979.45	475.81	1436.2	23.33	2125.95	1069.36	1400.55	443.43	1605.75	259.72	1469.95
M63623_at	138.55	84.07	221.85	256.33	49.05	12.8	571.55	414.01	173.85	105.01	630.35	229.31	233.55	229.17	241.65	15.06	207.95	116.74	224.95
S74728_at	1556.35	152.81	562.1	649.41	2041.35	307.24	1268.3	572.47	1806.65	630.39	1515.7	480.13	2042.75	330.57	1419.1	105.5	1961.05	190	1859.85
X86428_s_at	310.85	292.25	134.05	154.93	347.65	267.07	199.8	96.59	180.1	0.14	359.5	123.74	215.4	6.36	215.2	108.47	189.7	22.49	207.35

U85992_at	1050.4	251.31	296.8	342.95	1068.35	701.52	1122	874.13	341.1	311.98	929	523.82	1257.9	0.57	706.35	406.52	1055	87.4	705.5
U03486_at	1177.65	450.07	576.65	666.31	1494.2	1083.57	1546.4	1012.72	1135.4	69.58	1136.75	254.91	960.5	300.94	492	455.38	1085.75	214.04	1058.8
M22995_at	256.15	278.11	167.1	193.04	347.55	89.45	258.25	100.34	578.55	3.89	460.85	26.66	419.9	88.67	727.55	330.29	597.25	30.19	372.75
U52111_rna4	291.3	31.11	184.2	212.84	915.2	735.53	275.75	85.35	193.6	96.45	402	119.22	343.05	148.56	292.6	127.7	241.85	51.41	221.05
X59373_at	1313.7	406.44	126.95	146.72	1232.75	160.02	1747.6	1042.56	945.25	811.97	1555.55	415.71	869.15	500.7	888.75	806.88	711.55	191.13	1096.1
AC002115_c	1854.7	126.29	763.15	881.69	2683.85	1513.84	2467.65	771.81	2337.35	749.04	2723	1056.84	2363.15	747.62	2588.4	1397.38	2225.9	469.94	1506
AC000115_c	99	33.52	139.9	161.64	133.85	13.93	196.75	33.45	251.65	45.75	308.1	268.7	784.45	158.46	352.8	325.13	308.2	311.69	413.65
X02875_s_at	781.3	271.95	417.4	482.25	997.55	315.16	717	365.01	665.8	89.1	969.9	999	683.85	441.73	252.25	109.25	600.75	630.24	467.5

Z86000_at	2946	5.37	608.1	702.58	2803.15	86.9	1924.6	230.09	2555.65	537.19	2351.85	240.49	4817.6	302.22	1878.8	253.29	2102.35	337.5	1816.05
X15393_rna1	1839.1	163.77	1071.15	1237.65	1570.55	316.43	1946	1194.3	1221.75	188.3	2311.15	159.17	1840.85	287.58	1684.45	109.81	2260.3	621.12	2305.1
X04145_at	51.55	6.01	39.75	45.89	49	9.19	45.35	5.59	68	25.31	89.2	34.51	260.4	199.97	365.8	469.66	80.6	29.7	66
M30448_s_at	12543.75	4790.15	6232.65	7201.1	14762.75	5346.08	15433.3	2490.01	14004.4	1204.77	12412.55	824.27	11296.3	32.1	13101.3	6611.45	14500.9	4648.52	12335.95
X60655_at	2573.25	1495.46	1470.45	1698.97	2803.65	1992.84	3325.3	486.21	1300.8	920.51	1972.45	1925.1	2412.35	2526.85	1073.3	707.39	3202.25	666.02	2393.1
M76732_s_at	1623.15	513.29	587.85	679.18	1545.15	819.47	718	429.92	613.25	469.73	1133.75	1140.92	1401	869.18	1409.65	645.8	882.6	732.7	663.6
X81851_s_at	119.5	31.82	80.5	93.06	129.6	1.41	150.75	134.99	111.2	40.02	143.7	121.34	343.7	305.89	140.5	20.65	97.45	24.11	126.2
U32499_s_at	131.55	7.71	155.2	179.32	435.05	86.62	1155.25	1450.35	301.9	262.34	298.85	260.85	358	222.17	652.2	515.76	210.85	62.3	216.6
U77604_at	1616.75	233.56	778.45	899.37	2764.45	325.91	1799.45	1179.38	1325.7	238.58	1412.05	598.42	796.4	417.62	1737.2	947.24	1100.35	324.35	1573.45
D21235_at	3649.1	1293.86	1934.2	2234.74	3538.2	922.07	3408.7	1571.33	3104.6	542.21	3188.55	269.9	2752.8	171.69	3016.55	1455.16	3217.4	1651.66	3754.55
X69398_at	490.35	240.2	325.5	376.04	539.55	662.77	474.85	452.48	759.7	312.54	382.35	46.32	537.8	263.04	296.45	149.69	192.1	30.41	230.95
U89335_cds2	1318	564.13	631.8	730.02	1660.8	694.94	1057.65	46.32	722.35	268.91	1322	168.57	1706.5	191.34	1546	346.2	1701.4	104.51	664.25
HG2157-HT2	2819.6	1275.76	1231.2	1422.56	1145.05	411.61	1777.6	122.9	1199.45	101.75	1185.15	1102.73	2320.35	807.59	1384.1	340.4	1310.45	1049.28	1129.9
M14328_s_at	56623.2	8398.17	31914.75	36873.57	56525.25	6541.94	67559.9	12335.48	51381.95	340.33	43855.45	129.75	49955.65	5318.79	63149.7	15884.31	61281.8	4598.46	43835.4
X69978_at	256.4	136.05	313.6	362.32	874.75	477.51	410.65	68.52	510.05	28.78	565.5	182.43	352.6	214.54	286.5	157.83	331.65	140.08	369.35
L22569_at	76.05	40.38	34	39.32	447.2	456.65	68.9	39.88	110.05	64.84	69.7	3.82	398.7	457.78	62.8	11.31	115.15	47.59	99.95
X84746_at	779	464.71	101.75	117.59	335.1	11.74	1057.8	629.04	1285.35	1488.95	514.55	338.49	827.1	975.1	393.4	241.41	376.6	178.19	1396.25
AJ001047_at	98.7	68.31	180.4	208.46	433.45	466.05	149.35	135.55	220.15	103.31	161.85	50.42	154.85	11.95	119	52.47	104.85	8.98	140.65
HG2197-HT2	3628.9	65.05	2283.15	2637.86	2869.7	975.81	3299.15	384.03	3917.9	223.73	3661.95	787.36	4011.95	2043.89	3232.8	962.09	4173.2	2132.78	3302.05
L40933_at	993.1	556.21	179.6	207.47	763.95	590.36	391.8	153.3	477.15	365.22	1120.75	285.88	247.35	23.12	1926.35	263.11	602.4	364.87	710.1

U47334_at	672.6	54.73	521.05	601.96	486.5	334.18	1236.45	298.47	1108.15	366.78	562.55	479.91	1157.05	293.1	697.65	618.36	721.9	562.86	757.2
U34587_at	1014.85	288.99	1045.85	1208.37	1322.55	502.54	1331.5	1174.79	433.7	197.85	1271.6	647.43	1528.5	41.29	1079.35	326.33	896	100.97	1293.05
Y00083_s_at	505.25	68.94	277.6	320.74	591.5	206.19	196.15	72.34	731.8	200.96	197	161.64	464.3	82.73	248.95	17.18	359.85	412.6	602.7
S42457_at	22.35	9.55	28.55	33.02	192.9	135.34	418.2	311.69	144.75	82.8	39.9	11.74	81.55	50.42	299.75	395.06	122.55	94.26	153.6
M86699_at	328.95	74.32	279.05	322.37	67.4	2.4	155.2	149.34	382.5	321.31	420	70	509.8	236.46	44.15	25.1	390.75	42.21	253.75
X95632_s_at	558.7	226.42	560.7	647.85	950.6	403.33	253.3	141.99	750.9	914.57	253.55	197.21	548.3	504.17	601	7.35	679.7	4.1	423.1
X99728_at	5318	154.29	3157.6	3648.25	4572.9	769.62	5332.45	1958.61	4601.45	1723.01	5278.65	92.14	4054.5	121.34	4475.15	853.69	4240.3	478.43	4195.25
U77970_at	717.7	658.6	825.55	953.82	442.45	259.72	551.8	371.23	757.45	509.47	397.6	238.58	631.65	473.27	460.5	179.46	213.2	91.36	888.8
S77393_at	48.6	13.86	23.3	26.87	38.95	11.53	72.35	51.69	72.45	6.58	131.05	95.39	59.7	3.11	67.4	33.8	46.85	13.08	489.25
AF009368_at	2737.6	522.41	1090.9	1260.35	3644.05	518.95	4208.1	2008.32	3360.9	78.06	3422.6	263.61	2465.55	323.93	3311.9	980.33	3368.8	1746.7	3377.9
M71243_f_at	262.3	200.82	108.2	125.02	133.95	62.3	202.95	1.91	164.1	48.79	185.75	4.03	148.05	43.35	281.85	159.03	508	550.41	309.2
L34075_at	1407.05	213.05	924.05	1067.66	929.4	447.88	1023.6	281.43	1636.7	213.97	2106.8	846.83	1447.65	551.9	1465.75	634.49	1112.05	167.09	1128.85
D50931_at	143	122.19	318.55	368.05	305.35	194.38	595.15	51.55	316.75	16.19	205.95	143.47	544	335.73	326.65	103.45	438.35	334.25	163
D86970_at	960.35	34.72	297.3	343.51	532.35	95.95	1417.3	1224.99	796.45	694.73	1270.75	724.43	1579.25	717.78	1006.55	87.04	675.7	367.13	684.45
U00930_at	206.35	109.39	372.3	430.2	1053.4	769.19	1063.85	828.8	564.65	334.11	784.2	886.43	302.5	385.51	629.9	308.02	204.35	219.7	646.1
U40317_s_at	164.1	1.13	349.25	403.55	237.35	35.43	263.5	124.88	234.05	125.79	200.7	1.27	515.05	252.51	145.75	3.75	166.8	78.21	125.95
D16562_at	6304.4	1781.91	3800.6	4391.13	7187.75	92.28	7885.15	2850.14	6882	404.89	6251.75	2096.22	7041.25	2047.43	7161.3	61.24	5524.9	441.38	6132.45
M19888_at	4824.15	2014.62	2029.75	2345.12	4727.55	863.45	3695.5	1721.81	6355.8	1502.04	5703.4	566.96	8116.55	1600.54	6468.8	44.55	5205.85	339.48	5045.4

U72066_at	1123.45	146.3	917.25	1059.74	1731.7	669.06	1253.8	560.88	1233	374.34	1415.1	111.16	2495.8	271.1	1459.5	517.88	1389.6	656.62	1394.5
Z22548_at	3979.25	1811.68	2207.75	2550.75	4420.1	362.46	5275.2	417.33	3020.4	595.67	3599.2	724.5	3614.15	1837.42	3544.05	240.49	3369.55	544.68	4113.65
U50330_at	1544.8	994.62	709.65	819.89	664.7	313.39	1136.3	423.56	1123.5	527.5	1037.4	216.52	552.65	326.61	574.65	425.47	879.45	925.25	1723.45
M19722_at	1837.65	176.56	1188.9	1373.63	1180.6	610.8	1111.4	45.68	1330.2	588.17	1386.35	328.45	1712.15	566.04	1002.6	550.69	1567.55	32.46	1337.25
M21642_s_at	47.25	46.03	425.1	491.16	642.3	854.19	373.55	457.71	29.2	6.51	225.55	274	139.45	174.73	62.15	18.31	188.5	176.78	111.9
U41060_at	1277.05	113.77	795	918.53	1399	487.06	596.75	592.48	1599.35	414.15	691	671.19	1435.2	716.02	1547.05	152.1	1462.85	521.49	1195.1
D67029_at	760.6	128.69	260	300.38	919.4	103.94	782.55	340.75	487.6	390.89	623.3	623.81	649	64.21	429.9	383.25	908.45	748.19	769.4
M16938_s_at	1426.75	507.49	359.05	414.86	1497.3	593.12	1310.1	485.5	1150.35	140.36	1078.75	42.92	2125.1	558.19	1367.15	272.73	1153.75	186.75	1017.35
D16532_at	482.1	519.86	270.7	312.82	373.9	313.67	133.1	11.17	268.7	79.62	670.9	311.69	149.9	9.48	171.15	2.19	142.45	115.19	452.95
L14812_at	317.95	273.58	357.4	412.95	1138.75	1100.89	448.55	95.81	464.75	373.28	475.2	480.55	649.85	93.41	560.95	424.33	617.05	302.29	652
HG909-HT90	737.25	294.93	229.9	265.59	928.35	641.42	555.9	114.69	493.2	247.06	675.05	272.02	549.95	9.55	552.3	2.12	852.05	484.72	677.85
HG2416-HT2	64.3	22.06	58.25	67.25	73.75	38.54	143.65	122.4	74.25	4.31	126.3	74.81	493.95	101.05	68.25	38.11	64.25	29.06	87.7
U50136_rna1	2353	447.74	1040.65	1202.29	1126.8	602.88	1099.35	374.13	1688.7	336.16	2730	1393.28	1800	707.39	630.65	38.54	2129.2	746.42	1404.15
U26727_at	749.1	217.93	960.1	1109.31	781.65	633.21	1563.8	123.32	1261.9	32.24	1640	1.84	621.6	444.35	1153	1406.15	1150.45	223.23	841.85
L37199_at	1451.8	148.21	988.95	1142.61	1356.4	977.22	1303.75	186.75	820.7	101.26	1026.85	275.42	1639.5	381.84	1761.1	502.89	889	84.15	1270.4
U53347_at	8661.85	1541.99	5020	5799.97	6127.95	1353.47	7261.6	673.02	7085.7	456.51	7526.75	849.87	7229.7	1854.74	6447.6	4291.71	6796.95	2611.42	8230.7
D38305_at	1346.7	305.19	125.3	144.82	1893.55	541.15	1104.95	102.18	1798.5	318.34	1595	387.64	954.15	168.08	1405.25	813.53	2349.2	1380.7	2343.95
X62691_at	52472.2	32878.35	16454.05	19010.64	60995	14745.01	57591.05	18198.74	62033.4	5047.47	50253.45	7437.7	47322.75	23384.23	49215.85	14454.18	44868.2	21914.23	64154.2
J03810_at	296.8	47.52	8.9	10.32	146.4	41.44	228.25	34.29	205.35	121.98	322.7	97.72	268.6	221.75	248.15	285.88	106.05	50.7	127.8
X51521_at	6184	2157.81	3068.6	3545.43	5604.4	1115.11	5790.3	1180.59	6359.05	1057.05	5737.1	1471.77	4920.25	728.11	6192.45	1659.37	6153.9	1630.31	6032.6

X79204_at	458.2	117.52	322.8	372.93	534.6	342.52	610.95	418.11	271.9	206.33	386.5	311.98	542	114.98	304.7	180.31	98.05	23.69	291
U31501_at	1995.45	1675.77	2232.8	2579.67	1193.7	355.39	2406.4	369.53	1576.9	799.45	2451.3	305.61	2250.95	319.97	813.05	795.42	1235.2	122.9	2851.9
U27333_s_at	470.6	446.47	88.6	102.39	1726.9	2194.15	565.65	603.52	240.2	135.91	462.05	480.48	244.75	18.6	719.85	725.7	184.8	23.33	137.05
X05130_s_at	19790.9	3399.91	9570.95	11058.09	12701.45	4578.59	13202.5	1842.01	12248.9	6406.39	10880.5	2666.08	9889.95	2757.65	12614.65	2157.03	16032.85	1639	11523.35
U39231_at	246.05	242.61	105.8	122.19	156.55	56.36	232.2	51.05	387.15	99.35	430.9	455.8	294.5	235.47	144.55	35.71	78.65	20.01	105
U46461_at	3124.5	1827.02	726.15	838.98	662.55	165.11	2135.8	1231.36	976.05	972.63	2849.45	1016.75	2000.05	1547.5	2073.9	734.83	3018.5	881.48	1355.6
D38555_at	1566.75	318.69	1123.75	1298.32	1469.85	594.46	1804.25	41.51	1768.8	513.5	2094.5	16.26	1304.95	1016.18	2335.85	719.34	1566.7	381.13	1575.65
D16350_at	470.95	545.67	140	161.79	599.7	662.42	1068.1	1153.01	673.55	93.55	1148.7	894.63	545.8	358.22	288.8	263.75	460.2	336.3	264.9
X07496_at	884.2	764.38	301.8	348.75	436.15	143.33	625.8	164.9	1100.1	1044.96	1885.65	741.83	1926.8	854.61	2447.05	380.35	1009.9	390.18	941.05
L19063_at	88.6	26.87	36.2	41.86	427.1	427.09	227.8	171.83	186.85	80.82	133.2	131.95	94.25	55.37	112.45	87.33	40.95	27.65	119.05
U76764_s_at	251.75	27.22	148.65	171.76	434.9	374.91	738.6	417.76	351.9	115.97	199	10.75	296.45	52.96	137.25	9.4	830.25	691.76	273.15
X63380_at	1637.75	538.18	375.95	434.38	1626.55	978.14	1297.15	471.15	1048.65	452.48	634.6	147.36	1489.8	496.39	1527.15	1122.11	1953.3	1596.65	1361.1
U61374_at	2109.55	994.55	1518.5	1754.47	2275.85	897.25	2157.2	1120.62	1947.85	244.02	1838.05	363.95	1849.15	102.04	1649.8	74.95	1773.35	252.93	1911.45
D38128_at	420.3	156.55	482.8	557.77	576.4	242.82	374.35	124.52	386.35	333.68	493.1	151.75	383.8	208.74	416.9	43.42	947.85	791.18	569.65
X76180_at	2107.35	144.46	1218	1407.28	2027.05	41.37	2952.6	513.36	3201.9	8.2	2794.1	763.82	2008.4	1178.18	3200.55	846.48	3649.15	505.51	3450.15
X97674_at	228.2	89.66	104.9	121.2	518.1	36.49	210.6	168.43	135.7	47.09	596.35	679.74	182.45	25.24	250.25	99.91	137.2	26.16	274.5
D12485_at	281.9	51.34	598.4	691.41	893.4	875.68	1127.9	916.41	568.3	73.68	490.9	411.11	2157.5	1584.48	517.6	165.75	300.9	210.58	370.15
U79261_s_at	1763.05	458.98	1027.85	1187.59	982.15	304.13	813.15	97.37	1475.45	104.86	1558.05	1160.29	795.35	456.58	976.45	458.42	1127.85	226.06	1744.3
D87457_at	343.8	371.51	682.45	788.49	104.3	65.05	147.65	91.85	79.2	11.46	584.8	694.52	168.9	68.45	97.3	10.18	76	26.87	84.4
S78771_s_at	5046.35	1504.09	1557.15	1799.09	2215.4	978.21	3504.1	2688.7	3385	1480.68	3399.7	290.34	2461.6	172.11	2270.45	703.5	3204.6	918.39	3731.4

U89717_at	503.25	2.19	135.85	156.91	486.7	260.64	716	253.99	413.75	30.9	656.9	84.57	873.5	80.04	682.25	403.69	385.5	28	447.5
HG2480-HT2	571.4	645.87	469.75	542.7	556.5	0.57	1334.4	774.56	1103.35	735.74	607.8	510.11	627	360.06	492.85	225.78	864.95	199.76	1061.35
Z47556_rna2	33.3	16.83	41.95	48.44	52.6	1.13	38.15	15.49	59	60.25	103.5	64.06	353.05	372.29	283.65	177.27	107.8	94.19	178.05
X99584_at	709.6	421.86	398.3	460.19	411.1	317.21	412.2	33.52	457.8	115.82	357.65	397.75	367.6	387.78	600.95	38.11	741.25	30.05	300.25
J05200_rna1	74.85	5.87	75.35	87.04	73.85	27.51	174.55	164.12	147.5	4.24	292.25	116.04	128.05	81.81	89.85	9.83	95.7	47.52	77.55
L34035_at	63	11.03	102.25	118.16	202.25	190.85	114.15	75.02	213.7	212.98	129.25	88.32	194.6	56.29	103.7	88.67	67.25	11.53	78.65
D86966_at	3727.4	687.31	1858	2146.63	2861	707.11	3456.45	241.48	3978.35	695.72	4300.3	1113.98	4438.6	167.16	3829.8	897.74	3467.75	184.77	3782.9
HG371-HT10	141.45	146.3	36.5	42.14	626.3	836.22	368.7	489.04	269.9	133.93	91.9	90.79	94.95	45.75	269	199.26	371.9	352.85	292.05
X89894_at	593.35	95.39	225.2	260.22	565.4	80.19	762.6	267.71	277.15	238.22	457.4	133.78	894.75	20.15	661.65	328.31	776.95	159.74	307.8
L05424_cds2	476.2	422.43	613.75	709.16	636.7	624.66	552.5	241.97	172.95	81.67	908.45	1021.13	462.4	425.68	310.55	230.73	433.85	406.66	901.55
U46024_at	66.1	72.27	6.95	7.99	90.55	25.39	100.2	0.71	41.35	43.91	269	136.47	102.15	58.05	227	286.66	43.4	28.14	119.8
U68494_at	676.35	187.6	433.25	500.56	1097.45	552.75	759.45	400.15	1252.95	280.93	590	182.57	720.35	18.03	1341.4	61.94	875.9	107.9	756.55
L42025_rna1	323	137.74	270.4	312.4	507.9	95.88	631.8	119.5	441.9	186.53	600.55	162.99	554.2	152.03	705.15	323.64	639.65	317.99	503.55
M23114_at	2636.65	435.22	1743.85	2014.76	1238.85	1268.05	2011.4	411.54	2383.9	292.18	2046.35	712.41	2131.95	159.88	2677.7	1140.28	2821.9	47.38	3100.9
U79247_at	308.15	235.25	206.05	238.08	516.65	477.51	184.05	83.93	215.2	91.78	352.9	359.92	518.8	168.86	765.3	215.95	90.7	76.65	445.1
U57623_s_at	1010.75	759.08	471.7	545.04	934.25	567.31	968.75	355.32	1290.55	507.35	353.8	252.44	685.1	443.64	1037.1	58.97	976	288.78	982.95
X13973_at	7387.85	950.42	3394.25	3921.68	7513.1	1574.3	6275.4	231.51	5207.1	576.72	5611.1	533.02	5797.1	41.15	4969.85	564.77	5136.05	1531.95	5655.6
U07158_at	2428	406.16	1440.8	1664.67	1945.9	165.6	1652.5	1120.62	2012.95	930.91	1875.3	128.27	1858.4	335.88	1751.35	115.05	2050.4	678.82	2463.45
AF015913_at	4112.25	1248.4	2314.85	2674.49	4600.8	515.06	4422.65	985.78	3925.9	1560.3	4573.15	240.35	3212.75	710.57	3534.65	951.55	5250.95	1140.92	5250.8
M19645_at	7734.4	157.97	4046.35	4675.04	5793.15	1844.49	6677.25	4036.8	5726.75	2090.28	6485.1	2319.45	3676.25	507.21	3524.1	2808.77	5888.55	3314.42	6596
S77415_at	428.45	411.89	759.7	877.8	589.4	158.67	1118.6	1046.8	625.75	120.14	169.9	67.6	545	165.04	721.8	398.1	311.95	183.35	329.8

M37981_at	199.35	0.78	124.45	143.75	256.2	207.89	102.15	59.47	505.85	341.89	219.95	84.36	394.65	269.34	51.45	10.54	47	2.55	200.6
HG2228-HT2	461.9	8.06	277.8	321.03	604	53.46	548.55	268.21	412.7	12.73	651.9	174.51	539.05	40.66	482	67.03	415.25	55.23	476.55
L14856_at	5700.25	550.48	1757.2	2030.25	3997.9	1384.66	4779.55	748.33	4887.25	699.54	5201	1219.62	5849.4	710.5	3959.45	252.79	4602.8	134.35	5352.1
Z21488_at	759.55	196.65	297.65	343.87	834.85	130.18	308.9	266.3	734.65	370.31	364.9	170.55	155.45	31.89	421.25	21.43	468.7	193.61	340.35
M60858_rna1	8877.85	2045.45	4492.7	5190.73	8312.15	3490.35	5429.1	4591.39	11858.8	5011.41	9474.35	5823.8	10345.9	639.37	11350.15	3263.93	9263.8	116.53	10841.55
X00948_at	72.1	33.23	34.95	40.38	118.25	79.27	89.2	96.45	202.25	69.79	161.9	112.71	92.7	72.69	64.75	60.17	48.2	18.38	126.55
U49441_at	1106.3	168.57	796	919.66	1164.85	465.63	1133.2	263.61	1481.15	199.47	1155.2	290.34	1128	416.49	848.55	683.98	576.8	337.15	1272.45
J03915_s_at	2089.3	118.51	450.2	520.15	2204.65	300.59	644.2	352.99	1694.55	772.09	2546.8	2825.03	1356.3	804.69	1102.7	790.97	2045.7	301.79	2193.2
X77753_at	337.95	211.78	175.25	202.44	374.15	95.25	160.25	204	226.4	245.51	263.15	89.73	628.9	48.08	260.15	302.57	79.15	13.51	400.85
U79300_at	202.55	196.08	215.5	249.04	553.3	629.61	727.6	1004.66	211.5	256.4	414.3	74.25	40.55	13.79	197	240.13	188.15	235.68	278.9
L14754_at	1314.7	942.71	972.45	1123.52	2101.15	502.54	2462.1	2709.07	1140.8	461.18	2423.5	1406.86	1821.4	1290.75	1945.3	515.91	2310.75	25.39	1960.35
X95289_at	84.55	32.88	152.55	176.28	249.2	143.83	211.2	201.67	113.7	18.53	839.95	892.72	277.2	88.67	177.15	104.02	142.35	69.51	144.55
L10955_cds1	459.6	64.77	575.5	664.96	943.35	514.7	1162.2	957.42	358.95	71.21	231.15	73.89	792.45	648.49	639.1	700.6	807	260.92	824.95
U36621_cds2	87.75	26.66	266.85	308.37	81.7	25.17	353.3	369.53	155.7	15.98	485.15	183.21	212.2	74.1	276.3	236.46	122.65	22.7	149
L38933_rna1	136.9	50.91	237.3	274.22	300.7	38.18	296.65	74.74	600.4	175.93	522.05	112.5	207.45	23.83	200.25	19.73	501.3	554.37	839.1
X79200_s_at	93.65	86.62	26.4	30.55	309.1	392.59	74.55	54.8	31.15	5.59	18.85	14.78	32.9	15.56	178.15	181.8	61	52.18	222.25
X60003_s_at	1933.75	454.03	1765.85	2040.22	2510.9	694.94	2288.35	1244.15	1552.25	69.23	3391.75	922.85	2772.95	570.14	2092.3	20.93	1643.3	1304.89	1772

M91432_at	284.5	132.79	33.95	39.24	433.4	195.16	129.7	5.66	230.75	109.53	235.7	3.68	230.6	4.67	392.65	279.66	226.3	96.45	514
U01317_cds6	449	371.51	144.25	166.67	213.3	21.07	263.3	86.83	500.85	492.78	721.8	785.45	198.05	76.3	232.1	102.39	105.65	26.23	213.4
X85740_at	230.1	31.11	96.25	111.23	201.35	83.23	167.55	57.63	128.45	91.85	171.45	15.06	203.4	14.28	238.8	120.21	395.65	305.54	155.3
U66578_at	331.7	316.08	221.45	255.9	455.55	96.94	770.2	944.55	586.9	242.25	929	478.85	354.45	323.64	786.6	519.02	778.25	74.46	748.25
HG613-HT61	44838.05	24715.29	19620.6	22669.28	59513	10225.19	56136.95	28809.86	52536.7	9300.58	47623.1	13897.05	44131.65	1979.69	50545.95	3318.52	43167.8	5628.71	47995.7
U59913_at	121.35	37.12	120.05	138.66	325.05	192.12	278.2	342.66	403.8	156.27	306.6	348.32	305.45	106.42	365.85	478.5	376.8	114.55	244.45
X07732_at	204.5	66.75	393.8	454.95	940.15	1053.38	206.2	0.85	149.35	71.49	185.2	23.9	313.4	171.12	291.45	127.49	184.65	31.18	226.8
Y09912_rna1	112.25	25.39	43.9	50.77	268.4	236.46	257.55	241.62	116.7	14.14	180.8	1.98	146.55	15.77	163.4	17.39	116.5	69.72	145.7
HG627-HT50	2921.55	298.47	1809.95	2091.13	2483.6	572.05	2849.05	704.21	2396.95	755.12	3171.45	1000.49	3116.8	585.2	2530.9	73.26	3169.95	748.19	2451.2
X74142_at	200.15	192.69	58.4	67.46	236.75	272.73	48.5	11.17	154.2	108.75	106.55	118.44	87.65	62.86	150.3	188.51	204.55	232.43	45.75
X63097_at	180.55	47.16	49.95	57.77	261.65	271.74	702.55	581.45	243.35	252.51	133.6	12.02	104.25	93.55	284.35	209.94	242.1	86.83	199.6
X58199_s_at	598.5	91.64	273.65	316.15	1274.3	595.24	180.5	86.13	433.95	21.28	171	19.23	748.45	445.83	1001.75	387.99	823.15	329.87	765.4
L13738_at	1394.95	1415.84	702.85	812.11	432.3	41.58	1697.65	121.83	1366.5	1441.93	561.7	120.35	2227.25	1497.72	1403.85	1087.04	2004.4	961.52	726.25
X63527_at	70224	25473.67	33566.45	38781.91	82142.8	1358.07	106465.4	20283.92	76974.8	12700.21	75535.05	4418.22	72679.15	24873.68	71326.7	3739.46	69399.55	13213.63	74630.2
AFFX-HUMTI	925.4	77.78	368.15	425.32	1078	524.11	561.15	172.04	530.2	509.54	896.5	19.37	1158.85	311.34	753.95	144.46	1085.75	385.87	912.65
X85137_s_at	641.15	293.8	402	464.43	638.2	366.99	126.55	87.75	488.3	130.67	901.95	378.51	443	461.6	278.8	228.4	529.55	231.86	476
M74091_at	144.65	168.79	328.15	379.08	191.85	29.49	382	348.18	143.4	117.52	212.35	204.99	276	321.45	88.05	85.77	249.8	194.45	165.35
J04444_at	7019.9	2061.36	5023.85	5804.43	11191.6	1598.77	10195.25	1092.97	9410.4	1536.97	12779.75	3675.47	7309.5	2216.07	7259.05	250.1	8627.75	1942.35	9989.5

L49218_f_at	30.15	7.85	31.3	36.2	92.4	70.71	78.7	84.85	52.7	39.32	70.9	68.45	39.55	18.46	58.8	12.73	39.85	9.26	45.2
X07820_at	169.35	38.54	290.65	335.8	579.1	612.64	650.65	520.22	487.1	112.15	346.7	67.18	526.85	663.62	190.4	133.08	773.3	315.94	257.4
M37245_at	1629.3	565.69	1092.15	1261.83	1298.05	715.8	768.7	687.45	2127.75	633.21	1256.15	1010.67	1784.2	523.12	1516.85	721.18	1316.1	106.35	1606.45
M94547_at	373.3	160.8	50.05	57.77	619.4	592.27	380.5	354.97	336.9	202.52	422.5	386.79	255.35	127.49	427.55	303.7	444.5	154.01	235.65
J05272_at	7478.25	1247.55	2787.8	3221.01	5203.35	860.48	6773.15	34.01	5404.9	735.39	7010.3	1343.93	5805.05	2183.9	6828.6	514.35	5876.1	1262.47	6760.45
X14787_at	865.45	757.24	675.65	780.58	492.55	172.04	158.3	72.83	339.95	109.25	728.65	139.51	589	180.74	425.25	35.57	892.8	340.4	333.85
X16699_at	28.75	7.42	21.85	25.24	83.45	71.77	69.05	64.84	42.75	17.18	73.5	63.5	41.15	9.97	57.6	2.4	36.2	1.13	43.85
M60483_rna1	1040.35	133.43	889.35	1027.5	1102.2	343.23	988.45	171.76	1095.6	183.56	760	277.61	992.9	76.37	732.1	364.44	1107.7	249.33	1090.8
U77665_at	696.2	81.74	415.7	480.27	1286.45	202.59	683.05	581.31	648.35	71.77	919.2	180.17	681.4	455.09	569.7	30.55	450.75	394.21	850
L48728_cds1	739.35	148	708.2	818.26	900.85	250.95	1442.9	569.65	726.4	94.19	1244.65	829.22	502.1	577	1064.95	168.5	369.75	309.08	1145.7
S75989_at	298.95	66.68	689	796.06	314.45	113.91	444.8	272.24	764.05	240.77	343.8	38.61	873.4	663.55	244.85	154.5	868.4	679.25	553.85
Y08991_at	432.5	96.87	249.5	288.22	176.75	110.1	423.6	136.33	700.8	303.77	641.5	148.35	402.05	414.72	535.95	55.79	346.75	269.76	506.1
X99656_at	2823.5	2728.87	1025.65	1185.04	1277.05	1091.84	1171.75	170.77	1533.95	204.14	826.15	416.13	2777.1	67.03	1282.4	600.05	2017.45	218.99	1228.95
U90268_at	53.15	30.76	67.45	77.99	75.85	44.9	128.65	69.08	102.65	57.91	80.9	60.53	96.6	59.82	471.15	498.44	79.55	15.91	62.3
U79245_at	71.75	40.8	256.45	296.35	561.7	191.34	347.5	1.13	191.4	227.55	498.55	14.5	133.65	59.75	356.8	417.9	387.05	86.48	556.9
X58022_at	311.55	371.3	93.85	108.4	82.95	16.05	412.35	518.95	70.9	11.31	124.4	42.28	161.05	125.79	63.45	13.93	59.25	6.58	85.25
X99657_at	110.1	29.42	55.7	64.35	156.75	9.69	276.95	168.36	122.1	25.6	214.15	129.33	161.5	39.03	159.95	76.01	166.85	84.22	124.65
D88422_at	10770.1	8790.61	4831.55	5582.25	17077.55	970.36	9077.95	5134.94	16033.05	330.71	11480.4	3728.43	10165.25	460.96	15028.65	2200.02	12071.2	3615.72	12980.65

J05213_at	103.1	66.33	62.65	72.34	236.65	195.09	173.55	193.25	79.85	36.84	124.45	56.64	116.1	20.65	193.85	212.06	101.4	40.31	88.15
M19507_at	51.2	1.7	62.6	72.27	231.8	185.4	105.85	93.83	392.55	151.25	241.55	159.17	145.2	106.77	157.6	113.56	141.35	33.73	214.4
M37104_at	1553.2	123.46	835.95	965.84	1770.9	341.25	1503.4	326.68	1624.25	109.25	1234.1	452.27	1047.6	124.17	1815.85	537.05	1302.5	594.11	1706.75
X77307_at	150.95	127.92	10.75	12.37	364.1	400.22	158.8	193.46	230.1	141.99	274.75	132.02	311.8	109.18	132.55	34.01	48.4	16.83	158.2
X58234_at	379.45	243.6	105.4	121.76	256.2	170.84	995.5	635.55	231.55	249.96	538.85	270.33	296.25	194.1	248.35	36.84	617.1	34.22	173.8
X52520_at	381.65	131.17	304.45	351.79	414.9	106.49	942.8	860.27	438.85	315.86	513.65	220.41	581	24.61	598.75	181.51	339.55	98.08	711.7
J04449_at	139.55	51.41	79.95	92.42	269.2	223.73	122.4	102.67	221	93.76	184.1	54.31	133	99.7	611	566.96	129.15	53.39	105.6
U14550_at	1327.4	601.47	1161.1	1341.52	1863.85	32.88	1861.85	793.16	1240.25	147.43	2093.45	537.19	1512	152.88	1642.25	139.09	2062.65	299.74	2240.4
M34182_at	5091.1	2510.37	1900	2195.28	5090.95	2051.95	4857.1	1479.69	4598.35	1605.2	5794.8	1777.24	4794.3	316.93	5461.3	570.07	4283.35	516.26	4214.85
S78653_at	214.55	180.24	83.65	96.66	231.25	216.3	596.95	406.66	966.6	224.72	761.7	355.53	534.15	240.77	775.5	70.85	520.4	57.13	681.25
U72761_at	1762.05	162.85	1040.35	1202.01	1653.85	622.18	2446.3	658.74	1553.35	11.81	1697.45	432.68	1923.35	29.49	1842.6	418.61	2923.65	204.57	2114.1
U15642_s_at	217.1	56.29	398.4	460.33	420.5	1.7	230.5	58.41	338.9	31.96	771.25	596.73	430.45	343.72	147.3	164.61	329	21.78	543.6
U13045_at	224.15	96.66	104.85	121.13	58	23.48	55.5	30.97	481.4	480.83	353.85	30.62	92.35	76.3	200	164.76	324.65	22.27	352.85
X83107_at	148.65	56.07	16.2	18.67	130.7	58.83	94.4	56.14	65.9	34.22	367.7	61.09	168.4	171.4	131.8	7.64	52.15	6.43	337.65
X90846_s_at	338.85	82.38	192.3	222.17	343.3	84.71	331.35	39.67	312.85	94.54	344.8	166.59	308.4	5.52	422.05	92.56	343.15	164.97	235.65
X00038_at	216.95	79.13	454.4	524.96	202.9	38.89	181.3	125.16	118.85	33.3	344.2	180.6	375.7	253.43	308.6	285.25	177.45	92.98	206.7
Z21507_at	14968.4	5411.49	6411.95	7408.29	14695.95	6178.06	13279	5843.96	16257.45	4541.25	15613.9	1536.54	15077.5	447.32	14955.2	2228.09	16488.3	3540.48	16383
M21535_at	247.35	186.61	21.5	24.89	670.8	827.74	355.3	112.15	165.15	80.54	230.05	178.54	327.7	117.38	184.35	40.66	222.2	134.92	90.15

X02544_at	89.4	21.64	29.8	34.37	513.15	516.68	143.15	136.97	124.25	0.35	147.5	110.31	82.05	14.5	107.45	75.31	63	30.55	99.05
L49380_at	5213.85	2493.61	2982.7	3446.16	3917.15	709.02	4596.5	997.87	3619.6	989.53	3992.55	1404.53	4844.35	253.92	4460.75	850.86	5376.8	1190.34	4166.95
D63135_at	294.65	280.23	125.8	145.38	906.3	343.09	231.55	20.86	256.35	29.63	1109.55	136.12	996.05	136.54	502.3	143.68	927.7	151.32	622.05
Y10571_at	106.05	62.01	25.75	29.77	509.4	670.76	86.05	43.63	167.15	59.89	218.75	207.82	196.8	239.71	188	237.16	133.35	158.46	317.8
Y10506_at	182.75	100.76	102.25	118.16	55.6	7.78	723.15	352.78	274	237.59	688.35	656.69	302.6	339.55	271.2	310.28	233.65	5.3	240.8
L10403_at	212.5	159.95	60.45	69.79	190.15	41.65	186.55	85.21	286.25	41.65	155.1	46.39	175.8	9.9	208.7	114.69	317.75	3.46	147.4
HG3044-HT3	74197.95	35646.59	49090.65	56718.24	40931.8	18552.93	45489.4	7018.74	53854.1	7020.44	49467.9	13820.69	43474.3	2707.23	37215.2	8411.18	50490.15	28299.47	47236.85
M96759_rna1	416.15	141.92	380.5	439.68	610.5	632.86	320.85	170.48	176.95	29.2	116.05	38.11	253.65	174.87	273.05	160.3	374.2	195.16	513.1
X16660_cds1	114.35	13.65	111.65	129.05	226.7	165.32	98.4	57.13	102.45	22.84	250.75	162.99	155.6	34.65	189.7	158.82	78.3	4.81	255.1
HG2028-HT2	1315.9	38.04	399.4	461.46	1268.85	102.04	1996.1	1050.34	1557.85	42.07	2432.05	522.34	2132.5	253.14	2112.05	769.26	1682.45	28.07	1891.05
M21388_r_at	19155.55	3332.52	7458.25	8617.16	13115.75	5259.53	12711.9	2200.66	14593.55	1979.26	16516.85	5731.31	16796.2	1122.6	14641.3	9.76	14986.9	2442.91	12867.4
U32439_at	718.9	14.85	437.45	505.37	584.45	591.07	652.65	381.91	634.5	7.35	1183.65	771.1	463.85	390.96	309.5	217.79	268.1	294.72	456.8
D14043_at	818.15	96.8	601.3	694.66	905.05	428.72	530	553.52	970.45	442.44	1015.7	128.98	818.7	246.78	1201.9	859.98	475.2	293.87	487.1
M63962_rna1	1274.05	117.17	199.65	230.73	495.65	24.96	762.45	299.18	1010.9	350.3	2000.65	949.43	770.8	284.68	732.45	430.84	790.6	584.35	839.3
X01630_at	8585.45	1638.72	3348.35	3868.65	8060.8	1043.27	7525.2	3179.72	3852.2	280.44	5487.4	436.28	3396.3	351.86	3460.8	1457.91	4361.5	2649.53	5174.2
M24594_at	146.75	53.25	62.35	72.05	441.1	387.49	581.55	141.35	280.4	233.77	369.15	305.82	578.8	660.44	133.3	2.12	593.5	167.58	250.85
Z46632_at	210.15	215.6	37.2	42.99	331.95	366.07	1766.7	2258.92	750.85	312.75	209.45	253.07	148.2	84.85	419.65	97.09	942	336.3	399.35

M96982_at	3804.9	1623.66	1443.95	1668.28	3092.25	60.88	3022.4	579.69	4960.7	145.66	2996.6	1327.52	3484.45	480.48	3182.2	40.59	3774.45	823.99	3968.4
M23178_s_at	454.35	242.33	402.2	464.71	407	347.19	981.75	928.93	202.85	211.78	291.65	289.14	893.35	833.33	448.9	36.35	526.25	154.64	732.6
X83573_at	2430	481.26	1098.9	1269.68	2313.55	172.04	3058.15	1377.23	402.35	86.76	2323.3	416.77	2252.35	817.2	2022.35	565.19	1199.4	438.55	1685.5
U34877_at	1341.6	858.99	1065.1	1230.65	1738	275.06	1884.95	464.36	2430.55	118.44	2246.15	859.49	1224.45	912.95	2042.15	137.53	2308.35	245.72	2312.55
M18079_at	463.9	378.16	169.05	195.37	365.35	307.66	398.95	106	406.9	163.48	89.65	40.09	262.4	250.6	361.8	483.24	244.55	263.54	198.2
L41559_at	1475.4	15.41	783.7	905.52	2199.3	568.09	2256.75	606.49	1050.25	1192.68	2031.85	363.38	1288.8	660.58	1500.1	259.23	1179.9	145.24	1402.95
X00540_at	77.9	27.29	100.2	115.82	269.35	337.78	94.8	59.82	192.95	114.76	165.9	74.67	253.5	285.11	143.55	156.62	174.65	161.01	130.8
Y10262_s_at	186.75	64.42	448.65	518.38	178.7	156.55	151.65	91.29	118.05	39.24	360.6	115.97	150.55	6.01	107.3	35.07	417	276.62	280.4
U91985_at	1414.95	194.1	1088.2	1257.24	887.35	102.6	1438.65	134.14	1491.55	171.47	2223.15	400.29	1446.3	230.94	1317.2	469.66	978.2	575.73	999.6
X89426_at	32.65	19.3	32.5	37.62	48.75	16.9	88.55	98.08	348.45	107.98	274.55	185.76	211	24.61	66.2	26.3	160.5	169.99	58.15
U57316_at	1736.9	155.7	1238.2	1430.62	2309.3	1136.89	1822.6	325.69	1721.05	582.44	2253.25	508.48	1807.05	32.17	2557.5	908.07	2011.2	481.96	1277
AB000897_at	821.9	336.02	967.6	1117.94	3667.45	3385.84	4008.35	3267.05	2326.45	2039.23	2666.75	1467.32	2095.25	1222.09	2761.75	2496.16	1577.2	359.78	2170
D28118_at	398	175.5	39.2	45.25	900.9	494.83	346.1	17.96	304.15	269.05	731.95	71.21	540.85	25.24	671.6	354.54	137.85	48.3	736.3
X66360_at	182	226.13	36.65	42.36	161.6	175.65	100	18.95	45.8	12.59	297.45	374.7	23.55	9.55	42.7	21.92	21.6	7.35	39.45
J04599_at	2560.65	580.75	1001.2	1156.83	3053.75	3306.22	1615.6	569.79	2933	1798.6	1182.9	593.4	2939.1	336.58	2325.55	127.35	2184.4	965.63	2395.85
M27492_at	550.1	480.55	354.5	409.56	1117.9	707.96	923.4	786.59	1036.6	796.91	841.9	642.62	569.7	88.11	1199.75	666.45	401.05	200.61	951.7
U11732_at	416.55	473.55	77.25	89.31	481.55	226.91	928.7	1131.09	192.65	132.87	155.1	55.3	139.85	37.97	245.6	59.4	794.7	383.11	766.4
M18700_s_at	2899.85	294.37	1795.7	2074.65	2961.8	1089.09	3881.55	1805.46	2002.35	246.43	2575.75	90.16	3185	722.1	2489.5	291.61	2729.6	832.55	2552.5
X80343_at	362.25	178.4	142.75	164.97	585.9	9.9	322.15	115.19	291	82.59	401.5	123.46	324.3	51.62	370.95	145.03	281.95	49.85	260.9

X74795_at	3342.4	660.58	1961	2265.71	2501.35	622.61	3751.8	79.9	3803.55	129.47	3844.85	642.12	4584.8	530.9	4508.05	391.67	4426.4	868.04	3467.05
M36429_s_at	3286.4	1254.83	1754.95	2027.63	2191.25	1032.16	1288.5	300.52	1818.4	259.65	1880.4	104.51	2964.05	1331.84	2160.6	371.23	2735.35	515.69	2992.4
M63154_at	33.05	15.77	30.45	35.14	412.1	501.06	89.35	49.29	142.35	77.57	584.4	743.59	42.9	6.08	46.65	18.6	27.2	17.54	38.6
X54925_at	128.8	69.44	142.2	164.33	222.7	159.66	113.95	8.27	338.25	212.91	313.2	188.66	515.15	686.25	680.65	232.71	780.25	387.99	280.05
X04106_at	17023.75	1567.44	6562.65	7582.38	13587.3	5444.86	16918.35	5686.62	13087.4	4804.79	14106.35	3623.71	15556	2092.75	12228.25	2578.75	18147.35	4864.26	18807.2
M22919_rna2	1445.5	287.51	871.7	1007.2	3458.65	1742.81	2481.8	240.84	1612.05	296.63	1994.55	38.82	1322.6	257.67	1035.3	539.1	1570.85	290.55	1427.25
M95767_at	101.3	59.54	170.25	196.65	202.6	37.34	798.5	739.07	184.1	144.82	96.6	54.59	164.45	108.26	91.4	52.18	46.1	15.56	225.45
U43408_at	482.85	261.98	142.9	165.04	1028.4	786.16	950.7	506.15	437.15	35.85	923.3	63.78	483.35	289.84	1505.1	15.7	1122.85	449.37	465.45
U85767_at	179.4	121.76	170.8	197.28	250.4	185.83	1065.75	1303.98	105.65	5.73	314.65	90.3	154.9	61.66	318.9	131.66	251.05	49	183.8
HG2367-HT2	308.45	264.67	293	338.56	716	329.79	904.55	1119.28	536.75	517.81	351.2	45.96	86.6	2.12	293.1	149.77	514.8	299.81	294.05
L47276_s_at	719.4	185.69	668.3	772.16	348.15	164.12	846.6	876.95	825.95	94.82	739.25	410.9	795.1	407.43	897.7	154.29	662.75	327.18	210.45
X03473_at	136.8	108.33	375.8	434.16	460.4	430.35	112.8	42.99	456.75	450.78	152.1	48.93	327.5	227.12	408.9	483.24	93.1	32.1	287.5
D29012_at	11758.8	3478.68	5462.35	6311.14	11311.7	2124.43	10813.1	4143.65	9301	1341.1	11025.35	73.89	10970.45	1540.72	11590.15	1919.72	9114.55	408.5	9480.5
S69231_s_at	127.9	5.23	125.65	145.17	234.8	198.98	143.1	12.02	83.55	62.44	238.65	92.28	335.5	283.97	152.9	41.72	197.75	96.94	394.85
U43030_at	1343.7	512.37	288.4	333.19	930.5	441.66	820.3	368.4	856.25	480.34	1867.5	834.95	1773.9	642.62	1273.75	830.5	1078.4	74.95	1333.2
M55284_at	384.55	479.21	121.85	140.78	126.45	73.89	145.75	25.81	177.85	133.71	187.45	91.99	156.75	26.66	175.65	10.82	129.85	10.54	112.8
U07857_at	3420.4	330.36	1475.7	1704.98	3497.6	287.37	3467.6	1289.9	3592.6	246.21	3031.7	723.23	3301.1	113.99	3116.95	384.74	2951.3	621.26	3233
D49354_s_at	545.15	252.51	585.5	676.42	463.65	153.09	620.1	247.2	995	334.46	543.9	253.71	773	793.09	137.25	114.2	579.5	191.63	419.65
HG1034-HT1	295.75	244.45	693	800.73	256.5	188.66	213.75	70.07	284.65	192.55	518.2	502.47	172.8	10.47	276.3	22.63	165.35	75.73	130.15
U60975_at	962.8	77.64	657.65	759.79	882.15	682.85	1444.7	101.12	1069.9	403.9	878.8	128.13	1228.7	1166.02	1421.6	399.8	1066	258.52	859.9

M62840_at	178.3	127.7	79.65	91.99	150.25	54.24	176.85	88.32	198.25	12.52	334.8	268.42	170.75	14.35	188.9	59.96	156.7	13.29	190.5
L77561_at	447.35	354.76	499.15	576.65	484.8	97.44	462.45	237.09	323.55	176.14	398.65	169.21	198.75	129.47	505.55	33.3	501.2	22.91	65.8
M76729_at	83.9	2.55	33.2	38.33	173.65	138.24	169.35	113.21	123.1	0.57	162.65	94.26	104.8	16.69	104.75	24.25	176.9	131.95	123
L42621_at	1762.95	374.13	554.65	640.85	1303.9	379.57	1801.15	162.42	1374.95	90.3	2183.25	202.87	1215.75	678.33	1584.85	88.18	1273.25	254.91	889.1
M64358_at	176.4	204.07	73.65	85.06	235	226.42	758.25	958.48	386	7.21	287.2	248.76	324.65	351.64	194.8	74.1	60.4	49.07	103.15
L24804_at	2130.35	562.08	1060.9	1225.7	1915.4	170.41	2961.4	1069	1955.9	738.36	2224.3	644.17	1363.2	193.46	2827.8	166.59	2505.95	214.89	1997.3
M55422_at	1307.55	1108.81	478.55	552.89	1459.65	598.71	807.45	193.39	1087.45	181.8	714.15	20.58	1076.05	114.34	901.35	21.99	600.15	466.9	688.1
U18004_at	21	1.13	32.45	37.55	56.75	16.33	37.95	32.6	37.6	27.72	66.55	12.23	55.9	41.44	155.85	75.02	44.8	21.92	47.95
HG862-HT86	567.7	520.29	73	84.29	291.7	67.18	330.4	237.02	204.65	61.31	311.5	68.02	202.35	40.38	149.8	54.59	241.1	180.31	120.75
S52969_cds1	4141.6	1425.24	1659.45	1917.32	2434.6	870.87	3210.65	292.81	2872.65	942.93	3400	65.34	2857.9	1813.87	3084.5	664.54	3401.65	598.28	2807.15
X83857_s_at	50.15	21.99	68.7	79.34	91.25	76.58	86.1	102.39	36	23.33	93.8	104.51	221.8	136.47	66.85	27.93	274.1	78.63	107.3
HG1227-HT1	106.4	60.81	123.45	142.62	176.4	106.35	147.95	121.98	61.3	9.05	126.3	41.01	154.55	116.04	93.75	15.34	78	5.52	130.1
X75091_s_at	1668.05	437.63	1102.6	1273.92	1127.5	36.06	831.3	453.82	1091.05	199.62	1188.1	349.59	1536.45	438.62	1407.95	316.15	1562.7	502.47	1176.7
U62437_at	1595.7	957.42	254	293.45	2499.05	1564.47	2204.35	1731.07	1616.3	1007.63	668	295.15	1140.15	702.93	2411.6	659.73	1424.35	606.63	954.35
U13061_rna1	108.35	83.51	41.85	48.3	124.1	39.74	166.2	140.57	365.2	407.58	55.05	21.99	164.5	173.1	115.3	88.53	215.9	255.83	87.75
D86980_at	137.7	181.59	165.2	190.92	466.2	621.41	127.6	18.53	33.25	1.63	564.6	365.15	124.85	152.24	161.6	182.43	224.9	65.48	236.7
Z71389_at	269.95	155.21	772.05	892.02	528.9	358.08	463.25	328.45	368.7	53.03	477.15	74.88	554.35	357.87	216.05	84.64	256.2	32.53	265.95
Z29331_at	321.6	260.64	61.35	70.92	1075.7	999	681.2	303.63	824.95	217.86	529.3	374.34	100.2	48.37	167	13.15	120.45	6.58	858.55
U19718_at	1182.35	4.45	430.85	497.73	2060.45	1065.68	1032.3	42.28	588.2	78.49	678.1	49.36	657	47.94	1641.1	1330.63	601.55	49.43	1178.1
HG2604-HT2	238.7	131.38	155.55	179.68	233.1	27.72	214.2	8.34	199.25	62.3	196	43.98	149.85	8.41	273.7	55.15	239.8	77.36	307.45

U82468_at	1554.35	1272.16	242.75	280.51	466	95.32	1363.3	1003.38	470.25	104.86	636.25	475.95	516.05	4.6	1229.75	1200.03	678.1	96.59	1115.8
HG4068-HT4	189.1	19.09	50.15	57.91	286.35	237.94	324.5	88.39	308.25	130.74	902.95	1015.33	497.9	50.91	426.95	24.4	331.55	166.81	326.35
M60614_at	247.5	4.95	147.1	169.99	226.1	170.13	212.1	81.46	135.65	32.74	145.2	31.68	261.7	85.28	238	24.75	274.2	158.53	132.5
U18291_at	548.3	30.55	598.7	691.69	875.75	122.4	594.35	430.42	914.3	152.31	1032.6	362.32	159.55	44.48	535.55	534.5	944.8	392.73	1423.85
U57877_at	2352.4	111.02	1169.05	1350.64	2065.95	683.42	1851.25	530.26	2661.2	899.3	1909.85	768.55	2482.5	326.54	2561.35	339.62	2266.15	651.46	2650.95
L47738_at	2122.8	362.6	929.05	1073.46	2267.7	373.07	1347.25	776.19	1619.75	242.33	1299.6	468.39	1675.5	60.81	1005.2	684.06	1229.55	418.54	1119.4
HG3565-HT3	487.45	164.26	304.75	352.07	946.8	251.45	1092.4	1094.32	147.55	87.19	1362.75	989.03	697.15	166.1	1047.2	964.21	430.25	120.99	559.4
HG3992-HT4	255	17.25	73.75	85.21	194.45	175.72	294.5	69.01	230.35	38.25	684	693.53	479.9	386.08	170.95	107.55	194.1	168.57	158.55
S77154_s_at	148.1	92.49	295.6	341.53	427.55	536.2	471.35	491.93	254.45	130.74	211.5	66.47	360.35	326.75	187.55	87.47	304.55	205.27	398.8
X69141_at	2406.45	57.63	1589.5	1836.5	2148.1	323.85	2388.15	286.45	2470.3	1347.89	2091.4	688.01	2518.9	1525.51	2587.15	756.96	2810.8	1422.7	2327.05
HG2999-HT4	310.25	269.76	342.25	395.48	738.45	523.75	559.25	211.5	338.35	64.56	221.7	66.61	194.45	123.39	409.95	424.76	185.15	80.68	289.95
D50924_at	258.65	147.01	263.25	304.13	237.05	215.6	195.8	146.09	471.15	152.52	280.8	147.36	324.1	271.1	177.1	141.42	467.65	252.79	478.4
U14383_at	742.7	210.15	274.3	316.93	575.2	456.08	769.45	324.07	655.6	25.03	1817.6	436.71	878.6	68.31	981.05	206.55	688.65	231.15	1083.55
D14838_at	315.8	110.31	506.65	585.41	175.85	6.29	1071.75	601.11	447.2	349.03	703.6	369.11	413.3	307.45	200.4	9.05	626.3	271.81	1228.85
U55853_at	402.15	11.1	324.65	375.12	245.6	172.96	398.75	147.86	449.15	161.01	528.85	274.85	639	627.91	382.85	492.08	300.05	260.85	534.35
U22897_at	829.25	294.79	567.3	655.49	814.8	150.19	764.65	274	870.8	280.3	818.5	286.52	775.65	255.76	809	60.53	715.6	15.98	1101.5
D16105_at	3577.35	985.78	1559.15	1801.35	2964.65	841.95	3802.85	2147.13	3298.4	628.05	2678.8	889.82	3157.3	468.53	2818.45	145.88	2330.55	1161.71	3056.15
X77533_at	93.1	10.61	30.15	34.86	96.25	6.58	107.45	62.86	82.6	1.41	103.95	58.34	85.05	11.38	88.3	21.64	84.55	14.5	105.8
HG880-HT88	2190.8	1926.02	401.8	464.29	340.2	138.03	1549.1	792.81	783.85	779.16	3609.75	1352.62	2333.8	367.98	4247.75	192.26	2387.9	2411.38	2176.5
X17059_s_at	222.5	112.01	112.1	129.54	440.85	107.41	296.6	130.81	148.3	29.7	429.25	48.3	159.85	103.45	189.85	180.1	375.95	93.13	260.55
U09716_s_at	530.75	272.73	62.75	72.48	621.85	321.1	994.85	824.84	171.35	87.61	752.8	3.96	303.7	323.43	591.3	80.04	828.5	165.89	768.05
L24893_s_at	109.55	55.93	41.95	48.44	159.1	85.56	101.5	17.96	350.2	383.39	288.45	350.65	96.55	23.83	63.5	17.25	81.4	28.14	209.4

D49372_s_at	164.5	160.09	449.6	519.44	223.8	51.34	544.85	630.39	247.8	81.18	694.2	878.65	758.55	107.69	276.05	340.75	287.65	348.82	130.9
U77949_at	129.1	61.52	502.65	580.75	431.55	549.35	327.5	385.37	264.7	79.05	277.85	86.05	238.85	188.44	272.65	158.32	230.4	255.97	126.15
M19483_at	17225.85	5692.99	9827	11353.87	17501.8	4424.93	20219.45	8663.54	16009.95	570.85	16240.65	4128.44	14459.55	538.74	14328.65	1027.64	18343.75	5893.24	17524.25
U53209_at	926.9	198.7	832.8	962.23	709.95	206.12	762.15	631.52	1372.5	344.64	1305.05	583.86	1284.4	516.61	695.6	17.54	1416.9	178.76	937.55
U90910_at	59.1	20.08	89.75	103.73	112	14.85	266.6	218.92	117.95	43.2	144.8	127.28	103.85	41.37	94	55.72	69.05	3.46	71.6
M59916_at	2610.45	767.56	213.3	246.5	1981.5	248.34	1398.35	1312.6	2235.2	474.19	2585.8	546.03	2772.05	261.42	507	346.91	1656.6	892.37	2998.5
S79873_s_at	645.2	135.06	287.1	331.77	597.3	375.47	407.6	387.07	702.3	207.61	360.85	28.35	585.9	362.04	596.2	109.32	358.55	154.5	407.95
D63484_at	440.15	168.93	159.15	183.92	535.55	251.8	471.75	195.09	379.1	48.37	533.45	103.87	455.95	173.45	407.75	96.1	443.4	22.77	539.75
X89101_s_at	648.3	234.76	315	363.88	1078.65	912.1	562.45	274.29	1278.55	1022.26	699.9	113.99	780.5	597.79	1652.6	1190.2	640.85	189.72	557.15
U31556_at	483.15	173.17	1041.15	1202.86	492.15	300.59	533.1	262.76	646.5	504.45	1010.1	304.76	591.1	499.22	556.6	184.41	625.3	143.26	511.35
L13848_at	748.7	202.94	440.85	509.33	270	242.11	578.9	685.19	1355.15	381.63	1119.45	190.71	1178.25	482.88	1678.1	62.79	641.1	386.22	1055.4
Y10510_at	49.6	16.97	42	48.51	300.8	331.07	87.45	106.28	45.6	16.55	90.6	58.55	92.7	29.98	170.95	132.3	225.75	124.8	80.85
U66619_at	73.4	19.52	70.9	81.88	196.85	46.32	174.2	80.89	116.45	68.09	237.4	33.23	174.3	112.15	243.8	2.4	126.65	28.64	101.25
L12711_s_at	16932.4	4346.87	5958	6883.68	14489.6	6471.16	15701.4	2227.53	12031.45	8524.81	10775.5	4759.39	16020.05	1514.27	15423.3	1038.17	16369.5	384.24	11081.45
U28413_at	368.65	216.59	707.35	817.2	217.15	66.4	722.95	128.06	528.15	38.82	702.95	233.84	331.35	378.23	584.5	34.37	295.1	60.95	570.45
X79865_at	3734.1	737.65	2044.45	2362.09	3409.05	274.99	3748.45	437.77	2635.8	1749.95	3461.35	1099.76	3334.1	19.23	3246.95	1124.09	3860.9	794.22	2687.1
X04571_at	142.6	154.01	280.1	323.57	604.5	363.03	167.4	211	119.9	28.85	341.15	123.53	326.95	52.26	172.8	179.04	580.65	38.25	621.2
S38953_s_at	1030.4	268.42	587.7	678.96	936.75	267.78	586.35	321.1	1031.4	37.34	1109.5	144.67	1022.65	100.62	524.25	63.85	784.05	311.34	868.4
U18259_at	339.7	366.14	33.85	39.1	307.3	339.69	453.4	385.09	354.55	48.15	478.4	630.31	312.75	191.98	353.3	104.09	335.1	82.87	710.85

M60724_at	24.05	7	264.65	305.82	205.55	72.62	334.5	421.86	250.85	283.76	245.9	282.28	147.95	121.41	55.4	3.39	41.75	6.15	219.1
M21305_at	16.55	2.47	8.55	9.83	40.45	10.39	61.2	49.64	245.45	310.63	54.55	64.7	26.3	16.97	159	189.36	17.4	4.1	132.45
AF003743_at	383.05	41.79	263.45	304.41	635.45	250.81	491.5	148.63	426.7	116.25	476.35	105.43	349.85	152.1	407	57.98	421.25	3.18	376.55
U61836_at	1166.65	171.33	708	817.98	1416.2	466.55	2099.4	834.24	1113.55	1193.81	926.9	780.36	1172.95	1081.38	1228.85	232.99	1874.5	433.03	1370.45
L27943_at	2424.05	546.95	913.15	1055.07	2990.9	830.99	2909	252.15	2851.95	872.78	2913.55	455.02	3384.9	187.81	4245.7	2665.93	2543.3	745.86	1175.75
D79985_at	1194.05	705.62	311.75	360.13	966.85	370.59	624.85	90.72	1960.3	2026.57	571.7	82.31	482.8	0.71	1859	1836.21	845.9	375.76	1218.45
X52011_at	364.6	158.82	141.2	163.2	556.75	49.57	716.4	499.36	276.15	279.52	405.8	105.22	142.9	93.76	342.95	27.93	179.45	208.53	622.5
HG4120-HT4	434.6	97.86	1178.9	1362.03	597.6	474.47	1308.8	954.17	658.3	555.64	579.75	127.49	241.5	136.61	182.75	27.08	1088.5	126.01	640.9
L32164_at	327.1	175.93	185.65	214.47	705.9	22.77	591.85	326.47	719.3	93.62	1043.15	855.25	492	70.85	483.1	463.86	180	31.11	704.7
D86062_s_at	170.1	65.05	105.25	121.55	898.7	320.32	810.15	693.46	181.8	16.4	387.55	307.8	445.25	337.64	504.9	279.59	639.65	595.03	161.7
M62843_at	397.7	344.93	46.9	54.16	788.1	777.53	833.2	857.01	391.9	306.46	273.65	14.07	247.65	7.99	476.55	370.74	159.5	46.24	344.05
HG3033-HT3	1009.6	479.84	543.5	627.91	1006.1	98.01	2992.8	3149.03	1074.3	79.05	942.2	434.87	1296.3	241.97	911.7	100.41	1548.4	1336.71	647.45
HG2796-HT2	292.35	51.83	293.4	338.99	848	398.53	1056.9	397.96	773.4	658.18	781.85	74.03	1042	498.79	1471.95	328.17	822.2	662.13	1230.95
U44105_at	35.5	5.8	136.4	157.54	95.5	17.25	74.5	97.16	57.45	12.52	352.35	295.5	168.25	206.55	167.6	219.2	63.4	13.72	20.7
X75346_s_at	965.7	223.73	593.35	685.54	269.35	170.91	187.1	3.68	531.15	434.38	582.5	164.33	275.75	96.94	745.65	615.54	596.5	138.31	517.95
U62317_rna6	243.9	203.08	143	165.18	1059.6	1137.73	476.05	122.26	194.6	166.31	1009.2	1092.06	398.45	60.03	214.5	89.94	438.85	97.79	166
Z68747_at	284.9	100.55	209.95	242.61	679	12.87	514.15	651.17	362.25	287.3	296.65	85.06	380.6	195.73	370.5	84.99	373.6	71.42	332.7

X07767_at	2325	1041.14	1153.5	1332.75	1356.85	223.38	705.3	813.74	1580.4	20.22	1261.6	400.79	2143.7	39.74	1732.7	313.25	2135.3	763.53	1561.5
U10685_at	593.7	173.95	60.8	70.29	305.9	241.41	944.75	820.17	420.95	374.27	107.35	15.91	919.75	88.46	489.8	254.42	206.6	118.37	342.3
M54968_at	314.7	81.32	31.5	36.35	239.7	208.31	215.1	94.89	225.2	163.34	184.45	145.31	294.9	210.15	428.9	118.51	494.85	142.91	260.6
D14533_at	161.45	28.35	331.75	383.32	174	84	175.9	188.94	832.4	75.24	570	304.9	1224.1	111.02	672.4	40.16	430.45	80.4	523.05
D10922_s_at	161.35	128.62	38.55	44.48	155.05	177.98	195.6	214.68	369.45	167.66	401	303.49	362.45	469.59	164.95	136.97	165.75	196.79	316.45
Z68204_at	1167.55	143.61	1000.8	1156.26	1237.35	33.45	1131.85	595.03	601.7	103.1	1157.1	452.55	821.9	124.59	1031.15	300.87	1080.2	56.71	1053.15
L19605_at	5169.45	107.13	3069.5	3546.42	4912.75	378.51	5175.6	2027.28	5544	294.3	4262.15	1011.8	4985.8	186.96	3541.65	1004.16	4715.15	311.48	4399.15
U04520_at	435.9	215.38	371.95	429.71	915	490.17	417.55	319.4	695.95	728.67	588.5	470.65	916	649.97	946.25	570.42	635.5	135.48	256.5
M64571_at	6321.75	1995.95	3514.3	4060.35	5189.95	1285.03	5218.05	443.29	5268	707.53	6024.65	960.46	6538.35	1048.71	6367.85	893.57	6270.85	1114.9	5534
HG2649-HT2	1252.2	80.89	235.7	272.38	1452.45	514.56	819.2	304.62	790	609.95	1214.75	593.05	512.55	152.1	1062.5	958.98	1117.8	853.76	962.75
U57342_at	7149.9	1540.36	3636.05	4200.99	4407.55	845.91	6809.05	2763.3	4835.45	1083.78	6083.95	2584.4	6142.9	425.4	5572.15	1382.32	6346.55	411.61	4678.95
M31423_s_at	119.55	98.5	246.05	284.33	111.15	96.24	280.35	276.97	202.95	203.01	443.05	472.7	263.05	329.16	392.8	116.96	218.75	81.53	71.1
M13232_s_at	245.25	147.43	41.35	47.73	731.25	220.97	540.2	150.47	538.45	66.11	988.65	512.44	320.65	267.07	403.85	9.83	402.6	380.56	753.15
M64098_at	6574.35	789.63	3957.6	4572.58	8041.85	284.61	8179.9	1551.67	4905.55	750.03	5834.65	141.63	6813.45	578.34	7135.5	458.77	7465.45	232.57	6427.45
M91670_at	4046.3	699.47	2630.75	3039.5	4231.55	423.34	4357.4	1434.86	4403.45	586.12	4872.7	1478.42	4649	511.52	6433.35	1871.5	5346.6	516.33	4182.85
HG3731-HT4	104.05	49	64.5	74.53	246.95	189.86	194.15	170.62	458.35	534.08	437.85	538.18	160.75	167.51	146.45	43.2	79.6	19.09	166.8
U79241_at	658.85	227.9	914.85	1057.05	538.75	387.85	1030.8	1025.87	771.75	627.98	607.8	79.48	993.55	845.06	737.9	571.48	639.4	555.93	972.75
M75126_at	7313.35	424.76	4094.6	4730.83	6543.15	261.56	8544.6	86.55	6158.35	411.89	6212.8	307.17	6935.65	1136.82	7430.55	1195.08	7613.4	62.65	6201.65
M34276_at	1404.85	12.23	293.75	339.34	756.95	166.24	1146.9	669.77	1017.55	118.3	1639.95	931.61	1224.9	65.48	860.45	52.11	838.3	155.99	1095.75

M98045_at	3979.15	333.83	1109.3	1281.7	3037.3	902.98	3344.95	448.94	2537.35	1350.08	3392.55	205.41	2569.8	918.11	2971.25	1262.26	3196.4	474.47	2318.2
M11726_at	1489	76.51	582.5	673.02	1434.3	77.07	1900.6	948.09	903.75	374.27	1498.35	395.91	1541.7	440.24	1286.65	670.55	763.6	50.06	722.7
M27161_at	386.15	323.36	89.15	103.03	196.4	127.99	686.6	149.77	852.05	5.02	316.1	248.48	1100.45	359.28	268	13.01	389.35	272.17	508.55
L32140_at	77.2	26.02	110.7	127.84	280.05	260.57	353.75	107.69	297.75	392.37	639.9	135.06	107.6	3.96	442.05	119.15	186.4	150.47	153.8
D31887_at	3221.9	47.94	3370.5	3894.18	3055.3	1221.03	3301.75	1542.27	3822.55	474.26	3672.2	285.25	3074.65	418.82	2949.6	963.5	2769.45	329.02	3048.35
D13626_at	120	140.71	137.25	158.6	118	135.2	128.25	131.45	64.05	49.57	36.75	11.24	55.6	50.49	161.25	191.27	320.75	106.84	352.35
S60415_at	43.8	7.78	52	60.1	139.85	62.58	111	130.39	58.05	1.34	143.45	156.06	57.3	10.61	98.2	34.08	234.55	245.72	55.1
U65918_f_at	351.55	187.6	406.55	469.73	69.25	57.06	86.3	46.1	187.7	51.48	144.6	118.37	48.65	42.21	84.5	53.88	273.2	325.41	238.25
X98833_rna1	930.7	321.31	460.1	531.6	534.3	435.01	535.4	47.52	927.85	740.13	672.5	322.16	885.75	815.79	273.05	81.67	554.25	200.75	260.4
X54304_at	2509.5	173.24	1286.8	1486.76	3006.6	410.97	2267.95	686.67	2228.65	268.21	2587.45	16.76	1769.3	387.07	1270.95	716.37	2446.2	63.78	2896.5
D79987_at	219.75	77	139.8	161.5	515.9	357.65	208.25	75.45	349.05	154.79	338.7	101.54	457.7	263.04	403.1	193.18	209.95	57.35	308.75
D43950_at	5585.25	1539.16	3034.05	3505.48	5165.5	635.27	6601.85	583.29	5054.35	443.57	4423.85	699.68	4513.2	171.54	3826.3	655.63	5720	990.37	5802.85
D90209_at	11101.6	3725.46	13290.4	15355.39	17124.9	1240.69	21212.9	9533.07	17613.45	2148.97	22274.65	6268.57	18436.15	6774.58	17740.5	6030.91	9304.5	796.06	20949.2
M59807_at	2292.95	793.02	569.75	658.25	2307	546.31	1631	1283.96	1651.2	280.44	2876.45	692.89	282.15	79.97	2275.35	168.22	1219.9	657.18	2161.1
U07559_at	442.9	151.32	355.25	410.48	360.25	416.27	703.6	352.56	258.3	261.21	947.25	95.25	646.2	86.97	246.5	45.11	336.3	271.81	585.4
J04617_s_at	188856.55	72271.2	127537.55	147354.19	237719.05	19731.75	186920.8	98558.24	236625.25	87907.03	201427.5	117066.91	194747.95	32852.82	197563.69	118042.14	210526.5	33002.09	155143.91
U78793_at	804.3	445.34	1015.8	1173.66	2379.55	1871.92	1550.95	748.33	393.25	372.01	1532.4	615.32	1510.5	21.5	1064.85	479.21	694.2	123.32	1224.8
S77835_s_at	54.3	26.87	88.3	101.96	53.65	54.94	62.85	55.79	85.6	102.81	35.05	32.88	79.5	13.86	39.95	0.21	66.95	20.44	54.05
S67070_at	605.05	154.22	1161.2	1341.66	499.7	199.83	1929.7	2232.05	473.95	73.89	550.55	31.75	2338.3	266.58	1026.95	449.65	438.25	76.58	494
U36601_at	130.95	31.04	75.45	87.19	160.4	49.5	559.2	518.59	169.25	37.55	208.25	1.48	152.9	25.88	173	14.71	177.1	36.49	175.25

X77744_at	412.75	370.74	61.9	71.56	463.05	15.49	629.2	430.2	547.7	252.86	182.9	81.88	903.3	408.14	140.65	123.67	347.95	140.93	727.85
D14889_at	1089.95	423.49	477.35	551.47	710.2	34.37	1177.6	118.51	1596.25	317.56	1112.95	393.5	1563.8	70.57	1605	291.47	1237.4	218.35	1227.35
HG2161-HT2	2319.95	2075.99	502.9	581.1	1670.85	1115.18	1647.05	1522.19	1290	600.33	1710.5	418.18	1682.5	1548.28	1877.3	1484.78	2489.35	2153.78	1251.9
M23161_at	831.45	127.63	880.3	1017.1	463.8	504.73	877.2	112.43	1423.4	412.67	971.8	480.13	1086.3	439.54	1275	119.64	932.6	130.11	1262
X62466_at	426.6	292.04	592.75	684.83	1360.5	317.92	975.85	860.76	297.85	7.42	896.4	683.63	678.85	646.65	726.95	375.83	410.6	351.15	670.95
X79882_at	971.6	819.4	763.45	882.12	1300.45	876.18	2132.75	702.65	1327.15	1457.28	1655.8	123.32	1597.3	916.83	3315.85	1891.3	3015.3	209.59	2719.65
M58026_at	19147.75	5052.91	8790.8	10156.74	13259	813.31	15765.15	2887.89	25487.45	7728.61	18363.2	1339.83	33776.05	7972.13	29181.05	7013.16	44388.95	26367.09	24996.55
M93143_at	1803.15	415.28	318.75	368.33	1422.5	285.25	686.8	820.95	1576.6	69.86	1036.25	716.79	1230.85	73.04	302.6	340.54	1157.2	289.07	1187.05
X58529_at	524.85	479.35	303.35	350.51	590.35	633.5	317.2	103.24	741.15	683.56	526.5	137.04	537.75	312.19	647.05	516.12	354.9	86.27	324.95
L20316_at	3387.05	660.93	1183.35	1367.19	3756.3	2078.75	3228.8	1923.61	2427	985.99	2306.6	1736.23	2949.25	35	1874.8	1857.29	1301.55	802.35	2466.45
U50361_s_at	87.6	3.11	13.55	15.63	134.6	91.36	502.95	692.19	48.9	16.55	389.5	499.5	307.55	261.13	266.15	275.42	47.7	47.23	38.9
Z84483_at	607.55	419.24	521.1	602.03	404.25	232.57	846.75	620.06	945.75	180.81	885.6	311.69	454.2	460.89	249.2	29.84	775.65	574.52	661.05
Z26256_at	320.8	104.09	201.5	232.78	504.45	70.22	549.9	540.8	356.85	25.95	456.05	151.82	537.95	199.76	813.35	263.68	337.95	115.47	761.05
U87459_at	1753.75	568.87	943.8	1090.5	1739	471.78	2951.25	1155.34	2288.15	39.1	2070.65	740.13	1945.7	618.29	1893.65	935.43	2156.9	49.5	2307.75
S72493_s_at	17820.15	4330.11	11506.2	13294.03	18157.9	296.7	24197.2	8332.26	12364.55	1089.3	13388.85	1735.74	16844.1	1442.22	23005	12256.42	21048.85	9409.26	13367.45
AB000115_at	1033.25	231.15	1045.75	1208.23	3060.05	2633.62	2484.2	1864.64	2619.9	1343.36	2589.25	1734.6	1940.1	178.62	4458.2	2836.06	2013.15	10.11	2978.9
X52599_at	133.75	4.03	77.05	89.02	166.95	49.29	174.8	58.41	97.25	43.2	108.95	4.45	92.05	36.56	154.05	65.41	309.55	364.09	226.1
U72209_at	747.35	72.48	410.2	473.9	844.6	323.29	855.6	879.36	904.35	154.08	745.5	123.89	136.25	60.03	290.05	167.8	745.35	58.48	799.7
U35113_at	1617.8	1405.02	441.45	510.04	1341.55	517.67	1037.6	721.96	1185.7	475.18	1129.15	352.49	1488.3	81.6	1693.35	365.79	1919.4	210.72	1272.2
D79994_at	1124.7	268.7	965.95	1116.03	833.1	237.87	334.1	254.42	1270.4	94.61	1827.95	126.22	1279.1	750.1	729.15	150.12	1416.85	309.64	1191.35

M86667_at	1635.4	596.09	622.25	718.92	2977.95	798.54	2363.8	757.03	2579	752.08	2351.5	248.34	1710.3	328.1	2210.9	372.36	2543.15	1052.81	3568.4
HG3859-HT4	130.95	0.78	449.1	518.87	562	622.25	112.6	67.74	157.25	124.52	327.6	325.27	115.9	2.26	125.85	55.93	164.15	18.74	191.35
M16405_at	2547.4	756.32	693.4	801.15	2254.1	869.46	1777.05	179.82	1551.1	39.17	2567.1	828.73	2449.5	150.61	1442.6	38.33	1563.9	260.92	1024.2
Z69915_at	352.9	21.07	175.6	202.94	172	83.16	93.8	17.39	413.4	168.29	547.55	21.43	373.75	76.58	321.1	104.51	571.6	145.1	195.05
U18467_at	273.3	8.34	149.15	172.32	561.75	359.99	282.8	265.45	530.05	23.83	542.4	376.04	381.9	115.68	233.95	144.74	152.65	31.89	473.95
X96401_at	447.15	95.39	933.15	1078.13	645	125.02	510.95	194.53	1649.55	875.89	1492.35	1540.43	2115.35	264.95	2021.45	826.68	1362.45	778.88	784.45
D38500_at	220.55	51.55	135.15	156.2	122.1	83.44	237.8	28.71	258.55	43.49	221.2	52.61	759.4	800.73	454.1	381.13	182.6	20.22	153.25
U60061_at	655.25	391.24	652.75	754.13	451	211.42	1221	671.04	865.9	389.9	820.05	139.65	625	70.85	582.95	382.47	549.45	107.13	906.65
HG3638-HT3	155	93.2	499.8	577.42	358.55	77	362.35	288	628.8	582.66	825.55	620.77	277.45	81.39	183.25	5.59	942.75	127.63	246.2
L38929_at	282.4	281	175.05	202.3	706.4	815.29	74.65	9.55	211.05	51.41	173.2	23.9	161.7	28	108.6	30.41	213.5	150.33	211.65
M22324_at	775.15	355.75	456.45	527.43	1066.95	808.86	425.15	94.26	505.1	220.19	656.6	61.09	493.05	65.83	571	204.07	652.45	137.11	1902.3
AFFX-HUMIS	36.5	6.65	42.6	49.21	91.9	73.26	195.8	211.14	48.15	14.5	100.65	61.02	63.7	27.29	123.75	53.81	34.2	1.27	115.7
U84551_cds2	256.1	287.79	49.4	57.13	424.75	476.1	66.9	19.23	369.85	372.01	150.65	88.03	99.7	89.38	73.65	52.82	251.05	285.03	212.85
U82979_at	526.9	374.06	215.15	248.55	596.7	486.07	498	147.22	678.8	613.77	397.05	231.86	639.95	16.19	496.3	138.59	410.05	222.95	523.5
S68616_at	8608.4	683.91	3664	4233.31	6658.35	413.59	6224.85	347.4	5502.2	946.11	8221.2	996.88	5171.5	335.31	7472.45	748.61	7417.7	2787.42	7005.25
U68162_cds1	659.8	379.72	128.5	148.49	481.45	457	183.6	65.2	227.95	126.78	390.5	38.18	451.55	388.84	699.8	318.76	241	92.07	370.95
J04058_at	701.25	80.12	885.8	1023.47	1054.25	444.13	1295.35	247.7	1013.95	90.86	782.75	55.51	691.75	412.46	690.3	529.76	463.85	0.07	834.25
M30269_at	244.6	142.41	114.65	132.44	550.5	432.04	285.65	382.76	150.95	100.06	138.2	0.42	190.25	189.29	216.9	16.69	336	396.97	425.35
U60644_at	4127.35	1488.53	1807.2	2087.95	2247.7	1993.9	2966.05	470.86	3179.45	1099.76	3659.3	3443.04	3939.55	440.46	3672.55	277.26	4595.6	574.17	2974.95

D29675_at	320.2	310.99	49.8	57.56	98.75	6.15	194.35	15.91	196.7	214.39	480.6	427.38	245.95	181.09	180.25	57.06	1080.25	1120.13	513.3
Y10260_at	461.4	535.7	430.95	497.87	335.85	222.81	1662.4	1784.88	323.7	15.41	179.3	56	338.4	17.68	355.35	138.24	219.85	22.13	262.75
J02986_cds1	364.1	237.02	288.65	333.54	525.4	263.19	874.8	432.75	590.05	210.36	436.35	344.01	156.15	49.14	386.75	109.25	473.95	222.1	299.85
HG2036-HT2	2440.3	671.75	881.25	1018.16	1729.1	713.33	1933.9	1022.76	1586.7	431.34	2647.25	1062.99	1763.95	140.22	1769.8	973.97	2500.65	1128.61	2299.55
J05412_at	1777.65	507.63	1262	1458.05	1616.9	873.98	1147	156.13	1445	47.38	1811.85	85.91	1946.9	687.87	1495.5	103.24	1952.1	126.29	1288.35
M67439_at	122.45	15.63	130.15	150.4	233.8	59.26	254.2	163.34	167.55	30.9	519.65	484.16	205.9	37.19	273.3	28.43	256.05	95.25	234
D86479_at	930.55	474.54	473.1	546.59	1167.65	106.28	1357.9	386.5	1110.95	202.16	1041.95	120.28	1870.25	176.42	1655.9	434.87	966.85	264.39	1554.4
X78925_at	82.3	0.28	47.9	55.3	200.6	225.14	594.35	781.14	214.35	248.69	442.7	551.83	32.15	20.29	64.2	25.88	182	216.52	229.05
U37022_rna1	2120	296.56	1069.3	1235.46	2256.9	349.17	2611.65	224.65	2383.1	347.05	2075.3	103.94	2243.75	575.8	2241.95	50.42	2548.45	525.45	2693.25
K02054_at	112.9	99.28	142.85	165.11	692.35	496.04	175.35	49.85	236.85	186.46	236.85	36.13	142.15	23.41	526.15	385.59	116	9.05	224.8
D38293_at	1107.6	84.99	183.5	211.99	1633.2	360.91	893.4	68.02	1259.45	241.05	1548.15	59.89	622.55	436.5	822.15	458.84	1246.55	417.41	1202.3
X81892_at	192.1	151.6	217.95	251.8	369.55	207.11	497.4	272.66	392.45	40.8	721.95	708.45	290.8	336.58	388.35	289.14	354.3	182.29	502.15
X76057_at	918.5	271.67	589.8	681.51	948.05	393.65	1395.6	12.3	549.7	432.04	523.6	318.62	845.5	873.28	331.95	243.03	1232.2	688.72	1190
HG960-HT96	1296.05	281.36	1348.8	1558.32	638.7	255.27	1660.75	296.35	1637.95	663.62	1943.7	532.03	1183.6	384.1	364.4	105.5	1405.35	150.83	1206.55
X93499_at	2108.45	941.94	1242.2	1435.14	1758.45	195.66	1851.8	557.34	2119.5	678.82	1288.95	519.94	1708.4	645.02	1842.3	196.86	2591.7	170.55	1509
M33308_at	2353.85	592.2	1432.7	1655.34	2079.45	234.97	1856.9	710.78	2146.3	391.03	1879.4	714.46	2249.2	83.58	2182.9	311.13	2092.7	787.01	1400.3
U13021_s_at	131.1	82.73	124.55	143.9	98.85	11.1	185.85	174.3	186.75	85.49	70.3	17.11	166.3	30.26	82.9	14.42	95.45	43.77	168.35
U33448_s_at	711.55	227.05	264.35	305.4	614.85	79.83	881.45	762.05	565.95	69.51	1661.4	1768.9	654.75	309.92	1748.45	1733.76	2031.75	286.73	966.4
X59766_at	73.65	26.94	56.7	65.48	118.9	45.11	141	59.96	230.4	238.01	248.9	58.97	52.05	12.94	193.5	125.58	137.05	12.52	107.15

M16342_at	1702.65	141.07	1272.75	1470.57	1908.45	49.71	1769.95	143.05	1972.75	989.03	1961.55	171.61	1758.15	96.38	1657.75	468.03	1794.65	1256.32	2405.55
M77235_at	155.95	6.29	52.15	60.32	256.55	60.17	492.6	465.28	156.55	13.22	215.55	54.24	175.4	91.5	103.95	2.05	156.7	84.15	189.4
X86098_at	321.2	256.68	53.7	62.08	118.2	32.53	105.8	9.05	385	31.68	472.85	222.39	88.35	63.99	326.05	321.52	179.05	91.71	284.9
D14827_at	447.45	333.97	927.65	1071.76	786.6	658.6	707.7	367.55	1103.1	8.91	387.4	75.24	936.9	424.97	192.2	74.39	455.25	246.99	757.3
HG651-HT52	445.55	131.17	306.85	354.47	501.7	321.45	550.9	383.96	299.05	31.32	425.85	21.14	532	174.09	512.1	112.71	998.35	921.01	264.75
U43747_s_at	437.45	358.57	51.15	59.04	304.15	96.1	772.2	195.3	1231.4	1088.52	619.9	85.98	823.15	99.63	360.05	307.1	502.1	529.62	767.75
U82279_at	96.7	21.35	43.95	50.84	101.55	57.06	124.2	74.53	93.2	48.37	197.55	178.69	135.85	67.39	113.5	20.79	143.2	100.69	136.05
HG4683-HT5	2405.15	1390.24	1407.05	1625.71	3905.2	733.27	2324.6	443.36	3309.2	1291.04	2936.1	1038.03	3524.7	432.61	3759.4	509.68	3550.25	143.05	4068.05
D14497_at	28.55	16.48	16	18.53	167.8	41.44	270	363.88	102.8	84.15	62.55	42.07	81	86.55	81.1	9.9	326.55	416.98	28.2
L37360_s_at	272.05	92.28	666.15	769.69	947.1	1056.28	676.7	433.74	242.55	81.25	865.65	235.11	334.6	24.18	1306	1299.66	848.2	488.05	281.5
D13631_s_at	340.1	101.96	43.1	49.78	979	1284.81	483.75	301.16	64.7	16.83	709	627.35	302.05	319.68	77.65	52.4	191.5	205.34	284.65
M76559_at	327.75	322.65	307.25	355.04	390.4	206.9	547.25	392.23	101.1	20.22	445.55	171.9	368	120.49	382.25	71.49	145.3	144.53	87.4
X51953_at	179.75	1.77	63.85	73.75	194	75.38	284.5	266.3	125.6	99.14	560.55	609.88	132.2	52.18	167.8	28.14	224.35	53.67	321.8
L35240_at	3511.55	2596.28	1285.4	1485.07	2160.5	310.99	2360.5	477.86	1285	820.53	1394.3	1424.4	2918.65	413.45	2849.2	529.34	1587.8	326.82	1762.55
D63876_at	1402.55	164.97	1083.65	1252.07	899.3	709.23	1114.8	776.26	1299.8	448.31	2041.65	184.77	925.5	536.27	2189.65	494.34	1685.75	16.19	1047.8
L35475_at	1918.75	940.24	1037.5	1198.69	1227.8	272.38	1548.6	302.08	1423.2	31.11	2135.1	79.05	1542.2	240.84	2336.75	89.02	1764.35	487.13	1675.3
U28043_at	281.05	3.89	141.1	163.06	817.7	721.11	276.55	70.92	288.65	36.84	202.35	30.9	916.8	846.97	222.95	2.47	116.4	4.53	593.45
M74542_at	604.3	378.58	997.55	1152.51	1964.45	910.68	1159	541.5	1249.8	1387.63	1513.35	271.03	1480.75	254.49	1547.5	337.57	711.85	554.02	735.6
Z83803_at	139.05	23.41	54.4	62.79	120.7	23.05	86.65	41.93	121.1	0.14	118.05	15.06	90.15	8.41	99.85	24.96	112.25	20.86	130.65

U08191_s_at	321	76.51	443.3	512.23	1210.95	582.16	1188.05	591.78	599.35	452.9	1300.2	1300.51	512.15	348.11	569.05	451.35	797.7	721.11	686.25
Z37976_at	561.1	155.42	271.75	314.03	693.5	191.48	345.5	140.86	1285.45	102.6	1428.95	343.87	1392.35	2.62	1231.3	344.5	1306.75	123.67	610.5
D45248_at	3898.55	1242.46	2088.15	2412.58	2521.5	1382.68	3994.95	3221.37	4073.75	8.84	3527.9	1733.4	4090.3	969.73	3911.55	543.69	3405.25	961.31	3796.15
U43431_at	1086.2	152.74	370.7	428.37	1028.6	100.55	751.3	566.82	1356.95	506.36	1560.35	667.3	1755.4	54.45	859.1	207.32	1186.9	280.16	1404
M29610_at	635.55	310.91	369.95	427.45	660.05	476.24	243.45	308.09	444.4	267.57	310.25	323.78	416.7	110.45	219.75	213.05	579.65	363.95	347.15
L34657_at	185.35	5.16	47.95	55.37	178.75	115.05	303.3	68.17	172.05	68.94	107.5	37.19	230.9	147.08	158.8	45.54	290.7	296.7	258.35
AFFX-HSAC(	181684.16	62761.03	94679.85	109391.05	131294.8	10548.48	146993.3	5740.29	165667.95	42190.88	105076.84	3881.38	161707.2	10690.75	189101.84	69119.61	176076.45	50335.75	78284.2
X66894_s_at	715.95	134	167.25	193.25	1038.2	796.06	495.15	455.73	1201.35	265.66	490.1	10.18	466.95	82.8	529.35	324.77	994.5	113.99	675.7
D89377_at	1394.85	562.22	463.15	535.07	876.35	58.76	618.2	219.34	1100.65	1253.63	636.55	419.95	469.2	440.81	1075.2	968.03	1098.7	364.02	878.05
L08044_s_at	724.95	609.88	379.5	438.41	1175.25	1247.41	1336.1	366.71	604.85	172.46	1005.7	981.75	898.65	594.18	405.8	10.32	537.55	80.68	1086.95
X69878_at	979.5	185.97	982.9	1135.61	1374.85	862.32	887.9	347.19	522.25	109.25	1063.5	185.4	992.9	296.56	1073.15	496.74	1013.95	183.78	1069.65
U84573_at	877.7	69.44	521.35	602.38	805.8	7.5	740.6	251.31	852.65	121.13	627.75	68.09	568.95	363.81	778.35	227.19	511	1.7	346.9
U66052_at	1821.35	84.08	615.9	711.63	1402.5	541.22	2193.8	651.67	1704.3	398.95	1970.2	620.56	1480.25	1085.34	1671.5	551.83	1439.65	167.65	2182.9
HG4073-HT4	1446	556.92	1467.75	1695.85	1302.2	717.85	2102.3	803.41	2661.85	752.72	1838.6	332.48	3661.35	1898.79	3879.6	2931.66	3037.45	1278.94	1914.85
U35835_s_at	459.8	183.85	640.25	739.7	737.3	629.89	628.35	788.21	1071.65	317.14	374.5	156.13	592.9	656.34	386.45	291.54	875.9	590.86	626.6
M35252_at	185.8	142.69	201.25	232.57	247.65	147.86	125.65	110.38	294.9	299.81	115.95	36.13	295.35	162	490.6	19.09	230.9	228.68	218.65
U84487_at	149.7	62.93	80.95	93.55	222.8	53.03	168.3	113.99	831.45	627.27	291.65	88.46	236	49.64	198.25	71.77	194.8	12.3	229.35
M30607_s_at	119	57.42	405.75	468.74	263.2	221.47	127.2	4.95	232.15	151.39	138.95	70.5	235.9	174.66	93.95	16.05	102.9	24.75	242.8
S81221_at	88.45	33.45	76.9	88.81	199.05	45.33	207.05	65.55	111.35	45.47	349.8	166.03	103.3	6.22	305.65	195.52	123.8	63.07	195.25

M97639_at	182.8	42.99	143.6	165.89	419.5	98.85	265.35	95.67	383.8	250.6	968.9	208.17	294.45	122.68	254.85	74.6	562.45	460.11	904.3
U37143_at	229.75	235.68	255.05	294.65	314.65	84.92	116.35	39.81	531.45	42.5	465	528.07	93.15	27.08	307.5	285.53	144.85	36.98	199.75
HG4011-HT4	2697.95	1758.08	1234.55	1426.31	901.8	214.39	719	145.81	488.8	201.24	966.05	15.06	1344.6	600.62	1063.1	704.7	1657.15	943.78	560.9
Y11710_rna1	1219.2	604.01	167.9	194.03	991.9	70.57	887.8	371.23	1318.05	348.53	1387.2	532.45	1243.1	340.54	894.25	203.01	644.7	4.67	795.05
D86985_at	828.05	43.35	152.95	176.71	193.1	141.99	638.45	410.05	491	462.73	253.6	3.82	572.1	60.67	286.7	161.93	443.3	53.32	804.05
X14085_s_at	1335.8	941.16	913.75	1055.78	1020.55	660.23	1143.75	318.83	865.45	24.82	1995.3	466.41	1372.55	82.66	1020.15	925.11	865.05	705.76	1411.05
HG4541-HT4	9779.8	1136.89	4410.05	5095.34	8054.8	4717.39	8749.4	1957.98	7669.2	1134.62	8092.45	452.19	7956.15	2543.39	6686.15	2112.2	7611.7	3619.68	7427.5
U30999_at	975.15	199.47	541.3	625.37	1042.5	238.15	1297.25	635.05	1220.7	204.21	736.45	698.83	1369.8	146.51	1319.45	620.34	1043.5	504.59	1006.45
U17033_at	47.35	6.86	74.45	86.05	219.8	283.69	119.1	65.76	49	4.67	45.35	10.82	52.55	19.02	31.6	16.83	58.25	3.04	162.45
S69370_s_at	671.3	346.91	466.45	538.89	999.55	1072.47	718.5	853.62	736.05	409.34	218.15	68.38	198.1	1.56	337.6	219.63	119.45	73.19	237.6
U68488_at	658	392.02	334.5	386.5	1285.25	87.75	637.3	315.94	1283.25	156.34	978.65	32.88	1538.75	31.89	719.05	152.81	1508.65	12.37	727.8
S80905_f_at	259.35	115.61	193	223.02	256.95	182.65	1013.9	1159.51	164.2	1.41	252.6	38.47	142.4	71.28	785	794.08	355	62.93	267.2
Z49254_at	5338.2	3220.59	2435.95	2814.5	6678.2	624.38	6690.8	1008.9	4835	240.13	7209.55	3280.76	5729.55	2878.7	7874.9	996.31	5342.15	275.14	5401.75
HG1602-HT1	1102	249.61	555.6	641.91	1896.85	1143.04	930.2	708.66	911.5	234.05	836.95	869.39	1347.9	240.42	662.65	345.28	744.55	479.63	1049.85
M26041_s_at	866.7	10.32	457.85	528.99	1154.7	688.86	674.1	11.17	1173.45	608.61	1565.9	836.37	1291.6	442.79	716.5	395.13	618.95	83.09	713.3
U69611_at	153.25	200.46	89	102.81	521.2	721.96	37.95	22.27	272.05	214.47	165.95	142.62	144.8	147.22	57.2	28.99	456.6	157.54	231.9
X56465_at	68.9	50.06	130.5	150.76	48.15	44.76	177.75	25.53	47.45	20.72	197.6	206.19	54.7	35.36	36.9	2.69	39.35	9.12	272.4
L40392_at	276.2	240.56	268.8	310.56	405.2	253.85	278.2	189.79	574.8	549.85	661.25	25.24	935.15	58.62	882.65	57.35	862.35	179.53	324.65
U75276_s_at	2074.35	133.71	1227.35	1418.1	2400.8	160.09	2554.45	159.17	2509.8	105.36	3306.7	1683.06	3119.8	78.91	2088.75	1521.62	2815.1	89.66	3040.55
L78267_at	102	94.61	214.9	248.34	111.3	21.78	525.7	532.59	236.6	177.48	174.95	38.4	235.35	106.7	25.1	0.57	23.7	4.1	105.45

AB002365_at	121.1	42	308.05	355.89	239.65	219.56	84.2	93.06	709.45	75.31	113.6	29.7	229.05	169.21	424	533.16	51.8	4.38	338.55
HG4036-HT4	507.4	410.83	391.1	451.84	401.4	292.46	647.5	370.38	890.1	153.16	510	359.92	576.35	202.16	692.8	191.77	276.25	303.84	815.3
U14588_at	4863.8	375.61	3460.6	3998.26	4774.45	913.65	3268.3	2092.47	5800.6	183.85	4809	829.01	4661.9	2400.63	5307.7	425.82	6109.05	582.59	4325.15
L36861_at	549.4	573.04	196.35	226.91	185.55	33.59	461.15	535.63	197.3	17.11	506.1	499.5	189.9	13.44	141.65	89.17	474.85	476.52	291
M14016_at	2785.45	782.41	1099.35	1270.18	3142.7	232.78	3100.8	52.89	3422.15	266.65	3663.7	259.79	3471.6	459.05	2975.95	281.22	2989.2	170.55	3353.2
U38480_at	798.15	247.42	247.2	285.67	225.8	50.35	888.3	375.47	592.15	134.28	1595.75	1068.79	556.4	692.68	310.15	153.8	657.4	364.73	483.05
L19711_at	1560.95	57.06	781.15	902.48	1515.85	190	1959.4	99.42	1953.15	246.57	1421.4	648.7	2159.85	174.73	2633.1	1024.17	1565.3	210.15	1572.15
HG1327-HT1	46.1	13.15	18.45	21.28	578.25	421.93	310.1	200.68	352.85	186.75	285.45	35.43	512.4	196.72	169.8	209.16	93.45	4.6	246.35
X68561_at	344.9	137.74	390.1	450.71	1272.9	347.76	246.5	140.86	748.95	140.22	685.6	630.6	933.3	429.21	605.35	293.8	704.05	81.81	542.95
S79639_at	2476.25	512.86	1102.05	1273.29	2016.3	70.71	2529	152.31	2145.55	516.54	2470.8	58.41	1773.95	142.62	2110.4	1002.96	2621.6	296.56	2026.15
Y08136_at	136.2	76.08	44.05	50.84	259.15	280.08	64.7	19.94	52.5	11.74	342.25	108.82	305.45	56.07	328.6	45.11	100.9	80.61	183.8
U82671_cds2	1058.85	38.54	220.25	254.49	598.1	513.08	501	145.52	584.5	186.25	1190.55	608.89	1057.7	375.05	478.15	250.39	533.7	347.76	1430
U36341_rna1	5240.2	1111.57	2073.55	2395.75	4986.05	807.87	5242.75	517.25	4885.25	696.29	4848.7	885.3	5741	812.32	6259.25	1116.59	6805.5	1188.36	5818.55
L76937_rna1	353.7	55.44	76.85	88.74	229.3	210.15	1032	487.34	962.65	382.05	728.65	20.86	484.65	448.94	169.15	137.39	121.9	6.22	741.45
M91463_rna1	771.6	18.38	699.7	808.36	1421.6	1138.72	1087.8	1102.66	1234.95	231.58	1082.45	482.18	925.95	582.02	1105.2	51.19	1044.35	440.32	1131.1
X55079_rna1	3404	69.44	1268.3	1465.41	3517.6	65.9	3446.95	1578.05	1676.8	620.56	1817.75	603.09	2094.65	237.94	2157.6	23.76	1970.25	1415.42	3035.1
HG2815-HT2	6130	1611.35	2745.6	3172.22	5481.5	1782.62	5997.85	721.04	4534.85	2167.07	2857	84.99	3775.25	65.41	9265.05	6389.49	5126.5	2374.61	3780.15
S76617_at	142.5	46.39	111.85	129.19	106.9	1.27	178.55	130.6	113.9	6.08	119.25	30.33	290.45	3.89	247.9	107.48	243.3	47.66	168.3

Z34975_at	578.3	664.4	78.15	90.3	599.65	537.19	334.95	2.47	300.7	185.4	154.15	40.23	446.05	442.58	400.85	354.9	180	93.06	432.45
M90391_s_at	77.5	31.68	260.8	301.37	531.75	165.82	107.1	50.2	491.4	514.77	146.05	109.25	176	22.91	211.4	155.85	126.9	1.27	234.2
D16217_at	5429.55	1222.66	3239.35	3742.65	5959.35	1168.07	4627.65	1411.46	5162.8	1752.63	6198.5	1007.06	5201.15	1484.15	5214.4	263.04	5222.2	1677.96	5783.55
M11353_at	4874	3134.32	1942.2	2243.93	6934	1026.86	4716.6	1822.21	7394.75	1415.42	6473.85	2810.68	5573	984.29	5498.15	755.97	5055.3	1893.91	6844.35
D25328_at	4185.5	1099.27	2811.95	3248.8	4108.7	1057.12	4937.9	2125	4505.7	830.99	3438.8	430.77	2975.85	998.51	3515.85	352.63	3713.65	274.85	5413
M55682_s_at	215.35	132.3	137.35	158.75	155	27.58	217	100.69	180.45	8.84	678.25	710.43	534.4	338.85	630.25	602.95	413.9	353.55	264.05
AB000381_s_	132.8	24.47	72.45	83.65	678.25	508.9	338.6	391.03	505.2	513.93	274.4	158.96	235.4	175.65	124.1	80.61	160.05	163.98	167
HG2600-HT2	73.85	8.27	38.4	44.41	81.8	14.85	160.1	122.05	148.45	81.25	81.85	2.47	92.9	20.79	73.45	25.81	121.25	28.78	119.75
D13720_s_at	168.05	12.09	24.25	28.07	349.4	414.93	547	538.67	615.6	443.5	490.7	385.94	329.05	200.04	274.2	319.05	245.45	153.37	323.1
X77584_at	19060.65	3530.37	8838.75	10212.11	22072.45	2062.28	17340.85	4916.02	17142.85	2139.92	12713.5	785.74	12967.75	2606.89	14419.75	505.93	15468.85	3035.54	15789.55
U76369_at	1180.85	181.8	340.25	393.08	400.45	256.61	457.75	289.7	487.65	447.39	1311.05	826.96	182.95	15.63	264	136.33	717.8	389.33	204.8
M29877_at	925.1	176.21	418.75	483.87	1372	330.08	1025.95	469.45	1304.7	405.03	987.2	56.29	743.85	79.41	585.1	115.82	604.8	383.82	811.75
L07493_at	556.15	33.02	598.1	690.98	1112.15	653.86	477	483.1	1846.15	1014.2	563.3	387.07	710.9	610.37	1417.15	234.97	665.1	429.64	966.35
D87002_cds2	109.5	11.74	111.4	128.69	220.8	124.31	298.35	326.33	246	91.22	354.55	154.36	202.3	76.65	290.65	3.61	197.2	67.03	321.65
L19593_at	401.95	192.55	133.8	154.57	187.3	165.6	560.25	716.09	520.55	307.38	240.1	225.57	241.2	1.27	595.85	151.96	303.6	134.77	373.15
M60047_at	4220.35	2011.79	1818.55	2101.17	3807.6	287.79	3651.75	68.52	10186.8	2938.59	6550.15	2089.15	14133.35	10134.33	20366.15	14298.76	12569.5	10327.72	4647.55
U66618_at	1744.8	641.35	869.6	1004.66	1901.7	379.01	2068.15	938.4	1689.9	55.15	1705.45	15.34	2184.4	532.03	2966.05	682.85	1620.15	480.34	1701.2
U06863_at	2333.8	274.5	849.2	981.18	1875.8	204.92	2186.6	514.77	1759.9	25.46	1876.05	197.49	2124.4	47.94	1728.75	6.86	2377.9	157.4	2459.3
L41349_at	253.5	55.86	16.85	19.45	100.4	27.72	39.45	20.29	174.15	181.66	125.75	30.48	41.75	4.17	58.9	46.39	20.55	6.58	50.4

M93056_at	145.8	14.14	389.35	449.79	181.45	19.73	668.1	343.94	434.95	333.68	595.3	62.93	881.2	983.73	160.75	54.09	478.25	413.73	569.3
M89955_at	345.35	41.93	203.95	235.68	715.8	647.71	424.4	201.53	202.05	12.23	253.9	54.87	479.6	332.34	1160.75	275.42	928.55	197.49	437.8
U49785_at	2845.95	1095.8	1482.75	1713.11	3350.8	130.81	3421.35	167.09	3951.05	276.13	3304.6	830.14	4011.85	1273.85	3374.2	466.12	3149.15	188.59	3566.55
X77748_at	371.4	431.76	97.1	112.15	131.2	74.53	45.9	28.14	612.85	815.93	158.2	132.65	90.15	27.37	635.9	794.08	177.8	108.19	122.55
M22960_at	2785.35	577.64	632.9	731.29	2014.25	377.24	2049.6	227.41	1411.85	175.29	2295.35	413.45	2034.95	645.8	2240.95	458.13	2508.5	597.65	2224.5
U90902_at	114	46.95	68.15	78.7	362.6	366.56	364.85	373	167.75	95.39	256.55	0.78	312.8	25.03	506.3	435.29	199.15	178.4	205.3
U95090_at	293.85	137.67	130.5	150.76	486.65	185.62	272.75	9.55	199.7	5.23	510.75	224.37	700.45	541.71	225.7	90.65	268.1	192.47	769.1
X78669_at	255.7	90.37	271.65	313.88	275.55	268.63	271.95	273.86	381.1	149.62	583.5	612.35	466.6	102.95	53.8	38.33	411.45	52.96	129.2
U03056_at	133.6	32.67	234.25	270.61	366	361.19	119.8	37.76	550.9	628.48	464.25	440.32	325.75	386.43	434.35	491.93	180.65	175.57	733.95
L03427_at	177.75	146.44	170.05	196.5	321.35	380.92	140.55	36.84	80.7	61.38	146.9	118.79	57.1	0.99	120	79.05	189.4	133.78	189.55
X60382_rna1	80.3	81.03	180.7	208.74	162.2	88.39	34.8	29.84	536.6	282.28	465.45	483.45	469.15	141.49	84.45	21.28	225.2	8.2	263.4
U58681_at	866.5	894.91	158.05	182.65	274.25	133.57	1037.45	343.44	191.85	38.25	1110.55	65.69	876.65	790.33	496.7	428.79	341.1	165.46	415
M15353_at	118.45	133.29	13	14.99	59.05	51.12	384.75	374.41	278.45	79.41	184.1	41.15	117.9	118.65	384.65	93.83	245.9	181.44	198.55
M82962_at	91.35	12.8	81.35	93.97	82	31.11	176.25	135.27	187.2	28.43	154.4	29.42	188.1	10.18	226.5	51.34	104.5	2.69	140.65
X66785_at	2092.25	109.81	1423.35	1644.52	1903	874.83	1896.55	507.49	2354.8	47.52	2915.4	918.25	2957.15	527.57	2713.25	467.19	2081.7	208.03	2508.8
X75304_at	1080	464.14	658.3	760.56	969.7	458.49	1036.55	940.95	1258.25	228.32	762.75	459.55	1469.65	493.63	896.8	736.95	1095.55	745.08	559.95
D29640_s_at	697.2	191.48	563.85	651.46	924.1	277.19	568.35	134.85	767.35	452.48	607.85	87.89	381	459.05	1026.2	56	1012.4	31.25	1231.55
L04510_at	377.4	54.45	253.85	293.24	692.05	427.87	219.25	240.2	511.25	187.31	180.2	175.36	344.1	49.21	281.35	316.29	286.45	115.89	250.1
X53961_at	233.35	82.66	125	144.39	247.65	20.15	317.05	226.91	237.35	164.26	474.4	208.03	162.1	12.87	316.95	18.03	173.75	18.17	188.4

M96789_at	721.6	124.45	896	1035.2	1355.5	511.38	726	375.05	326.9	132.94	1333.9	382.54	691.1	419.03	889.3	528.07	710.7	595.81	1139.3
D21260_at	5125.75	1283.75	3338.2	3856.84	4015.45	1559.81	3014.6	1709.78	5504.4	1883.59	4895.8	2188.35	5292.15	2482.72	4370.8	57.56	4404	552.25	5554.15
L34600_at	493.95	100.76	338.75	391.38	765.3	48.65	601.5	143.97	1073.4	202.37	994.2	503.74	910.8	113	705.15	136.97	697.8	238.72	1160.15
U73167_cds7	154.6	67.32	327.8	378.73	310.45	125.37	118.1	22.49	144.5	31.96	210.7	156.69	424.55	264.95	414.75	157.19	187.55	125.37	255.55
X78549_at	1874.7	1252.57	793	916.27	1418.95	437.2	2165.9	350.02	1475.7	961.67	1046.95	142.34	2156.25	214.89	1159.3	450.29	929.2	461.74	1051.35
X58964_at	187.9	65.62	88.85	102.6	348.2	202.94	143.75	52.82	147.4	8.06	247.6	207.18	131.05	51.41	205.75	26.8	203.9	134.07	204.6
L43575_s_at	934.15	139.94	695.5	803.56	622.55	693.74	786.8	74.25	1285.2	149.62	470.1	574.88	384.8	385.09	644.9	245.51	718.45	144.04	611.6
X16316_at	2490.35	96.38	1190.95	1375.96	2338.75	379.22	2185.9	168.15	2068.9	131.24	2275.25	250.1	2626.1	770.6	1870.8	330.22	2371.15	451.49	2430.5
M76424_at	1939.95	1015.19	1139.25	1316.28	3052.1	2357.21	1657.85	1375.25	1867.15	554.16	3613.35	710.43	2219.8	119.08	1772.4	1349.02	1516.7	299.53	3427.55
L35035_at	414.75	391.38	393.4	454.53	281.7	106.49	223.2	89.66	628.4	91.36	841.2	504.59	746.05	823.85	561.8	152.59	537.1	18.38	342.5
X90872_at	5470.1	549.99	2870.45	3316.4	4846.95	2810.25	3685.35	603.37	4912.9	1564.4	4963.35	137.67	4655.7	410.12	4419.2	127.99	5574.5	668.78	5409.9
D90084_at	3100.8	169.14	1737.65	2007.69	3065.5	130.11	3105.45	682.29	3373.4	708.38	2666.25	177.55	2658.2	890.53	3133.35	608.32	2760.3	552.11	2471.4
M76482_at	2353.95	186.89	1918.25	2216.28	3367.95	856.09	2436.8	243.24	2757.25	2698.67	2839.55	965.55	3307.7	635.12	2559.45	1082.79	1678	582.37	2833.65
S78234_at	371.25	183.35	426.4	492.71	485.65	460.96	172.85	2.33	572.55	235.11	254	166.17	251.45	218.99	304.45	289.98	363.45	275.42	350.2
D26528_at	410.1	219.49	475.1	548.86	986.25	594.18	946.55	523.61	862.35	205.27	1040	757.03	1009.55	34.72	936.85	4.6	772.3	438.97	799.1
X79780_at	1076.05	1192.82	108.65	125.51	684.15	699.26	323.65	328.31	311.6	182.43	156	52.18	355.1	221.32	266.65	28.07	312.4	323.43	334.55
L14269_at	328.75	366.07	31.25	36.13	74.75	9.97	804.85	958.06	332.05	353.77	556.9	219.77	203.95	163.98	464.7	511.38	101.35	98.22	592.15
Y07829_xpt3	338.75	359.71	207.3	239.57	1127.65	248.41	357.35	238.65	271.4	27.86	852.95	733.62	468.9	93.34	550.1	443.92	538.25	508.2	346.6

AC002486_at	695.15	89.59	132.2	152.74	743.7	431.76	1337.05	1283.33	591.55	26.38	684.3	242.82	611.75	420.52	500.65	524.46	423.85	128.48	694.4
X95425_s_at	56.8	2.97	248.55	287.16	91.2	89.1	83.75	44.19	143.4	111.86	80.85	9.55	171.1	81.18	128.7	16.97	73.8	41.01	334.8
X63717_at	749.1	320.18	250.55	289.42	806.95	377.81	295.05	133.15	945.3	154.43	719.55	187.03	1095.25	69.79	675.4	275.35	755	23.62	271.6
L13977_at	1643.5	141.42	1001.15	1156.76	2184.1	604.29	1115.35	113.63	1224.05	46.88	1054.75	603.94	945	408.99	642.55	510.04	1038.1	322.72	796.5
M96684_at	277.45	150.83	124	143.26	273.5	83.58	249.25	66.96	233.2	166.17	191.2	113.84	206.6	5.09	216.25	84.36	211.9	133.64	153.45
M11058_at	818.65	113.49	323.6	373.92	626.1	510.67	674.65	329.72	907.8	677.97	566.75	29.77	792.75	377.52	567.2	112.01	1378.15	784.25	990.5
X79536_at	583	284.26	411.35	475.25	138.6	155	637.85	19.59	493.05	541.15	652.1	37.76	264.9	171.83	820.85	14.07	729.1	517.18	499.15
U44975_at	1453.2	1283.82	1518.75	1754.69	1150.2	669.63	1827.25	405.38	2119.1	1430.48	1315.05	524.46	1139.8	152.31	2432.4	1478.56	1986.5	1199.68	1440.25
X03656_rna1	2102.9	193.32	1013.15	1170.62	1712.55	51.27	1942.5	165.32	1949.55	283.76	3135.75	771.81	2033.35	1312.32	2463.15	4.17	1805.6	434.87	2628.8
D32050_at	5757.95	640.71	5119.05	5914.45	7271.2	656.48	6058.15	383.89	6672.3	704.28	7254.9	205.06	5666.4	700.6	5877.7	3537.94	5731.2	3264.43	6186.1
U36759_s_at	2110.1	387.07	1342.6	1551.25	1842.2	1693.52	1648.7	739.21	2798.85	1467.6	2512.5	444.91	3759.95	663.2	2309.8	984.58	2392.15	301.86	2224.7
U40271_s_at	1394.5	511.66	752.6	869.6	1294.85	1278.94	852.65	390.96	430.05	34.44	813.85	373.28	960.25	641.56	446.7	165.75	1405.5	252.44	841.55
X69550_at	12704.05	4696.39	5310.15	6135.21	7646.95	3291.37	8861.85	269.34	7530	1153.43	9050.1	886.15	12257.75	848.17	10010.8	2346.04	13949.8	1052.88	11549.5
K03515_at	11591.35	6601.76	4897.9	5658.98	11837.05	2071.75	12148.6	1687.58	9931	4694.48	10762.85	1531.1	11905.25	3036.53	11887.4	2099.97	13484.85	349.95	13419.15
U70370_at	131.4	16.97	77.25	89.31	113.6	6.36	216.9	135.06	158.8	121.06	334.1	333.9	134.1	81.03	156.85	69.08	177.35	53.39	115.8
S78569_at	177.3	178.05	11.4	13.15	477.15	284.61	55.05	54.24	293.8	126.57	252.75	325.06	158.1	139.58	71.35	64.7	207	33.09	43.95
J03779_at	903.05	451.49	635.05	733.76	786.25	660.51	450.95	167.51	1100.3	1357.08	862.4	129.26	1005.5	462.17	892.85	972.91	470.1	122.05	955.55
D86981_at	36.5	14	26.65	30.76	51.4	38.89	186.8	17.82	217.8	267	272.95	101.47	30.1	0.14	300.5	298.96	232.55	296.07	97.15
L40387_at	207.75	2.47	78.35	90.58	527.25	234.26	394.8	333.33	1258.5	224.29	425.85	170.2	485.15	389.97	291.7	44.26	818.25	827.81	1173.35
X53683_at	395.95	66.11	1069.95	1236.23	981.55	1003.6	816.25	831.35	1710.4	271.67	570.45	298.33	1409	1073.67	458.95	297.2	1345.45	1013.5	793

X80818_at	2284.25	567.31	1125.6	1300.51	2899.7	265.59	3015.85	1113.06	2209.25	583.86	2774.15	735.04	2278.15	49.57	2819.25	915.49	2256.45	529.98	2077.1
X06956_at	753.8	725.77	622.25	718.92	122.6	40.31	492.05	509.61	676.7	860.69	606.95	505.37	597.95	718.77	456.3	93.76	1211.85	6.43	378.2
J03600_at	203.45	43.63	96.35	111.37	252.6	37.05	827.25	946.89	271.4	71.42	192.1	127.42	173.85	111.51	157.25	54.38	169.65	91.99	236.95
X90999_at	1592.35	523.61	869.35	1004.45	1349.6	1210	1830.05	64.13	1078.2	663.97	963.75	551.61	1612.55	600.26	1277.55	184.63	1240.9	435.58	825.55
U93091_at	21.9	10.47	12.7	14.71	217.6	203.79	55.05	3.04	55.35	19.02	88.6	11.03	25.25	7.14	31.65	9.12	150.3	20.79	103.5
U90912_at	372.05	3.61	372.3	430.2	382.8	113.56	105.65	34.29	658.9	339.84	209.25	238.37	329.35	39.53	348.4	367.41	416.1	62.23	897.95
HG3162-HT3	1218.2	147.08	527.35	609.31	1178.25	354.33	1200.75	636.89	1453.05	355.6	1616.3	738.22	1471.35	244.87	844.8	580.96	860.35	411.18	1054.95
M65134_s_at	343.9	280.16	119.45	137.96	454.55	500.7	155.2	15.27	281.75	350.09	312.05	245.86	532.45	126.08	295.4	151.46	417.85	331.14	373.3
U32645_at	245.2	225.71	508.25	587.25	292.1	213.69	739	369.68	1131.65	746.63	180.65	129.19	383.15	119.43	865.7	151.75	495.65	312.33	314.9
HG732-HT73	828.45	3.89	362.95	419.39	1336.9	482.39	550	488.05	969.55	164.4	673.45	63.85	879.25	137.11	1040.55	76.01	641.75	307.66	884.55
L22454_at	994.05	1107.12	221.4	255.83	1215.85	1098.21	1105.2	203.36	1148.5	209.73	668.25	354.61	1248.45	1173.44	601.15	90.58	630.25	534.22	469.15
L13698_at	305.6	126.29	384.5	444.2	329.4	303.63	541.75	403.55	406.3	109.18	615.05	307.66	698.5	100.13	283.4	280.16	488.45	26.09	605.25
X04347_s_at	10241.7	3812.86	4968	5739.87	11071.1	736.38	10931.5	1349.16	15705.55	321.24	11703.25	1189	13616.75	342.17	14326.7	1210	14739.25	2984.06	13106.6
U24105_at	3556.1	553.95	1779.45	2055.91	3740.85	715.24	4377.2	386.93	3434.5	201.95	3889.25	836.44	3090.05	1311.75	3922.5	884.45	4697.6	2023.88	3884.05
HG3299-HT3	151.85	54.52	91.25	105.43	172.5	4.1	237	64.49	120.45	17.18	304.15	133.01	161.95	2.62	327.15	175.15	241.9	41.44	186.35
D38462_at	59.6	47.38	97.1	112.15	93.4	42.99	548.95	282.91	207.9	162.92	73.6	47.52	61.05	7.42	157.75	86.48	193.95	143.33	176.6
X60787_s_at	209.85	8.98	368.9	426.24	251.7	29.13	151.4	1.56	537.3	360.2	484.85	460.96	210.3	26.73	237.8	59.26	289.7	44.69	292.25
D87469_at	1760.7	151.89	487.3	563	910.15	1033.01	1844.8	428.51	2226.95	474.96	2330.9	298.12	1458.95	226.34	2419.95	238.79	2126.05	650.89	1826.2
X61072_at	802.25	232.14	426.35	492.64	841.65	175.72	1328.2	1010.17	947.45	284.61	1234.45	477.08	1496.95	38.68	901.3	145.24	783.95	548.93	914.1
X56687_s_at	461.85	335.95	132.3	152.88	228.8	131.95	226.35	37.69	837.15	115.75	786.2	653.08	324.75	189.29	311.4	17.39	643.3	385.09	420.6
X67337_at	48.3	26.16	191.35	221.11	53.3	6.22	272.25	220.83	120.6	16.55	372.2	136.75	50.8	24.04	93.7	81.03	123.35	126.36	46.6
L76200_at	7665.5	2856.71	4561.8	5270.63	11847.2	3068.84	11368.9	724.36	10156.2	359.78	9543.25	789.63	8986.2	1143.39	10793.45	510.74	10481.2	992.35	8696.2

M98833_at	32.2	18.53	164	189.5	475.1	438.55	165.5	35.78	49.95	33.02	176.3	49.64	90.95	93.97	277.85	271.88	351.05	27.08	105
AFFX-BioC-5	53939.75	8279.87	27241.6	31474.31	276739	296362.03	280747.94	329889.44	126036.95	99184.52	191653.11	226727.72	87610.8	71160.4	98926.25	6948.38	67172.85	3897.22	88029.5
HG3477-HT3	157.4	86.27	80.4	92.91	161.05	48.86	140.75	94.68	138.8	27.44	169.1	73.96	112.35	6.86	135.55	38.54	110.15	28.35	116.75
D11428_at	534.45	342.17	543.95	628.41	775.9	201.95	597.65	210.22	1056.3	194.03	951.95	936.28	921.7	733.13	1044.9	650.82	520.4	298.26	494.05
U10991_at	1234.75	314.73	780.65	901.91	779.45	230.73	1316.35	709.16	739	641.06	1232.6	85.42	869.4	319.33	1300.8	174.66	1071.75	339.62	457.15
HG3132-HT3	330.3	257.67	32.25	37.26	253.35	30.19	170.1	151.18	135.45	14.21	361.25	344.01	224.95	15.77	192.6	19.37	550.65	546.38	113.95
U73379_at	3648.5	1011.73	2166.05	2502.66	2962.35	980.97	3322.7	390.89	4951	793.09	4975.1	279.73	5578.65	623.31	4513.65	117.03	4286.95	868.54	3809.65
M34057_at	127.1	28.43	218.7	252.72	607.05	182.65	410.85	465.06	932.4	38.47	225.1	139.02	558	603.44	463	592.84	1072.4	179.46	939.7
D21089_at	2210.55	188.16	1865	2154.84	2965.35	169.49	1721.4	195.44	3072.25	158.46	2620.3	1645.15	3197	36.49	2758.8	871.58	2332.6	99.42	3772
M84424_at	127.35	66.96	194.9	225.14	143.75	3.18	63.75	17.47	366.1	229.67	495.35	169.49	328.85	393.79	169.45	189.86	162.95	193.39	188.45
X54232_at	9138.7	2129.66	4902.8	5664.63	9022.95	1051.11	9892.6	1884.72	9731.35	966.69	8313.35	1605.77	8688.95	10.11	8696.85	516.4	12216.05	612	9391.95
J05252_s_at	364	144.39	38.6	44.55	785.9	315.09	460.7	426.53	169.75	50.98	517.9	11.88	501.6	210.86	661	237.73	937.15	364.8	478.75
M93119_at	87.3	54.31	24.6	28.43	266.7	141.99	670.15	792.31	301.9	269.97	110.1	117.8	202.2	196.29	280.15	329.16	424.1	64.77	105.15
U68723_at	116.25	13.65	49.9	57.7	178.05	130.18	110	46.24	141.35	32.88	236.15	118.44	206	28.85	211.1	29.27	97.25	14.5	108.3
HG2147-HT2	4333.45	2762.88	2164.45	2500.82	2352.35	277.68	3921.15	1126.92	3373.15	703.64	2459.15	1423.34	3580.6	316.36	3158.25	686.11	3850.5	1416.19	2708.1
X05299_at	502.9	5.94	284.05	328.17	490.25	222.67	511.5	2.55	541.75	50.84	589	49.36	731.85	210.79	1069.95	808.15	3182.8	810.49	1963.05
M16707_rna1	819.15	390.82	411.75	475.67	499.45	370.59	367.15	382.47	770.65	127.07	729.2	887.56	1155.5	145.81	744.2	985.14	162.45	88.46	1034.05
M58297_at	825.45	685.26	137.9	159.38	840.45	308.93	855.5	806.38	982.45	858.22	426.3	141.99	645.25	446.11	318.65	20.01	390.85	180.95	337
M33600_f_at	169.7	44.83	64.05	74.03	184.9	113.99	157.5	29.27	249	106.91	406.7	270.26	162.7	26.73	188.45	46.6	181.35	62.44	257.85

M21984_at	1804.45	85.49	1122.7	1297.12	2396.55	654.14	1024	85.28	1171	395.84	2216.3	738.22	1916.15	270.47	1520.9	111.02	1577.45	195.23	861.35
M38591_at	23400.2	7428.86	8517.4	9840.81	24366.7	1085.27	22479.15	10984.13	26257.85	1311.05	19813	4209.13	23984.45	1458.97	23644.85	4533.47	26128.45	1117.16	27208.85
D32202_at	843.65	76.3	324.95	375.4	1800.8	1169.7	1194.2	900.43	414.9	290.76	972.1	81.88	1018	184.98	688.8	172.53	506.5	287.93	852.05
S76638_at	1237.5	324.28	773.75	894	432.2	223.3	833.2	91.92	775.75	232.14	1394.85	60.46	1164.25	444.7	1485.75	601.68	1155	58.27	816.1
HG4319-HT4	48179.6	8401.98	31186.2	36031.9	54064.2	3717.26	51856.45	42437.37	71695.95	23735.24	57555	20807.32	55852	9703.48	56190.75	8273.08	60181.45	3190.68	66367.09
D00763_at	2568.25	433.67	1586.5	1832.96	3423.85	829.51	2959.4	1163.05	3348.5	53.46	3464.4	1542.62	2930.45	251.66	3084.55	63.14	3254.95	489.81	3608.1
Y08409_at	200.65	155.63	121.55	140.5	140.25	118.72	354.4	230.94	320	201.53	506.65	624.16	186.3	145.24	108.4	16.69	157.5	41.01	160.9
U90907_at	1229.2	546.03	580.6	670.76	529.25	239.64	1522.9	548.01	1266.2	409.41	976.15	317	1369.55	46.88	1118.6	180.74	1299.4	428.37	861.75
U20240_at	912.1	165.75	427.2	493.56	1134.3	197.14	1084.65	30.48	1024.6	570.35	1577.45	223.38	984.5	403.05	1072.4	532.59	953.85	114.9	941.9
U03905_at	501	164.47	55	63.5	79.1	12.16	208	176.07	196.35	122.26	201.45	114.9	124.5	2.97	253.4	298.82	56.45	6.43	142.8
M26692_s_at	431	238.44	502.3	580.39	961.65	862.18	660.9	283.27	1132.15	331.85	258.3	183.14	1192.55	221.11	660.85	125.23	650	128.69	598.95
X81882_at	706.7	271.39	232.85	269.05	325.1	254.7	336.1	296.56	478.55	1.34	231	198.7	106.55	59.33	672.8	296.84	307.3	15.7	561.5
X77548_at	651.2	38.89	237.8	274.78	142.45	116.04	936.35	401.57	747.25	445.12	604.05	103.59	803.2	85.84	234.35	63.99	398.35	267.07	725.8
L09753_at	50.95	4.88	77.2	89.24	113.4	70.71	53.85	36.42	112.3	32.67	78.25	16.19	49.85	20.15	43.7	0.85	41	12.87	315.15
X13461_s_at	14236.4	6351.94	4383.4	5064.44	7475.45	1873.2	9860.5	2176.19	13284.55	7358.79	10484	1194.44	22330.75	6322.74	12196.05	1611.99	25095.05	13410.63	13619.55
U09002_at	718.1	368.26	267.95	309.64	837.6	507.7	470	9.33	733.25	158.89	580.45	201.74	381.05	52.54	783.7	194.45	561.5	218.78	320.9
D26600_at	9527.55	3659.91	5148.25	5948.11	14267.9	250.03	11659.7	4473.44	16325.9	4277.15	7856.75	3177.53	12283.55	5724.24	12057.3	1955.29	10184.5	602.6	10159.8
M54915_s_at	3489.65	126.64	2419.25	2795.12	3021.5	780.5	4009.55	114.9	3355.85	926.8	3602.05	403.12	2683.1	949.5	3217.75	2768.82	3458.6	278.74	4399.35
HG3527-HT3	6311.6	470.51	2795.4	3229.78	5334.3	170.98	4556.7	1062.07	4001.5	1304.47	5218.1	2218.34	4868.2	2333.59	4629.6	257.1	5823.15	973.33	5117.95
X63575_s_at	341.1	14.99	271.7	313.96	273.05	67.95	840.5	898.87	312.2	177.63	1049.6	962.23	387.7	291.05	441.3	437.98	118.5	101.26	152.9

Z33642_at	874.2	105.92	490.85	567.17	1380.95	5.87	179.9	24.32	335.7	26.59	537.35	19.16	750.3	641.06	969.4	93.48	762.3	184.98	516.1
M38690_at	1882.95	381.91	952.7	1100.68	1814.2	386.5	2275.75	96.24	2335.75	604.65	2021.4	113	2757.45	359.28	2311.1	1491.71	2806.85	1213.32	2476.25
HG855-HT85	206.05	205.13	256.6	296.42	568.7	564.84	397.15	485.29	543.8	282.98	588.15	294.51	410.9	285.25	463.7	547.02	188.3	97.72	382.65
U22970_rna1	1005.7	791.68	162.15	187.31	2062.1	514.35	304.95	88.74	505.05	488.26	344.05	143.61	373.4	117.24	287.5	41.44	552.4	493.42	1116.3
Z29090_at	47.3	22.91	12.2	14.14	69.55	4.03	109.55	90.16	419.55	412.17	460.7	370.52	92.85	47.16	97.2	36.63	75.6	53.32	292.5
X04828_at	4002.85	2439.45	1005.95	1162.27	1641.65	1146.72	2718.3	620.98	1619.25	1768.4	1317.55	827.81	2859.3	408.14	3446.3	853.9	3298.7	1092.34	2028.05
M35296_at	1125.05	88.74	711.2	821.66	1815.45	635.19	1668.85	865.15	1009	921.08	1678.85	247.84	1397.55	180.52	2201.4	983.3	1763.25	463.23	1964
M27396_s_at	5745.8	936.63	3796.9	4386.89	4320.95	1825.68	5836.4	2180.43	4689.1	56.14	4931.35	368.05	2305.6	6.93	2232.05	2709	3120.45	3667.13	4406.75
Y07596_at	265.65	64.28	324.85	375.26	271.3	328.95	158.55	86.62	247.35	126.36	333.55	132.44	192.1	138.59	152.4	139.72	163.45	128.34	132.95
U90651_at	478.4	188.23	280.3	323.85	583.1	643.75	859.6	504.73	741.7	586.47	444.6	407.01	502.25	10.54	654.55	459.83	885.55	98.22	876.05
Z35491_at	499.2	184.98	397.5	459.2	844.85	447.81	465.9	145.1	479.65	100.2	342.95	47.87	443.6	296.56	700.8	416.63	362.65	237.66	824.8
U24056_s_at	1052.85	844.5	1116.9	1290.47	1748.6	471.78	1155.35	985.35	1768.25	258.16	1025.45	351.93	1852.25	561.8	776.6	411.11	938.85	127.92	1068.75
L36642_at	224.65	73.75	32.1	37.05	50.35	36.27	305.6	409.27	126.3	21.92	262.1	100.69	165.5	175.79	158.95	193.82	149.75	174.02	56.05
S67325_at	921.25	104.02	712.6	823.36	1549.7	825.62	1710.65	33.59	881.1	695.37	1287.35	373.28	907.3	257.67	676.3	297.55	1081.3	844.71	1299.1
AFFX-BioDn-	38811.5	738.93	17878.75	20656.78	188774.06	192632.52	204896.45	236484.58	90936.1	65514.29	149265.25	182504.33	56959.4	62418.16	67773.3	1124.73	52112.2	1438.11	67334.15
M31520_rna1	9344.25	3884.21	3531.7	4080.43	9682.05	267.36	8209.65	4471.96	11021.15	371.58	7415.65	2339.6	6599.65	644.67	7404.35	947.88	9326.85	242.61	8893.15
U90716_at	1002.25	655.42	656.5	758.44	700.25	655.42	608.85	70.64	1006.35	113.49	496.15	419.67	941.7	0.99	1050.4	377.03	845.9	394.71	579.1

M91556_s_at	307.5	290.34	12.9	14.85	66.9	29.98	61.95	49.43	67.15	20.01	517.8	665.25	101	56.14	247.85	254.77	204.5	206.05	229.9
M83941_at	130.65	86.05	14.2	16.4	84.55	9.26	382.2	439.11	61.65	31.89	443.1	316.64	60.6	38.89	71.85	40.52	47.8	2.97	52.55
U50950_at	302.2	156.84	67.8	78.35	283.4	319.75	516.25	155.78	408.7	97.3	325.35	44.19	366.05	130.04	44.15	17.61	242.6	38.04	183.9
X54938_at	2759.3	895.2	1346.9	1556.2	2214.35	1292.38	2496.7	259.37	2848.2	1097.29	2516.2	121.76	3588.45	517.81	3260.95	889.47	2579.35	218.28	2903.75
D55638_at	203.45	26.23	404.65	467.47	265.65	70.92	572.9	429.64	388.6	221.32	416.5	205.34	235.05	75.31	332.65	276.41	215.4	146.37	744.85
HG2797-HT2	52.45	22.42	9.5	11.03	30.95	18.03	59.35	63.43	40.55	38.68	78.1	85.84	64.65	12.8	85.7	0.85	96.6	65.2	37.2
D49396_at	479.1	162.07	595.5	688.01	365.6	283.69	426.7	475.74	966.35	409.2	654.8	136.75	1034.25	497.03	741.4	529.76	619.05	224.79	831.8
U65404_at	900.75	951.13	102.85	118.86	1117.05	1243.16	602.45	404.25	1027.35	1075.01	315.95	92.7	332.9	123.46	365.95	173.88	697.05	583.43	1419.25
D12625_at	592.05	188.02	687.65	794.43	2430.25	2548.63	2426.25	3112.47	1424.4	453.54	738.75	466.05	988.05	357.3	4508.5	4647.11	866.65	198.63	1390.7
AF005887_at	923.65	103.17	190	219.49	773.45	77.29	2044.5	1207.17	736.95	684.83	1265.15	391.53	960.95	169.35	501.8	282.98	1247.6	195.3	950.65
U87223_at	709.5	379.15	131.85	152.38	490.55	69.08	754.25	725.7	521.65	41.51	429.4	233.77	207.5	64.63	240.5	66.75	241.45	11.38	570.8
M36067_at	325	275.06	96	110.87	475.75	381.2	244.65	20.44	453.3	144.11	381.05	292.25	332.15	83.51	739.1	594.39	640.8	556.92	442
U79260_at	1604.5	201.81	1087.15	1256.03	1623.2	85.98	731.3	302.64	1745.95	553.03	1600.3	76.65	1306.2	613.91	1442.7	8.63	1629.5	363.03	1857.5
L78833_cds1	170.4	205.2	224.2	259.08	215.9	116.53	218.95	98.36	435.15	409.49	620.95	621.62	481.8	198.7	331.05	138.95	132.55	97.51	250.85
U07424_at	2314.45	691.2	1899.9	2195.14	3264.35	321.66	3040.4	650.82	3599.15	243.17	3604.5	213.97	2913.8	427.94	3237.05	381.2	3532.8	1384.37	3237.7
M25322_at	574.8	357.09	515.45	595.6	144.5	0.85	218.9	99.7	105.15	41.79	465.9	313.67	660.85	185.76	179.85	181.23	173.1	22.63	102.55
D17516_at	263.8	265.73	99.65	115.19	860.45	462.66	550.1	53.17	1019.45	64.56	1087	327.25	519.25	260.57	216.7	157.68	668.55	172.32	657.55

M75110_at	71.45	15.49	36.7	42.43	175.3	178.05	146.8	117.52	196.8	6.22	251.65	8.27	154	123.6	261.05	163.98	179.45	174.44	337.9
U78798_at	841.05	471.71	367.9	425.11	992.6	947.66	391	149.34	345.35	194.53	1019.3	850.79	605.2	704.28	563	357.23	385.6	186.25	501.45
L42373_at	630.25	224.22	753.2	870.17	1101.7	675.29	1022	293.59	1041.2	243.53	1302.1	96.45	1319.3	195.02	1483.15	100.62	1011.4	272.66	791.2
M77829_s_at	436.8	103.38	129.05	149.13	687.9	560.74	189.1	49.07	188.65	75.02	668.75	760.07	307.6	31.4	264.5	57.98	266.5	12.45	230.85
HG2280-HT2	2473.95	592.63	1003	1158.81	3165.1	1134.48	2539.1	1268.83	2914.75	1066.67	3061.6	82.17	3078.7	44.97	2792.4	168.29	1979.45	262.27	2269.8
L29218_at	640.75	347.68	198.2	228.96	348.2	191.77	179.7	57.56	175.6	39.32	579.8	312.4	205.7	88.81	562.5	520.85	443	393.43	372.55
D14530_at	74984.1	39039.37	25309.95	29242.62	90311.4	6675.94	81422.1	31667.21	83046	20101.34	60781.15	12832.5	56930.1	19445.86	55867.8	13149.22	54139.35	12418.42	65868.2
M15517_cds5	1339.4	688.3	293.2	338.7	552.45	577.49	388.05	317.56	730.85	254.35	660.55	289.98	811.6	104.93	127.15	33.45	621.2	551.68	593.5
U88892_at	597.55	216.16	108.2	125.02	332.35	1.2	434.2	102.39	548.8	222.46	387.15	168.93	651.55	620.2	494	153.3	435.95	95.95	328.3
M80783_at	1190.85	245.86	631.25	729.38	1348.45	359.71	833.6	539.24	1266.15	199.33	1060.05	251.23	1114.6	465.42	827.55	329.87	1394.15	59.61	1069.55
U59289_at	320.25	145.88	343.5	396.83	881.85	349.81	920.3	702.72	551.2	119.93	463.15	365.22	378.8	420.45	618.05	118.3	672.7	469.94	711.55
D17408_s_at	154.25	49.43	240.65	278.11	57.5	32.95	309.8	159.24	356.15	368.19	174.55	190.28	154.05	115.19	180.3	18.95	116.65	0.64	86.2
X95095_at	92.65	43.77	28.65	33.16	117.45	11.53	130	115.4	127.95	34.29	469.05	616.39	68.55	32.46	156.75	88.88	98.75	82.24	88.55
S80335_at	306.85	327.32	116.8	134.92	261.25	118.58	245	53.32	366.85	329.3	572	190.78	179.25	72.76	300.85	89.87	151.95	1.2	154.3
U43519_at	562.25	389.97	84.75	97.93	255.55	113.63	294.45	190.14	347.5	237.45	947.15	59.75	1143.05	161.43	342.1	139.87	239.5	25.74	593.8
X62534_s_at	321.55	151.11	211.75	244.59	291.85	115.05	284	9.05	545.5	44.83	767.35	269.76	709.15	154.93	610	80.19	310.8	417.9	651.35
U43292_at	695.15	668.71	83.9	96.87	285.2	90.79	441.95	280.65	188.6	18.81	707.9	491.16	1161.85	199.47	943.25	1125.93	269.25	71.21	363.1
HG358-HT35	2725.05	363.38	1205.2	1392.43	3050	1100.97	3672.25	2463.91	2321.45	692.05	2908.5	340.97	2775.8	41.86	2487.35	579.05	2207.85	314.73	2716.7
D49410_at	716.6	61.94	452.85	523.19	988.5	232.07	2242.25	2498.42	855.3	364.87	1879.2	414.36	1017.95	53.39	565.3	6.65	829.2	365.86	1231
U24266_at	1491.8	124.73	942	1088.38	826.3	297.55	1700.65	197.49	2338.9	478.15	1411.65	1499.14	1218.2	312.97	1274.1	185.69	1830	724.93	1948.55
HG1515-HT1	15313.8	7137.39	7505.7	8671.96	13319.4	6210.66	16256.1	9575.92	11215.6	1262.89	9013.1	2869.44	8585.55	2144.58	11255.7	2900.27	10610.55	2355.44	12130.45

M11437_cds:	48.6	46.81	32.05	36.98	147.7	151.46	147.1	164.05	352.95	383.18	140.95	45.47	89.1	38.47	83.45	26.94	40.75	11.95	59
D38498_f_at	444.35	293.38	119.05	137.53	274.5	131.24	318.25	103.45	996.85	610.59	989.95	1215.02	236.6	32.95	463.6	454.39	522.4	629.47	358.95
V00594_s_at	301472.56	163568.58	195311.75	225659.06	385376.88	169126.58	302362.06	199730.73	443291.66	323712.28	257018.91	114845.02	398219.88	46866.27	510271.91	228426.03	299596.34	109810.5	237577.44
X95648_at	756.95	183.21	331	382.4	513.75	135.98	540.75	166.95	493.95	572.97	640.9	318.48	768.4	299.25	481.5	156.13	558.65	39.1	454.35
M63256_at	383.5	98.15	35.45	40.94	108.95	53.81	318.05	264.25	180.4	215.1	689.5	392.16	271.85	303.42	199.2	81.32	254.7	272.09	259.5
L77886_at	4016.2	441.23	1806.95	2087.73	3523.95	98.92	3171.7	1118.22	3223.45	953.11	3054.1	965.06	2374.2	848.25	2606.35	633.5	2281.8	427.23	2702.35
X76498_at	91.25	8.13	45.45	52.54	181.75	29.49	379.7	317.49	263.05	35.85	229.6	203.51	118.85	62.72	149.65	78.42	194.65	154.79	173.35
D38535_at	3211.45	337.22	2482.7	2868.45	3103.3	909.76	3446.2	1106.48	2337.05	296.35	2675.4	115.4	3505.75	238.51	4146.7	1661.42	2944.7	336.87	2460.4
M73482_at	81.15	33.16	44.25	51.12	125.75	60.74	99.2	62.08	56.8	10.89	75.55	5.87	301.55	301.86	345.95	348.11	132.9	29.13	176.1
X73478_at	2915.9	1097.71	1884.1	2176.9	2203.15	59.61	2634.3	517.6	2885.65	1346.83	2578.75	854.4	2838.85	1055.92	2955.85	404.96	4474.2	44.12	3221.6
Z79693_s_at	265.65	204.85	37.3	43.13	642.6	653.08	115	36.06	405.85	92.7	259.65	324.92	347.2	221.18	513.85	6.72	591.35	7	742.6
U80669_at	154.35	85.91	259.65	300.03	766.9	154.57	460.65	538.89	325.2	270.54	257.9	230.52	724.15	871.37	370.05	83.09	158	0.14	361.15
AF000545_at	1211.6	84.85	315.4	364.44	1287.25	377.81	1190.5	763.25	993.5	35.07	1592.85	461.81	1355.6	407.15	1534.5	29.13	1085.7	379.43	1174.4
U15008_at	11658	6476.81	5760.25	6655.22	12872.65	1447.8	14049.35	2256.73	12286.45	1101.6	12010.35	73.89	14380.85	3702.2	12832.9	1155.41	10608.05	3999.75	13839.55
U47928_at	186.45	4.74	86.3	99.7	522.8	213.83	607	627.49	508.35	443.71	241.8	84	240.3	55.01	621.55	674.08	194.7	34.22	282.5
X63337_at	1004.15	83.79	621.3	717.85	1228.5	496.25	755.75	66.4	470.65	277.4	1076.35	582.3	1156.4	194.45	1264.4	18.38	1245.85	528	537.65
X52001_at	113.45	61.73	65.1	75.24	183.4	143.12	226.7	209.45	123.95	33.59	425.25	178.4	45.3	22.49	78.15	4.45	212.5	196.01	102.6
Z21707_at	161.4	27.44	242.15	279.8	209.75	133.29	118.4	59.54	110.7	13.15	306.05	277.12	403.15	244.59	108.65	32.74	174.25	80.26	234.4
X59932_s_at	4530.95	909.69	2515.1	2905.93	3873.5	1483.79	3833.95	265.38	3622.35	231.86	3154.95	18.88	4229.75	533.23	4025.75	531.81	3741.95	206.55	4187.45

AFFX-BioC-5	347.15	69.79	149.15	172.32	1455.5	1111.85	1443.5	1678.11	363.8	4.95	524.3	214.11	378.25	113.91	1762.6	599.49	428.15	217.86	390.45
X04707_at	136.5	98.71	491	567.24	221.05	164.4	394.45	78.42	118	58.12	223.95	207.54	140.45	24.25	403.4	395.41	367.1	352	538.55
L05779_at	559.1	18.38	615.55	711.14	882.15	175.57	609.6	397.68	901.05	494.9	535.45	294.23	1405.75	959.47	1479.4	831.7	610.9	222.6	967.35
K01396_at	3381.7	566.11	2337.25	2700.37	4132.1	1084.14	3903.35	218.57	2018.75	95.39	1548.15	13.51	2569.85	1230.44	2292	335.17	5518.35	214.18	5348.35
J03133_at	904.55	70.5	916.4	1058.82	1141.05	748.76	699	394.99	1006.55	352.49	900.35	123.53	1312.3	832.12	619.55	423.06	1065.9	262.76	1069.1
K02882_cds1	1059.05	702.09	302.1	349.03	1551.05	595.31	1487.4	1079.89	1162.95	405.95	1331.9	40.16	1392	295.15	543.7	491.72	695.2	288.64	528.3
U00921_at	142.65	74.18	190.3	219.91	98.4	88.11	171.55	162	65.25	10.39	621.5	413.52	329.85	344.01	72.6	6.79	282.4	231.22	173.3
X79568_at	216.5	79.9	104.7	120.92	301.05	46.32	199.3	36.2	425.4	238.72	174.75	91.15	198.45	88.03	298.95	93.69	301.75	64.42	146.05
HG1869-HT1	5304.85	1632.5	2588.3	2990.5	5705.5	960.11	4377.4	2396.67	4354.9	780.93	4379.25	1527.85	4802.6	607.69	3897.85	682.71	4567.75	1018.73	4495.55
M74297_at	193.45	44.48	134.3	155.14	544.65	542.56	300.6	77.78	413.5	325.69	777.6	86.55	572.75	570.85	765.05	903.47	270.4	194.74	330.35
U84540_at	64	37.05	73.45	84.92	128.4	98.99	80.65	46.74	163.3	100.27	138.25	1.91	439.4	172.39	245.4	135.76	239.1	206.05	184.85
U06155_s_at	32057.55	16066.67	14455.1	16701.15	25810.95	8588.73	32345.2	19772.83	36260.5	2578.54	27074.25	11691.38	30828.9	2575.28	30228.75	158.75	34157.35	12898.69	42289.4
D50640_at	423.1	109.88	232.2	268.28	332.35	27.93	122.65	78.84	474.15	78.84	318.8	10.47	245.25	55.93	474.3	516.05	242.4	295	663.3
U32519_at	2036.15	324.21	1395.1	1611.92	1961.25	301.3	1958.9	89.24	2039.85	2.62	1773.25	686.67	2412.6	509.82	3102	243.39	2670.55	30.76	1018
S76965_at	276.2	279.73	204.65	236.39	923.4	355.82	352.1	123.6	324.05	215.74	360.2	55.15	373.85	362.11	432.95	47.59	315.7	37.9	713.45
L17328_at	236.35	203.72	45.75	52.82	298.4	32.53	119.65	56.64	542.9	82.02	467	316.64	104.75	44.34	511.05	492.22	303.5	129.26	363.1
HG162-HT31	1948.75	122.12	1611.9	1862.38	1320.5	163.62	1718.9	233.77	1555.9	175.65	1479.3	1333.18	1359.7	308.16	1605.3	317.35	1601.45	515.83	1645.8
M92432_at	753.95	573.68	181.65	209.94	549.85	189.29	392	106.35	401.85	67.67	703.5	410.97	1230.15	1039.52	901.7	409.98	431.4	50.06	515.4
X89986_s_at	407.9	100.83	260.25	300.73	658.05	284.61	90.6	51.48	509.5	0.42	858.15	301.44	454.85	142.91	579.1	97.86	221.05	41.08	78.6

D82346_at	339.4	111.02	422.3	487.9	969.45	938.97	402.3	161.64	405.1	463.58	238.4	90.09	285.9	93.62	290.6	37.19	227.35	0.92	220.95
X90761_at	856.6	734.54	560.55	647.64	1259.65	554.3	1197.55	1353.9	1402.65	170.91	734.85	164.97	946.15	129.75	1514.45	935.57	1076.65	1.34	371.35
X14850_at	3059.1	479.7	1955.35	2259.14	2351.85	820.03	3078.2	604.72	4457.1	1219.62	4105.95	1336.93	4722.65	435.22	3777.2	351.15	4446.9	332.48	3264.95
X02158_rna1	90.75	48.86	42.7	49.36	159.85	82.66	255.5	80.61	294.75	225.07	120.15	87.19	257.15	213.33	476.7	282.14	148.6	54.73	138.2
X82125_at	71	16.69	74.75	86.34	101.1	4.81	193.3	129.4	561.45	321.8	254.75	124.38	362.15	358.43	100.65	35.14	402.45	412.17	324.8
D63486_at	2717.1	877.1	2365.15	2732.61	1369.05	807.73	1712.45	574.95	2452.6	845.7	2322.65	1859.9	2483.8	1590.42	2352.9	1517.88	3942.9	153.87	3330.1
X97675_rna1	305.2	69.58	270	311.98	163	94.05	606.1	788	435.5	30.41	217.85	53.25	683.45	179.96	163.85	77.57	271.9	224.01	365.45
S85963_at	345.8	326.68	443.85	512.86	113.55	0.64	270.2	235.04	398.3	408	650.25	374.41	458.15	228.32	699.1	763.39	448	407.43	217.7
U07223_at	253.5	179.18	213.95	247.13	104.55	49.85	222.95	121.69	131.1	33.23	386.4	397.39	112.2	44.55	169.55	75.17	111.05	69.08	220.6
Z48199_at	24930.3	2311.25	10539.8	12177.51	16330.4	4746.1	15223.95	1177.26	20332.35	439.18	19650.95	717.64	24033.2	8092.69	21790.5	3452.52	29050.2	7257.04	20961.6
M62958_at	315.15	328.59	136.5	157.68	401.25	409.06	240.55	129.75	300.05	229.74	514.35	312.33	153.35	111.37	126.6	114.13	497.85	574.81	256.6
U71087_at	947.35	652.73	79.05	91.29	439.15	52.11	801.05	258.31	946	218.64	1170.7	679.67	516.5	148.92	472.75	415.71	612.4	111.44	994.45
U64520_at	773.15	0.35	131.75	152.24	627.85	351.5	412.95	331.14	494.3	534.86	405.3	416.2	141.85	114.34	464.85	250.25	400.9	81.46	817.65
U19252_s_at	288.4	262.9	158.9	183.56	293.7	157.26	132.25	107.69	460.7	454.95	263.8	262.76	770.65	91.71	206.35	35	439.35	479.06	168.1
D83783_at	2079.8	518.59	838.85	969.23	1338.15	444.56	1376.15	685.4	1534.4	297.13	1649.2	41.3	1211.95	855.95	1267.95	410.33	1674.05	50.98	1617.85
Z11933_at	67.3	33.23	111.4	128.69	391.4	458.49	87	87.26	112.35	27.93	108.65	67.95	121.05	20.44	104.25	63	122.75	31.47	142.25
HG2465-HT4	669.8	126.43	606.15	700.39	1172.8	262.48	1232.55	668.71	1084.5	97.44	716	35.92	1192.3	162.49	938.25	427.3	1305.95	39.39	1241.2
X16135_at	4011.65	1058.75	2200.65	2542.54	4260.15	764.88	3725.85	755.97	5031.2	263.18	4757.4	156.27	5253.75	1039.66	5402.45	1043.48	6469.7	1643.6	3989.4
X17093_at	411.55	169.78	646.2	746.56	786.25	779.3	846.45	556.14	436.8	17.39	674.8	284.4	359.1	346.06	604.4	218.92	429.75	460.11	621.45
Z48475_at	247.25	6.15	136.7	157.97	260.2	9.19	210.5	12.87	189.15	39.67	227.2	121.06	252.5	79.62	308.7	36.49	262.8	35.07	240.85
M13560_s_at	161.25	48.72	99.25	114.62	188.65	8.98	177.55	41.65	230.5	79.05	202.55	175.15	147.6	51.05	212.05	144.04	226.35	40.23	359.75
X77922_s_at	156.25	148.85	65.15	75.31	175.7	88.53	629.55	328.59	660.7	320.32	105.6	34.65	30.4	7.5	194.55	189.29	271.65	337.08	244.65

U26398_at	495.5	494.83	131.05	151.39	754.95	535.77	371.35	267.36	584.85	97.09	222.5	87.12	395.1	240.42	542.1	64.49	186.5	30.55	418.1
U69645_at	1512.4	481.82	883.6	1020.92	1603.55	278.11	2039.15	461.67	1619.85	299.18	1510.6	45.96	1579.45	342.73	1405.9	585.91	1461.7	33.09	1656.35
HG2981-HT3	10994.4	4946.21	5924.4	6844.94	5497.45	4192.22	4983.75	1339.9	5787	4674.26	6364.4	2535.97	7701.1	2658.01	5331.9	2172.23	10474.95	143.61	10416.3
L35545_at	975.85	562.93	872.25	1007.84	1150.65	557.55	1073.2	282.7	980.45	397.75	820.2	911.18	1349.8	29.56	1561.25	210.93	989.85	250.81	1250.2
U86529_at	4400.55	183.49	1300.55	1502.67	4513.35	1807.15	3583.75	362.11	3095.6	597.08	3519.8	749.25	3014.55	782.56	3696.5	228.96	3600.85	989.74	3952.2
U57796_at	197.65	238.37	256.45	296.35	151.15	137.96	161.8	140.15	247.85	231.86	60.15	15.91	169.6	196.86	76.8	4.24	193.75	145.45	194.4
X99897_s_at	111.1	38.18	68.1	78.63	291.45	268.49	94.75	42.36	122.5	4.53	143.9	133.08	100.75	3.18	151	80.04	107	46.1	109.3
HG371-HT26	283.4	259.65	821.85	949.57	244.65	89.31	283.95	226.77	302	147.22	528.55	573.68	210.95	108.82	204.95	143.75	168.9	56.43	587.6
M86873_s_at	44.55	7.85	24.15	27.93	64.7	18.24	85.1	46.1	221.3	111.3	106.25	77.43	78.55	72.76	73.5	28.14	81	20.51	225.7
M60165_cds:	2197.5	29.7	848.45	980.26	1848.25	370.74	1412.6	183.71	2476.65	87.19	1392.85	334.53	2536.85	123.53	1748.5	523.54	1930.7	358.22	1275.85
U28049_at	680.3	258.09	625.45	722.59	516.6	369.53	1077.1	184.98	534.65	212.91	708.35	218.43	1135.15	289.56	1442.6	318.06	1140.75	222.53	708.45
D87075_at	788.55	6.86	99.95	115.47	933.45	486.7	159.45	66.82	607.15	697.99	1432.85	477.65	212.8	46.53	826.45	475.39	559.85	64.56	726.2
M32886_at	1768.9	234.19	1036.85	1197.91	1771.6	352.14	2226	603.44	1671.65	401.14	1840.65	96.38	1677.65	141.49	1509.75	9.12	1663.1	23.33	1440.05
M14199_s_at	74664.5	22922.99	47836.15	55268.81	106681.05	24295.55	112395.35	3557.24	94137.35	53195.01	75529.7	12602.06	76630.45	6595.26	93452.59	17395.25	78178.5	8995.25	73688.85
M74525_at	220.75	247.84	344.95	398.6	104.1	67.18	155.85	137.96	191.75	198.34	72.8	13.86	204.7	231.79	404.9	200.96	299.8	378.3	367.55
D10656_at	1058.55	133.15	1122.15	1296.48	1153.45	52.96	642.4	7.35	783.5	701.45	1401.85	101.05	1101.95	191.41	622.6	625.08	658.25	231.58	1278.5
X79483_at	1344.2	108.61	651.45	752.72	671.7	212.98	1391.4	290.76	607.5	344.08	1587.55	1617.79	1298.45	931.05	1774.1	1016.82	795.5	427.23	1242.4
D26018_at	95.25	66.26	77.6	89.66	329.2	219.91	101.9	41.86	231.3	144.53	136.75	37.55	300.65	169.78	112.85	5.87	201.95	15.91	321.1
M26311_s_at	3652.4	676.28	1319.6	1524.66	4188.6	590.01	2927.7	127	4099.65	1146.86	3225.75	913.37	5512.65	1440.59	6892.05	1906.71	8919.15	490.94	9619.95
Y09616_at	3717.85	510.04	1846.9	2133.91	2772.45	2179.37	3469	11.31	3794.8	507.42	3385.5	698.76	4074.9	1230.08	3496.95	1068.08	2347.5	931.26	3579.65
Z29083_at	3399.65	1220.4	1756.85	2029.89	3612.45	559.25	2428.35	766.72	2966.55	582.02	3076	522.98	2939.25	370.17	3644.3	1563.27	4364.45	650.33	3210.7
M77140_at	117.2	104.93	22.75	26.23	93.55	7	61	44.41	354.55	280.65	95.9	42	131.35	79.13	275.9	335.88	156.35	7.85	434.3

U23850_s_at	527.4	659.02	375.5	433.88	756.5	528.77	244.05	213.05	125.9	61.66	234.15	242.75	166.1	90.37	378.55	134	713.55	230.3	740.35
M31211_s_at	1255.85	760.92	612.15	707.32	1500.95	723.72	1381.05	94.82	1608.8	452.27	1285.25	449.37	1467.4	618.29	1437.5	115.26	1612	151.75	913.4
U31903_s_at	3749.65	41.79	1163.8	1344.63	2600.4	384.24	2893.3	888.41	2657.3	303.07	2628.95	282.49	3099.15	87.04	3044.6	243.67	3731.6	956.15	2559.2
HG4668-HT5	2068.8	102.53	966.05	1116.17	1959.45	318.27	1534.45	287.16	1820.85	307.95	2131.15	129.33	1363.2	95.18	1283.6	806.53	1882.55	277.68	2305.9
M21154_at	681.75	238.79	811.05	937.13	928	690.7	411.75	381.34	1119.05	392.94	628.85	117.73	875.4	488.19	795.4	440.95	699.5	131.38	949.45
X61123_at	1393.6	615.47	1285.55	1485.28	2447.55	177.41	2254.35	907.15	1991.9	248.48	2666.55	1182.21	2061.75	59.04	1661.7	913.16	1663	1360.05	2421.4
D88795_at	85.35	21.99	121.4	140.29	382.7	417.05	411.1	145.1	509.45	270.04	176.55	138.66	129.5	46.53	579.4	677.41	241	141.56	183.25
V01516_f_at	269333.81	135676.53	135146.95	156145.94	279614.13	46259.92	280963.56	176869.53	177229.69	22991.01	172003.75	13728.26	216858.75	36428.38	233759.56	52882.19	218159.5	34135.87	108403.65
X52282_s_at	83.55	39.1	90.65	104.72	558.55	656.27	341.35	242.61	533.65	333.4	158	4.1	472.5	453.4	127.1	12.3	251.6	170.41	285.35
L33881_at	500.95	89.73	237.8	274.78	579.55	579.62	273.45	63.85	1124.45	344.71	1345.6	574.17	626.15	396.33	657.6	725.63	421.75	157.19	612.75
U18062_at	418.55	107.83	436.8	504.73	429.65	203.01	731.8	254.28	698.95	302.01	476.65	154.64	480.05	55.79	682.75	542.85	519.8	209.16	680
M96739_at	2879.3	590.01	1228.7	1419.59	2761.1	137.46	2371.5	434.02	2164.7	726.48	2941.05	621.48	2208.35	73.04	2501.05	371.16	2489	452.83	2444.55
M21064_at	607.7	719.98	281.05	324.77	185.55	170.34	57.6	31.82	248.75	256.18	180.05	160.3	211.65	72.48	29.5	4.1	492.3	278.74	645.1
J03060_at	254.15	223.8	131.65	152.1	189.15	55.65	342.6	91.92	235.3	89.8	208.85	59.89	390	207.47	209.75	43.06	120.3	49.78	261.1
U22961_s_at	550.4	109.46	117.05	135.27	646.6	335.59	716.05	426.74	235.4	198.56	778.35	452.05	396.4	55.15	559.75	259.3	566.75	261.28	313.05
X83228_at	44.5	24.47	11.5	13.29	131.35	113.49	98.9	44.12	201.45	190.71	35.15	21.14	46.4	12.16	38	14.28	77.15	15.34	48.9
AF009674_at	1263.2	1064.9	1257.35	1452.75	1022.75	177.27	2004.9	1827.16	3267.85	1401.98	1958.1	1489.87	2422.5	1661.56	1353.9	502.33	1698.25	281.36	1614.7
HG511-HT51	1517.4	18.81	1184.45	1368.46	1242.65	652.45	2172.1	59.11	1049.65	102.88	1145.7	535.28	1414.15	93.41	1611.45	62.86	2007.75	111.79	1730.05
U33429_at	1471.6	798.75	231	266.86	954.2	357.94	815.8	170.55	431.6	49.07	1431.8	121.76	706.05	86.9	605.25	86.76	762.5	2.83	808.2
D85418_at	646.25	659.8	107.2	123.89	882.45	136.97	492.75	196.65	964.3	65.76	1154.25	621.76	874.85	773.22	286.1	49.36	420.7	119.64	735.35

Z24680_at	964.75	263.4	700.3	809.07	1491.4	381.98	1010.4	264.18	590.4	626.36	606.9	226.84	930.25	492.36	503.95	383.46	1336.45	232.85	924.05
X97302_at	1249.9	224.01	629.6	727.47	1302.75	315.02	788.4	487.06	1483.5	66.47	1450.3	163.34	1204.7	65.34	1189.4	221.04	980.1	248.19	777.35
AF014958_at	90.6	6.51	65.8	76.08	191.55	131.59	191.95	75.59	132.7	26.73	207.7	81.32	754.8	851.07	201.55	9.69	99.25	23.55	133.35
X86570_at	274.15	207.82	149.15	172.32	815.45	791.61	584	593.4	453.25	233.27	393.25	266.65	323.75	24.96	260.15	155.78	326.05	29.06	347
X76942_s_at	494.05	211.07	55.85	64.56	536.8	72.55	743.8	565.69	900.7	104.09	825.8	327.81	739.1	9.76	391.4	425.4	682.05	232.71	394.85
U68019_at	486.5	76.08	101.25	117.03	387.85	373.56	184.6	49.21	611.4	610.8	387.55	221.54	347.65	166.95	695.65	555.29	761.95	391.81	638.9
V01510_rna1	154	0.85	79.6	91.92	221.65	124.66	265.95	33.45	172.2	116.67	257.15	90.44	207.5	8.49	169.9	0.14	164.9	47.38	243.05
U15306_at	131.95	67.53	293.8	339.41	138.4	1.41	211.5	133.93	134.75	0.49	131.25	29.77	156.05	42.21	74.1	8.63	263.55	194.38	88.55
D26561_cds2	81	17.82	151.2	174.66	428.25	553.74	134.05	152.38	471.85	556	184	116.67	330.4	12.59	279.55	191.41	402.9	220.62	231.55
D79993_at	409.45	44.48	334.6	386.65	565.5	142.69	385.8	411.11	879.1	53.74	810.4	103.8	226.5	287.23	816.3	86.27	669.1	105.92	557.95
X85373_at	3386.85	1131.44	1234.2	1425.95	3196	1191.9	3659.65	1310.06	3884.1	328.95	2727.75	182.08	2978.25	715.95	4170.75	115.75	3014.3	618.15	2792.6
D25215_at	227.5	179.61	41.1	47.52	458.9	240.7	518.5	402.77	350.2	352.85	442.8	372.65	73.6	16.4	117.4	38.89	377.4	72.27	636.15
X90978_at	734.7	490.73	377.55	436.21	1097.7	877.8	539.25	306.81	890.55	908.28	486.45	170.62	607.7	576.86	936.75	782.27	1150	312.68	1195.35
D76444_at	326.95	25.24	120.75	139.51	556.95	561.94	828.25	233.42	572.35	133.86	622.55	588.67	718.2	179.18	221.1	118.94	671.45	141.49	904.05
M64269_s_at	719.3	69.01	454.75	525.45	1161.55	834.46	439.15	99.49	618.4	102.11	1117	522.98	810.2	113	521.05	452.34	711.3	152.59	846
M24766_s_at	2328.25	1646.64	811.3	937.34	1178.85	557.98	1043.65	557.13	1074	583.22	694.2	345.92	912.05	511.73	1533.35	724.43	1803	440.53	968.25

L06499_at	133059.91	44352.84	77599.3	89656.61	318855.56	109782.94	265584.69	174915.73	270242.09	93336.11	269834.56	179677.61	260821.84	113404.02	240190.41	63014.67	197195.34	72936.86	154703.36
U18422_at	91.25	5.73	133.15	153.8	141.6	35.64	157.55	137.53	58.5	49.36	112.35	21.57	478.15	281.07	389.7	535.56	274.3	237.73	45.35
X91247_at	2295.15	694.45	1805.1	2085.54	2078.85	118.86	2733.95	702.93	3007.15	335.38	2196.75	267.22	2251.2	281	3870.55	1888.75	2472.15	272.87	2725.65
S59184_at	1025.95	606.34	603.15	696.85	1132.3	215.24	673.8	338.28	1217.7	149.34	1107.75	232.43	675.6	170.7	1018.05	88.74	561.8	324.7	1064.75
L13689_at	537.75	41.51	361.05	417.12	370.65	94.12	524.7	288.92	951.5	539.81	955.8	132.79	409.45	143.19	319.55	136.4	525.5	211.28	618.8
U01160_at	409.95	437.91	463.95	536.06	634.05	168.36	904.15	594.46	493.95	452.62	968.35	94.82	1004.5	536.27	452.15	443.85	775.4	19.09	880.9
X04445_rna1	1973.55	612.99	545.5	630.32	3751	2314.64	3504.45	640.57	590.2	243.1	1706.15	1532.37	2680.55	253.22	1436.35	1001.05	1493.7	1357.36	1184.2
X99142_at	1954	791.96	1333.6	1540.79	1260.1	1075.65	1503.25	356.17	1721.05	603.8	1730.4	236.74	1563.15	922.85	3080.9	542.35	2112.45	981.96	1139.75
U88726_at	928.05	131.17	176.35	203.72	641.5	363.31	298	214.54	498.85	168.5	741.2	97.3	369.75	313.04	720.5	181.87	710.8	364.3	510.35
D89052_at	7416.3	1656.61	4055.05	4685.08	9647.9	191.48	10310	3971.39	5693.4	1729.16	7531.6	1573.03	7478.95	2756.8	7332.45	1895.68	7440.9	1446.46	7566.75
HG742-HT74	663.9	480.27	143.55	165.82	809.25	376.82	859.3	241.41	154	47.94	764.35	9.69	903.15	181.51	810.45	70.92	848.1	495.26	850.6
M60721_at	263.95	51.97	135.9	156.98	278.05	17.18	430.5	224.58	347.8	73.26	379.1	92.63	381.15	37.83	591.9	559.89	688.05	840.68	746.55
U78575_at	966	387.35	515.6	595.67	975.95	142.76	933.2	269.12	1296.35	25.53	974.6	24.89	852.2	183.56	1028.55	4.17	1222	45.4	1395.75
U10117_at	1380.5	374.63	779.25	900.36	1503.35	390.25	1258.3	336.44	1266.6	271.67	1440.7	311.55	986.15	570.28	1372.2	389.05	1187.9	142.98	1449.95
D25278_at	220.65	199.47	410.2	473.9	336.3	363.45	852.6	592.13	614	66.33	204.4	205.63	664.9	416.2	904.45	414.01	456	421.58	549
U94319_at	493.35	37.41	232.45	268.63	267.7	61.52	402	352	440.9	37.76	463.15	313.46	505.4	379.29	60.6	18.38	223.65	274.71	192
U17195_at	364.85	98.5	120.7	139.44	596.35	312.47	90.7	4.1	200.9	197.71	265.8	5.66	114.05	20.72	591.1	377.03	90.25	13.36	328.7
L42611_f_at	60380.4	4768.73	22267.65	25727.59	50995.65	3773.33	60228.85	4775.16	34655.8	5211.8	33840.4	2983.71	37970.85	4759.47	39033.75	2796.4	39989.65	304.27	31763.4
X12447_at	58292.5	34085.23	31191.45	36037.91	66194.65	8628.75	81883.8	16039.87	30072.05	8834.52	45390.65	10903.09	56453.7	12553.55	59700.45	13255.35	62933.6	23571.69	52077.9
D13789_at	619.95	35.99	191.35	221.11	671.95	58.34	711.55	521.35	548.25	4.31	417.5	24.32	658.05	104.58	668.25	490.8	500.9	0.14	436.95

D50370_at	245.1	189.36	155	179.04	224.9	201.38	78.85	51.97	308.8	166.17	259.25	12.66	67.4	26.16	286.35	302.15	297.45	340.75	111.75
U15197_at	547.95	290.83	112.6	130.11	1081.1	271.95	1357	503.04	371.2	11.31	2430	1335.3	1456.9	255.97	567.85	310.21	936.95	275.84	784.25
L01664_at	61.15	10.39	17	19.66	194.3	217.22	64.15	35.85	79.7	55.3	529.2	50.35	156.3	148.63	249.8	303.21	76.35	40.52	78.7
M88468_at	4755.4	539.81	2504.65	2893.83	5079.05	1102.45	5285.3	290.62	5512.35	201.03	5655.6	743.73	5317.95	716.09	5422	1011.3	5517.85	355.46	4786.4
M61176_at	200.7	261.91	51.25	59.18	119.85	76.01	166.45	74.74	210.9	105.92	444.15	58.9	142.1	80.47	241.3	78.06	509.7	184.98	229.65
U21551_at	679.35	19.45	240.2	277.47	401.65	121.98	1055.3	830.71	299.65	331.56	325.1	245.65	210.3	204.07	667.2	726.48	533.5	182.01	360.1
U41654_at	2806.3	983.16	816.35	943.21	2827.1	337.43	2223.75	714.53	2009.55	13.65	1750.4	78.35	1935.85	760.21	2300.05	72.76	2380.15	390.11	2015.15
D42138_at	393.5	138.17	210.7	243.39	606.15	436.36	439.45	3.04	387.1	229.67	459.05	562.93	500.65	196.36	727.6	840.18	275.6	10.47	845.75
U77594_at	186.8	115.82	52.3	60.39	149.05	45.89	302.65	306.81	123.9	65.34	187.3	77.5	172.15	58.34	159.95	73.19	170.1	78.77	153.4
X99894_at	193.85	17.04	110.3	127.42	296.1	75.94	275.3	147.22	236.95	112.08	307	153.73	529.8	334.46	207.75	27.22	173	50.35	218.45
HG2815-HT4	25471.25	15518.09	11445.1	13223.46	32920.15	7741.62	28916.25	7362.89	27843.05	5295.45	20002.05	4513.25	19891.8	3361.87	25628.5	1999.56	23891.15	6929.86	22123.9
U20536_s_at	116.3	14.57	24.55	28.35	89.85	4.6	597.25	780.01	197.45	203.86	85.7	34.08	244.05	224.08	161.5	148.35	258.65	217.58	78.05
U44754_at	414.15	112.36	626.55	723.87	319.1	19.66	696.95	536.91	557.45	15.49	556.2	11.17	297.75	284.61	225.65	205.7	397.6	67.74	582.75
M37825_at	50.35	3.89	246.65	285.03	308.4	305.61	187.1	127.42	262.15	76.72	231.4	234.34	151.9	50.35	69.55	15.77	168	187.81	433.65
D14664_at	329.1	423.7	44.95	51.97	116.4	126.01	99.6	123.89	369.75	434.23	70.75	29.06	294.4	378.3	281.95	326.61	102.1	114.69	116.5
U72511_at	6983.75	1044.04	5058.7	5844.66	5479.8	1867.75	8741.85	6542.22	5464.6	925.89	7370.1	3372.05	5909.1	171.12	7421.15	227.62	8622.4	1710.49	7671.3
X71490_at	3550.2	1148.06	1735.25	2004.86	3429.9	850.37	3016.95	431.97	2982.5	1774.56	2040	903.12	3339.1	597.79	3469	531.46	2401.5	296.42	1522.6
X66142_s_at	919.65	203.58	95.1	109.88	906.95	770.25	933.45	503.25	897.25	468.6	616.35	429.14	1442.65	109.67	246.1	84.71	919.7	164.33	1434.9
U91616_at	260.5	12.73	65.7	75.94	573.25	625.72	448.05	386.15	187.3	122.9	238.7	170.27	209.65	23.69	101.45	2.19	206	45.54	153.95

U51240_at	319.45	38.11	94.5	109.18	333.95	196.93	393	196.86	677.95	700.11	220.65	126.64	403.25	191.27	158.95	9.4	516.1	408.14	340.7
Z46376_rna1	1090.95	91.85	523.6	605	846.1	123.89	380.2	374.63	1119.65	181.94	1265.75	246.57	756	23.33	448.35	512.02	1062.95	347.68	1006.15
U24389_s_at	556	489.18	23.8	27.44	586.45	208.95	676.6	445.76	493.15	585.98	766.35	171.05	557.7	65.9	524.05	641.28	458.25	382.33	526.05
S39329_at	146.6	8.34	269.4	311.27	283.3	310.84	512.2	573.04	307.6	240.84	496.65	168.22	110.9	13.01	255.95	75.31	215.8	181.02	121.25
U45878_s_at	147.4	148.63	543.85	628.41	230.7	76.79	249.8	202.37	441.2	428.51	978.9	808.22	154.05	92.84	194.55	148.28	429.7	480.27	530.35
U25265_at	987.75	381.06	728.8	842.02	651.45	13.93	651.05	407.65	1469.15	353.48	910.8	357.09	1078.3	61.38	574.85	528.7	738.45	47.59	1009.45
M17466_at	2934.2	0	1201.8	1388.47	3070.7	569.08	4038	1520.7	1917.9	869.88	3168.2	919.1	2534.15	331.14	1913.6	1009.61	2441.55	550.62	2525.25
L19493_s_at	362.6	227.41	388.85	449.22	118.85	8.56	469.1	489.6	283.35	46.03	153.5	4.53	160.2	196.72	383.15	161.57	201.9	51.9	217.25
U60205_at	1224.15	681.72	591.4	683.35	603.2	64.21	765.1	123.46	1621.05	1048	476.95	443.85	1376.4	31.25	1470.95	85.77	1125.1	739.78	455.2
U68105_s_at	9744.9	3906.2	3794.7	4384.34	8254.7	5011.12	6876.95	4098.74	7475.1	1270.67	7109.5	487.76	6418.65	571.13	6533.35	1385.43	8758.3	2938.03	8064
U60666_at	1159.8	395.98	402.7	465.28	740.3	8.49	1385.45	775.91	636.1	621.26	1331.1	552.96	791.75	201.17	414.55	148.99	857.2	62.37	812
S81264_s_at	2305.55	1158.59	1276.4	1474.74	1307.95	131.88	944.6	782.06	1327.4	262.9	1579.2	452.41	1463.35	784.96	1786.55	175.29	1350.25	1202.58	1357.2
M74002_at	650.6	109.74	404.7	467.54	378.8	267.99	602.8	44.97	1217.05	49.43	1120.95	365.08	747.35	80.12	935.4	40.02	451.4	208.31	851.15
M21259_at	1897.4	337.01	1086.85	1255.75	1744.7	282.28	1507.7	932.67	2746.85	276.13	2149.65	534.5	1794.75	103.03	2557.4	290.2	1424	554.23	1965.25
D87460_at	3211.05	118.86	1263.75	1460.1	4083.35	992.42	3377.9	942.15	3978.9	170.84	3798.2	376.75	4058.6	34.37	3952.8	1138.16	3226.8	429.64	3903.55
U22029_f_at	1265.6	640.21	261.15	301.72	431.05	385.16	544.05	428.86	756.2	466.97	865	675.29	1180.9	976.51	732.3	542.07	963.65	198.06	382.25
Y07846_at	229	121.91	110.45	127.63	260.8	110.73	139.9	49.64	185.6	1.7	215.1	154.71	262.7	51.62	165.4	27.01	169.35	1.63	209.9
X68742_at	166.25	84.36	418.35	483.31	283.7	303.49	92.75	77	37.05	8.98	216.95	193.96	493.7	31.96	264.3	82.73	204.8	254.56	735.4
X56411_rna1	61.6	18.1	22.8	26.3	150.9	89.1	359.4	157.4	292.65	342.31	204	189.36	77.2	20.93	168.25	39.1	153.75	119.71	72.85

X52611_s_at	1172	1017.53	1490.95	1722.58	357	50.77	294.1	28.14	691.35	424.19	787.9	361.33	645.6	218.64	1020.45	574.67	1022.6	725.21	889.65
L13943_at	104.8	37.62	75.65	87.47	489.1	254.13	112.2	125.16	249.3	224.44	750.6	537.12	56.2	40.87	102.6	60.53	57.25	44.48	270.25
L20433_at	156.3	93.34	105.05	121.41	121.6	7.64	191.4	157.4	92.25	2.9	137.95	68.38	193.55	29.49	129.9	13.86	148.15	17.18	100.55
U77129_at	228.3	50.35	185.65	214.47	732.1	451.98	172.65	16.48	658.85	369.75	461	155.85	111.65	58.05	248.95	12.09	360.5	279.02	145.85
L05597_at	18.95	2.33	214.7	248.05	421.9	499.78	123.75	106.84	148.15	147.43	132.8	139.44	170.15	204.57	31.05	4.17	24.95	4.74	264.9
Z48579_at	797.55	228.75	422.2	487.76	861.1	56.99	545.4	409.7	361.55	309.64	1273.85	681.58	286.35	215.03	870.05	16.62	603.15	92.84	828.3
M73077_at	195.85	38.4	602.05	695.58	335.9	153.58	122.25	21.85	473.25	239.64	260	99.84	238.75	27.37	144.35	0.78	400.6	128.41	279.75
J03242_s_at	689.1	186.68	453.6	524.11	1151.25	287.44	1439.3	653.37	552.9	385.94	611.5	477.01	818.3	15.41	867.8	209.16	619.6	645.73	1146.2
HG4188-HT4	147.55	40.38	105.85	122.26	155.8	45.82	88.25	19.87	148.35	33.45	298.25	189.01	363.4	259.79	483.3	542.49	291.2	240.27	306.25
D63476_at	1980.2	374.91	1216.2	1405.16	1873.55	639.72	1700.85	247.28	1611.05	121.83	1559.05	389.97	1966.25	626.14	1977.9	127.7	1471.35	136.12	1592.4
S43646_at	687.15	240.63	690.2	797.47	1105.8	604.43	925.75	93.55	994.35	249.26	903.9	44.69	1054.2	188.09	865.85	236.24	829.85	451.06	1356.2
HG415-HT41	361.1	30.55	534.85	617.94	750.1	153.44	1089.35	650.04	1569.4	1076.64	800.85	5.02	680.6	509.54	1322.45	837.71	390.8	41.58	461.95
J02923_at	1824.6	805.25	1503.3	1736.94	1533.3	213.55	1331.3	454.25	963.45	288.57	1615.45	160.02	988.7	70	1031.2	326.4	981.4	228.68	740.05
X74104_at	5060.2	508.98	2474.75	2859.33	4224.4	1021.06	5590.75	1044.61	4915	188.09	3952.9	408	3956.7	902.27	3772.7	1374.62	4614.7	522.41	4562.5
X70040_at	1383.15	1189.57	1289.25	1489.52	1474.6	795.21	1484.05	373.14	1037	839.62	792.25	375.97	2206.85	1724.28	1592.95	1432.53	2237.15	473.55	2286.55
X58288_at	279.8	283.69	317.8	367.13	266.9	307.87	261.35	246.29	493.85	125.65	169.4	20.36	468.25	219.27	517	220.19	203.25	214.61	124.5
U07231_at	397.35	125.51	543.35	627.84	126.05	64.13	250.6	73.54	191.1	23.05	675	149.48	324.5	213.12	370.3	185.69	615	239.28	825
AFFX-CreX-3	290246.88	166945.02	173993.11	201027.92	662228.06	606850.75	688132.69	778190.75	369613.66	236731.09	489178.16	489249.53	317990.94	42701.12	423904.16	239675.91	286238.56	67725.77	222371.34
HG142-HT14	92.5	59.54	74.15	85.63	163.15	34.44	154.45	121.27	88.9	13.15	299.85	192.26	63.6	16.55	127.8	9.76	125	1.41	160.2
D83407_at	284.4	164.05	236.1	272.8	169.7	6.36	220.8	43.98	672.7	147.93	527.25	208.1	282.15	202.16	573.6	96.59	301.55	179.11	450.35

X62078_at	3119.05	836.15	2139.6	2472.05	3943.1	466.55	3628.85	457.85	4471.75	97.65	3590.35	1328.3	5297.3	1859.55	5030.45	1635.33	5434.45	2036.96	4938.45
HG3570-HT3	2148.95	989.03	896.6	1035.91	2742.7	931.4	2739.75	360.41	2028.8	6.22	3067.8	813.17	2990.9	285.95	3319.3	500.49	2797.2	646.15	2914.2
M60751_at	23.25	4.03	26.45	30.62	34.55	27.08	72.6	84.57	133.75	173.59	251.25	338.21	44.05	9.83	43.4	32.24	128.7	130.96	79.75
L11573_at	144.25	108.54	165.55	191.27	579.5	753.78	663.55	408.35	205.5	104.51	487.25	270.04	455.1	7.64	172.75	168.79	562	541.5	677.85
X95384_at	249.9	195.16	56.05	64.7	400.1	402.63	1324.25	894.7	286.3	63.36	657.3	140.01	352.7	262.9	373.55	65.69	62.8	5.66	708.3
U18244_at	385.45	78.28	351.95	406.66	268.3	59.96	391.45	24.82	412.25	151.67	446.05	146.58	905.4	223.59	1474.1	1434.72	1821.5	1111.43	1165.6
X97065_at	1715	242.68	2049.4	2367.82	1017	821.23	1688.65	465.49	2414.9	916.83	2076.6	617.87	1549.3	525.24	1846.2	1152.73	1717.05	165.96	1745.65
U58516_at	4915.3	1506.7	3852.7	4451.38	3955.1	139.58	4265.9	2129.81	3414.25	189.01	1970.85	2111.35	1308.45	1420.65	3298.75	831.63	3590.75	1737.01	4132.35
AFFX-BioB-5	16152.75	3411.01	8604.95	9941.99	69571.1	73395.7	73094.5	84461.92	34160.3	25062.27	55802.05	67666.09	25526.55	22735.11	26140.25	2130.44	19627.95	1279.37	25817.05
X52213_s_at	594.2	424.55	223.6	258.38	286.05	72.76	518.5	335.59	365.9	75.09	447.45	292.95	492.7	340.12	969.05	198.91	296.3	29.13	220.5
J02621_s_at	6814.4	2207.45	4550.95	5258.12	6272.2	2800.28	5794.2	2000.69	5715.35	527.57	3823.85	303.42	3440.7	471.5	4545.25	1853.11	4204.95	31.89	5092.45
L20859_at	3327.75	1469.86	1882.5	2175.06	1769.25	365.64	1889.6	1023.89	6053.45	1837.13	2837.1	1200.38	4466.85	2836.13	4228.95	1038.24	4528.7	1945.96	2911.7
M22403_s_at	158.55	44.48	105.85	122.26	593.6	252.15	354.2	153.87	231.75	169.07	464.3	537.83	337.85	242.33	250.65	117.45	172.1	97.02	272.85
X77737_at	767.25	273.3	114.65	132.44	455.85	130.46	314.25	134.56	233.45	14.07	796	780.5	422.6	126.29	834.6	570.92	504.5	273.37	373.45
D85527_at	3738.6	922.07	1609.65	1859.76	3536.65	269.34	4094.05	398.88	3215.95	401	4074.4	327.53	4200.8	1899.29	3608.55	223.8	3906.45	1191.83	3397.05
Y08639_at	710.4	587.32	233.8	270.11	343.75	155.35	1178.4	761.55	800.65	229.74	329.2	73.96	348.8	198.7	369	291.19	939.35	378.23	971.25
M13577_at	122.5	34.08	117.35	135.55	76.6	32.1	247.95	213.76	63.8	22.49	166.25	139.23	167.85	187.45	82.1	31.96	160.75	24.25	142.35
U21556_at	393.75	137.67	124.4	143.68	245.75	77.15	986.1	1129.39	214.5	47.94	535.7	423.27	537.85	91.43	763.15	457.14	371.3	103.1	595.25

HG2414-HT2	142.05	49.99	101.65	117.45	160.65	43.77	224.65	136.54	153.75	44.05	218	171.54	171.3	14.28	147.6	13.44	196.4	7.21	210.05
AB002356_s_	1971.9	114.13	1484.6	1715.3	2199.8	243.67	2986	1122.6	2679.65	404.39	2900.8	656.05	3064.65	1370.44	2499.8	208.17	2963.3	332.91	2790.75
X55668_at	374.1	233.91	173.2	200.11	268.05	79.97	939.85	473.27	182.2	0.85	434.1	74.25	309.95	3.89	164	7.5	230.15	159.17	208.95
S77361_at	155.9	43.84	35.35	40.8	158.1	41.3	862.45	1088.31	79.55	13.36	272.65	306.11	209.75	171.47	345.65	209.94	219.55	157.9	81.25
X57152_rna1	11428.95	2639.42	5874.7	6787.52	11185.35	2713.8	12207.55	5138.33	10888.05	1550.19	10523.7	1176.06	8987.55	847.89	9252.45	4415.39	10956.1	4483.34	11990.9
Y10936_at	1264	816.85	901.95	1042.06	1394.05	756.96	416.5	118.65	415.9	228.4	757.5	273.08	1035.8	59.96	1679.75	1051.82	1353.05	444.7	684.05
J02843_at	257.8	256.26	259.1	299.39	363.95	107.55	122.55	26.94	100.5	37.9	309.1	93.48	340.3	105.92	255.8	264.88	616.55	324.07	157.35
M57730_at	561.1	239.28	465.95	538.32	666.15	361.4	1927.1	1065.61	1085.25	813.81	694.5	395.56	1415.65	786.23	1774.9	1981.6	858.2	325.83	524.35
D80011_at	86.9	59.54	32.25	37.26	126.3	101.12	78.05	43.35	76.15	3.61	115.6	105.36	57.35	16.62	38.45	10.11	64.4	2.55	89.75
HG831-HT83	505.8	436.71	367.35	424.48	125.9	104.79	202.05	190.57	720.95	2.33	803.8	120.77	262.95	198.91	343	333.47	516.15	305.97	874.05
U14972_at	46897.45	25511.92	27050.2	31253.27	71311.15	6512.1	80493.4	29078.92	73134.4	29377.46	54926.8	2764.51	57917.5	29829.16	69507.95	6450.58	54011.2	6588.54	58231.3
D86968_at	858.6	48.65	872.85	1008.41	712.45	263.68	637	316.22	953.25	327.89	1095	17.39	855.75	270.89	1023.05	286.31	980.25	116.6	727.65
X61118_rna1	159.6	81.74	328.9	380	84.2	27.01	102.2	67.74	293.8	23.05	434.3	319.33	186.2	170.98	92.05	47.02	161.85	164.26	220.7
X52008_at	613.05	454.03	477.75	552.04	1325.9	384.67	312.4	47.52	1085	541.64	1269.65	150.4	1293.5	1428.78	1495.25	144.89	1202.9	812.89	554.25
M34309_at	380.95	144.46	146.5	169.28	1162.6	909.76	147.1	45.54	406.3	0.28	297.35	210.36	414.45	341.89	232.9	101.26	482.5	254.42	241.55
AF001359_f_	6272.6	181.58	2490.25	2877.15	6850.65	2238.06	5154.55	342.45	6683.45	1165.81	7345.8	2263.45	7160.55	66.11	6850	355.67	6293.45	386.86	5745.15
U32581_at	263.5	303.77	593.35	685.54	121.8	115.4	159.2	85.84	102.25	45.47	169.9	118.23	192.1	204.07	97.65	48.15	77	31.82	251.2
X87613_at	149.05	72.34	49	56.57	65.5	34.51	169.65	206.83	74.95	65.97	211.4	182.15	137.95	115.61	138.85	130.04	107.85	99.77	168.55
U64197_at	93.75	3.32	65.5	75.66	291.75	122.54	527.2	638.94	215.9	183	113.65	77.71	148.75	92.84	576.05	336.79	125.05	77.15	581.05

U09584_at	3278.1	498.51	1568.4	1812.03	3829.45	1144.17	4315.85	1655.41	3026.15	583.01	3431.8	743.88	3471.75	703.08	2236.5	85.28	3030.3	309.43	2998.15
U77845_at	345.05	272.59	308.85	356.88	245.15	151.25	414.8	120.07	402.75	293.38	237.65	14.35	369	333.9	538.6	595.38	459.1	256.4	243.9
HG3991-HT4	2534.6	317.49	988	1141.55	2496.8	836.51	2538.55	797.55	2240.05	200.46	3762.8	2518.15	2857.45	658.95	2491.2	1744.01	2512.8	2460.31	4128.85
X92521_at	726.9	169.14	568.2	656.48	821.3	266.44	785.35	303.28	562.05	466.76	661.9	495.82	295.2	278.03	989.9	264.6	613.95	11.81	399.5
M88338_at	3058.35	2739.4	1530.35	1768.12	1535	420.87	2644.35	246	2742.4	1262.47	4711.85	1114.19	4219.35	1450.63	4241.1	174.09	3803.5	720.4	3612.2
M81780_cds4	112.7	18.81	90.1	104.09	214.55	132.02	199.05	1.77	215.45	55.37	182.4	101.82	344.25	321.38	186.2	16.4	117.8	61.94	151.2
X12517_at	1085.45	688.23	491.85	568.3	916.2	169.99	1352.3	674.16	1378.65	1237.51	968.6	221.32	724.2	140.86	772.4	146.23	1276.85	882.96	560.6
U33053_at	558	315.79	124.6	143.97	793.15	300.03	498.45	482.6	436.8	302.64	768.25	92.28	630.45	477.65	771.3	65.9	632.35	403.83	847.1
D44466_at	2418.15	389.97	1859.75	2148.69	2406.95	52.82	1775.2	2134.9	3009.65	1064.97	3081.3	154.29	3363.2	485.78	3375.95	454.03	3863.9	2253.41	2534.1
L10381_at	53.45	4.31	26	29.98	80.75	27.65	118.8	137.74	88.1	63.5	103.85	24.68	85.05	12.8	326.75	229.74	69.05	11.53	63.4
X77777_s_at	158.65	117.17	83.7	96.73	328.15	136.97	120.15	69.93	269.1	132.51	164	51.9	72.65	50.56	103.55	10.54	145.15	15.63	144.5
L38961_at	1520.35	215.31	1264.55	1461.09	1372.35	761.48	1108	309.85	863.55	598	1667.6	1163.33	1251.15	734.61	1088.85	121.13	1929.7	450.99	1787.2
D50487_at	577	182.15	410.15	473.83	675.75	260	323.05	134.7	431.55	305.12	1042	141	706.35	558.69	328.1	271.81	600.75	412.88	554
X95073_at	246.1	212.98	285.9	330.36	915.1	589.3	137.45	116.32	592.4	347.05	343.4	431.9	256.15	322.94	208.05	236.24	267.5	330.5	166.75
U57627_at	612.95	171.9	420	485.22	699.65	595.45	418	135.06	901.25	333.26	422.6	496.39	125.8	18.81	210.45	157.9	604.7	148.78	357.5
U94586_at	3644.95	803.63	1094.75	1264.8	3641.75	1747.05	3448.85	1658.24	2849.15	464.64	2596.15	398.03	2547.65	74.6	2721.05	91	2681.55	432.11	2423.5
D21853_at	6002.2	86.41	3417.65	3948.7	5689.1	1157.53	6672.45	2754.82	5408.3	956.15	5504.55	570.99	5845	160.23	5836.7	410.97	5828.75	254.06	5259.55

X00351_f_at	197359.69	73217.52	66900.35	77295.19	103946.25	43754.29	119525.95	11246.61	93770	34318.02	85827.75	2645.64	131204.2	44577.14	112691.1	27645.61	137487.2	29148.07	96670.1
M21119_s_at	202.45	230.73	31.8	36.77	135.25	132.58	749.2	1005.79	151.6	94.75	63.6	53.32	85.2	39.03	63.25	29.34	211.1	270.68	160
D28114_at	701.1	36.91	538.9	622.68	712.7	29.84	444.9	86.83	880.25	315.58	710.95	18.74	503.7	470.93	690.75	277.12	775.05	408.5	632.45
L11284_at	2557.4	577.99	1999.5	2310.12	2746.7	66.33	2285.4	15.13	2273.5	14.14	1913.7	55.15	2335.85	759.22	2637.2	206.9	3202.7	274.92	3123.4
HG2874-HT3	1668.8	738.5	1211.15	1399.29	2384.2	597.08	2443.7	777.11	3534.8	824.35	3791.9	31.11	2839.2	1950.2	2706.55	798.54	3155.9	2006.06	3082.25
X53002_s_at	2741.05	400.01	1572.85	1817.19	1875.4	1666.37	1997.2	121.48	2997.4	1017.24	2773.2	215.38	3035.9	1154.56	3397.8	1442.07	4538.4	485.08	4621.2
M87503_at	1363.1	1480.68	1149.2	1327.81	2288.85	80.82	2112.5	1164.46	1010.05	892.72	2428.1	1185.39	2258.75	602.38	1204.2	1111.57	1850.65	696.85	2838.2
U28758_s_at	127.15	58.19	44.85	51.83	167.1	118.37	355.4	288.78	391.1	23.48	1308.95	821.3	565.3	115.4	466.75	286.02	423.9	364.02	262.6
X81625_at	1587.65	387.71	963.4	1113.13	1311.2	203.93	1381.3	162.35	1504.15	32.17	1567.6	685.89	1931.9	89.66	1738.9	31.11	1668.6	443.5	1737.5
M37583_at	4541.65	96.52	2427.6	2804.81	4052.85	213.62	3727.1	2574.58	6957.1	1674.29	4237.6	2045.52	5295.4	664.26	7155	2863.36	6038.3	3093.17	4202.95
L29277_at	1175.25	170.2	753.05	870.09	1931.3	1.56	612.3	734.26	1189.35	403.26	1019.35	17.75	1258.05	68.38	1461.8	287.51	1729.3	290.48	1633.65
X66436_at	2214.4	193.04	1537.65	1776.61	2857.05	177.84	2966.6	122.19	1783.75	1050.83	1500.25	1025.38	2980.5	139.87	1749.8	478.85	2311.9	308.86	2615.65
D00017_at	92255.85	23316.35	72798.45	84109.72	107318.9	15078.2	98548.95	19008.23	123167.5	6105.72	88796.3	14901.71	114716.25	9929.83	126655.7	23666.45	116975.1	4706.65	94072.25
U24183_s_at	1594.9	506.15	1126.4	1301.36	1742.05	163.98	1773.55	66.96	1775.65	66.26	1897.95	388.56	1648.45	378.8	1847.35	767.99	2218.95	805.47	1543.9
D31883_at	2820.45	588.24	2120.15	2449.63	3274.05	478.22	3705.4	825.62	2951.2	552.96	4120.65	1166.23	4015.4	46.39	3506.35	1248.82	2868.5	681.79	2852.65
M18731_at	145.4	103.24	80.75	93.27	148.25	86.62	136.65	90.72	83.1	19.23	199.35	32.88	160.5	3.96	236.45	53.67	86.1	11.88	174.55
D83542_at	3625.25	1101.18	1254.7	1449.71	4060.55	2265.5	4549.5	2200.94	2666.1	141.99	4809.65	1706.18	4030.15	786.23	3829.9	237.45	2388.85	458.84	2917.95
U25975_at	557.5	321.88	268.7	310.42	517.05	338.49	511.6	85.56	654.8	309.15	682.7	130.96	741.45	137.25	167.25	199.62	503.85	18.88	370.05
Z50781_at	3413.25	824.42	3016.75	3485.54	3124.2	1116.66	3293.8	737.23	2863	1337.99	3936.2	1014.84	3039.3	2310.82	3402.95	622.32	3056.65	444.7	2938.05
X51417_at	1087.95	871.23	271.45	313.6	736.6	19.23	1209.85	1146.43	430.9	60.39	1638.05	690.91	832.25	449.51	577.9	243.67	874.7	676.56	378.65
M86849_at	1145.75	274.71	353.7	408.71	986.05	216.87	626.55	720.75	999.15	146.16	531.65	476.38	1044.75	376.96	836.3	360.48	1138.3	285.95	747.95
HG3729-HT3	377.5	158.53	297.3	343.51	488.1	492.15	318.35	114.9	540.15	625.72	523.55	508.48	1178.1	479.28	469.9	467.96	313.2	20.22	893.1

U73167_cds4	394.75	479.49	28.65	33.16	339.25	213.76	80.85	39.53	469.6	428.93	102.85	4.17	105.05	75.87	98	6.93	78.15	12.37	272.45
U06643_s_at	1183.4	1242.81	663.25	766.29	1960.3	1715.02	2595.35	573.53	3521.4	1084.84	4736.75	14.92	3064.6	2387.9	4127.5	2884.71	1139.8	1032.38	3328.85
M90516_at	1218.7	8.06	592.75	684.83	1051.05	56.36	648.95	342.88	930.05	1.48	658.5	781.78	536.15	450.07	572.15	135.13	811.8	272.09	1076.95
D50916_at	298.1	187.24	585.7	676.7	237.25	163.13	1225.65	569.57	1174.5	48.37	340.6	266.86	262.75	145.88	290.4	233.35	667	246.78	660.2
Z74615_at	1397.35	157.76	666.35	769.83	2115.4	1281.84	1907.15	1382.04	1312.3	328.1	2060.9	550.69	1284.35	318.41	1141.75	1025.8	1429.2	45.25	1597.65
X51408_at	442.05	98.08	248.65	287.3	199.25	143.33	538.5	540.8	802.7	487.76	462.1	69.3	487.6	338.7	284.1	288.78	776.85	145.17	233.85
AB002366_at	53.6	25.31	173	199.83	627.5	627.49	122.15	79.83	172.5	164.19	94.8	51.9	97.95	58.9	254.55	281.64	465.05	46.46	296.05
D88378_at	418.95	99.91	440.2	508.55	425.55	396.33	471.9	273.23	491.4	263.19	625	332.91	300.5	273.08	387.3	120.77	304.95	349.24	562.85
M74290_at	119.35	62.58	49.95	57.77	208.5	108.75	128.2	80.47	240.4	93.34	166.05	87.89	186.7	96.17	150.95	2.47	113.15	35.57	166.65
U55054_at	3848.55	1750.3	2750.15	3177.53	2811.95	764.31	2239.25	1928.21	2631.7	196.58	2984.7	201.24	3996.65	117.87	3167.5	411.54	4061.5	411.68	3442.6
M80333_at	640.3	414.65	471.1	544.33	902.95	284.61	435.7	44.55	1464.45	944.06	860.25	333.83	560.75	469.02	605	211.42	346.35	106.14	792.3
X14346_at	255.25	92.56	123.15	142.34	119.9	51.48	546.7	653.79	104.4	64.35	150.1	74.81	120.5	31.96	215.5	93.9	129.95	24.96	149.1
X12791_at	1443.25	308.93	1171.25	1353.19	1290.25	63.71	1022.15	372.43	1579.1	74.39	1641.2	561.73	1463.95	427.02	1690.9	225.43	1673.6	642.05	1496.05
X82676_at	144.85	17.47	82.55	95.39	141.35	49.85	144.85	115.89	101	20.22	295.75	23.97	338.25	208.38	282.9	178.05	258.35	112.22	391.3
D14695_at	3367.65	1404.38	1898.45	2193.37	1611.05	586.55	2617.85	218.43	2664.5	552.11	2578.45	536.62	2409.9	195.59	1830.6	1.13	1908.55	44.19	1890.75
M57230_at	52.55	23.41	11.65	13.51	60.4	50.77	118.75	103.73	54.8	14.42	32.95	13.93	180.05	203.29	42.8	11.31	15.3	5.09	62.7
U40279_at	815.8	328.8	617.55	713.54	1208.35	953.67	1019.1	531.32	1143.9	635.26	957.55	112.92	942.35	335.81	921.35	474.4	988.25	269.9	529.45
D63878_at	2153.85	1214.6	1058.15	1222.52	1789.75	31.61	1982.9	37.19	3412.55	474.82	2649.4	767.21	3114	387.35	2447.4	5.37	2263.3	1095.59	2116.25
X82634_at	657.75	90.86	318.65	368.19	1207.95	322.09	605.25	95.67	984.8	799.45	935.9	521.42	918.1	56.29	1106.55	132.02	1177.35	460.82	955.75

X05232_at	153.1	149.34	48.85	56.5	189.1	156.55	243.85	7.99	359.65	361.26	378.6	299.25	429.65	166.81	84.05	6.86	124.4	5.37	77.25
D31767_at	2940.7	1233.62	1385.3	1600.61	3352.3	999	3000.35	1037.11	3304.9	493.28	2754.5	432.61	2988.75	523.9	3280.6	1254.97	2797.15	598.85	3075.3
X15729_s_at	4594.3	97.72	1885.95	2178.95	3058.3	1206.47	2772.5	2140.84	4092.15	793.44	2589.65	546.24	3287.15	1254.34	3884.75	835.45	4207.15	867.12	3230.3
M99701_at	419.65	100.34	359.85	415.71	460	422.99	778.15	416.27	1059.15	521.77	600.25	248.97	836.1	214.82	758.8	390.61	634.5	23.19	664.75
U04285_s_at	479.15	352.35	597.8	690.7	445.9	319.33	467.7	296.14	911.35	217.15	273.95	138.81	506.65	521.21	1073.75	394.49	927.35	389.4	524.6
X04085_rna1	542.15	62.86	456.55	527.43	927.3	109.6	1040.55	639.86	886.85	189.01	406.05	478.64	792.6	218.78	805.2	281	1190.45	582.59	711.8
U91327_at	3551.05	12.66	1098.8	1269.54	2983.6	593.83	2749.25	1054.23	2475.45	1694.72	2953.8	371.8	3363.3	1282.97	2600	617.87	3175.85	838.98	2767.4
U49973_xpt2	285.7	373.07	27.6	31.82	34.7	25.03	83.1	65.9	173.3	155.28	53.35	34.58	184.9	130.25	576	19.94	23.25	10.25	42.3
AF015910_at	186.15	33.3	214.55	247.84	443	173.81	761.15	90.44	394.95	114.62	241.3	10.32	238.1	48.65	249.05	44.05	203.3	1.84	473
U18235_at	440.95	215.17	228.6	264.18	527.5	73.68	769.55	639.15	311.2	66.61	668.9	516.47	402.3	28.71	523.55	237.8	433.1	54.59	370.05
M21812_at	2124	562.29	430.85	497.73	1889	68.73	1948.35	784.39	1860.85	774.35	1564.65	374.98	2058.55	351.22	1854.85	250.39	1429.5	779.37	1783.15
M19154_at	598.3	311.98	243.1	280.86	135.6	102.25	151.4	12.3	271.45	281.22	408.2	319.05	169.35	124.24	141.45	137.53	364.5	198.41	116.15
J05614_at	4629.85	425.61	1907.2	2203.49	4602.3	1171.82	3242.95	320.11	6875.25	784.11	6369.6	250.03	8000.3	1412.38	8572.45	2226.33	7615.35	4150.36	7030.1
Z48054_at	1104.15	33.59	677.1	782.34	964.8	322.02	798.05	403.69	764.05	483.31	771.25	185.19	802.85	31.75	411.9	284.4	530.75	364.65	1015.95
M28879_at	392.3	140.71	102.35	118.3	368.75	93.83	164.3	151.6	119	24.89	74.25	15.49	276.55	294.79	231.4	237.16	351.2	219.2	398.2
U26174_at	87.1	92.21	20.75	23.97	151.25	62.01	146.3	124.31	521.55	680.31	695.1	853.05	130.85	22.98	101.8	60.53	44	3.82	55.2
M11722_at	190.6	11.31	444.35	513.43	167.15	195.8	134.05	76.72	541.2	424.12	221.6	46.24	370.6	175.5	1572.05	444.42	166.9	138.88	434.5

U60665_at	37.8	4.81	46.45	53.67	48	40.02	40	30.83	37.5	24.75	46.6	27.86	32.2	2.83	75.8	47.38	60.55	34.44	36.5
Y07707_at	40.05	28.21	14	16.12	73.75	8.13	46.75	31.61	312.75	45.04	84.15	77	112.15	105.01	65.9	13.58	96.85	92.84	87.3
V00535_rna2	42.7	26.16	35.15	40.66	942.7	990.94	144.85	111.09	126.1	109.74	261.7	310.42	81.1	2.26	406.6	523.54	174.8	147.93	87.65
X97335_at	474.85	87.61	87.4	100.97	324.3	53.74	280.65	1.77	585.5	435.01	378.7	310.84	338.05	109.11	675.2	72.55	263.65	43.91	383.9
U83601_at	226.5	64.63	155.35	179.53	215.25	58.9	322.1	110.31	179.65	39.81	281.95	150.68	843	842.45	317.5	136.61	290.95	108.82	429.5
U81523_at	102.2	15.41	71.05	82.1	329.9	166.17	54.75	27.93	164.3	48.79	418.6	84.29	501.5	73.82	104.95	25.95	314.35	34.01	59.95
HG721-HT48	1132.25	494.62	571.1	659.87	441.1	192.62	758.65	279.52	784.85	778.74	598.6	97.3	974.65	190.57	1342	351.71	1269.1	206.9	803.55
U13616_at	129.55	147.71	90.2	104.23	153.35	12.09	310.3	417.33	55.2	52.47	35.4	27.86	38.75	27.08	61.25	62.15	129.95	83.65	31.25
M13485_at	1369.55	326.33	275.85	318.69	1040.6	965.48	1549.2	833.25	913.95	417.12	1285.4	561.44	1017.3	633.43	1045.8	213.83	1977.25	272.59	922.25
Z29678_at	487.85	119.43	436.45	504.24	866.55	444.56	781.9	29.98	271.8	93.34	803.05	337.36	618.35	14.5	1034.35	207.54	724.35	187.88	957.6
L23333_s_at	3241.95	2322.21	940.65	1086.75	3929.35	191.13	3176.95	1289.97	3028.3	80.75	2051.4	1340.39	2236.35	566.18	2549.6	44.26	2760.95	458.56	2578.3
Z97074_at	1307.2	759.72	981.4	1133.92	2184.8	330.22	1720.1	93.76	1538.1	513.36	831	426.53	1380.8	198.27	1085.2	458.91	1428.35	425.32	1030.55
D63478_at	2665.25	383.61	1533.7	1772.01	2371.75	88.32	2220.65	12.37	2998.5	8.91	2815.6	503.18	2665.15	463.79	2718.2	958.41	2733.2	675.71	2918.3
Z49995_at	151.85	66.26	36.75	42.5	188.05	68.94	266.45	241.48	88.45	9.55	192.05	50.98	135.95	21.28	399.45	352.63	150.9	61.09	131.25
X64072_s_at	193.65	16.76	127.35	147.15	201.5	141.7	467.25	531.25	233.4	35.64	194.85	13.93	135.1	40.31	195.9	60.81	213.15	104.86	219.65
L39061_at	188.15	32.88	184.55	213.19	636.1	808.22	27.55	15.34	552.7	491.86	322.75	248.41	386.25	457	372.85	103.59	379	43.7	346.3
D38024_at	4525.9	120.49	1454.4	1680.37	5130.35	1998.5	4581.55	244.59	4311	1705.26	5348.1	2636.24	4728.8	823.64	4707.75	1011.94	4642.7	158.25	4355.25
U76272_at	120.55	29.06	39	45.11	242	166.88	617.95	562.22	112.1	66.75	405.1	399.94	86.7	38.04	53.7	18.38	145.65	94.68	102.65
L41913_at	181.65	221.96	32.2	37.19	260.3	313.96	481.1	488.33	127.3	55.58	114.75	58.9	45.1	12.02	188.15	173.31	34.6	4.1	101.1

S58733_at	1288.25	80.96	752.7	869.6	1679.2	1002.11	888.6	257.1	1482	108.75	1025.25	1007.98	1363.1	123.46	320.8	165.89	695.85	503.39	926.85
HG4243-HT4	859.2	659.31	169.6	196.01	774.05	323.5	251.75	62.01	649.6	496.25	1106.8	72.27	266.05	58.9	1086.65	645.38	923.3	823.78	627.85
L12468_at	170.6	173.24	126.95	146.72	127.8	59.26	83.8	11.46	475.85	183.49	219.75	2.05	335.05	125.79	291.75	239.21	174.8	135.34	48.15
M14306_at	52.2	1.41	11.1	12.87	60.7	3.39	132.9	51.9	90.25	39.95	45.25	7.14	93.1	35.21	248.25	332.13	223.3	159.52	243.9
M23263_at	930.15	227.9	741.9	857.15	435.85	95.95	491.35	86.62	784.85	853.27	369.9	278.88	620.75	496.04	767.2	311.27	848.9	19.23	636.2
D28588_at	651.7	74.25	404.75	467.61	593.95	165.82	232.7	214.39	349.8	246.36	397.1	386.65	259.85	178.54	469.95	148.56	794.4	3.25	581.35
S83308_at	511.7	43.56	346.5	400.36	1293.9	348.04	299.6	330.64	541.3	484.79	585.95	44.34	527.4	525.8	208.15	42.78	105.35	41.08	555.4
X04500_at	6465.7	1880.34	9973.9	11523.58	4112.15	202.3	5229.75	2308.49	11202.1	4160.19	11477.75	5314.4	7643.8	6549.65	5462.9	3663.38	3661.75	2118	3154.25
D87437_at	1368.25	210.65	1116.35	1289.83	1434.8	229.24	1256.3	289.91	1513.6	223.45	2160.7	575.02	1581.8	886.15	1509.25	486.84	1664.05	674.08	1646.75
Y08417_s_at	343.15	184.91	401.9	464.29	132.45	66.4	267.05	148	326.3	101.82	241.25	1.77	498.6	391.17	287.05	317.84	453.15	140.93	168.35
M86400_at	14544	1849.23	6401.2	7395.77	11202.4	4363.27	10936.7	7540.3	12336.2	683.91	14168.45	8947.94	12071.8	2145.93	15525.95	1150.82	15028.65	3404.65	17891.5
J02943_at	366.7	269.83	54	62.37	81.1	2.69	408.2	209.16	387.95	372.86	528.7	585.48	775.9	449.44	531.15	117.73	532.55	322.51	721.8
U09607_at	728.05	186.61	771.3	891.1	2244.05	1996.37	449.75	126.08	1555.05	1585.55	1280.35	720.61	1980.25	746.49	1069.65	829.79	1372.1	1218.2	1386.6
U02683_s_at	117	91.36	103.35	119.43	80.85	41.79	102.45	98.92	108.45	95.39	167.35	78.98	114	42.28	201.15	68.38	88.3	24.61	174.9
X52541_at	616.1	95.32	1953.25	2256.73	1635.55	246.71	346.65	147.43	1010.3	151.6	2128.3	933.81	1210.15	39.95	635.6	316.36	555.5	21.78	2044.4
U32849_at	390.15	130.74	189.65	219.13	632.9	366.85	846.9	399.23	481.45	68.52	354.2	118.09	447.6	118.09	645.1	67.74	566.5	221.75	277.9
X83543_at	247.2	58.41	409.4	473.05	316.7	9.9	197.15	86.62	850.4	780.22	410.05	164.12	578.7	499.08	970.5	220.76	661.45	552.75	336.9
D50923_at	322.75	169.07	530	612.35	366.75	62.15	282.4	3.54	239.4	45.54	336.35	125.23	387.95	202.44	319.25	26.94	440.15	214.32	338.95
HG1723-HT1	151.5	23.33	294.4	340.12	528	543.34	370.65	137.39	782.8	582.66	274.3	136.75	720.15	750.45	161.85	7.42	324.85	298.19	474.65
X58723_at	359.95	26.66	339.8	392.59	355.35	132.87	333.3	290.48	367.7	74.95	501	290.48	501.55	50.98	280.7	2.97	461.45	56.21	438.8
U10690_f_at	717.65	58.62	557.8	644.46	546.95	279.38	521.15	229.74	923.85	12.09	1129.35	224.79	707.35	741.83	850.05	504.1	745.1	319.75	520.6
HG3925-HT4	1288.05	607.05	511.35	590.79	1464.65	342.17	1548.5	721.53	1979.25	1311.47	1513.75	112.92	1927.9	650.4	1374.9	296.28	950.4	342.38	1673.2

M13150_s_at	307.5	71.13	317.8	367.13	165	61.94	387.65	14.78	543.75	185.62	677	100.97	399.35	362.68	482.25	315.58	527.5	388.77	745.4
U26710_at	417.2	335.88	645.75	746.07	727.6	540.65	391.5	290.06	312.3	52.04	436	344.93	362.6	287.79	457.15	114.48	502.9	60.25	407.8
U18242_at	522.4	144.11	145.05	167.66	972.85	307.66	474.15	30.9	973.25	263.54	1005.55	606.77	435.1	459.9	952.15	472.14	406	5.8	1010.05
U79280_at	295.5	166.59	694.1	802	295.95	136.54	369.75	400.29	205.75	19.02	271.3	45.4	299.25	32.31	2489.9	3194.28	294.7	45.4	276.55
U31201_cds2	16937.25	8301.5	12950.6	14962.8	9425	4444.87	9446.85	2057.04	9341.65	950.99	7321	2219.61	10005.45	3023.52	7374.35	585.7	16399.85	3933.71	14307.25
X51730_at	37.1	14.14	72.15	83.37	54.35	61.16	38	46.67	105.8	140.01	49.65	20.29	120.25	130.46	190.75	87.19	122.25	158.32	10.75
L37378_at	372.85	395.34	357.9	413.52	132.15	88.03	160.7	62.65	702.1	45.25	302.5	70.85	146	20.79	130.75	22.56	142.95	17.32	279.6
Z67743_at	1812.15	1115.04	551.9	637.67	1656.35	1151.52	1855.15	998.36	1289.2	656.2	1651.25	302.85	1471.25	935.43	1640.75	1030.18	1411.65	478.64	1148.4
X75958_at	376.4	48.79	133.5	154.29	408.1	71.13	542.75	467.19	190.4	95.74	612.45	119.15	312.65	131.45	104.35	82.1	36.5	6.22	178.6
U59752_at	2056	776.83	945.2	1092.06	2349.15	827.67	2022.8	858.85	1832.5	2007.05	2800.75	2343.85	2843.35	582.16	1815.9	134.49	2093.15	286.17	3554.55
Y08612_at	562.85	6.01	352.8	407.58	513.6	77.07	665.25	336.37	364.65	178.69	485.85	189.01	707.55	118.3	413.75	18.03	980.75	195.52	752.2
L20941_at	23229.35	10963.48	13851.2	16003.38	28938.5	708.66	30938.4	7276.98	21199	7234.13	19418.5	5810.3	14445.25	4066.64	18861.3	3115.94	13551.8	1695.78	16257.9
M37485_cds1	84.25	31.61	49.55	57.2	109.25	20.15	112.15	101.89	74.7	11.03	144.3	36.35	137.8	69.01	75.75	4.03	139.1	26.73	87.75
L38932_at	2953.4	155.14	1619.55	1871.22	2995.95	776.05	3059.75	259.86	3931.45	132.16	3661.85	695.86	3589.6	843.01	3056.15	1648.48	2674.85	154.93	3437.1
L18920_f_at	43.8	21.92	16.75	19.3	428.3	562.57	61	52.75	140	162.78	175.3	108.75	286.4	209.02	51.1	13.29	74.5	70.71	69
D50920_at	1169.1	305.61	531.55	614.12	1924.3	970.86	1472.95	62.01	2012.35	1008.83	1866.1	204.21	1976.75	239.21	2461.15	714.53	1641.2	922.35	2202.35
HG1071-HT1	458.6	55.3	432.25	499.43	206.75	202.73	477.25	202.59	254.05	237.94	612.2	234.76	642	590.72	510.9	149.77	494.25	190.28	398.85
U30246_at	350.45	203.29	59.4	68.59	461.85	166.52	105.95	127.92	362.8	385.51	651.3	363.31	107.3	61.8	59.65	12.23	99.75	33.16	782.1
X55448_cds1	6612.35	2293.64	2599.85	3003.86	3436.25	1551.75	5858.9	1934.08	4718.9	413.66	5651.65	380.21	7225.5	1948.65	5632.3	2527.48	6971.5	1319.89	5274.9

X02761_s_at	68940.4	14336.59	48309.15	55815.41	50752.4	9351.77	59316.35	1209.37	72485	26387.1	51324.45	13878.6	52023.9	3883.72	45342.4	10448.49	44434.6	21923.56	45140.7
X68264_rna1	256.7	51.62	243.65	281.5	712.15	2.62	202.2	10.47	192.45	109.96	552.25	542.28	616.6	360.34	403.2	412.38	181.15	97.37	860.35
U56816_at	4391.1	395.84	2553.85	2950.69	4403.55	487.41	2235.4	1688.01	4173.55	1159.73	3669.2	1403.89	3575.65	1105.7	4649.35	1834.45	5893.75	2403.1	3908.95
D30758_at	1499.9	1300.23	1652.8	1909.61	1337.8	763.53	3273.75	1779.15	1900.9	788.99	2276.45	323.08	1378.35	722.03	1848.8	1995.46	1342.8	323.71	2108.7
U96113_at	781.5	306.46	171.4	197.99	991.15	192.97	627.25	41.22	1094.85	111.79	1146.05	28.64	714.85	77	615.7	798.32	988.25	289.14	1200.1
D45399_at	112	35.64	369.35	426.74	86.2	29.56	254.7	270.4	90.9	100.97	263.4	10.04	51.6	32.53	100.65	60.6	394.75	239.5	448.15
L07541_at	182.4	12.02	35.2	40.73	229.9	11.74	264.8	235.04	567.05	169.63	292.05	240.35	307.55	215.74	522.2	249.61	438.45	435.22	347.2
M17254_s_at	155.15	5.87	146.25	168.93	158.75	2.19	947.1	1043.83	207.7	83.58	206.1	155	599.1	624.94	294.6	36.35	618	582.94	302.6
D21163_at	824.4	202.52	409.85	473.55	533.35	228.04	1296.25	2.19	1048	278.88	1500.4	203.08	1046.2	784.46	882.95	619.21	1212.75	232.43	1071.55
D21852_at	700.1	222.88	520.95	601.96	813.65	97.79	1244.55	774.78	899.75	41.22	974.55	105.85	1233.55	533.79	1072.9	254.42	1124.1	252.3	602.9
J03077_s_at	21769.85	5378.47	14312.3	16536.12	14759.15	4287.54	18531.75	5261.79	19660.4	5321.83	15263.75	3633.04	16102	1498.08	18372.2	2123.3	15764.2	2163.32	18695.65
U58089_at	1040.6	10.32	738.6	853.34	758.25	281.5	786.7	9.33	1215.2	73.4	851.9	91.78	1160.35	121.98	926.3	103.24	1181.15	174.44	1350.8
U49187_at	84.75	30.76	80.55	93.13	230.15	42.21	114.4	102.81	153.1	148.21	74.3	13.58	109.85	81.67	80.25	11.1	469.85	85.77	358.45
U16258_at	193.1	38.75	45.85	52.96	174.55	45.61	303.2	102.25	231.75	111.93	501.95	297.48	666.95	522.06	551.15	488.54	159.6	109.04	199.25
X84194_at	841.5	392.44	397.05	458.7	842.9	78.21	306.95	77.15	871.95	304.83	978.7	661.71	670.15	112.08	493.2	132.51	533.05	225.5	249.2
M62762_at	4774.4	1138.44	3540.7	4090.9	4540.35	47.31	5445	480.27	4760.85	1247.69	4180.3	1912.02	5156.95	732.07	5136.95	213.62	5363.5	98.57	3895.25
L13197_at	734.1	111.58	165.7	191.48	380.1	168.57	1166.55	493.49	900.3	69.72	1114.35	425.89	1242.65	798.39	572.35	323.08	580.6	283.13	883.6
L00022_s_at	301	144.53	50.7	58.55	214.75	46.46	408	348.46	302	182.72	147.5	2.97	213.65	192.26	374.35	114.76	242.2	155.14	156.25
M94172_s_at	2837.6	372.22	892.2	1030.82	2699	269.41	2413.55	201.17	2044	615.75	3637.95	1685.25	3017.6	115.12	2319.45	334.67	2248.35	946.32	2602.45
HG825-HT82	461	188.51	93.8	108.33	175.45	63.14	292	238.58	242.95	30.33	131.9	45.4	141.85	9.12	119.75	15.63	145.9	48.65	158.8

M85220_at	441.8	85.84	332.6	384.24	597.6	225.43	767.25	412.46	1123.2	1060.09	681.15	25.81	1047.55	612.57	574.1	69.58	465.15	114.2	1442.45
X12458_rna1	1753.1	916.98	900.9	1040.86	1553.9	153.02	1544.3	1034.5	1525.35	1119.28	2104.05	1150.67	583.85	149.55	905.1	687.73	1193.15	815.08	1890.6
L40384_s_at	103.35	72.34	155.55	179.68	159.7	39.74	105.5	63.78	219.25	52.54	248.75	166.1	677.9	80.04	92.2	23.33	100	10.89	67.45
X63187_at	1268.2	331.77	592.2	684.2	545.9	482.95	752.2	734.12	616.6	265.17	1165.35	261.7	835.95	545.96	996.8	159.1	773.6	640.07	1248.3
X54816_at	286.3	7.5	332.25	383.89	285.1	38.47	210.35	153.8	568.3	188.23	917.65	777.89	536.35	321.52	462.6	421.72	618	631.87	337.55
D89859_at	337.15	228.47	283.5	327.53	157.15	124.66	152.7	20.65	237.8	153.58	165.2	9.62	110.2	120.77	108.8	9.33	204.45	127.63	526
S80050_at	560.55	289.28	357.1	412.53	566.5	580.96	911.3	380.42	736.6	542.92	1119.8	984.43	304.75	104.16	741.05	805.75	850.1	496.53	802.35
U12622_at	891.35	544.26	341.65	394.78	377.8	144.67	701.8	673.73	1154.3	414.93	341.8	372.79	1013.55	808.86	478.05	12.8	354.5	320.74	404.8
X05323_at	325.05	175.29	79.7	92.07	240.95	212.63	226.65	22.7	343.25	371.58	865.45	194.53	500.3	290.9	290.55	21.28	334.8	252.86	246.2
U33838_at	436.35	177.98	139.95	161.72	177.45	218.71	58.15	58.05	18.25	12.52	232.7	264.6	71	81.32	446.5	70	39.15	46.32	72.15
Z49989_at	4403.8	2672.72	1653.35	1910.25	1280.3	1284.53	2488.85	357.44	1095.85	1298.32	1047.05	729.38	3529.35	988.18	2365	2379.7	5273.65	2157.74	2542.2
X97074_at	6409.75	2642.67	3135.3	3622.51	5209.5	590.15	6848.05	1902.05	5140.45	136.97	5346.45	85.07	6116.4	1791.1	6140.85	1770.95	5864.85	618.93	5810.55
U46116_at	42.1	8.06	121.8	140.71	356.15	420.52	86.65	62.01	85.6	14.14	444.95	552.89	122.8	4.81	48.7	29.7	180.3	185.83	190.4
J04177_at	532.15	83.09	342.7	395.98	887.6	86.13	345.35	229.17	993.35	532.8	802.1	404.04	803.8	13.44	718.4	194.45	782.7	371.37	563.25
D86962_at	936.45	369.32	416	480.69	719.4	227.41	300	252.86	828.35	299.32	1430.3	200.54	516.05	288.99	902.45	625.15	821.3	187.67	1209.3
X67683_at	5445.95	437.63	2088.3	2412.79	6065.2	638.09	5303.55	1279.23	3772.55	2544.95	5375.8	745.86	5217.35	981.53	5566.15	455.59	5429.95	1614.54	5220.45

U79272_at	270.25	140.22	180.45	208.53	63.8	7.64	87.8	45.68	197.15	13.51	268.8	131.8	180.75	161.01	233.15	5.87	245.75	260.43	454.1
M16653_at	5430.15	272.59	4790.45	5534.74	12480.85	12742.14	12285.75	11765.48	5217.75	1863.86	10035.5	7554.16	6031.45	1658.24	6030.35	1410.89	4253	275.77	3958.9
M88458_at	2876.15	345.56	2127.4	2457.9	3601.95	278.67	3020.9	2011.86	4053.95	52.26	3602.55	704.77	2713.45	691.62	2934.1	725.07	4677.25	1044.33	3702
D13988_at	2901.75	928.36	2674.35	3089.84	3958.75	1118.29	3565.55	1240.9	2693.5	1225.56	3396.6	527.22	2715.45	445.97	2856.65	1822	3200.45	756.25	4017.95
X14675_at	1097.1	349.17	710.95	821.45	731.6	17.39	1467.5	756.75	1162.95	9.69	1771	68.17	1202.2	1071.97	1774.7	700.74	1503.65	114.76	1440.05
AB004884_at	729.95	401	692.8	800.44	1382.7	684.48	1288.05	575.23	1348.35	170.48	1255.55	123.81	1242.5	204.07	669.55	546.95	813.6	33.38	1276.15
X57351_s_at	12104.45	2813.51	5118.95	5914.31	13232.6	5129.21	10819.85	3681.41	18739.7	898.73	15699.9	7683.85	12818.75	3906.84	12237.45	4511.27	13814.95	4830.46	18633.55
AF005037_at	527.85	101.75	432.8	500.07	1847	1774.56	1325.6	1474.03	1162.35	437.77	939.45	639.72	897.75	456.01	649.2	216.94	571.4	18.95	652.6
U66048_at	761.9	833.11	101.5	117.24	716	35.21	323.1	158.96	909.3	58.27	592.15	617.94	625.2	627.63	791.7	794.79	903.15	412.88	622.4
X51956_rna1	173.85	67.39	54.5	62.93	119.2	67.32	141.15	38.11	145.55	143.19	163.7	135.2	99.25	8.41	80.25	51.83	111.2	19.23	72.55
X98311_at	362.3	259.37	158.9	183.56	848.15	128.06	616.65	265.52	872.25	182.93	646.15	236.1	1224.05	75.45	793.8	855.88	1410.4	211.85	505.25
Z48804_at	267.3	216.8	37.4	43.27	187.15	132.72	109.25	20.58	597.8	44.83	322.1	339.69	192.9	117.95	274.6	179.18	201.7	83.3	341.55
L25851_at	282.3	113.28	405.6	468.67	477.75	447.67	388.2	444.91	1463.55	557.98	560.6	581.38	877.3	61.24	1115.35	176.99	484.3	605.85	902.2
U69263_at	235.85	201.74	202.7	234.19	124.3	67.88	739.4	957.85	308.5	326.68	858.3	131.8	452.05	302.43	487.45	366.92	666.35	160.16	287.55
M25164_at	302.25	309.36	283.15	327.18	137.75	86.76	491.6	174.66	142.25	87.19	669.2	343.23	647.25	219.42	569.9	163.2	276.35	260	222.85
X73460_at	67627.55	27051.29	40078.25	46305.53	83478.6	14474.48	94908.45	32557.95	80923.85	9455.93	73848.2	5001.08	73678.6	24770.37	76577.6	13778.68	75388.1	21165.97	82115.25
X68149_at	319.6	152.74	171.25	197.92	517.4	521	856.35	74.32	217.2	88.11	872.45	950.42	673.65	351.36	699.55	102.88	512.3	167.87	419.95
U03642_at	502.8	142.41	297.55	343.72	579.65	192.83	533.1	37.48	408.65	310.35	890.3	738.22	689.2	20.79	891.5	395.41	674.8	199.12	629.25
L38490_s_at	2527.9	252.01	2236.9	2584.48	2564.7	360.62	2659.6	735.96	2741.85	673.94	3677.05	545.82	2674.4	332.91	2906.5	364.73	3324.6	764.38	2953.4

L38487_at	2361.95	309.5	1265.8	1462.44	2596.45	256.47	2714	758.16	2486.7	38.18	2308.75	379.79	1260.4	186.68	1775.6	316.5	2572.8	579.83	3196.65
L13203_at	2249.35	129.61	778.9	899.86	2196.85	498.72	2377.4	950.78	2305.5	1119.92	3148.5	13.72	3158.65	513.85	2433.4	923.2	2024.05	147.86	2403.6
U52426_at	1238.3	299.11	1087.6	1256.53	1658	184.55	1105.5	213.4	1261.7	821.23	1589.25	155.21	422.65	332.55	1386.1	494.97	1523.3	86.13	1482.6
M55905_at	957.8	72.69	439.8	508.13	1012.45	376.25	512.7	432.04	905.2	22.77	1050.95	262.12	693.4	395.13	512.9	166.03	763.75	91.85	693.8
X59770_at	523.75	443	226.15	261.28	696.15	614.55	495.3	130.67	596.8	84.99	149	76.08	425	152.45	604.8	134.35	573.2	298.96	218.8
HG273-HT27	49.05	7.57	73.8	85.28	95	63.5	400.55	533.94	69.05	83.09	119.7	57.7	153.45	151.25	62.35	11.24	34.1	13.01	54.95
U11036_at	171.95	114.62	45.25	52.26	269.6	204.07	145.6	42.14	164.35	46.74	206.15	133.86	97.65	37.55	113.1	33.38	130	72.55	174.1
M91036_rna1	512.65	581.6	155.2	179.32	1164.45	1156.33	287.5	94.33	499.65	133.29	932.75	43.06	831.25	289.7	130.55	49.43	299.95	262.97	205.4
U66828_s_at	875.75	543.13	1263.75	1460.1	1070.9	1055	519.75	14.5	757.3	368.12	638.4	119.64	649.9	152.17	749.3	533.02	923.3	394.57	431.65
X67325_at	173.65	24.54	42.3	48.93	304.8	187.95	125.7	3.54	165.95	35	165.75	9.55	128.85	1.48	149.5	73.68	245.6	95.6	241.75
U28488_s_at	57.8	10.18	73.8	85.28	233.8	127.14	107.8	80.61	235.65	55.79	52.2	11.74	120	32.81	83.05	17.04	104.05	0.49	93.25
X69116_s_at	121.75	38.82	149.55	172.75	142.6	3.11	92.5	52.18	529.3	74.95	115.7	53.32	82.25	54.38	207.6	177.48	114	88.67	362.3
M27281_at	1336.2	562.29	1040.5	1202.22	491.05	370.74	855.4	529.2	681.45	129.75	1581.9	248.05	602.1	435.29	526.9	556.63	730.1	360.34	1162.85
U34044_at	457.6	45.68	357.4	412.95	590.15	54.38	959.95	361.83	963	286.1	926.7	519.44	591.5	83.86	1271.95	629.54	776.45	136.68	189.65
HG3039-HT3	6720.7	1033.65	3145.4	3634.1	8230.95	1885.92	6851.15	260.71	5366.9	5.23	7143.75	1296.34	7692.4	1338.69	5554.8	268.42	5893.05	1583.71	6439.2
M36200_at	889.05	992.99	364.3	420.87	1234.45	747.2	596.8	323.71	710.8	55.01	914.2	146.51	1094.1	508.13	746.4	535	934.1	92.91	758.35
S70585_rna1	122.85	26.94	246.2	284.4	222.75	134.14	145.15	47.59	339.5	1.27	401.1	59.11	131.85	14.5	421.85	238.93	140.6	7.92	438.4
S80267_s_at	393	274.07	238.35	275.42	345.95	294.51	332.65	404.96	198.55	138.24	316.15	8.98	274.25	264.53	145.5	141	333.5	97.72	322.65
X87767_at	79.4	0.42	79.6	91.92	252.4	143.26	170.65	151.39	350.85	71.35	651.2	800.59	152.5	70.71	119.55	99.35	279.75	211.5	152.55
X14766_at	904.55	26.94	543.2	627.63	1361.05	798.39	1198.5	400.22	835.9	668.78	1165.55	37.97	1497.2	304.62	1117.4	231.79	1301.25	191.41	801.45

D87024_at	997.75	233.84	616.8	712.62	1726.7	103.52	988.15	453.33	758.75	719.91	1266.25	728.81	1278.65	239.36	1362.7	331.21	1587.65	153.94	745.65
L49054_at	430.65	86.34	511.45	590.93	297.35	317.56	897.9	209.87	463	83.44	530.7	645.16	635.85	73.61	452.65	6.72	735.5	168.86	733
M94856_at	1904	446.47	829.95	958.91	2081.25	81.95	1744.95	342.88	2213.15	74.18	2125.3	442.08	1798.85	284.89	2008.4	753.35	1294.05	32.88	1811.2
M17252_at	324	108.05	255.85	295.64	439.15	318.41	1534.45	916.62	347.7	37.48	434.15	201.6	343.1	69.01	909.55	852.13	460.65	259.01	277.4
U31248_at	628.15	258.02	85.2	98.43	1029.2	311.27	289	115.12	699.9	277.04	1031.3	536.84	569.3	32.81	685.35	429.71	365.2	225.43	774.85
M22489_at	269.95	292.53	33.35	38.54	250.5	174.23	242.9	18.1	131.5	39.74	220.85	143.19	109.35	13.79	106.95	69.51	73.15	4.6	639.05
D79995_at	389.25	233.27	62.15	71.77	725.1	454.67	374.35	91.43	739.75	197.49	493.7	283.41	744	519.86	763.7	822.79	598.1	553.24	488.8
Z35402_rna1	3956.35	505.51	2464.9	2847.94	2751.3	1160.22	2863.4	618.01	3951.8	1165.74	3059.55	71.49	3396.5	1041.43	3542	284.82	4128.6	540.09	3495.65
HG2591-HT2	271.8	16.83	228.85	264.39	364.4	64.06	439.4	405.17	496.15	70.22	504.1	130.39	241.2	75.66	303.75	44.48	448.3	25.6	359
U53445_at	120.1	6.08	40.9	47.23	360.7	393.72	394.8	288.92	142.15	91.43	292.1	265.87	175.65	176	155.35	89.59	240.8	271.53	244.45
U94855_at	7526.25	4182.32	4310.6	4980.44	9588.1	198.27	11058	3980.3	5635.2	1385.08	5914.8	714.88	6312.95	240.06	7793.7	801.58	8928.4	3171.09	8446.9
M28983_at	685	189.5	475.4	549.28	425.5	355.53	406.15	167.37	1040.7	244.8	719.35	146.72	654.8	220.48	550.4	9.9	591.7	53.03	1239.5
S71018_at	884.15	220.69	680.45	786.23	935.2	569.5	622.4	639.79	862.6	210.29	897.85	104.3	208.75	109.96	385.3	340.26	229.35	45.18	660.2
L11672_at	2897	30.26	826.25	954.66	2506.3	538.82	1931.55	949.43	2531.35	66.54	2871.4	770.04	3693.05	1492.07	1976.9	608.54	2409.8	199.26	2459.65
L48513_at	1390.75	168.22	1090.45	1259.85	1098.1	198.98	789.55	1029.62	1289.45	429.57	1149.75	71.49	831.7	319.9	722.55	312.05	1350.75	289.14	1328.35
X93921_at	5181.1	2702.99	1301.25	1503.38	2088.85	2442.42	3440.65	616.81	2352.15	1300.02	2249.45	2046.58	3027.9	1373.48	4300.85	1657.39	5335.75	1255.18	3689.3
U60276_at	842.6	522.83	611.5	706.54	934.9	732.7	1302.65	1116.59	641.55	395.91	681.95	532.8	1315	1285.38	1327.7	1377.16	2129.35	434.38	1059.85
D16581_at	3137.15	565.33	1796.1	2075.22	4272.75	349.81	4371.75	197.07	3721.9	522.55	2829.75	1019.15	4312.85	750.45	5059.3	361.19	4049.55	384.03	3561.5
X59434_at	2620.5	327.25	1562.75	1805.6	2067.25	659.52	2577.05	343.58	2209.05	11.95	3136.9	65.05	3006.4	131.66	3095.2	322.58	2310.4	1259.22	2592.85
U22816_at	201	0.28	245.9	284.12	82.85	21.28	103.45	89.59	449.3	197	228.3	28.14	260.1	14.85	414.3	496.39	181.45	199.62	174.05

U86136_at	381.7	264.46	518.7	599.34	518.1	414.36	152.05	37.55	413.45	529.98	752.55	767.42	418.45	431.83	157.75	58.34	120.9	27.58	110.35
X83441_at	496.05	250.1	191.7	221.47	445.2	247.63	179.4	218.5	609.2	116.67	368.6	169.56	312.9	360.91	398.55	435.37	106.4	81.88	168.5
J03798_at	325.8	146.23	354.6	409.7	803.1	554.09	811.25	434.8	501.25	34.15	560.2	167.02	341.25	383.61	542.25	62.72	379.75	82.66	737.35
Y09615_at	495.3	138.59	115.2	133.08	728.8	598.64	329.85	215.6	216.65	149.13	325.65	144.6	484.05	566.46	246.8	280.72	157.4	62.08	190.5
Y13115_at	159.7	99.56	274.35	317	275	244.94	197.1	107.62	628.25	157.76	703.05	514.56	392.3	323.43	279.95	370.03	400.1	2.4	198.55
S73205_at	303.7	193.89	227.75	263.11	79.65	47.59	327.6	404.89	93.85	57.91	137.2	30.55	216.8	197.57	43.45	24.4	322.5	245.51	158.15
D50919_at	319.9	345.35	279	322.3	293.4	20.79	233.15	173.59	299.6	273.93	141.35	68.52	325.3	273.08	418.45	503.25	564.35	270.19	826.9
D83776_at	249.4	0.99	200.65	231.86	584.65	278.39	369.9	454.53	434.25	267.78	249	122.47	276.5	77.92	487.8	27.29	212.45	34.44	488
D49958_at	238.45	230.02	259.4	299.67	133.45	148	83.45	0.92	166.9	186.82	462.7	347.9	222.1	236.74	225.4	180.88	395.3	151.32	232
HG1862-HT1	8093.55	2820.72	3401.95	3930.59	7011.8	3671.44	6009.95	2872.48	6615.7	2793.21	7556.4	2364.14	8643.55	285.6	7860.6	94.61	9993.05	309.36	8427.65
U12140_at	245.9	263.19	87.6	101.26	711.95	623.17	122.9	72.27	70.55	19.59	574.9	485.36	547.1	69.72	718.45	214.18	551.25	82.38	472.1
X15088_at	224.2	89.24	233.5	269.83	213.7	62.51	384.5	436.43	120.45	68.66	356.5	207.18	145.3	16.69	200.25	111.37	86.45	58.19	143.6
S67970_at	176.9	30.83	271.8	314.1	339.7	129.82	88.9	11.03	191.3	55.44	269.75	35.14	61.25	36.56	301.95	323.36	196.45	2.9	247.2
Y10812_at	159.55	40.38	61.75	71.35	219.15	5.02	142.8	89.8	266.4	99.28	149.3	23.48	166.25	37.83	178.2	22.91	200.9	56.14	124.05
U46767_at	257.5	175.79	41.45	47.87	410.8	77.22	671.3	412.1	124.25	112.08	135.65	97.51	258.15	226.63	712.25	384.74	402.35	435.08	643.55
HG1400-HT1	961.65	78.42	576.25	665.74	1079.6	419.31	600.3	171.12	1351.15	448.66	1011.35	84.78	970.2	82.87	970.7	142.41	1094.85	162.42	747.2
D79988_at	76.4	26.45	388.3	448.59	109.85	37.97	151	159.95	147.9	68.73	149.25	84.07	271.6	205.49	242.7	221.04	251	249.33	419.9
U61397_s_at	1107.35	274.43	524.95	606.49	669.15	111.09	1060.75	483.17	1153.55	77.29	1299.65	223.09	913.8	295.57	1266.7	216.52	1129.7	142.41	963.25
Y07566_at	293.45	279.52	240.8	278.18	111.75	59.33	395.4	274.07	482.6	101.82	198.8	120.49	277.15	301.72	359.4	141	243.05	252.79	98.6
M33493_s_at	756.85	147.86	693.75	801.51	937.75	229.31	688.95	547.23	300.3	154.15	715	165.89	153.6	0.42	482.2	233.91	457.7	364.02	1022.4
X06825_at	192.8	3.96	191.55	221.25	596.65	435.93	532.05	106.7	154.2	24.61	231.7	147.5	153.5	70.99	272.95	77.85	230.4	21.35	222.25

Z78291_at	95.55	16.9	50.35	58.19	77.9	58.83	321.75	373.56	73.55	23.55	89.3	41.01	77.55	56.92	138.35	117.59	77.1	2.83	72.1
X79888_at	109.2	104.79	217.05	250.81	549.8	254.84	111.3	94.19	620.05	127.49	456.5	311.69	282	337.86	320.3	255.97	124.3	75.66	417.2
M63391_rna1	830.9	350.72	522.45	603.66	2630.2	500.07	1928.7	337.29	813.5	525.66	1393.15	1278.52	2360.6	422.99	860.9	269.41	1884.95	1243.02	783.95
X59812_at	1377.45	130.6	529.2	611.36	2874.55	2106.82	4441.75	2573.94	2261.95	1126.35	742.35	14.92	1838.7	699.33	2833.2	437.42	1849.5	1599.33	1448.05
M99063_at	778.15	206.4	704.6	814.02	367.45	6.29	996.75	292.95	859.7	208.17	737.5	33.66	1076.15	67.1	925.15	214.75	362.85	19.3	493.95
M96954_s_at	1910	85.14	841.4	972.13	1264.5	602.6	1180.65	192.83	1659.75	363.38	1525.05	292.39	1787.3	112.43	1167.45	206.97	1888.4	207.47	1470.4
M37190_at	980.7	130.53	250.15	288.99	673.55	553.31	503.5	208.31	547.55	297.06	1200	355.39	376.7	148.21	896.35	197.49	1002.65	324.77	421.85
U28687_at	620	47.94	205.8	237.73	286.05	226.91	773.2	520.71	425.25	337.78	829.8	10.61	781.7	191.91	557.7	48.93	775.85	447.53	856.3
U45880_at	538.1	572.05	500.4	578.13	826.95	973.33	749.15	1007.98	235.2	36.91	723.25	766.86	393.8	187.67	382.95	240.77	197.9	22.34	692.9
X82103_at	1473.1	169.14	724.75	837.43	908.9	195.3	1091.9	40.16	1375.65	52.26	967.8	276.2	1369.95	347.26	1140.7	66.89	1216.35	211.5	1009.5
L10717_at	183.05	88.46	52.65	60.88	219.35	172.32	68.75	8.7	41.85	14.64	150.85	134.7	374.9	51.19	37.85	25.95	20.55	2.19	223.1
J00277_at	5534.7	1508.4	3334.2	3852.32	4984.7	2723.35	4702.2	3845.25	5215.6	676.42	3552.1	124.73	3565.8	392.02	3884.15	405.81	5471.25	1204.7	3178.3
D87463_at	249.9	165.04	139.9	161.64	271.45	60.6	240.6	137.04	150.1	15.98	175.7	1.7	246.45	1.63	243.5	46.95	165.45	101.47	211.6
X90828_at	263.55	50.42	133	153.73	200	99.7	324.25	4.45	661.8	750.52	167.55	0.07	295.8	9.33	580.8	593.69	335.9	269.83	324.35
HG2348-HT2	2151.6	296.42	1328.7	1535.13	2741.75	250.39	1206.9	952.33	1362.95	1288.56	1861.75	742.67	1472	1197.27	2613.9	1434.86	2150.75	460.4	1800.8
U77948_at	871.2	285.39	443.45	512.3	488.4	415.21	993.7	514.21	875.65	6.43	581.3	601.18	985.2	7.07	701.15	582.73	1224.7	71.7	1056.4
S69369_at	833.35	249.4	713.55	824.42	692.2	199.4	1611.35	1212.62	231.45	165.53	1167.3	612.07	369.75	69.08	329.15	100.62	817.25	910.68	233.65
X15306_rna1	521.35	112.22	162.5	187.81	390.6	430.77	551.3	508.98	297.65	128.91	693.5	571.91	543.05	470.44	658.8	39.88	829.35	396.47	212.2
HG3417-HT3	375.7	138.73	101.85	117.73	467.65	415.28	75.15	22.13	147.1	117.24	109.9	7.64	187.5	132.65	199.85	183.92	540.6	120.35	171.45
X92689_at	786.9	204.35	173.5	200.39	219.9	171.54	397.2	282.56	373.35	276.13	471.4	361.47	580.4	347.9	131.55	12.8	276.75	258.73	426.15

AF007551_at	1118.15	37.26	669.65	773.65	1121.1	251.45	698.95	537.61	986.5	1065.19	1345.85	550.06	1148.3	48.65	1118.2	131.24	666.9	260.78	1515.25
X59798_at	2917.7	509.82	1244.1	1437.41	1912.5	2178.03	2609.55	387.71	4205.4	705.83	3022.1	178.76	4113.3	2297.81	4055.45	299.74	4344.55	2800.64	3712.2
X66503_at	423.75	81.53	93.25	107.69	191	133.08	125.55	79.27	361.25	264.81	432.05	67.67	101.1	14.85	174.2	106.21	108.5	62.08	436.8
D31891_at	1072.5	41.44	671	775.27	855.4	652.24	1229.55	299.6	1463.35	182.36	1249.5	383.68	1841.55	235.82	1677.25	850.3	1089.35	490.38	1436.2
L40391_at	1091.5	95.88	765.7	884.73	1094.4	465.42	959.55	53.25	1224.5	150.76	1297.8	13.29	1237	1374.33	714.6	37.62	1321.65	126.93	1553.95
J02854_at	772.85	596.3	654.9	756.6	859.1	612.64	1119.85	964.14	657.9	441.66	596.55	291.26	423.2	32.81	1046.9	899.58	293.65	150.68	412.95
L76703_at	568.4	165.04	466.6	539.1	468.95	234.97	984.65	567.45	643.75	109.11	411.45	431.12	453.2	570.92	521.9	126.15	603.85	73.89	614.05
U82306_at	511.35	107.41	318.6	368.12	144.75	42.78	831.9	436.28	259.1	123.6	293.05	104.02	711.15	652.16	212.2	141.99	370.25	129.47	444.35
M80899_at	2914.45	742.53	715.95	827.24	463.85	498.58	501.25	392.8	1067.7	383.68	1666.8	1810.33	1618.95	977.43	721.6	340.83	2491.05	496.88	1537
U03057_at	29394.3	1865.63	11757.7	13584.65	25232.35	14691.77	19817.85	4999.32	28095.1	6937.28	23366.7	8077	27517	16135.33	22641.7	4568.76	27121.95	5824.51	18592.05
M29581_at	1164.2	317.49	606.05	700.25	1098.35	664.89	904.5	132.23	1771.3	601.47	1270.2	697.63	823.35	122.97	1162.95	213.9	533.4	67.46	1004.95
AB005535_s_	280.3	136.47	36.5	42.14	214.85	77.29	139.3	145.38	60.65	0.78	89.2	54.16	245.8	129.54	171.15	191.13	148.45	81.95	86.8
D80006_at	738.6	5.09	765.45	884.38	436.45	178.69	540.85	73.75	905.55	47.87	399.75	387.85	857.35	432.4	716.8	38.75	752.5	432.18	750.8
D30755_at	4941.55	10.54	4654	5377.12	4080.15	639.15	6006.85	497.45	2937.4	751.09	2846.35	2070.76	3111.1	273.65	2977.95	708.73	4407.35	1572.82	4198.3
V00563_at	781.05	1.2	435.7	503.46	343.95	80.82	1046.1	298.96	808.3	349.73	1264.25	568.73	1004.15	396.62	1122.55	99.21	680.7	149.77	1106.3
X76059_at	136.05	69.51	502.8	580.96	307.3	0.28	290.05	260.43	510.85	455.73	1227.85	461.25	640.6	533.16	534.35	395.63	173.2	145.66	337.9
U90544_at	110.4	34.79	60.55	69.93	159.5	72.69	171.25	1.34	99.05	63.57	91.3	29.13	344.15	256.61	129.6	82.17	85.75	17.75	71.3
U34962_at	1695.6	196.15	791.55	914.5	2071.4	64.35	2056.6	459.48	1782.95	83.93	2036.9	827.31	1908.5	87.12	1753.05	1928.07	2035.4	110.45	1252.9
S83549_at	64.15	40.38	63.15	72.9	104.65	30.76	111.9	22.2	122.2	56.29	86.2	51.05	326.45	348.96	79.4	66.75	52.9	6.93	75.15
U78628_at	1030.8	108.61	356.6	411.96	645.9	215.81	818.15	503.25	888.9	227.69	743.8	10.89	1061.25	41.79	498.35	302.43	1144.8	211.57	798.6
U90336_at	161.1	37.34	150.4	173.81	341	113.56	1507.1	1894.48	165.95	41.79	340.75	175.15	557.9	522.41	202.3	58.27	202.45	70.22	720.15

L32606_at	178.95	4.88	138.35	159.88	208.55	132.02	180.5	135.91	200.05	53.81	252.2	150.61	352	86.97	218.3	3.25	310.25	60.17	134.4
U96629_rna2	2680.95	264.1	1107.7	1279.86	3042.6	222.03	2692.05	590.08	3047.9	964.35	2374.35	27.08	2399.55	34.01	2283.05	2.19	2882.45	358.86	2887.5
D63813_at	279.75	40.66	662.9	765.94	546.95	140.08	946.8	539.95	850.85	226.49	272.85	77.29	322.65	200.46	678.2	343.37	725.5	291.47	718.8
U94333_at	94.6	4.95	70.85	81.81	324.65	195.94	87.45	15.91	163.5	35.21	173.45	50.84	213.95	16.76	109.7	60.53	124	19.94	109.95
X80198_at	3653.6	1444.19	2592	2994.74	1719.95	1110.51	3953.65	1291.25	4071.75	12.8	3159.45	493.91	4671.75	447.67	3773.5	1626.77	4229.6	1164.75	4044.05
U21931_at	1872.15	559.39	792.85	916.06	1212.2	219.06	971.4	959.83	871.55	926.95	1363.2	126.71	1271.35	1028.35	1342.65	1042.35	1418.9	513.08	1363.7
U67092_at	15	3.11	14.05	16.19	45.75	41.22	106.1	126.57	29.6	19.66	236.7	312.82	19.85	5.44	27.95	6.86	21.05	8.41	43.2
X05608_at	816.95	133.29	433.5	500.91	693.95	8.84	1327.25	948.87	650.45	54.24	945.9	262.9	610.65	6.86	692.7	46.53	518.45	121.98	511.7
U66561_at	165.4	105.78	355.75	411.04	455.85	341.46	578.8	735.11	134.7	34.51	176.15	166.52	131.35	133.43	138.75	157.47	249.85	282.35	212.75
M32011_at	470.6	16.55	405.05	468.03	319.65	275.84	336.9	34.51	219.7	204.35	365.3	159.95	377.5	483.1	297.2	332.06	212.4	125.02	206.7
M20642_s_at	37.45	16.62	39.35	45.47	122.7	116.25	130.25	139.65	56	21.5	442.55	608.75	262.25	82.94	78.1	4.24	85.55	26.52	208.85
HG3914-HT4	3153.65	680.87	1656.4	1913.71	1650	457.78	2954.05	1578.19	2170.05	743.66	2731.05	46.74	1907.6	743.31	1021.1	744.58	1358.65	378.09	1758.35
X13546_rna1	4083.25	2193.66	1682.7	1944.12	3775.95	1986.05	3957.35	1899.64	4709.45	1641.69	3393.55	702.79	3726	151.6	4250.65	819.61	3877.95	330.71	4825.6
X62744_at	191.65	20.29	84.3	97.44	335.6	159.38	177.85	54.09	281.25	13.08	377.65	153.65	250.9	53.88	209.35	55.08	254.4	37.76	189.1
U71092_at	99.4	25.17	46.7	54.02	156.45	19.73	195.8	128.69	144.85	17.18	170.9	39.03	371.3	420.45	181.65	178.12	190.4	41.86	203.8

U24685_at	1044.2	38.61	596.85	689.64	2041	1186.67	1200.95	1440.73	861.9	943.85	654.9	286.66	1203.8	349.73	1647.7	198.27	820.55	113.21	480.5
HG2855-HT2	9224.6	179.61	1230.4	1421.57	4207.9	1448.72	6544.9	1110.58	7608.4	1308.57	2674.85	924.54	7775.5	3845.81	8082.2	5193.42	13090.75	8167.15	5230.05
U40571_at	1491.85	49.14	599.9	693.11	1014.55	258.16	806.1	608.11	530.35	461.53	1295.65	92.7	821.7	165.18	877.1	56.29	697.2	447.46	445.05
X85753_at	968	290.06	875.95	1012.08	959.55	177.41	672	58.55	709.85	558.83	1180.55	381.06	913.8	39.17	358.05	351.5	512.05	377.81	637.6
L33799_at	472.65	263.82	405.25	468.18	653.8	292.04	798.85	192.55	223.4	64.35	500.8	257.25	1069.6	329.65	757.2	319.19	519.7	201.1	549.1
X53390_s_at	2728.1	735.25	1429.8	1651.94	1787.65	216.3	2178	165.04	1445.8	200.68	2261.6	96.03	2701.5	987.4	2625.5	177.48	2556.55	575.37	2652.45
U17032_at	203.65	193.68	71.15	82.24	670.75	302.29	310.45	156.62	345.05	225.64	521.1	702.72	313.7	195.02	252.2	23.62	133.5	53.46	89.6
M63838_s_at	525.4	136.61	455.05	525.73	913.95	28.5	255.7	211.99	1141.4	405.31	845.7	133.08	864.9	20.65	625.2	7.78	800.95	225.78	769.95
U41737_at	79.95	11.24	79.95	92.42	193.45	44.62	119.1	20.93	201.4	61.52	761.8	901.56	117.45	33.16	776	4.81	131.9	0.28	290.75
X95404_at	28475.7	3402.17	15782.65	18234.94	19291.35	5080.77	26510.8	10041.2	18139.45	365.64	23338.65	5066.21	20158.45	12800.12	25745.25	109.67	30941.75	861.04	23938.6
X68487_at	1109.95	163.98	518.85	599.41	1135.9	295.99	547.45	418.11	1260	486.35	813.25	28.21	1266.3	544.76	647.7	12.3	844.5	9.33	804.45
M29474_at	41.65	16.48	44.2	51.05	168.3	37.76	229.8	221.18	131.5	39.03	59.9	3.96	109.85	81.81	468.15	318.13	142.75	94.82	228.7
D78361_at	40143.75	12684.44	22775.9	26314.84	46565.85	7026.87	49517.1	15826.46	46499.2	2680.92	34917.85	5312.42	36261.55	6096.75	41356.15	2717.48	36557.3	2044.95	34382.65
Z29574_at	290.7	172.39	342.55	395.77	268.5	276.34	863.55	972.48	72.2	6.22	479.45	421.22	161.25	35.71	325.85	104.44	186.4	34.65	238.3
M63180_at	1562.35	401.85	1453.5	1679.38	1928.75	265.52	1598.25	625.15	2085.95	106.99	1927.65	310.21	1493.8	315.79	2347.4	307.17	1588.4	193.32	2089.25
M84332_at	11751.2	3545.15	5934.95	6857.17	12497.9	2716.42	12417.95	2409.89	11736.45	339.62	13012.6	1212.69	9181.4	630.46	10849.7	433.03	12227.85	3685.23	14273.4
U23803_at	794.05	555.86	237.65	274.57	576.8	240.7	224.5	94.47	824.6	30.26	434.45	276.27	1155.2	776.54	544.7	368.97	707.85	138.1	807.5
U08998_at	1320.4	947.66	318	367.41	2314.6	196.29	2201.4	253.43	1777.9	141.7	1184.8	693.39	1250.25	351.64	1142.6	989.81	865.35	533.94	1253.35
X03342_at	63023	38396.18	25239.9	29161.65	90380.15	637.46	88929.6	14843.44	80725.55	8948.93	66743	10003.58	62184.35	34076.54	65000.55	18709.55	64650.75	34454.27	72083.16

U18543_at	716.9	115.4	169.05	195.37	1279.4	842.59	668.65	411.18	810.15	121.41	300.45	145.03	636.45	720.05	452.15	271.32	885.55	247.28	1051.4
L05624_s_at	1943.2	701.31	1648.9	1905.09	931.6	800.3	1857.95	436.92	1430.05	75.17	953.45	297.76	2066.95	1455.58	1998.25	1118.01	2102.25	612.14	1427.6
U14518_at	298.1	105.08	211.95	244.87	392.45	455.31	368.45	206.12	594.4	25.46	497.05	153.23	413.45	140.36	652.65	283.05	622.7	380	420.6
U67369_at	280.55	42.92	299.35	345.85	526.6	571.77	504.25	233.56	306.9	206.9	638.75	669.84	845.1	326.12	803.95	573.11	427.6	253	559.85
J04621_at	96.8	34.08	15.5	17.96	160.8	123.89	77.8	35.21	166.8	36.77	123.95	113.63	195.4	95.6	213	55.86	31.6	3.96	138.95
Z23091_rna1	128.35	142.48	65.45	75.59	252.95	6.43	82.2	83.16	67.45	61.59	129.7	102.25	106.25	7.14	137.45	62.3	61.6	17.82	77.05
X83378_at	375.9	193.46	658.4	760.71	584.35	340.75	1232.85	1391.23	872.2	171.4	1214.4	649.55	1156.15	344.01	1389.65	644.39	1651.65	1279.79	709.85
HG3576-HT3	179	52.75	122.9	141.99	264.8	261.35	211.2	43.27	157.3	73.11	251.05	149.55	225.8	22.34	219.45	140.08	276.85	4.74	123.45
U24577_at	47.25	0.07	17.5	20.22	65.95	64.42	210.3	227.69	421.55	269.2	68.55	42.5	202.9	215.95	205.1	201.95	57.5	53.6	79
X54489_rna1	605.25	244.31	973.7	1125.01	666.95	314.45	524.4	358.36	560.2	187.95	357.85	201.31	383	237.02	805.65	312.47	638	635.69	468.25
X14474_at	491.05	402.84	62.8	72.55	184.35	104.3	160.55	54.38	650.7	319.9	781.05	141.63	569.65	410.9	107.4	41.01	506.95	37.97	259.8
L38810_at	4193.75	2116.16	2009.4	2321.57	3779.8	625.93	3375	2272.92	3352	1029.12	3507.5	216.8	2969.6	580.53	4122.75	941.09	3904.6	208.6	2777.3
L34409_at	77.15	14.35	52.6	60.81	195.8	189.08	196.25	259.58	42.95	13.36	72.85	24.11	71.85	11.24	248.2	147.64	214.5	169.85	103.1
X54457_s_at	166.95	86.05	133.15	153.8	155.5	70.57	130.2	60.67	136.45	60.17	145.15	130.18	89.5	11.46	78.85	24.11	86.45	3.04	81.2
U52154_at	617.25	161.86	194.15	224.36	561.5	434.87	447.5	164.05	642.2	508.27	994.35	593.9	293.85	11.38	401.6	97.44	435	217.79	604.2
D42055_at	338	33.23	371.55	429.28	873.75	907.15	668.25	419.95	352.75	418.96	695.85	270.61	307.05	34.15	385.25	34.44	513.1	610.8	730.25
U93867_at	510.45	224.79	597.4	690.28	755.45	401.57	771.05	231.58	165.15	145.31	1195.55	805.61	478.5	239.57	309.25	47.02	801.45	409.49	605.3
U46025_at	7780.6	1228.24	4956.3	5726.43	7006	1073.25	8286.45	4598.81	13400.85	5953.34	15452.65	4286.55	13473.45	2965.68	16993.8	1895.33	19920.45	3936.67	15739.5
U51333_s_at	1549.5	403.48	628.45	726.13	2076.75	647.92	1500.1	1316.49	947.5	355.39	827.2	262.2	1339.65	167.23	1504.5	88.81	2143.95	114.34	1533.1
X57025_at	421.1	186.82	216.35	249.96	405.2	374.63	522.2	706.26	209.3	32.24	269.95	88.03	331.85	276.27	396.65	142.34	215.9	196.43	394.9

Y00067_rna1	170.95	199.19	248.25	286.87	131.2	33.8	126.05	128.62	223.5	192.05	535.75	398.31	230.7	285.39	321.6	430.35	100.95	40.09	87.85
U28368_at	98.4	99.14	71.25	82.38	69.85	55.65	50.6	50.49	159.2	192.05	267	20.51	102.7	84.43	106.5	106.07	169	215.53	97.55
HG3543-HT3	1593.55	679.46	531.3	613.91	2185.5	393.15	2263.15	1247.41	1132.35	11.81	1556.4	94.05	1806.85	492.92	1500.85	84.64	1248.95	291.96	1578.05
U49742_at	803.85	342.17	47.2	54.59	1130.8	633.85	893.1	642.62	1077.75	65.83	999.8	781.21	953.45	407.93	498.1	413.66	497.65	369.75	170.6
HG3635-HT3	463.35	363.81	374.2	432.33	986.45	265.8	397	269.27	318.7	383.39	339.5	371.37	300.7	208.46	436.55	222.1	838.95	5.87	618.7
J03890_rna1_	328.95	41.79	729.2	842.45	354.8	122.05	278.25	36.84	287.6	27.58	534.15	205.27	367.05	175.43	344.95	25.39	383.3	158.11	333.9
D84239_at	346.3	100.27	131.05	151.39	949.75	388.84	277.55	274.29	254	28.43	450.9	247.35	301.6	38.75	223.5	90.51	339.6	180.45	244.1
S72008_at	845.35	34.29	798.95	923.13	1348.75	519.51	392.35	324.92	1230.65	406.94	920.45	380.07	1140.6	112.71	1269.7	314.24	856.35	547.09	857.25
U10492_at	158.25	148.28	211.6	244.52	37.2	0.28	165.15	13.36	172.25	175.72	613.7	538.82	227.15	151.11	213.85	269.48	240.25	178.83	103
U20350_at	481.75	274.57	57.25	66.11	303.15	7	880.05	741.97	476.65	170.34	109.95	85.35	490.8	13.29	206.25	156.77	312.8	147.79	307.55
AB001325_at	5299.75	44.9	3487.75	4029.73	8607.5	415.78	8827.45	6779.81	7461.25	1930.9	9530.4	6986.5	5349.95	832.9	5174.55	2342.15	4629.35	202.02	8917
M64174_at	731.85	46.88	199.6	230.66	584.55	661.36	801.35	119.43	715.7	319.9	471.95	416.84	1111.7	338.7	563.85	300.17	603.9	275.77	799.1
V00572_at	13268.35	1636.46	5656.25	6535.15	11141.9	107.62	13295.5	4211.81	8251.45	4394.03	6572.8	1921.07	11919.6	5803.51	15858.3	9964.41	15172.7	8175.29	8707.05
U90304_at	392.45	155.07	75.6	87.4	317.95	188.87	131.1	14.99	442.8	187.1	327.85	152.38	895.6	172.25	388.65	349.95	342.55	278.25	207
X83425_at	3360.2	142.98	1529.3	1766.92	3166.55	612.14	2271.45	843.51	2749.45	371.87	2475.2	668.92	2541.15	494.9	2437.25	777.75	1889.2	499.78	2329.3
S62696_s_at	160.25	16.76	52.95	61.16	280.7	183	217.5	96.87	156.5	12.87	238.7	152.03	154.15	36.56	115.85	0.92	353.95	295.78	137.3
U10473_s_at	259.1	24.47	411.1	475.03	219	52.89	152.4	3.82	99.9	93.48	305.85	365.5	288.15	187.6	99.25	94.96	85.2	28.14	58.05
U43959_at	720.5	619.71	198.15	228.89	399.1	89.94	307.35	47.45	390.95	40.8	317.5	22.06	340.25	77.85	567.95	151.53	314.1	169	330.05

D13146_cds1	2232.9	669.63	1132.45	1308.36	2761.2	1043.69	2648.3	34.93	2702.2	867.9	2238.3	238.15	2306.1	196.86	3185.95	1439.17	3071.7	857.3	2549.9
D50927_at	461.15	103.73	389.15	449.65	307.85	174.87	124.3	55.58	763.25	266.79	392.95	229.17	457.75	119.01	326.85	21.14	380.65	137.67	439.5
M31651_at	385.75	215.74	100.1	115.68	171.25	58.62	755.2	990.94	114.4	58.27	422.8	450.99	384	180.17	446.5	489.18	120.75	18.74	429.7
M16424_at	1302.75	345.42	729.25	842.52	2123.7	1394.98	1979.15	865.85	1106.55	532.1	1633.45	949.15	1551.95	1331.55	1419.3	1386.21	1265.1	813.46	942.85
U12259_cds2	205.6	76.79	117.85	136.12	161.45	2.33	857.15	1103.86	286.25	171.76	176.8	34.37	302.15	190.99	159.05	18.6	351.15	88.6	165.65
X93512_at	144.4	59.26	79.1	91.36	66.25	10.96	281.9	79.62	108	57.28	303.1	95.46	81.25	8.7	96.6	37.34	410.35	458.28	156.75
X67247_rna1	73047.35	38238.71	37625.45	43471.58	85426.9	19558.57	82782.66	35718.86	94358.8	7849.02	73648.65	12924.71	82862.8	12493.3	83971.6	5624.47	73654.5	21319.98	76984.8
U73377_at	11786.65	324.92	7241.45	8366.56	10041.8	4162.17	11480.35	4925.49	13262.9	538.96	10623.7	2033.5	15768.55	5282.3	11735.15	961.03	14948	3769.3	15997.1
U77413_at	491.8	464.14	400.05	462.24	282.2	264.88	257.45	247.98	168.75	27.93	335.35	232.99	579.8	111.44	716.55	202.02	115.6	20.93	219.45
Y08836_at	861.35	353.06	466.5	538.96	1320.75	483.31	1582	973.26	1473.85	402.7	987.85	172.32	395.3	57.28	609.95	440.6	845.05	181.37	1032.95
Y08265_s_at	3871.25	887.49	2375	2744	3566.8	617.59	3680.9	1162.06	4212.3	605	3804.5	999.14	6188.05	672.1	4938.1	826.04	5577.55	1428.43	2899.2
U07804_s_at	108.55	77.43	104.25	120.42	146.45	112.92	167.05	31.89	119.8	11.17	99.25	0.64	243.15	208.81	329.25	34.86	71.8	12.59	247.35
X95097_rna1	1150.15	299.18	876.05	1012.22	1744.25	869.25	1255.75	1003.46	1138.9	21.92	1365.05	1440.45	1597.45	13.79	631.4	495.96	2261.75	401.99	1195.65
X78678_at	224.25	137.39	73.2	84.57	226.25	18.31	192.75	19.02	227.05	161.86	316.05	262.41	267.9	71.42	334.55	281.64	279.35	49.85	381.85
M93651_at	2553.15	298.89	1808.35	2089.29	2062.95	222.53	2072.75	87.47	2488.35	66.4	2122.85	808.01	1880.35	138.95	2699.7	898.03	3269.35	110.8	2790.4
U21936_at	278.3	138.31	28.2	32.53	225.1	127.99	172.2	203.93	309.8	200.82	186.25	180.1	244.1	190.78	38.4	0.57	66.3	33.94	193.55
M76665_at	393.1	152.59	638.75	738.01	382.2	378.44	399.9	239.57	511.2	512.51	360.55	260	259.5	275.63	524.65	595.6	470.65	242.33	267.1
X57129_at	53.45	8.7	82.3	95.04	113.3	100.13	177.45	212.91	71.75	48.01	78.15	52.54	106.15	15.63	118.8	75.09	285.7	210.72	100.65
HG1800-HT1	74224.2	46256.1	31455.1	36342.6	106843	622.25	87719.95	25909.59	90428.1	14687.17	81273.25	14771.53	73310.7	17751.49	71718.8	12604.32	63156.5	10570.96	71453
U73936_at	531.9	64.21	333.9	385.8	1098.5	215.24	1055.6	169.28	683.15	565.05	962.15	246.99	756.35	341.32	465.7	514.21	796.6	285.95	784.95

M88579_at	133.2	82.59	284.8	329.09	128.15	119.15	439.1	227.83	326.2	154.43	189.8	73.26	670.8	200.25	446.7	348.89	134.7	45.68	380.75
J05158_at	694.95	74.88	891.1	1029.55	492.2	123.46	785.9	182.01	1167.8	526.94	1893.45	958.48	691.75	253.5	1168.05	116.46	1126.95	203.58	1407.3
D17570_s_at	34.15	17.18	44.65	51.55	90.15	53.39	182.9	139.72	272.3	207.32	81.35	57.77	61.85	50.98	61.85	6.29	70.15	12.37	164.35
U09646_at	107.9	46.24	71.65	82.8	153.2	37.19	184.25	158.75	456.3	290.76	262.15	12.23	160.3	39.17	337.15	394.64	267.5	255.27	556.05
HG4518-HT4	137.95	48.3	159.35	184.06	381.95	113.49	146.7	96.87	575.6	680.52	451.15	513.29	281.05	130.74	115.85	29.34	105.75	41.37	736.4
J04970_at	67.1	13.01	178.15	205.84	31.35	6.29	61.5	27.72	68.45	52.68	36.3	14.85	70.55	56.64	59.35	39.24	133.75	62.58	106.25
U27333_at	1953.55	1260.42	2446.15	2826.24	3430.75	1826.1	2767.15	188.02	1960.9	1811.89	3519.85	970.5	4074.35	2198.75	2613.45	1418.53	2763.3	1013.99	3631.8
J00268_s_at	458.4	117.8	195.45	225.78	481.75	255.34	934.05	875.33	579.4	14.42	591.3	182.01	669.95	32.88	1354	70.43	429.85	72.34	402.05
S81944_at	42.25	7.42	29.05	33.59	87.4	58.83	143.15	167.09	104.35	25.81	41.95	11.67	89.05	89.87	203.4	175.22	58.85	32.6	50.8
M81933_at	1354.95	428.01	637.3	736.38	642.35	519.79	794.3	578.98	498.45	424.9	1437.7	380.42	387.95	5.73	1329.1	766.08	1004.2	481.96	1212.8
D84557_at	1407.8	141.28	823.05	950.99	868.6	670.34	1797.7	1010.88	2069.55	191.84	2004.55	606.2	2558.15	581.31	1430.7	1044.68	1618.4	1191.33	1710.25
Z19554_s_at	43350	13290.5	17890.05	20669.79	43674.9	12592.3	44742.4	15101.82	41007.9	7318.27	33556.45	11202.2	37545.65	1357.29	37786.55	1302.84	52281.2	11337.61	52048.45
M15958_at	200.4	103.1	107.05	123.67	176.5	3.68	342.6	106.07	204.3	3.96	230.1	56.29	213.8	41.01	286.75	78.28	211	70.85	322.65
U14391_at	2401.8	319.47	1103	1274.35	1606.9	129.12	2467.4	101.68	2405.2	570.49	2783.85	96.52	2669.1	270.96	1940.35	598.42	3054.35	128.62	1740
M55153_at	2218.45	971.21	341.15	394.21	1158.7	432.18	1531.2	972.7	1289.3	1063.49	2097.95	734.61	2212.5	81.88	1757.6	308.02	3187.4	3439.08	1464.1
X83412_at	159.65	29.91	143.35	165.68	254.75	173.59	233.65	101.05	208.75	98.36	164.3	90.09	236.75	34.58	254.3	73.11	139.2	4.95	136.9
M86528_at	1349.5	364.3	187.4	216.52	1174.85	160.44	964.15	427.16	878.1	111.16	621.7	477.44	1147.25	15.91	907.15	285.32	840.8	286.38	927.3
J05125_at	235.65	250.81	98.05	113.35	118.3	53.6	82.45	59.89	81.35	4.17	542.95	576.08	105.5	32.95	126.6	44.41	185	93.9	332.7
X77909_at	1085.15	402.56	664.8	768.06	445.25	207.39	1036.5	88.11	880.45	700.53	2295.9	1044.82	1963.7	81.32	1007.8	757.59	900.6	497.24	1431.85
HG2992-HT5	1581.45	335.95	368.75	426.03	1827.35	672.39	1484.3	156.84	1625.6	569.65	1619.5	612.21	1342.6	261.63	1881.55	713.97	1367	124.59	1632.3
J04031_at	1333.25	195.09	822.3	950.07	1985.9	363.31	1293.1	424.83	1935.45	849.87	1863.45	398.31	1939.1	446.04	1421.2	53.17	1788	136.19	1728.55

U09510_s_at	10444.3	1531.73	5019.8	5799.83	7862.55	5449.32	7023.95	1343.57	8886.8	688.16	8254.75	2225.05	7335	1311.54	6706	4690.24	8105.65	5531.34	8660.65
L37368_at	4270.55	128.76	3700.95	4275.95	3849.75	298.05	5167.95	501.41	5369.45	2350.78	5922.25	1152.37	4779.45	1628.82	5153.4	354.4	6946.45	3008.39	5685.9
X55889_at	498.95	508.06	144.9	167.44	473.5	262.05	238.3	128.98	165	23.76	691.85	87.61	334.6	204.07	202.6	72.55	166.95	140.78	185.75
X65784_s_at	3072.5	683.35	1845.4	2132.07	2538.1	695.37	2753.5	790.26	3130.25	1353.76	1272.3	1152.58	2230.75	1379.21	2648.7	891.66	3611.45	825.83	3099.15
M64710_s_at	168.45	141.49	86.15	99.49	186.75	103.03	198.45	81.11	188.35	11.95	138.8	56.14	208.95	9.83	189.45	8.13	208.6	35.64	106.85
U66061_cds3	393.65	154.64	340.7	393.58	409.1	108.47	393.35	109.81	677.4	156.69	778.3	442.51	829.6	207.89	492.85	418.4	458.5	67.6	343.45
U64105_at	2877.5	838.06	1032.5	1192.89	1990.3	574.17	2945.55	331.28	2287.45	933.45	3135.95	293.66	2993.65	234.83	2740.25	60.32	3075.35	583.01	2556.15
Z22555_at	3283.5	1177.76	1421.9	1642.89	1940.95	7.42	2234.3	37.34	2894.95	906.44	1965.4	58.27	2407.05	924.12	2928.3	950.49	2600.85	407.65	2099.05
X78711_at	885.15	155.07	381.25	440.46	811.9	211.99	352.05	19.59	443.8	121.62	215.6	57.56	465.55	566.18	405	227.69	776.4	100.27	460.1
S77893_s_at	352.65	329.02	59.35	68.52	380.85	298.05	89.65	14.78	128.2	65.34	616.9	738.93	439.7	388.63	169.85	74.88	177.05	7.85	343.2
U30930_at	340.95	209.37	233.75	270.04	350.7	32.81	577	444.35	517.05	42.64	307.85	123.67	247.25	165.53	410.9	287.37	531.6	102.11	471.45
X99268_at	89.65	3.46	42.3	48.93	335.1	323.29	105.25	62.58	119	39.03	72.5	8.77	84.1	24.61	745.45	932.6	83.15	7.42	97.55
D88155_s_at	897.1	432.18	658.8	761.13	1552.35	80.26	234	141.7	1043.1	22.63	981.2	898.31	1146.95	286.73	1335.6	320.04	618	105.64	1024.05
U90543_s_at	522.7	247.49	347	400.93	429.25	430.84	468.45	3.04	630.9	617.02	646.2	313.67	302.45	46.6	680.8	474.89	515.6	591.99	263.55
J04093_s_at	65.55	15.63	233.25	269.48	100.8	42	144.55	36.7	255.2	189.08	108.3	58.41	275.85	120.84	122	101.4	54.2	13.15	561.65
D79998_at	611.35	142.34	487.35	563.07	354.7	130.96	554.75	63.99	1079.2	21.5	894.65	694.45	397.15	27.08	936.85	498.3	440.3	24.61	352

D79996_at	1729.4	50.91	1141.7	1319.04	1947.55	4.74	1665.4	670.05	2338.6	267	1439.7	1078.48	2126.15	314.45	2078.1	526.09	2575	226.98	1756.45
X13766_s_at	152.75	29.49	86	99.42	789.9	851.07	319.45	238.51	190.4	60.67	445.75	399.3	642.25	172.6	610.2	601.32	293.5	26.16	296.95
Z46788_at	56.4	45.25	30.75	35.57	341.4	421.44	165.1	200.39	105.8	87.12	272.75	11.38	73.7	52.89	43.3	7.21	79.95	18.46	123.8
J05249_at	1844.75	138.52	1294.45	1495.6	2231.15	195.94	2412.3	668.64	1799.95	117.73	2417.3	492.15	1779	821.8	2147.85	877.73	1869.05	448.38	2253.2
X01057_at	26.2	13.86	37	42.71	297.05	397.61	58.9	22.34	269.65	96.8	104.25	107.98	169.15	195.66	51.8	16.69	245.75	100.62	37.75
U47677_at	265.05	44.9	172.1	198.84	192.85	33.45	192.55	18.88	291.85	177.41	269	230.52	210.35	82.8	185.05	54.52	334.9	12.87	147.4
M93036_at	579.3	172.68	539.3	623.1	922.85	95.39	588	324.28	656.55	126.08	467.9	141.28	687.8	117.38	571	635.97	546.45	197.78	983.25
U41898_at	148.85	15.63	75.35	87.04	382.7	279.59	159.35	58.9	134.7	10.75	135.4	64.63	357.85	305.97	408.3	492.15	305.25	257.74	135.15
Y12670_at	837.3	194.88	488.45	564.34	992.1	143.54	217.7	51.62	874.35	473.13	350.55	174.16	761.95	259.01	156.55	54.24	456.65	404.68	393.5
Z19574_rna1	210508.06	66282.28	121755.15	140673.31	219196.81	55317.53	210591.64	80439.12	172667.8	38446.81	152268	58522.85	180740	5705.51	189444.09	12354.29	159358.06	9665.65	119947.95
M15990_at	845.4	500.21	553.5	639.51	1133.6	618.15	1079.65	668.29	1159.1	254.28	1096.85	107.13	1424.95	132.3	1268.25	38.82	829.6	263.89	1119.2
U28131_at	722.8	569.93	774.3	894.63	1453.5	310.28	1873.9	1655.62	1224.5	1194.44	919.55	575.66	765.15	453.89	1295.65	1273.29	652.7	352.85	531.65
X51698_s_at	315.3	105.22	127.3	147.08	260.65	204.57	615.5	629.47	288.5	136.33	247.45	25.81	348.6	64.77	654.65	638.16	274.05	71.35	384.2
U69141_at	1002.35	177.55	576.2	665.67	956.75	296.35	1449.65	796.7	663.75	669.98	740.5	386.79	1095.9	668.78	791.7	356.52	1053.2	118.23	882.35
HG881-HT88	1157.7	40.31	479.9	554.51	866.65	90.16	1585.65	643.26	597	211	1440.2	1299.8	706.8	105.5	1779.95	404.96	1530.15	960.75	727.7
M80244_at	74035.59	8164.54	45033.95	52031.25	51569.8	27073.14	57859.1	27205.65	51849.35	4449.19	53093.45	20313.41	34009.35	8974.39	43802.95	28475.69	39507.1	17616.57	36966.3
S82470_at	1801.35	480.05	749.85	866.42	1171.05	525.17	1301.85	347.97	1887.85	9.55	1710.95	336.09	1534.55	443.57	1190.65	104.86	1932.65	75.59	2003.3
X67098_at	105.2	9.05	55.65	64.28	265.7	95.18	472.3	579.97	566.65	134	400.8	421.15	227.1	236.46	347.25	268.91	383.6	402.91	284.85
U78521_at	3500.5	445.9	1392.1	1608.39	6353.35	1076.57	4612.9	1119.49	3587.5	210.86	3523.1	830.57	4894.75	131.03	4371	877.52	4009.65	1319.96	4391.8
S62539_at	488.95	112.78	394.45	455.73	545.3	267.85	292.3	200.82	645.7	169.28	376.25	318.41	735	422.99	453.05	117.17	772.95	213.48	553.5

M83308_at	1506.5	1097.57	380.2	439.25	473.7	70.43	469.8	18.24	475.65	130.18	579.45	256.18	638	109.18	410.5	97.3	403.95	148	810.4
HG2479-HT2	667	163.06	406.55	469.73	95.05	1.77	1456.8	970.57	377.85	364.65	226.05	50.56	477.45	313.04	602.7	358.5	264.85	179.39	975.75
U40371_at	295.7	96.03	329.25	380.35	561	579.4	644.75	256.04	292.3	225.85	421.35	115.05	286.3	88.95	536.2	292.04	586.25	19.16	555.05
U37359_at	126.45	30.48	424.7	490.73	276.25	281.22	204.55	50.84	453.1	81.74	223.45	86.34	342.65	415	171.05	184.34	201.1	100.97	292.75
HG4310-HT4	86	14.71	47	54.31	205	41.01	102.7	16.97	119.2	26.87	136.5	37.48	139.6	1.98	98.45	12.09	113.1	45.82	148.55
X63522_s_at	392.1	249.75	77.05	89.02	600.35	276.69	823.6	995.46	169.7	40.45	174.55	52.26	216.5	43.56	134.9	60.67	108.3	25.74	162.7
L38616_at	648.9	638.09	515	594.96	717.5	339.55	1346.6	558.61	1573.55	45.04	585.65	495.19	1120.2	150.19	1162.45	122.4	1164.9	174.51	998.4
Y10313_at	202.4	81.32	527.1	608.96	297.1	305.89	548.15	364.51	542.35	355.32	807.7	528.07	354.2	357.65	511.7	4.53	628.65	196.79	384.35
J04456_at	48885.05	35043.72	20873.55	24116.79	68017.15	8491.86	63130.85	25939.43	91690.9	22478.22	66508.8	12337.32	70532.5	31129.81	66084.1	14020.23	56271.15	17699.94	71190.45
U48213_at	599.35	3.46	658.8	761.13	697.2	91.22	760.25	348.67	786.75	55.08	911.45	239.5	651.45	5.59	686.7	1.84	1089.85	681.01	547.8
X98248_rna1	1347.15	66.4	579.85	669.98	858.2	327.67	961.7	141.7	1321.9	460.04	649.5	511.38	1124.2	268.13	596.55	389.97	1127.65	271.88	1346.55
X17648_at	40.45	4.31	402.7	465.28	722.5	99.42	146.1	83.01	278.35	306.25	327.65	204.14	263.9	251.59	425	496.11	472.4	101.54	209.5
U24152_at	2138.3	828.59	1079.55	1247.27	1256.65	135.69	2174.45	339.48	1928.2	458.77	1780.1	78.06	1246.65	623.03	1774	227.41	2285.2	0.28	2271.75
Y13896_at	228.75	162.56	51.95	60.03	140.8	71.84	375.05	441.31	526.1	16.83	589.75	720.33	357.4	212.13	470.75	449.37	339.8	308.72	548.65
X97160_rna1	2336.4	1263.88	1780.45	2057.04	1095.3	356.1	1765.6	91.78	623.8	177.34	2021.75	250.53	2027	525.38	1461.55	16.05	1694.35	54.8	2335.35
U80628_at	158.45	72.05	46.3	53.46	423.75	315.44	157.4	20.65	811.3	153.02	476.35	495.75	243.6	56.85	900.65	954.1	506.75	455.73	297.8
U23435_s_at	714.8	6.93	359.75	415.71	1102.35	47.31	1539.8	453.82	744.2	275.35	889.85	181.66	843.6	1.98	1082.5	184.41	925.95	6.15	912.85
X05360_at	226.65	170.06	148.25	171.33	88.65	64.13	148.95	155.35	410.5	179.18	435.05	40.52	256.85	234.41	216	162.35	300.6	163.06	253.55
D79983_at	99.35	45.04	23.8	27.44	479.55	606.06	192.95	19.73	84	17.54	220.55	37.12	116.3	42.71	298.75	18.74	288.1	311.27	330.1
U52828_s_at	597	326.4	486.5	562.15	683.75	882.82	1337.7	1198.4	159.05	37.26	191.05	102.46	438.15	50.56	561.2	560.59	294.05	135.13	399.55

U79271_at	113.45	49.43	298.2	344.5	96.65	37.41	144.85	101.05	238.85	64.56	315.65	286.45	348.9	408.57	49.7	9.33	298.8	318.48	113.25
X99325_at	4017.6	1356.8	1746.4	2017.8	2766.8	1436.56	3183.55	412.03	2013.2	1168.71	3238.65	593.33	2921.85	745.93	2335.75	1098.21	3479.65	302.43	2991.9
X74614_at	1762.45	330.29	453.4	523.82	1731.8	182.72	1857.75	1556.98	1102.55	823.57	1326.95	448.38	1987.85	743.38	1056.35	296.49	1336.95	91.71	1105.5
U51127_at	5629.7	382.97	1256.7	1451.97	4610.7	193.75	4131.75	1481.18	4210.2	989.81	3677.45	1903.32	3684.65	1210.07	3307	1155.98	3955.85	933.31	4696.75
U77846_rna1	1568.1	161.36	1385.9	1601.17	2604.45	1079.54	3214.55	2578.04	1780.35	232.71	2162.75	1322.64	903	579.4	3647.4	1362.03	2177.55	1047.72	3523.3
D63940_s_at	297.2	250.88	76.25	88.03	200.8	142.98	110.2	100.27	257.5	153.58	283.05	201.6	469.2	528.35	253.55	152.66	340.7	71.7	403.35
HG2662-HT2	256.05	102.88	125.5	144.96	404.6	239.28	412.8	183.42	224.2	112.57	364.6	87.4	231.8	13.15	485.4	304.62	282.55	54.8	263.05
HG1471-HT3	918.35	381.2	513.2	592.98	648.85	785.38	672.7	26.02	744.25	321.24	482.6	115.54	777.6	0.28	708.7	339.55	798.5	337.57	787.1
U52830_at	831.55	67.53	380.65	439.75	518.05	315.3	510.8	238.44	451.9	63.92	1119.55	319.4	901.7	312.68	623.2	606.41	854.65	184.77	611.45
Z00010_at	164.4	41.86	34.25	39.53	140.8	104.23	94.4	111.16	495.15	261.98	83.4	28.28	112.15	66.68	96.45	40.38	91.25	67.1	132.65
M83221_at	1884.95	0.07	1425.55	1647.06	2413.65	288.43	2239.85	476.66	1824.3	389.05	2357.8	191.2	2672.55	1237.93	1784.9	615.61	1620.45	42.07	1980.65
D63412_at	105.85	90.58	69.9	80.75	88.3	45.11	366.7	95.18	493.95	388.41	104.95	84.36	123.65	72.62	271.9	246.78	151.4	89.52	398.15
AC000064_c	768.55	388.13	580.1	670.2	483.6	289.07	637	102.67	670.55	494.06	818.85	372.43	837.6	134.77	552.95	121.69	754.85	268.21	997.65
X85750_at	150.9	110.87	250.45	289.42	152.05	7.71	93.45	85.77	185.25	195.66	80.05	15.77	172.5	184.84	95.4	2.26	410.8	509.26	111.3
L23808_at	94.15	35.71	25.85	29.91	131.3	32.24	93.8	87.4	132.9	30.97	94.75	26.38	138.9	103.38	153.4	0.71	91.4	8.49	190.05
AF006087_at	3015.3	1512.08	1356.05	1566.74	2777.65	167.23	3312.8	201.95	2482.3	1427.51	3247.95	388.98	2588.25	974.89	2755.8	45.82	3067.75	1754.54	3629.5
M91368_s_a	1262.65	329.44	239.9	277.19	904.85	196.79	1040.6	755.47	916.3	733.13	389.1	132.51	303.1	39.32	704.4	468.95	392.65	52.26	292.8
U19765_at	461.75	141.35	389.45	449.93	646	776.69	412.15	73.19	1008.8	494.41	421.7	344.93	655.55	120.7	521.05	120.84	572	92.49	472.2

3 15 - Stdev	3 60 - Average	3 60 - Stdev	3 120 - Average	3 120 - Stdev	4 dmso - Average	4 dmso - Stdev	4 15 - Average	4 15 - Stdev	4 60 - Average	4 60 - Stdev	4 120 - Average	4 120 - Stdev	Description
531.46	2407.5	665.1	2702.25	1444.83	2621.8	599.91	902.15	759.08	871.45	129.33	847	803.56	S78187, class A, 20 probes, 20 in S78187 2548-3064, CDC25Hu2=cdc25+ homolog [human, mRNA, 3118 nt]
202.52	793.9	327.67	1009.3	758.02	556.1	409.84	915.85	25.95	679.45	163.98	1038.6	419.88	X69636, class C, 20 probes, 20 in all_X69636 1268-1951, H.sapiens mRNA sequence (15q11-13)
11.24	21	2.4	194.15	241.05	331.65	118.3	122.45	0.35	242.65	172.75	69.2	7.21	X04688, class C, 20 probes, 20 in all_X04688 227-798, Human mRNA for T-cell replacing factor (interleukin-5)
23.12	346.25	78.98	104.85	28.5	236.1	241.69	118.6	7.92	1338.2	1815.43	64.45	21.57	U16306, class A, 20 probes, 20 in U16306 10722-11142, Human chondroitin sulfate proteoglycan versican V0 splice-variant precursor peptide mRNA, complete cds
1703.84	2422.6	1334.31	540.95	56.92	2156.85	1081.38	1162.9	454.1	977.65	398.74	868.8	456.93	M36542, class A, 20 probes, 20 in M36542mRNA 1437-1832, Human lymphoid-specific transcription factor mRNA, complete cds
18.95	228.5	37.9	198.75	48.58	2464.25	2961.43	395.85	121.69	431.1	95.74	230	18.95	12-Lipoxygenase
636.47	2075.75	734.47	1994.75	768.41	1745.25	1172.88	1767.45	950.28	1893.95	31.47	1274.5	1247.76	M35531, class A, 20 probes, 20 in M35531 2832-3318, Human GDP-L-fucose:beta-D-galactoside 2-alpha-l-fucosyltransferase mRNA, complete cds
167.37	302.55	132.87	343.3	428.37	391.85	441.16	58.1	31.54	249.55	130.74	236.3	245.79	X97671, class C, 20 probes, 20 in all_X97671 31-566, H.sapiens mRNA for erythropoietin receptor
32.6	803.5	643.61	1928.7	1303.48	737.95	735.74	1563.85	610.45	1760.05	118.3	759.25	660.79	AC002477, class A, 20 probes, 20 in AC002477cds 444-988, Human PAC clone DJ327A19 from Xq25-q26, complete sequence. /gb=AC002477 /ntype=DNA /annot=CDS
74.81	352.35	281.07	1005.25	162.85	340.15	204	322.35	32.6	1415.65	1852.97	313.7	103.8	L76528, class A, 20 probes, 20 in L76528exon 146-615, Homo sapiens presenilin 1 (PS1; S182) gene
17.89	289.7	113.42	182.45	111.65	643.45	211.92	210.35	115.33	501.95	446.82	438.9	5.8	U33267, class A, 20 probes, 20 in U33267 1613-2081, Human glycine receptor beta subunit (GLRB) mRNA, complete cds
2260.2	2054.2	2531.02	3307.5	4271.77	13812.45	14324.64	3814.45	2627.96	4836.25	4629.36	2134.45	474.96	J04423 E coli bioB gene biotin synthetase (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
821.52	4502.8	447.88	5465.3	4143.36	5000.25	478.78	5351.15	2405.65	2616	2675.13	2713.8	1624.65	M64673, class A, 20 probes, 20 in M64673 1541-2084, Human heat shock factor 1 (TCF5) mRNA, complete cds
272.66	3503.75	102.74	3174.1	663.97	1853.85	48.15	2015.15	1335.51	2211.25	1332.97	1813.8	259.37	U02031, class A, 20 probes, 20 in U02031 3660-4182, Human sterol regulatory element binding protein-2 mRNA, complete cds
14.07	229.35	212.49	229.3	134.63	705.3	184.27	214.45	155.35	374.4	254.56	416	338.28	U81607, class A, 20 probes, 20 in U81607 6007-6535, Human gravin mRNA, complete cds
299.25	353.9	456.23	82.5	77.78	123.55	120.99	451.8	264.6	146.2	90.79	276.7	290.34	B. subtilis /GEN=lys /DB_XREF=gb:X17013.1 /NOTE=SIF corresponding to nucleotides 387-658 of gb:X17013.1, not 100% identical /DEF=B subtilis lys gene for diaminopimelate decarboxylase (EC 4.1.1.20).
172.6	396.65	366.63	1827.5	2136.74	1504.35	1082.09	863.8	855.17	199.45	124.95	344.7	212.56	M33317, class A, 20 probes, 20 in M33317mRNA 1158-1693, Human cytochrome P450IIA4 (CYP2A4) mRNA, complete cds
225.35	1314	586.47	423.5	343.09	816.35	338.92	704.3	55.01	1180.8	1016.82	1553.45	685.68	D10704, class A, 20 probes, 20 in D10704 1969-2365, Human mRNA for choline kinase
266.44	1163.5	54.73	1133.55	131.31	538.85	426.31	536.7	535.85	689.7	683.63	1620.65	319.97	D86978, class A, 20 probes, 20 in D86978 5648-6086, Human mRNA for KIAA0225 gene, partial cds

2648.19	7069.65	1274.84	5661.7	1781.34	4706.95	1277.95	4229.7	621.12	5507.8	2380.83	4666.9	310	Z37986, class B, 20 probes, 12 in Z37986cds 409-631: 8 in reverseSequence, 826-1000, H.sapiens mRNA for phenylalkylamine binding protein
101.89	80	60.95	541.6	217.79	387.65	392.09	486.25	92.56	1084.05	817.06	517.75	325.76	U51903, class A, 20 probes, 20 in U51903 5202-5712, Human RasGAP-related protein (IQGAP2) mRNA, complete cds
91	2422	558.19	2456.2	85.98	1262.75	1149.83	1970.95	480.9	2803.4	1179.17	1937.05	247.28	D50914, class A, 20 probes, 20 in D50914 1543-2077, Human mRNA for KIAA0124 gene, partial cds
1170.05	1511.75	487.69	750.3	474.19	1756.15	650.89	1180.85	271.32	1219.8	320.04	1064.65	363.38	M13207, class A, 20 probes, 18 in M13207exon#2-4 6-288: 1 in reverseSequence, 2583: 1 not in GB record, Human granulocyte-macrophage colony-stimulating factor (CSF1) gene, complete cds
20.15	241.8	16.97	481.95	339.06	259.9	48.65	591.7	274.07	818.55	733.06	236.3	170.98	U25956, class A, 20 probes, 20 in U25956mRNA 1470-2046, Human P-selectin glycoprotein ligand (SELPLG) gene
251.02	412.35	241.05	463.45	3.32	419.85	315.44	523.95	195.23	974.1	812.61	778.65	780.86	X12953, class C, 20 probes, 20 in all_X12953 859-1130, Human rab2 mRNA, YPT1-related and member of ras family
22.34	133.7	81.46	113.7	95.32	107.9	49.64	153.55	41.65	456.8	412.95	111.3	21.07	M81650, class A, 20 probes, 20 in M81650mRNA 1200-1566, Human semenogelin I (SEMGI) gene, complete cds
444.7	732.7	398.1	354.85	51.55	276.1	193.46	801.55	405.95	164.75	17.18	466.05	414.15	X52005, class C, 20 probes, 20 in all_X52005 476-969, H.sapiens skeletal embryonic myosin light chain 1 (MLC1) mRNA
30.41	60.75	57.2	392.7	505.86	74.5	38.89	101.95	3.04	93.95	1.63	106	70.29	X86816, class A, 20 probes, 20 in X86816mRNA 4-193, H.sapiens estrogen receptor cDNA, 5 splice variant. /gb=X86816 /ntype=RNA
304.9	265.85	246.43	427.95	104.3	384.7	39.88	606.8	534.43	482.5	94.33	356.35	241.62	U88666, class A, 20 probes, 20 in U88666 3113-3653, Human serine kinase SRPK2 mRNA, complete cds
601.32	653.25	715.52	257.75	160.02	1034	635.12	950.6	888.55	395.85	380.07	263.15	176.28	D89289, class A, 20 probes, 20 in D89289 1431-1947, Human mRNA for N-Acetyl-beta-D-glucosaminide, complete cds
56.29	49.05	55.08	79.9	103.94	133.45	118.58	86.4	25.6	126.65	89.17	66.4	14.42	Z78285, class A, 20 probes, 20 in Z78285 3-137, H.sapiens mRNA (clone 1A7).
334.39	452.6	493.7	489.1	191.06	298.2	333.05	374.4	372.22	1028.45	1316.7	396.15	305.68	U54804, class A, 20 probes, 20 in U54804 2444-2912, Human Has2 mRNA, complete cds
120.77	917.7	288.64	942	13.44	1309.55	286.02	959.25	345.28	982.3	554.51	667.1	333.05	Homo sapiens /REF=M97935 /DEF=transcription factor ISGF-3 mRNA, complete cds /LEN=3970 (_5, _MA, MB, _3 represent transcript regions 5 prime, MiddleA, MiddleB, and 3 prime respectively)
271.18	168.25	98.78	350	147.22	109.2	61.8	67.35	26.38	516.8	294.02	553.3	18.53	U76992, class A, 20 probes, 20 in U76992 2137-2533, Human Tat-SF1 mRNA, complete cds.
5560.55	6571.45	2970.06	7250.55	3450.19	8316.45	7060.96	7661.9	4873.24	7268.55	468.88	8748.25	681.16	Nuclear Factor Nf-Il6
350.8	1945.25	168.93	2640.75	997.37	1157.85	778.31	1410.45	826.96	2297.7	383.96	1283.2	736.95	D82070, class A, 20 probes, 20 in D82070 285-843, Human aC1 mRNA, complete cds
137.67	4533.75	78.14	1663.85	1503.52	1835.4	1391.44	4600.15	3197.18	4684.75	1843.92	3923.5	831.42	U56998, class A, 20 probes, 20 in U56998 1628-2048, Human putative serine/threonine protein kinase PRK (prk) mRNA, complete cds
55.86	108.65	61.87	197.65	114.62	173.85	115.19	321.55	275.7	624.3	823.21	329.45	308.93	D49488, class A, 20 probes, 20 in D49488 640-1138, Human mRNA for alpha-tocopherol transfer protein, complete cds
1373.91	1617.85	1202.58	2336.7	683.91	3159.4	722.52	1839.5	927.02	3087.5	1652.79	1752.85	551.9	D11094, class A, 20 probes, 20 in D11094 973-1345, Human mRNA for MSS1, complete cds

1007.7	939.4	750.1	1403.35	450.36	878.95	803.91	1067.05	1072.04	1180	925.89	766.15	357.73	U31814, class A, 20 probes, 20 in U31814 1366-1876, Human transcriptional regulator homolog RPD3 mRNA, complete cds
33.59	375.8	72.69	576	384.24	223.4	18.38	300.55	304.41	562.75	379.93	173.9	197.71	M17219, class A, 20 probes, 20 in M17219 758-1286, Human brain guanine nucleotide-binding protein alpha-i subunit mRNA, 5 end
36.35	175.15	62.01	103.85	38.11	279.55	38.82	362.55	109.67	685.15	925.53	151.45	81.53	M25393, class A, 20 probes, 20 in M25393 1686-2253, Human protein tyrosine phosphatase (PTPase) mRNA, complete cds
1507.98	5736.1	1285.24	5909.15	2507.19	3235.1	2893.76	3488.65	802.5	3920.05	837.71	4646.15	45.61	Z37166, class B, 20 probes, 11 in Z37166cds 1006-1252: 9 in reverseSequence, 1432-1570, H.sapiens BAT1 mRNA for nuclear RNA helicase (DEAD family)
323.78	250.3	26.73	874.15	959.61	481.25	371.02	1320.25	880.7	1410	1706.67	608.6	541.64	U40434, class A, 20 probes, 20 in U40434 1561-2071, Human mesothelin or CAK1 antigen precursor mRNA, complete cds
4089.69	3632.25	894.14	5886.15	2891.43	4335.45	785.67	5372.65	1731.35	3646.85	1463.5	4464.2	106.49	U23143, class B, 20 probes, 10 in U23143cds 1258-1426: 10 in reverseSequence, 3604-3844, Human mitochondrial serine hydroxymethyltransferase gene, nuclear encoded mitochondrion protein, complete cds
359.21	962.1	307.17	1009.25	572.97	1806.55	1329.43	931.55	99.07	729.5	365.57	1549.5	138.45	J00129, class A, 20 probes, 16 in J00129mRNA#1 1282-1552: 4 not in GB record, Human fibrinogen beta-chain mRNA, partial cds
1062.5	4012.8	1510.52	2314.1	1519.43	4746.05	175.57	3886.95	2723.85	5061.6	4595.63	3019.3	508.55	Y07604, class B, 20 probes, 10 in Y07604cds 291-483: 10 in reverseSequence, 572-776, H.sapiens mRNA for nucleoside-diphosphate kinase
615.96	556.65	685.96	348.85	91.71	1382.05	796.84	473.85	412.03	288.1	11.46	272.6	108.61	L32179, class A, 20 probes, 20 in L32179mRNA 1028-1556, Human arylacetamide deacetylase mRNA, complete cds
1028.77	7079.7	402.77	4524.75	475.39	10279.25	1549.06	7301.6	1447.73	8277.85	3311.73	8802.6	650.26	D31846, class A, 20 probes, 20 in D31846exon#4 179-713, Human gene for aquaporin-2 water channel, exon1-4, complete cds
862.46	836.6	765.09	1126.15	1254.2	1349.4	846.83	389.6	141.14	868.2	579.69	548.1	404.89	D25217, class A, 20 probes, 20 in D25217 2864-3410, Human mRNA for KIAA0027 gene, partial cds
133.22	125	65.62	736	441.23	332.15	222.53	432.75	417.12	1036.7	1256.39	247.6	99.84	X05153, class A, 20 probes, 20 in X05153mRNA 264-678, alpha-lactalbumin precursor gene extracted from Human alpha-lactalbumin gene
1279.16	17971.35	769.97	19593.3	1040.86	19573.8	10770.37	17668.4	2333.59	16136.55	774.21	21604.95	2734.45	J04823, class A, 20 probes, 20 in J04823mRNA 61-391, Human cytochrome c oxidase subunit VIII (COX8) mRNA, complete cds
712.48	584.75	660.37	82.05	1.06	370.3	120.77	184.3	135.62	849.3	1095.45	358.8	93.76	U79295, class A, 20 probes, 20 in U79295 817-1345, Human clone 23961 mRNA sequence
44.12	1187.9	588.74	1604.8	1035.35	1630.4	957.42	861.3	443.36	650.1	87.12	266.4	92.35	X96849, class C, 20 probes, 20 in all_X96849 49-323, H.sapiens 5 mRNA of PECAM-1 molecule. /gb=X96849 /ntype=RNA
279.52	558.8	62.51	862.15	364.51	1840.2	1502.88	1240.8	410.83	708.1	11.03	1629.8	674.44	U85658, class A, 20 probes, 20 in U85658 2310-2736, Human transcription factor ERF-1 mRNA, complete cds
13.01	97.55	12.37	99.9	51.62	145.1	117.1	61.55	53.95	120.8	117.66	279.6	218.78	M59499, class A, 20 probes, 20 in M59499mRNA#1 3549-3891, Human lipoprotein-associated coagulation inhibitor (LACI) gene

2478.9	6194.8	1479.83	2353.2	589.44	4974.4	2340.66	3731.3	271.95	3139	1005.79	3812.95	952.4	X51441, class B, 12 probes, 11 in X51441cds 28-65: 1 in reverseSequence, 228, Human mRNA for serum amyloid A (SAA) protein partial, clone pAS3-alpha, Human mRNA for serum amyloid A (SAA) protein partial, clone pAS3-alpha
235.68	100.9	63.64	506	198.13	100.1	99.7	436.8	475.6	678.4	197.85	759.9	115.12	M31241, class C, 20 probes, 20 not in GB record, Human complement receptor 1 (CR1) gene
69.44	1619.25	16.9	2243.05	663.62	2121.9	1106.2	2022.3	804.4	2562	844.14	2163.7	1044.54	X73113, class A, 20 probes, 18 in X73113cds 2973-3339: 2 in reverseSequence, 3430-3520, H.sapiens mRNA for fast MyBP-C
439.54	944.8	61.09	1242.95	1181.36	1942.7	2188.07	2735.35	3076.83	702.3	290.76	517.7	127.56	Z38026, class A, 20 probes, 18 in Z38026cds 63-483: 2 in reverseSequence, 536-542, H.sapiens mRNA for FALL-39 peptide antibiotic
321.03	4798.95	32.6	4711.1	326.54	3405.25	3758.06	4693.15	194.81	5254.15	604.79	3917.65	749.32	D64154, class A, 20 probes, 20 in D64154 786-1254, Human mRNA for Mr 110,000 antigen, complete cds
109.25	1508.4	349.59	1488.35	1139.08	2183.9	2191.47	1909.35	999.07	2946.25	256.04	3702.15	36.13	U05861, class B, 20 probes, 15 in U05861exon 50-243: 5 not in GB record, Human hepatic dihydrodiol dehydrogenase gene
611.36	1945.5	403.33	734.2	105.5	3618.35	3234.66	2602.8	1244.51	1838.85	1179.1	1266.7	876.81	X59350, class C, 20 probes, 20 in all_X59350 2679-3220, H.sapiens mRNA for B cell membrane protein CD22
637.46	466.65	263.26	1284.55	174.87	945.1	486.07	765.1	41.44	1364.85	99.07	835.5	687.17	Proto-Oncogene Met, Alt. Splice Form 2 - Also Represents: J02958, U08818
379.43	792.6	998.01	393.8	234.9	855.15	476.38	332.2	40.02	1543.1	494.13	864.3	212.98	X62515, class C, 20 probes, 20 in all_X62515 13260-13708, H.sapiens mRNA for basement membrane heparan sulfate proteoglycan
332.13	1082.1	506.43	1632.25	134.56	1109.65	1008.69	951.65	479.06	858.95	389.83	396.35	54.94	X76223, class A, 20 probes, 20 in X76223exon 2-540, H.sapiens MAL gene exon 4.
97.72	3429.25	1091.84	3337.4	736.81	1473.55	391.1	2159.15	1802.63	2153.85	2202.99	2472.4	786.3	Poliovirus Receptor
78.21	152.2	57.84	216.25	106.7	394.8	279.59	126.15	48.58	413.6	481.54	660.1	359.63	M58583, class A, 20 probes, 20 in M58583 989-1487, Human precerebellin and cerebellin mRNA, complete cds
210.79	852.35	218.85	1134.4	72.12	880.7	1070.14	1710.2	124.88	1973.75	877.45	716.3	39.74	X53065, class C, 20 probes, 20 in all_X53065 85-462, Human SPR2-1 gene for small proline rich protein (exon 2)
290.55	81.5	22.2	195.95	163.41	264.5	216.52	185.5	93.76	255.75	161.01	227.55	79.69	X60484, class C, 20 probes, 20 in all_X60484 683-858, H.sapiens H4/e gene for H4 histone
1189.35	8576.95	1260.14	6132.1	695.51	4737.1	1183.7	6942.2	660.01	5126	4897.7	3612.8	2371.92	U66711, class A, 20 probes, 20 in U66711mRNA 556-1102, Human Ly-6-related protein -9804 gene, complete cds
647.99	2516.4	357.23	2406.1	296.14	2215.5	522.13	1958.85	387.42	3638.9	2100.67	1742.6	490.73	D50911, class A, 20 probes, 20 in D50911 3298-3718, Human mRNA for KIAA0121 gene, complete cds
779.44	1245.3	672.46	428.45	122.68	273.05	92.56	791.3	632.01	1780	1964.34	1630.7	594.11	U11870, class A, 20 probes, 20 in U11870mRNA 1901-2411, Human interleukin-8 receptor type A (IL8RBA) gene, promoter and complete cds
160.51	534.2	227.55	419	83.44	776.9	649.97	829.1	228.82	425.4	251.31	664.75	432.4	M25809, class A, 20 probes, 20 in M25809 1358-1817, Human endomembrane proton pump subunit mRNA, complete cds
428.15	953.45	209.8	784.25	873.77	1018.9	1107.75	391	305.61	991.95	247.13	426.65	502.68	M55210, class A, 20 probes, 20 in M55210mRNA#1 7322-7844, Human laminin B2 chain (LAMB2) gene
13373.65	15576.45	9118.92	23461.35	14536.77	8806.15	3290.8	10927.8	6257.47	10597.8	3837.89	9529.2	1256.39	J03827, class A, 20 probes, 20 in J03827 970-1438, Y box binding protein-1 (YB-1) mRNA

1593.54	2281.9	2099.68	2287.85	2734.31	5789.9	6986.78	9289.75	11391.28	8543.95	6207.34	4687.9	420.73	AB000462, class A, 20 probes, 20 in AB000462 6801-7227, Human mRNA for SH3 binding protein, clone RES4-23A, complete cds
135.76	779.05	273.58	1508.95	1135.54	761.65	647.64	935.9	335.88	557	149.06	1285.85	238.79	M90657, class A, 20 probes, 20 in M90657 581-1163, Human tumor antigen (L6) mRNA, complete cds
288.78	623.45	199.33	349.85	165.39	572.75	533.09	116.2	30.26	494.85	362.25	205.9	221.47	L07956, class A, 20 probes, 20 in L07956 2402-2930, Homo sapiens 1,4-alpha-glucan branching enzyme (HGBE) mRNA, complete cds
442.58	897.35	581.88	1858.2	884.87	1937.1	916.83	1154.55	299.88	1216.55	282.63	958.15	252.22	Z47087, class C, 20 probes, 20 in all_Z47087 1065-1438, H.sapiens mRNA for RNA polymerase II elongation factor-like protein
121.27	156.3	98.29	73.15	6.29	447.8	561.02	102.85	21.28	157.4	129.68	111.9	61.24	L26081, class A, 20 probes, 20 in L26081 1959-2487, Homo sapiens semaphorin-III (Hsema-I) mRNA, complete cds
27.72	56.4	30.69	51.1	3.96	518.8	698.62	42.9	16.12	362.9	464.29	40.75	28.35	K02402, class A, 20 probes, 20 in K02402exon#8 1383-1875, Human coagulation factor IX gene, complete cds
9157.88	51902.65	20548.03	47440.4	18505.41	53644.25	45927.93	44094.85	20345.09	40625.3	15823.21	46529.45	11100.52	
4.6	21.8	18.53	139.6	159.95	142.85	63.57	72.15	78.56	181.35	207.68	673.85	492.92	U19147, class A, 20 probes, 20 in U19147 34-66, Human GAGE-6 protein mRNA, complete cds
93.06	1374.85	417.83	1015.25	1044.47	1011.9	759.15	1538.9	38.75	1995	393.29	1367.2	1396.25	X82207, class B, 20 probes, 10 in X82207cds 876-1098: 10 in reverseSequence, 1166-1412, H.sapiens mRNA for beta-centractin (PC3)
251.31	3673.4	450.29	3575.2	720.68	3095.5	830.28	2677.35	943.21	2973.95	113.07	2310.75	48.86	X80199, class C, 20 probes, 20 in all_X80199 3718-4223, H.sapiens MLN51 mRNA
643.18	1649.45	453.75	800.85	800.23	861.05	783.26	1464.2	657.61	2041.65	991.72	1198.95	987.33	U53786, class A, 20 probes, 18 in U53786 6024-6432: 2 not in GB record, Human envoplakin (EVPL) mRNA, complete cds
1235.95	4015.35	368.76	1670.3	430.2	2044.6	684.48	3330.65	386.86	3003.45	474.4	1965.5	434.02	S57132, class A, 20 probes, 20 in S57132 3108-3615, COL16A1=type XVI collagen alpha 1 chain [human, placenta, mRNA Partial, 3720 nt]
237.09	1340.95	187.31	2250.6	901.56	3623.8	3407.97	2299.55	236.1	1303.3	319.61	2029.95	482.88	X13967, class C, 20 probes, 20 in all_X13967 3247-3806, Human mRNA for leukaemia inhibitory factor (LIF/HILDA)
55.01	412.75	171.19	412.8	7.78	452.2	27.58	1167.8	940.59	678.55	400.86	178.9	72.41	M63262, class C, 20 probes, 16 in all_M63262 161-540: 4 in M63262cds 231-340, Human 5-lipoxygenase activating protein (FLAP) gene
119.71	289.35	70.36	807.15	772.09	1495.65	1434.93	548.5	640.07	264.25	216.87	126.95	86.9	X80062, class B, 20 probes, 12 in X80062cds 1187-1268: 8 in reverseSequence, 1430-1463, H.sapiens SA mRNA
47.52	208.4	64.77	411.5	486.77	1216.9	1141.13	335.15	307.24	316.75	203.29	805.05	384.6	U25771, class A, 20 probes, 20 in U25771 1383-1821, Human ADP-ribosylation factor mRNA, complete cds
361.26	343.8	42.99	1095.8	334.89	274.75	80.54	1451.8	1599.62	444.8	150.05	728.15	477.65	U15641, class A, 20 probes, 20 in U15641 760-1283, Human transcription factor E2F-4 mRNA, complete cds
37.34	138.9	54.16	275.15	260.57	341.5	402.49	238.45	109.81	124.9	5.52	107.9	16.83	S78774, class A, 20 probes, 20 in S78774 411-717, Na+/Ca2+ exchanger [human, neuroblastoma x glioma hybrid NG108-15 cells, mRNA Partial, 760 nt]. /gb=S78774 /ntype=RNA
135.34	181.4	85.42	593.15	415.85	134.2	29.13	223.15	3.61	70.15	77.29	323.85	363.81	L40586, class A, 20 probes, 20 in L40586 1123-1255, Homo sapiens iduronate-2-sulphatase (IDS) mRNA, complete cds

185.97	1128.2	27.29	1263.25	555.29	1252.95	110.38	1730.7	88.95	716.45	883.11	1659.85	436.36	D83703, class A, 20 probes, 20 in D83703 2605-3169, Human mRNA for peroxisome assembly factor-2, complete cds
107.55	189.3	106.35	194.35	7.57	178.75	40.8	857.15	899.79	608	613.06	266.05	130.89	U28727, class A, 20 probes, 20 in U28727 7835-8357, Human pregnancy-associated plasma protein-A preproform (PAPPA) mRNA, complete cds
35.85	202.45	61.87	207.05	17.75	355.95	99.07	303.05	227.19	858.1	976.51	391.65	16.19	U20230, class A, 20 probes, 20 in U20230exon 7-127, Human guanyl cyclase C gene, partial cds. /gb=U20230 /ntype=DNA /annot=exon
408.35	678.95	117.87	1124.2	9.05	726.1	727.19	247.7	122.61	1412.2	1000.7	725.7	6.08	Tyrosine Phosphatase, Epsilon
49.64	75.1	32.53	44.95	10.25	634.25	847.33	781.1	592.56	394.5	30.97	66.4	32.1	U49957, class A, 20 probes, 20 in U49957 5044-5581, Human LIM protein (LPP) mRNA, partial cds
347.83	2566.4	439.4	1937.75	10.11	3272.95	1672.52	2192	322.87	1338.65	1147.7	2704.35	803.49	X52425, class A, 20 probes, 20 in X52425mRNA 3032-3536, Human IL-4-R mRNA for the interleukin 4 receptor
178.9	1893.3	132.23	1372.3	776.4	649.2	722.66	1187.15	20.86	1118.45	1358	744.95	118.02	S80343, class A, 20 probes, 20 in S80343 1609-2077, ArgRS=arginyl-tRNA synthetase [human, ataxia-telangiectasia patients, EBV-lymphoblastoid cells, mRNA, 2120 nt]
102.39	290.7	36.77	143.65	102.04	305.8	185.12	106.5	35.64	113	50.35	195.6	70	D86960, class A, 20 probes, 20 in D86960 5652-6168, Human mRNA for KIAA0205 gene, complete cds
165.25	385.6	19.23	695.3	697.49	1470.65	1763.17	305.05	167.66	1517.2	840.33	425.1	196.58	U27326, class A, 20 probes, 20 in U27326 1666-2123, Human alpha (1,3/1,4) fucosyltransferase (FUT3) mRNA, major transcript I, complete cds
243.53	70	31.54	372.2	309.71	215.65	81.53	295.05	93.55	303.3	310	311.6	57.84	U52077, class A, 20 probes, 20 in U52077cds 428-982, Human mariner1 transposase gene, complete consensus sequence. /gb=U52077 /ntype=DNA /annot=CDS
158.46	361.35	73.61	286.8	50.2	327.15	97.79	308.4	81.46	260.7	32.81	237.45	17.61	X99802, class A, 20 probes, 20 in X99802 1983-2463, H.sapiens mRNA for ZYG homologue
8607.54	14749.25	7264.32	15667.25	14915.36	8855.05	6944.71	6697	3009.16	8602.2	4864.33	13009.55	2338.47	X57959, class A, 20 probes, 20 in X57959cds 264-714, H.sapiens mRNA for ribosomal protein L7
3159.85	6135.05	2625.84	6398.65	2382.17	5433.05	4433.77	4095.55	1378.5	5244.45	874.62	5895.8	798.32	U38846, class A, 20 probes, 20 in U38846 1294-1732, Human stimulator of TAR RNA binding (SRB) mRNA, complete cds
554.23	2302.6	342.1	3274.95	270.33	1923.25	1299.03	2544.1	672.46	1179.4	651.39	3074.55	711.7	J04029, class A, 18 probes, 18 in J04029 1450-2007, Homo sapiens keratin 10 type I intermediate filament (KRT10) mRNA, complete cds
1618.36	8942.1	350.16	8261.3	3498.34	12542.3	2838.47	12853.3	4060.21	8037.2	3458.32	10780.25	1648.48	Cpg-Enriched Dna, Clone S19
1069.71	3603.35	919.73	2764.55	465.77	2564.1	659.16	2476.3	308.44	3086.25	400.86	2662.9	537.54	X93920, class C, 20 probes, 20 in all_X93920 1520-2031, H.sapiens mRNA for protein-tyrosine-phosphatase (tissue type: foreskin)
200.32	597	199.55	775.3	566.11	579	283.13	704.6	224.29	1000.15	1130.45	728.95	648.2	X63131, class C, 20 probes, 20 in all_X63131 1996-2179, H.sapiens My1 (PML) mRNA
572.83	966.25	446.82	310.6	69.44	731.5	488.19	480.7	398.24	567.6	581.95	822.5	390.61	M92269, class B, 20 probes, 13 in M92269cds 6175-6626: 7 in reverseSequence, 6958-7053, Human L-type calcium channel HFCC mRNA, complete cds
30.9	392.6	114.69	802.3	624.23	1099.25	692.89	274.05	60.32	1169.15	975.74	138.35	26.23	Z50053, class C, 20 probes, 20 in all_Z50053 2518-3047, H.sapiens mRNA for alpha2i-subunit of soluble guanylyl cyclase
1008.05	3363.85	336.51	3816.55	2016.74	3017.45	2327.3	3743.2	25.6	3472.75	543.84	3620.3	748.26	AB003177, class A, 20 probes, 20 in AB003177 604-1096, Human mRNA for proteasome subunit p27, complete cds

71.13	557.85	153.09	1044.25	52.54	875.8	209.59	534.9	234.48	243.25	4.6	905.2	158.67	X63563, class A, 20 probes, 20 in X63563cds 3176-3500, H.sapiens mRNA for RNA polymerase II 140 kDa subunit
151.89	444.25	319.26	309.6	173.38	1838.15	1556.55	695.6	224.15	981.6	1059.25	1480.05	673.94	M59216, class C, 20 probes, 19 in all_M59216 1586-2163: 1 in M59216cds 1091, Human gamma-aminobutyric acid-A (GABA-A) receptor beta-1 subunit
11.53	240	227.69	203.5	139.72	147.05	35.85	560.2	148.78	116.75	53.81	298.7	198.56	X07743, class C, 20 probes, 20 in all_X07743 2156-2679, Human mRNA for pleckstrin (P47)
9257.09	30659.9	12814.47	32719.3	11533.48	40695.05	27302.45	29136.35	7202.66	36287.35	4600.93	39355.75	5653.39	D87735, class A, 20 probes, 20 in D87735 127-643, Human mRNA for ribosomal protein L14, complete cds
129.97	55.4	16.97	599	525.38	319.15	165.11	453.05	188.87	1239.1	1413.79	430.35	178.83	S66896, class A, 20 probes, 20 in S66896 1272-1638, squamous cell carcinoma antigen=serine protease inhibitor [human, mRNA, 1711 nt]
440.17	1222.9	669.35	848.95	12.66	1145.95	1006.99	820.6	141.14	970.1	685.75	1635.3	701.59	Homo sapiens /REF=M11507 /DEF=Human transferrin receptor mRNA, complete cds /LEN=2826 (_5, _M, _3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
157.76	281	70.29	675.3	769.05	1004.5	1258.79	797.15	449.37	1265	1318.33	990.7	1244.51	L22214, class A, 20 probes, 20 in L22214 2335-2857, Human adenosine A1 receptor (ADORA1) mRNA exons 1-6, complete cds
263.75	1073.45	632.93	901.45	602.38	2453.2	836.08	2434.65	2922.83	854.9	259.37	3302.15	1431.96	X68688, class A, 20 probes, 20 in X68688mRNA 111-134, H.sapiens ZNF33B gene
1738.78	3026.45	2175.7	2734	2363.57	3049.75	2857.06	2661.45	248.55	1724.8	337.29	3515.3	213.12	U14187, class A, 20 probes, 20 in U14187 824-962, Human receptor tyrosine kinase ligand LERK-3 (EPLG3) mRNA, complete cds.
127	86.45	27.51	674.5	789.27	301.45	277.12	753.1	952.76	857.95	707.74	53.85	14.78	X55330, class C, 20 probes, 20 in all_X55330 1609-2120, H.sapiens mRNA for aspartylglucosaminidase
0.92	208	88.95	362.55	226.63	164.6	121.06	166.6	119.5	179.15	0.64	148.25	127.63	U60805, class A, 20 probes, 20 in U60805 3576-4146, Human oncostatin-M specific receptor beta subunit (OSMRB) mRNA, complete cds
28.78	225.35	33.3	607.35	442.44	327.3	28.43	249.9	62.23	607.05	206.97	247.6	30.97	M85276, class A, 20 probes, 18 in M85276exon#2-5 5-92: 1 in reverseSequence, 5295: 1 not in GB record, Homo sapiens NKG5 gene, complete cds
136.68	430.85	279.94	516.15	396.05	573.6	13.58	508.75	178.69	576.65	308.37	330.15	1.2	U79255, class A, 20 probes, 20 in U79255 760-1180, Human X11 protein mRNA, partial cds.
316.5	1203	472.35	708.25	422.07	559.9	631.59	666.75	104.44	476.15	76.86	1024.95	759.93	X66401, class B, 20 probes, 11 in X66401cds#1 327-615: 6 in fullSequence, 45931-47208: 3 not in GB record, LMP2 gene extracted from H.sapiens genes TAP1, TAP2, LMP2, LMP7 and DOB
646.72	1315.2	765.09	824.65	503.67	909.9	678.12	1655.85	280.37	685.55	166.67	639.95	541.43	X97230, class A, 20 probes, 16 in X97230cds 782-1274: 4 in reverseSequence, 1290-1353, H.sapiens mRNA for NK receptor, clone library 4M1#6
130.25	943.55	145.73	1485.1	980.19	2484.75	1365.49	1072.25	597.01	926.1	246.36	1033.55	463.93	Serine/Threonine Kinase (Gb:Z25427) - Also Represents: X97630
288.71	396.15	264.1	214.8	234.19	1530.95	1554.86	730.1	991.51	176	95.74	240.85	67.53	U61500, class C, 20 probes, 20 in all_U61500 6060-6577, Human GT334 protein (GT334) gene mRNA, complete cds
13652.11	94108.55	5562.32	94601.84	30947.16	109571.95	43682.16	129024.4	11334.78	106502.25	1294.07	116079.05	24879.77	M60854, class A, 20 probes, 20 in M60854 19-373, Human ribosomal protein S16 mRNA, complete cds
216.16	1879.05	184.77	2233.7	649.41	1114.85	1043.34	769.45	393.36	2160.1	1066.46	1533.45	18.74	U15174, class A, 20 probes, 20 in U15174 224-746, Human Nip3 (NIP3) mRNA, complete cds

70.07	1875.4	335.17	1736.5	967.46	1103.25	10.25	1261.4	250.46	1047.2	99.14	780.9	769.76	L28010, class A, 20 probes, 20 in L28010 1292-1748, Homo sapiens HnRNP F protein mRNA, complete cds
59.82	281.7	0.42	227.9	194.31	766.4	618.29	325.2	11.6	386.15	149.98	598.75	569.01	D31628, class A, 20 probes, 20 in D31628cds 781-1132, Human gene for 4-hydroxyphenylpyruvic acid dioxygenase (HPD), comlete cds
15.06	844.4	10.61	558.5	391.31	570.65	106.28	325.3	111.02	770.7	428.79	479.4	249.18	U09410, class A, 20 probes, 20 in U09410 1481-2003, Human zinc finger protein ZNF131 mRNA, partial cds
38.4	332.25	50.42	239.9	73.96	351.6	202.66	683.85	351.79	331.3	171.83	329	25.03	X51755, class C, 20 probes, 20 in all_X51755 8272-8537, Ig light-chain, partial Ke-Oz- polypeptide; Author-given protein sequence is in conflict with the conceptual translation gene extracted from Human lambda-immunoglobulin constant region complex (germline)
81.18	2194.3	349.17	2950.8	18.24	1261.2	940.88	1406.7	298.4	1118.2	366.85	663.3	229.81	X59303, class A, 20 probes, 20 in X59303cds 3274-3773, Human G7a mRNA for valyl-tRNA synthetase
593.26	693	751.51	276.8	228.4	196.75	183.49	553.7	390.32	586.05	489.53	395.35	114.06	X83705, class A, 18 probes, 18 in X83705mRNA 67-535, H.sapiens mRNA for c-sis proto-oncogene.
690.77	828.8	743.59	238.55	91.85	218.15	85.06	787.8	30.41	763.75	198.77	938.65	429.71	L37362, class A, 20 probes, 20 in L37362 1009-1579, Homo sapiens (clone d2-115) kappa opioid receptor (OPRK1) mRNA, complete cds
41.72	959.1	226.13	608.15	574.1	397.35	281.22	1226.15	868.11	738.95	300.17	652.3	496.81	U38964, class B, 20 probes, 10 in U38964cds 1096-1146: 10 in reverseSequence, 1291-1359, Human PMS2 related (hPMSR2) gene, complete cds
98.22	93.75	109.39	468.5	564.13	203.25	18.6	186.05	67.67	196.95	54.09	134.8	13.15	V01515, class C, 20 probes, 20 in all_V01515 5300-5550, Human gene encoding preproglucagon. Glucagon is a 29-amino acid pancreatic hormone which counteracts the blood glucose-lowering action of insulin by stimulating hepatic glycogenolysis and gluconeogenesis. Also included in the proglucagon sequence are two regions (GLP-1 and GLP-2) which are homologous to glucagon itself but not identical.
269.48	571.2	302.78	950.25	13.93	1004.05	998.65	512.75	216.3	674.05	35.43	882.2	791.25	U40343, class A, 20 probes, 20 in U40343 657-1119, Human CDK inhibitor p19INK4d mRNA, complete cds
265.17	994.35	152.95	1208.15	894.14	2738.5	2926.71	3416.9	378.3	4429.2	163.62	3878.15	269.62	Mac25
33.16	202.65	112.08	213.3	242.11	276.75	57.06	817.2	987.12	812.3	639.37	401.9	270.4	U79251, class A, 20 probes, 20 in U79251 1285-1747, Human clone 23878 mRNA sequence
50.84	242.9	264.18	439.25	346.84	371.55	230.87	794.45	239.07	401.9	447.74	497.85	396.19	X95592, class A, 20 probes, 20 in X95592 558-1122, H.sapiens mRNA for C1D protein
210.58	5533.1	1112.7	4131.6	2093.32	4719.9	427.38	4045.9	603.59	3888.4	852.35	3568.8	1017.53	U67171, class A, 20 probes, 20 in U67171 229-697, Human selenoprotein W (selW) mRNA, complete cds. /gb=U67171 /ntype=RNA
68.8	399.45	305.97	232.95	72.05	132.65	75.87	398.45	441.31	338.35	34.86	577.25	666.02	X63417, class A, 20 probes, 19 in X63417cds 170-524: 1 in reverseSequence, 560, H.sapiens irlB mRNA
146.09	559.65	104.86	285.9	119.93	566.55	304.69	798.05	166.81	1422	1681.64	836.05	499.15	K01383, class C, 20 probes, 20 in all_K01383 2141-2388, Human metallothionein-I-A gene, complete coding sequence
1405.94	3415.75	1202.58	2534.1	730.72	1219.85	630.95	2068.15	722.03	2947.05	1620.05	1337.45	1032.59	M31303, class A, 20 probes, 20 in M31303mRNA 933-1407, Human oncoprotein 18 (Op18) gene, complete cds
472.49	7027.25	148.42	5069.1	1043.83	5990.25	1037.96	5976.55	915.35	5499.85	3394.61	4789.8	1967.03	M57567, class A, 20 probes, 20 in M57567 491-953, Human ADP-ribosylation factor (hARF5) mRNA, complete cds

393.22	403.55	505.37	314.5	143.4	246.55	268.91	294.95	190.85	422.5	376.89	241.85	230.16	U73704, class A, 20 probes, 20 in U73704 1248-1764, Homo sapiens 48 kDa FKBP-associated protein FAP48 mRNA, complete cds
216.66	319.65	413.02	202.25	30.33	340.1	417.9	68.4	39.03	265.35	273.01	135.05	115.05	U50708, class A, 20 probes, 20 in U50708 946-1384, Human branched chain alpha-ketoacid dehydrogenase E1 beta subunit mRNA, complete cds
360.91	782.4	333.75	530.85	91.85	1225.85	791.04	1744.15	1643.81	1197.7	709.09	702.95	211.07	U92457, class A, 20 probes, 20 in U92457 2832-3375, Human metabotropic glutamate receptor 4 mRNA, complete cds
1650.74	932.3	502.75	2195.35	724.86	1237.2	169.14	1811.35	877.73	1123.15	1254.62	1148.9	473.76	Cystathionine Beta Synthase, Alt. Splice 3 - Also Represents: L00972
86.83	449.05	2.62	596.7	268.42	985.7	1239.13	202.65	106.56	103.7	10.32	1007.45	552.89	X04327, class A, 20 probes, 20 in X04327mRNA 1084-1564, Human erythrocyte 2,3-bisphosphoglycerate mutase mRNA EC 2.7.5.4
607.33	600	565.83	396.55	419.39	739.8	774.99	579.35	298.75	647.5	280.86	106.65	14.07	D13168, class A, 20 probes, 20 in D13168exon 2254-2800, Human gene for endothelin-B receptor (hET-BR)
487.48	903.6	114.98	538.15	10.96	1159.4	1086.96	92.2	38.04	693.7	382.4	256.1	217.51	M76766, class A, 20 probes, 20 in M76766 679-1129, Human transcription factor (TFIIB) mRNA, complete cds
104.16	116.75	93.97	620	798.75	1298.1	1663.96	146.4	89.52	94.75	81.95	165.8	169.14	U05012, class A, 20 probes, 20 in U05012 2204-2720, Human receptor tyrosine kinase TrkC (NTRK3) mRNA, complete cds
72.05	169.85	30.76	412.7	425.68	196.15	145.88	246.65	129.61	142.3	63.5	159.35	44.9	M97016, class A, 20 probes, 20 in M97016 1231-1803, Homo sapiens osteogenic protein-2 (OP-2) mRNA, complete cds.
678.68	1178	210.86	1213.5	1221.31	1167.15	593.76	1377.35	856.66	1179.1	394.71	1668.85	642.41	Macmarcks
10121.31	23065.8	4632.11	15590.7	3123.71	30660.55	2787.06	24766.1	3714.86	26586.45	362.25	34974.05	3919.14	X93036, class A, 20 probes, 16 in X93036cds 88-163: 4 in reverseSequence, 34-37, H.sapiens mRNA for MAT8 protein
142.91	842.45	524.32	520	273.08	1712.05	1558.96	1140.45	268.91	777.7	87.26	914.25	461.53	U20760, class A, 20 probes, 20 in U20760 4534-4966, Human extracellular calcium-sensing receptor mRNA, complete cds
157.05	204	229.67	155	162.07	1185.9	1619.42	91.5	53.6	95.8	64.21	383.95	175.43	D13666, class A, 20 probes, 20 in D13666 2630-3072, Homo sapiens mRNA for osteoblast specific factor 2 (OSF-2os)
630.74	1277.7	697.91	1030.45	894.28	2646.55	3024.51	1277.45	251.09	1377.65	825.41	1150.25	58.05	D56495, class A, 20 probes, 20 in D56495 1102-1600, Human mRNA for Reg-related sequence derived peptide-2
2.4	450.15	36.84	457.55	19.73	123	35.5	169.55	178.83	558.55	5.44	711	647.29	U15173, class A, 20 probes, 20 in U15173 1781-2303, Human Nip2 (NIP2) mRNA, complete cds
0.71	533.45	116.46	683.15	248.27	1129.45	347.83	564.3	390.04	925	653.08	770.15	821.3	Xe7, Pseudoautosomal Gene, Alt. Splice 2 - Also Represents: L03426
116.6	110.1	130.53	144.4	119.64	203.65	217.29	94.8	74.67	330.75	370.17	64.9	41.86	M31774, class A, 20 probes, 20 in M31774 1858-2392, Human thyrotropin receptor (TSH) mRNA, complete cds
176.42	485.8	114.83	868.25	544.68	1338.7	1032.8	621.1	286.52	507.6	1.27	684.5	324.84	U32576, class A, 20 probes, 20 in U32576mRNA 19-535, Human apolipoprotein apoC-IV (APOC4) gene, complete cds
2688.35	9201	2132.78	6462.35	402.27	3782.25	3625.12	5240.9	1423.26	10053.3	5610.61	7149.05	652.16	D42123, class A, 20 probes, 20 in D42123 655-1135, Human mRNA for ESP1/CRP2, complete cds

117.38	479.1	13.01	570.5	199.4	928.1	525.24	1998.25	1149.26	1389.15	1814.51	761	362.04	U06155, class C, 6 probes, 6 in all_U06155 512-660, Human chromosome 1q subtelomeric sequence D1S553. /gb=U06155 /ntype=DNA /annot=CDS, Human chromosome 1q subtelomeric sequence D1S553. /gb=U06155 /ntype=DNA /annot=CDS
192.47	668.05	302.43	463.45	26.09	961.05	353.34	488.7	97.02	368.2	239	853.8	84	U71374, class A, 20 probes, 20 in U71374 617-1157, Human HsPex13p mRNA, complete cds. /gb=U71374 /ntype=RNA
27.65	132.7	61.24	195.25	144.18	281.2	167.87	679.95	632.79	166.9	19.37	492.6	9.19	M11313, class A, 20 probes, 20 in M11313mRNA 3966-4522, Human alpha-2-macroglobulin mRNA, complete cds
366.35	363.35	472.14	725.15	842.09	970	317.21	211.9	135.06	136.5	8.49	528.3	670.34	B. subtilis /GEN=dapB, jojF /DB_XREF=gb:L38424.1 /NOTE=SIF corresponding to nucleotides 2020-2580 of gb:L38424.1, not 100% identical /DEF=Bacillus subtilis dihydropicolinate reductase (jojE) gene, complete cds; poly(A) polymerase (jojI) gene, complete cds; biotin acetyl-CoA-carboxylase ligase (birA) gene, complete cds; jojC, jojD, jojF, jojG, jojH genes, complete cds s.
566.96	835.25	658.81	423.05	257.74	891.2	261.06	1321.85	1291.25	515.5	529.48	688.4	572.47	M21574, class A, 20 probes, 20 in M21574mRNA 5807-6293, Human platelet-derived growth factor receptor alpha (PDGFRA) mRNA, complete cds
418.04	656.65	264.67	453.45	111.79	1009.95	922.7	186.4	42	173	30.26	707.3	132.79	L32832, class A, 20 probes, 20 in L32832 11295-11853, Homo sapiens zinc finger homeodomain protein (ATBF1-A) mRNA, complete cds.
220.62	648.3	225	468.6	191.77	409.9	168.29	448.1	381.84	428.4	101.54	270.1	67.6	U13666, class A, 20 probes, 17 in U13666cds 671-989: 3 in reverseSequence, 1329-1413, Human G protein-coupled receptor (GPR1) gene, complete cds
8.06	210.85	40.23	536.6	215.95	465.4	207.75	896.95	962.16	203.2	19.37	595.6	401.35	S70609, class A, 20 probes, 20 in S70609 1781-2339, glycine transporter type 1b [human, substantia nigra, mRNA, 2364 nt]
288.78	2137.45	60.46	1598.45	338.35	2301.05	275.56	1886.7	839.48	4876.3	3570.18	2539.2	1071.55	Colony-Stimulating Factor 1, Macrophage, Alt. Splice 3
280.72	720.75	271.32	1961.65	715.66	2124.2	336.58	1267.05	735.32	1576.3	202.09	1843.1	910.33	M81695, class A, 20 probes, 20 in M81695 4051-4597, H.sapiens leukocyte adhesion glycoprotein p150,95 mRNA, complete cds
301.79	443	501.48	539.55	560.24	255.65	39.95	246.25	172.04	770.75	1053.94	191.9	206.05	U18271, class C, 20 probes, 20 not in GB record, Human thymopoietin (TMPO) gene, Human thymopoietin (TMPO) gene
219.42	239.35	297.9	368.2	427.8	392.2	115.82	426.65	476.94	648.15	814.09	120.7	25.03	M26004, class A, 20 probes, 20 in M26004 3326-3894, Human CR2/CD21/C3d/Epstein-Barr virus receptor mRNA, complete cds
36.77	942.75	323.5	739.5	12.3	601.15	93.97	853.25	391.1	591.6	331.07	971.7	237.16	M36653, class A, 20 probes, 20 in M36653 1448-1663, Human 2-Oct factor mRNA, complete cds
48.65	389.05	303.42	1271.65	274	945.6	400.79	1305.45	495.19	1692.6	316.36	830.7	493.84	X53296, class C, 20 probes, 20 in all_X53296 1099-1657, H.sapiens mRNA for IRAP
25923.1	92400.05	9040.01	82149.95	24394.26	99048.6	57257.13	146577.16	19129.57	111027.09	54650.45	151394.3	35325.21	L04483, class A, 17 probes, 17 in L04483 39-272, Human ribosomal protein S21 (RPS21) mRNA, complete cds
1957.63	7266.95	120.7	6569.65	1660.22	5153.75	1322.36	6569.1	1378.43	6323.4	310.14	6284.45	742.53	D13900, class A, 20 probes, 20 in D13900 748-1234, Human mRNA for mitochondrial short-chain enoyl-CoA hydratase, complete cds

894.99	5697.2	935.93	3847.95	1370.73	7179.55	4243.14	6615.25	3115.87	6573.3	4548.11	10134.9	3077.89	U09937, class A, 20 probes, 20 in U09937mRNA 1176-1581, urokinase-type plasminogen activator receptor gene extracted from Human urokinase-type plasminogen receptor
468.18	980.7	217.51	2273.9	50.91	904	552.39	556.9	103.1	1245.95	301.86	774.85	87.19	M97388, class A, 20 probes, 20 in M97388 786-1332, Human TATA binding protein-associated phosphoprotein (DR1) mRNA, complete cds
219.13	305.1	387.64	682.7	677.27	68.3	52.89	424.1	263.61	140.75	155.78	464.3	535.28	X64624, class C, 20 probes, 20 in all_X64624 2881-3429, H.sapiens mRNA for RDC-1 POU domain containing protein
293.45	235.7	281	70.2	3.11	112.35	75.17	559.4	723.65	682.85	896.96	292.4	315.94	U32989, class A, 20 probes, 20 in U32989 1109-1559, Human tryptophan oxygenase (TDO) mRNA, complete cds
635.83	7146.7	851.36	3612.4	2763.37	8690.85	1222.09	7403.65	2117.57	10101.85	2135.53	5729.2	1495.11	L37033, class A, 20 probes, 20 in L37033 1039-1480, Human FK-506 binding protein homologue (FKBP38) mRNA, complete cds
18.17	7561.5	1143.25	6161.9	1775.26	6138.8	2980.88	5984.75	1651.16	7443.05	166.38	7320.35	1371.86	D26598, class A, 20 probes, 20 in D26598 187-571, Human mRNA for proteasome subunit HsC10-II, complete cds
310.99	1304.1	1418.03	1435.7	256.4	1443.05	41.37	2939.5	1499.63	2019.5	954.59	2134.55	258.87	U15637, class C, 20 probes, 20 in all_U15637 1727-2315, Human CD40 binding protein (CD40bp) mRNA, complete cds.
8.84	48.45	36.42	33.4	12.73	789.5	374.77	58.85	6.15	625.35	507.91	30.1	9.76	S75313, class A, 20 probes, 20 in S75313 1197-1387, MJD1=MJD1 protein {CAG repeats} [human, brain, mRNA, 1776 nt]
94.05	549.05	70.22	1319.2	1064.2	462.6	250.74	579.55	635.19	1104.35	1026.22	1114.7	82.73	U77664, class A, 20 probes, 20 in U77664 417-891, Human RNaseP protein p38 (RPP38) mRNA, complete cds.
185.76	298.45	366.92	223.7	24.04	507.35	682	212.45	89.87	662.85	633.92	62.7	26.16	Protein Kinase (Gb:M59287) - Also Represents: L29219
1556.13	3809.3	1202.65	3906.2	2530.45	3131.4	1351.28	4068.65	875.19	4415.2	960.11	3611.7	914.43	U65416, class A, 20 probes, 20 in U65416mRNA 1983-2316, Human MHC class I molecule (MICB) gene, complete cds
151.25	494.6	159.66	566.55	477.51	213.75	87.33	420.5	88.67	705.8	311.41	588.6	188.66	X91868, class C, 20 probes, 20 in all_X91868 915-1378, H.sapiens mRNA for SIX1 protein
18932.64	53918.85	15249.96	45569.55	3842.77	44193.4	21716.24	53764.45	4008.23	43076.05	14822.87	62183.75	8862.1	M24485, class A, 20 probes, 19 in M24485cds 109-604: 1 in reverseSequence, 3966, Homo sapiens (clone pHGST-pi) glutathione S-transferase pi (GSTP1) gene, complete cds
99.84	222.3	100.41	416.7	94.89	241.75	201.74	720.3	219.91	655.4	619.57	490.4	78.35	M57731, class A, 20 probes, 20 in M57731mRNA 617-1032, Human gro-beta mRNA, complete cds
217.93	55.8	1.7	673.95	794.58	60.6	45.82	457.6	531.18	288.85	352.35	451.05	142.06	M63896, class B, 20 probes, 10 in M63896cds 927-1197: 10 in reverseSequence, 1866-2022, Homo sapiens transcriptional enhancer factor (TEF1) DNA, complete CDS
361.12	1207.2	776.4	956.8	557.48	225.75	189.29	958.25	459.69	179.35	132.3	296.3	171.69	S78825, class A, 20 probes, 20 in S78825 523-667, Id1 (Id1-b)=transcription regulator helix-loop-helix protein {alternatively spliced} [human, glioblastoma cell line U251, mRNA, 1146 nt]
204.35	271.3	232.64	197.8	83.3	460.5	520.43	194.25	106.99	233.5	165.04	175.3	40.87	X64559, class B, 20 probes, 14 in X64559cds 202-580: 6 in reverseSequence, 679-823, H.sapiens mRNA for tetranectin

517.18	1263.85	597.15	503.35	451.06	1569.35	1287.29	1363.7	1307.16	184.4	2.26	413.55	263.68	L05425, class A, 20 probes, 20 in L05425 1712-2162, Homo sapiens autoantigen mRNA, complete cds
458.91	231.65	184.63	353	260.07	493.4	20.79	236.7	173.24	408.7	73.26	617	403.62	Z29064, class C, 20 probes, 20 in all_Z29064 3656-4251, H.sapiens AF-1p mRNA
1057.76	12530.8	3743.42	9661.05	1793.01	10467	8837.7	8588.2	3637.5	10492.75	938.83	12833.1	3325.1	X15822, class B, 20 probes, 11 in X15822cds 18-222: 9 in reverseSequence, 13-379, Human COX VIIa-L mRNA for liver-specific cytochrome c oxidase (EC 1.9.3.1.)
429.99	11773.65	2716.21	12004.8	1366.98	9139.3	6991.31	14445.85	508.62	16275.1	3491.83	15775.95	1086.61	U37690, class A, 20 probes, 20 in U37690 31-355, Human RNA polymerase II subunit (hsRPB10) mRNA, complete cds
14.07	48.6	5.52	656.35	880.14	223.15	270.61	231.35	159.17	686.45	907.57	126.35	46.17	S50223, class A, 20 probes, 20 in S50223 197-773, HKR-T1=Kruppel-like zinc finger protein [human, MOLT 4 T-cells, mRNA, 798 nt]
7468.75	19161.05	7641.91	21733.1	6837.02	17373.25	8288.78	23265.05	6690.43	17684.15	11886.82	21439.95	3835.7	X15341, class B, 20 probes, 13 in X15341cds 13-235: 7 in reverseSequence, 374-500, Human COX VIa-L mRNA for cytochrome c oxidase liver-specific subunit VIa (EC 1.9.3.1)
173.03	574.2	247.63	358.55	255.76	466.1	155.28	1048.25	97.79	527.45	362.11	541.4	339.41	X94232, class C, 20 probes, 20 in all_X94232 2035-2528, H.sapiens mRNA for novel T-cell activation protein
391.45	4956.25	186.75	4831.4	1191.19	4525.6	899.02	3639.95	832.05	4299.25	936.85	4106.1	1077.21	X57398, class A, 20 probes, 20 in X57398mRNA 3503-4007, Human mRNA for pM5 protein
313.96	3194	507.14	2721.45	400.01	3356.4	657.33	772.45	237.09	1628.85	665.88	3208.7	225.28	S34389, class A, 20 probes, 20 in S34389 1166-1586, heme oxygenase-2 [human, kidney, mRNA, 1627 nt]
996.38	2073.45	946.6	1382.1	702.72	1253.75	337.5	1616.95	346.27	762.4	529.06	2059.1	548.29	D10537, class A, 20 probes, 20 in D10537 1359-1876, Human mRNA for major structural protein of myelin, complete cds
38.54	46.05	33.59	229.6	17.96	168.05	161.15	99.8	21.35	142.15	91	73.65	5.44	X83863, class A, 20 probes, 20 in X83863cds 1151-1241, H.sapiens mRNA for prostaglandin E receptor (EP3f)
128.41	143.65	142.06	183.45	199.62	709.45	794.58	323.95	13.36	478.2	327.25	454.7	289.49	X72879, class B, 20 probes, 12 in X72879cds 3-52: 8 in reverseSequence, 76-208, H.sapiens 14A2AK DNA sequence
1248.61	1735.45	1407.5	1202.1	131.66	2996.3	1655.76	1089.8	745.15	717.6	881.76	2698.5	642.48	M17262, class C, 20 probes, 19 in all_M17262 16806-26862: 1 in M17262cds 1666, Human prothrombin (F2) gene, complete cds, and Alu and KpnI repeats
210.29	1989.85	360.55	3046.55	41.93	969.8	31.68	2454.05	294.37	1794.45	97.09	1718.8	119.78	Y00264, class C, 20 probes, 20 in all_Y00264 2984-3321, Human mRNA for amyloid A4 precursor of Alzheimer disease
111.72	169.55	191.7	99.35	13.08	197.35	181.66	400.2	62.93	276.35	195.23	143.45	83.51	X02612, class A, 20 probes, 20 in X02612mRNA#3 656-1184, Human gene for cytochrome P(1)-450
265.94	1190.45	20.29	1724.6	855.6	2796.95	2614.1	1900.4	1054.58	1415	1439.39	1596.05	67.95	X96969, class C, 20 probes, 20 in all_X96969 1470-1759, H.sapiens mRNA for urea transporter
651.32	591.5	773.43	1049.25	1173.73	147.85	3.18	595.45	173.45	280.75	7.42	595.2	267.14	Z80787, class C, 20 probes, 20 in all_Z80787 563-700, H.sapiens H4/j gene.
496.74	847.7	684.76	1491.2	916.98	984.6	336.02	739.75	371.44	2162.4	1719.54	460.9	86.69	U06088, class A, 20 probes, 20 in U06088exon 220-730, Human N-acetylgalactosamine 6-sulphatase (GALNS) gene
113.28	941.2	245.93	1147.35	489.39	1508	1098.84	686.15	145.88	1338.7	521.14	1001.25	369.46	M86917, class A, 20 probes, 20 in M86917 2402-2972, Human oxysterol-binding protein (OSBP) mRNA, complete cds
2233.4	10055.45	2359.97	9476.85	4104.12	11742.6	9961.86	7892.75	348.39	7941.7	2609.37	7294.35	96.94	U90878, class A, 20 probes, 20 in U90878 693-1179, Human LIM domain protein CLP-36 mRNA, complete cds.

69.93	150.05	108.82	150.95	125.23	242.55	209.94	155.5	102.25	142.4	64.21	153.15	43.63	D26135, class A, 20 probes, 20 in D26135 3247-3619, Human mRNA for diacylglycerol kinase gamma, complete cds
831.2	1974.65	1081.94	1472	1812.46	1196.7	1272.65	1315.45	158.75	1028.7	395.41	1143.85	998.79	D10522, class A, 20 probes, 20 in D10522 2000-2546, Human mRNA for 80K-L protein, complete cds
35.14	102.05	77.99	636.3	668.22	611.05	608.89	650	290.48	437	159.1	311.95	296.63	U10693, class A, 20 probes, 20 in U10693exon#2 1196-1553, Human MAGE-8 antigen (MAGE8) gene, complete cds
102.95	534.6	12.16	721.45	484.58	1518.05	1029.19	512.9	137.04	845.5	109.74	843.25	684.97	D87467, class A, 20 probes, 20 in D87467 5371-5857, Human mRNA for KIAA0277 gene, complete cds
204.14	819.8	278.46	395.5	231.65	904.95	222.1	1081.95	1189	534.6	109.74	845.6	477.01	J05582, class A, 20 probes, 20 in J05582mRNA 3910-4100, Human pancreatic mucin mRNA, complete cds
196.93	2109.65	281.64	3497.45	1705.75	5233.6	5189.88	3721.8	891.8	3514.95	89.73	3105.1	1180.02	M19045, class A, 20 probes, 20 in M19045 907-1414, Human lysozyme mRNA, complete cds
255.05	635.65	289.84	1011.75	391.81	1081.35	871.65	772.95	575.37	824.7	482.11	111.2	21.78	Y07867, class B, 20 probes, 11 in Y07867cds 643-787: 9 in reverseSequence, 1087-1237, H.sapiens mRNA for Pirin, isolate 1
51.62	2023.05	83.79	2572.9	127.84	2937.55	1183.48	3350.1	441.23	2006.9	61.94	2779.55	358.01	Z71460, class C, 20 probes, 20 in all_Z71460 2546-3033, H.sapiens mRNA for vacuolar-type H(+)-ATPase 115 kDa subunit.
4316.04	12394.25	1979.4	16460.65	6232.09	12946.7	7327.61	14112.95	878.86	12872.7	5079.71	15158.1	1333.6	U78095, class A, 20 probes, 20 in U78095 942-1434, Human Placental bikunin mRNA, complete cds
419.67	3526.35	54.09	2982.25	572.4	5429.9	2049.2	1992	929.28	4756.9	1995.46	3977.6	177.48	D86974, class A, 20 probes, 20 in D86974 5077-5308, Human mRNA for KIAA0220 gene, partial cds
356.95	458.7	495.82	1270.95	1367.62	543.55	428.15	740.1	366.99	799.3	603.3	429.1	324	U92027, class A, 20 probes, 20 in U92027 524-1028, Human clone 61501 defective mariner transposon Hsmar2 mRNA sequence
419.67	816.2	385.66	1333.5	1393.57	1138.2	154.71	658.4	10.47	964.15	773.08	948.2	386.93	X66397, class A, 20 probes, 16 in X66397cds 6605-6977: 4 in reverseSequence, 7352-7442, H.sapiens tpr mRNA
512.58	595.65	547.09	500.3	404.47	1009.9	791.68	573.9	629.89	949.8	4.67	664.9	3.25	U28249, class A, 20 probes, 20 in U28249 919-1405, Human 11kd protein mRNA, complete cds
464.99	3354.85	726.13	1751.8	182.86	3710.2	2855.16	1191.5	773.86	2480.9	221.04	1808.2	331.63	M98343, class A, 20 probes, 20 in M98343 2695-3163, Homo sapiens amplaxin (EMS1) mRNA, complete cds
1001.4	2483.65	1124.94	1852.55	1763.45	1777.45	712.41	2467.15	581.6	555.75	98.78	2578.95	603.37	X15331, class C, 20 probes, 20 in all_X15331 813-1194, Human mRNA for phosphoribosylpyrophosphate synthetase subunit one
178.05	410.55	49.43	593.6	567.81	746.1	1000.13	274.5	262.34	352.85	91.15	333.8	441.23	M30257, class A, 20 probes, 20 in M30257 2214-2709, Human vascular cell adhesion molecule 1 mRNA, complete cds
109.74	335.45	191.13	408.35	133.15	431.85	230.3	421.4	31.4	967.85	809.14	582.9	299.95	U71300, class A, 20 probes, 20 in U71300 1307-1703, Human snRNA activating protein complex 50kD subunit (SNAP50) mRNA, complete cds
101.89	173.5	125.16	214	37.76	439.15	122.54	1038.9	1286.93	765.8	892.23	458.45	300.73	M86757, class A, 20 probes, 20 in M86757 2-372, Human psoriasin mRNA, complete cds.
11.46	66.2	11.74	82.3	58.27	542.2	704.42	70.05	13.08	519.2	612.35	49.4	11.31	S82472, class A, 20 probes, 20 in S82472cds 3-73, beta #NAME? polymerase beta {exon alpha to exon VII region} [human, Genomic, 124 nt, segment 1 of 2]. /gb=S82472 /ntype=DNA /annot=CDS
67.1	215.55	117.73	416.45	469.59	776.7	926.17	128.65	32.31	766.45	462.24	112.05	44.19	X89430, class C, 20 probes, 20 in all_X89430 1828-2333, H.sapiens mRNA for methyl CpG binding protein 2

1187.8	1235.4	1212.83	1174.75	1189.14	573.75	225.5	1738.3	469.52	1020.6	952.61	1783.75	199.62	U28811, class A, 20 probes, 20 in U28811 3404-3866, Human cysteine-rich fibroblast growth factor receptor (CFR-1) mRNA, complete cds
2816.9	4166.15	804.05	1365.8	73.4	3108.85	2578.75	3595.35	1224.92	1401.25	625.44	1226.5	188.37	U73799, class A, 20 probes, 20 in U73799 34-265, Human dynactin mRNA, partial cds. /gb=U73799 /ntype=RNA
316.43	919.85	562.79	2141.3	1150.46	536.7	211.42	1625.45	15.91	1079.8	535.14	1546.25	334.25	X99140, class C, 20 probes, 20 in all_X99140 1407-1822, H.sapiens mRNA for hair keratin, hHb5
217.15	640.9	203.08	554.4	193.04	1324.2	1192.61	206.35	71.49	354.35	209.66	544.55	200.32	J05032, class A, 20 probes, 20 in J05032 1721-2153, Human aspartyl-tRNA synthetase alpha-2 subunit mRNA, complete cds
266.44	201.65	255.76	264.15	271.88	90.65	68.09	549.8	469.52	639.7	647.43	39.25	13.36	U25029, class A, 20 probes, 20 in U25029 1010-1556, Human glucocorticoid receptor alpha mRNA, variant 3 UTR
601.96	1158	1009.32	1276.35	145.45	1195.8	942.57	1137.65	1.48	516.75	292.67	867.55	544.97	U38175, class A, 20 probes, 20 in U38175 719-1205, Human HuR RNA binding protein (HuR) mRNA, complete cds
313.96	1242.95	504.94	960.05	34.29	1076.35	238.79	488.25	436.78	694.8	214.39	876.6	153.44	M37721, class A, 20 probes, 20 in M37721 3297-3705, Human peptidylglycine alpha-amidating monooxygenase mRNA, complete cds
47.02	175.3	50.06	253.95	90.86	204.1	156.55	214.1	135.2	306.35	36.42	211.4	116.25	M12783, class C, 20 probes, 20 in all_M12783 3467-3780, Human c-sis/platelet-derived growth factor 2 (SIS/PDGF2) mRNA, complete cds
851.64	1925.9	863.52	1346.45	681.44	632.7	492.85	1368	284.68	1022.6	1255.82	1041	502.89	D79984, class A, 20 probes, 20 in D79984 5275-5836, Human mRNA for KIAA0162 gene, complete cds
284.47	5062.4	990.8	4916.75	1867.68	7687.7	2528.19	5594.55	1004.73	4210	214.39	5472.55	425.32	U19796, class A, 20 probes, 20 in U19796 406-760, Human melanoma antigen p15 mRNA, complete cds
217.15	245.55	258.45	70.25	64.56	492.6	357.65	336.55	232.57	238.9	69.3	151	124.31	U30998, class A, 20 probes, 20 in U30998 43-166, Human (nmd) mRNA, 3 UTR. /gb=U30998 /ntype=RNA
120	1825.6	613.34	2797.15	655.84	1615.5	915.56	3119.45	1507.2	2519.25	211.35	3054	234.05	U34880, class A, 20 probes, 20 in U34880 1699-2179, Human DPH2L mRNA, complete cds
280.79	952.95	246.85	1079.05	303.99	349.95	222.53	1325.85	534.5	1102.9	344.22	505.4	445.48	X54637, class B, 20 probes, 11 in X54637cds 3359-3539: 9 in reverseSequence, 3857-4133, Human tyk2 mRNA for non-receptor protein tyrosine kinase
181.16	1420.6	423.13	1250.95	726.55	2647.4	3415.18	1741.65	584.28	1275.5	236.46	743	299.95	U90065, class A, 20 probes, 20 in U90065 615-1178, Human potassium channel KCNO1 mRNA, complete cds
154.22	824.15	54.52	1052.45	56.5	829.6	504.17	323.9	217.08	509.9	325.55	806.85	311.62	D64158, class A, 20 probes, 16 in D64158 415-668: 4 not in GB record, Homo sapiens mRNA for ATP binding protein associated with cell differentiation, partial cds. /gb=D64158 /ntype=RNA
472.63	473.4	327.39	152.6	53.74	233.6	210.15	757.15	703.78	318.3	382.12	270.95	173.17	L11695, class A, 20 probes, 20 in L11695 1767-2247, Human activin receptor-like kinase (ALK-5) mRNA, complete cds
546.74	1099.8	560.03	1103.1	1003.24	793.7	429.07	765.75	174.73	1046	323.01	1360.5	117.95	L14813, class A, 20 probes, 20 in L14813 856-1147, Human carboxyl ester lipase like protein (CELL) mRNA, complete cds
2180.01	5905.95	1609.02	5802.05	3168.76	5259.9	4221.99	6262	2331.05	3100.4	3490	5585	976.23	U90913, class A, 20 probes, 20 in U90913 659-1157, Human clone 23665 mRNA sequence
597.08	3136.9	231.22	2063.7	122.61	3111.25	112.36	3120.95	467.89	4168.65	2385.57	1749.05	1528.69	U37012, class A, 20 probes, 20 in U37012 3868-4372, Human cleavage and polyadenylation specificity factor mRNA, complete cds

35.64	211.35	162.71	190.85	137.11	289.05	357.02	258.6	307.59	332.4	179.46	544.95	202.87	U41740, class A, 20 probes, 20 in U41740 7119-7635, Human trans-Golgi p230 mRNA, complete cds
9.05	16.6	10.04	364.45	359	90.05	50.84	197.7	248.76	65.65	51.12	55.75	41.22	D00003, class B, 12 probes, 12 in D00003 1681-1981, Human liver cytochrome P-450 mRNA, complete cds, Human liver cytochrome P-450 mRNA, complete cds
576.43	2382.7	1154.42	3644.15	2955.78	7429.9	7266.23	3115.4	809.07	4741.65	753.28	4063.95	1368.89	AB000896, class A, 20 probes, 20 in AB000896 49-391, Homo sapiens mRNA for cadherin FIB2, partial cds. /gb=AB000896 /ntype=RNA
496.88	1326.1	97.44	519	455.94	439.1	33.52	1163.75	484.01	1233.55	958.06	743.85	256.47	L11353, class A, 20 probes, 18 in L11353 1698-2202: 2 in reverseSequence, 2208-2214, Human moesin-ezrin-radixin-like protein mRNA, complete cds
0.49	207.35	94.96	97.45	18.03	398.9	238.86	154.85	113.21	849.3	997.16	342.8	45.82	M91438, class B, 20 probes, 13 in M91438cds 55-181: 7 in reverseSequence, 300-540, Human kazal-type serine proteinase (HUSI-II) gene, complete cds
233.98	399.6	290.62	632.85	106.28	819.6	545.89	580.65	104.16	782.2	91.08	405.2	442.08	U12387, class A, 20 probes, 20 in U12387 1468-1722, Human thiopurine methyltransferase (TPMT) mRNA, complete cds
250.74	1289.6	176.92	1902.15	129.75	2266.95	690.07	1124.4	317.92	1510.6	323.29	1564.95	329.44	U41515, class A, 20 probes, 20 in U41515 61-397, Human deleted in split hand/split foot 1 (DSS1) mRNA, complete cds
1566.95	7171.55	1208.37	5254.6	656.34	5537	6566.05	4067.65	4872.6	3802.55	4018.42	9001	1246.49	Tubulin, Alpha 1, Isoform 44 - Also Represents: X06956
49.92	86.4	97.44	477.65	470.58	267.35	89.59	318.5	225.57	598.8	692.54	540.25	119.15	M18255, class C, 20 probes, 19 in all_M18255 18-408: 1 in M18255cds 47, PRKACB gene (protein kinase C-beta-2) extracted from Human protein kinase C beta 1 and 2 genes, next to last
265.17	283.4	266.72	260.7	53.46	362.85	103.03	438.7	18.95	446.9	29.42	361.15	73.75	D21205, class A, 20 probes, 20 in D21205 1715-2279, Human mRNA for estrogen responsive finger protein, complete cds
648.63	751.35	486.42	229.9	191.2	74.3	41.01	700.45	158.18	146.65	30.9	188.85	178.54	U71207, class A, 20 probes, 20 in U71207 1846-2224, Human eyes absent homolog (Eab1) mRNA, complete cds.
52.75	197.55	166.95	151.1	16.69	210.85	81.81	307.45	236.67	176.35	98.78	542.95	409.91	U10687, class A, 20 probes, 20 in U10687exon#10 1056-1412, Human MAGE-4a antigen (MAGE4a) gene, complete cds
744.44	5218.75	1052.25	3335.65	986.2	1335.1	1191.9	2159.75	2224.77	4192.2	3284.51	2380.65	161.72	S80437, class A, 20 probes, 20 in S80437 1601-2185, fatty acid synthase {3 region} [human, breast and HepG2 cells, mRNA Partial, 2237 nt]
50.49	128.45	64.7	157.05	173.59	211.35	138.95	135.85	110.95	110.4	38.75	138.05	39.81	X99374, class B, 20 probes, 15 in X99374cds 1846-2174: 5 in reverseSequence, 2211-2318, H.sapiens mRNA for fertilin beta
96.31	767.7	240.98	776.65	436.92	643.2	68.17	812	634.27	519.45	151.96	514	277.04	U12775, class B, 20 probes, 10 in U12775exon 156-342: 10 not in GB record, Human agouti gene
41.65	63.45	14.07	171.85	126.93	175.1	149.2	268.2	144.11	647	671.04	186.2	89.24	L41680, class A, 20 probes, 20 in L41680 1371-1869, Homo sapiens alpha-2,8-polysialyltransferase (PST) gene, complete cds
344.86	733.85	272.73	378.9	2.69	525.1	113	593.65	195.94	514.15	72.9	614.15	106.28	D38550, class A, 20 probes, 20 in D38550 3195-3735, Human mRNA for KIAA0075 gene, partial cds
216.23	413.85	313.46	477.1	140.71	695.9	194.6	310.3	87.96	496.15	457.57	503.8	164.19	M29874, class A, 20 probes, 20 in M29874 2457-2977, Human cytochrome P450-IIB (hIIB1) mRNA, complete cds
585.91	375.35	81.11	978.45	998.22	295.45	156.91	340.9	80.33	279.15	240.77	347.8	98.71	Y09836, class A, 20 probes, 20 in Y09836 999-1335, H.sapiens mRNA for 3 UTR of unknown protein

79.05	707.85	516.26	1388.95	1029.9	2275.35	1393.92	668.8	558.76	818.35	171.19	697.65	77.29	U63541, class A, 20 probes, 20 in U63541mRNA 431-977, Human mRNA expressed in HC/HCC livers and MolT-4 proliferating cells, partial sequence
481.82	1774.25	1297.75	3085.95	691.48	3523.95	37.55	4990.15	1077.98	4060.55	4397.57	6641.9	1413.93	K03189, class A, 20 probes, 20 in K03189cds 2-404, Human chorionic gonadotropin beta subunit gene
43.27	202.75	167.09	142.2	119.78	525.15	713.82	211.6	204.92	125.75	113.77	590.65	339.34	U58091, class A, 20 probes, 20 in U58091 427-883, Human Hs-cul-4B mRNA, partial cds
109.67	1052.1	108.75	1060.9	301.93	1835.65	1326.46	1060.25	312.75	556.5	621.69	1378.35	376.96	U23028, class A, 20 probes, 20 in U23028 1702-2236, Human eukaryotic initiation factor 2B-epsilon mRNA, partial cds
835.73	2814.95	1342.3	1536.7	260.36	1157.7	840.18	2771.7	149.2	2966.8	772.87	3692.85	1089.02	Guanine Nucleotide-Binding Protein Hsr1
854.4	2013.75	716.79	984.5	965.06	2526.2	1770.88	1063.6	222.46	2157.1	1541.63	2229.2	426.39	D30036, class A, 20 probes, 20 in D30036 1743-2283, Human mRNA for phosphatidylinositol transfer protein (PI-TPalpha), complete cds
188.23	908.55	851.14	1879.35	1305.96	2636.8	1223.29	887	740.2	669.4	84.29	1990.95	1651.59	X54667, class A, 16 probes, 16 in X54667cds 110-326, H.sapiens mRNA for cystatin S, H.sapiens mRNA for cystatin S
27.72	56.3	50.91	206.4	229.67	502.85	560.81	109.8	85.84	227.95	185.19	224.5	101.26	U07856, class A, 20 probes, 20 in U07856cds#5 1300-1846, Human endogenous retrovirus in complement C4A gene, A3 allele, HERV-K(C4) (gag), (pol), reverse transcriptase, integrase and (env) genes, complete cds. /gb=U07856 /ntype=DNA /annot=CDS
58.55	88.8	38.61	121.35	104.86	286.85	178.54	257.65	192.26	246.45	32.74	532.75	262.97	M88461, class A, 20 probes, 20 in M88461 993-1538, Human neuropeptide Y peptide YY receptor mRNA, complete cds
106.7	111.6	104.37	245.2	215.1	321.5	304.62	210.25	244.16	128.25	105.29	148.6	18.38	X99350, class A, 20 probes, 20 in X99350mRNA 1863-2424, HFH4_cds gene extracted from H.sapiens HFH4 gene, exon 1 and joined CDS
440.46	389.55	412.6	531.55	693.88	204.25	181.23	368.55	196.36	99.1	29.13	446.15	525.45	U31116, class A, 20 probes, 20 in U31116 1448-2012, Human beta-sarcoglycan A3b mRNA, complete cds
696.92	838.85	480.62	1321.8	1149.9	2055.25	1573.67	1959.85	1390.1	1589.1	653.37	1207.95	239.5	U71364, class A, 20 probes, 20 in U71364 1097-1553, Human serine proteinase inhibitor (P19) mRNA, complete cds
256.4	361.1	210.43	881.45	963.86	1634.55	1892.29	341.65	93.83	1552.25	1860.04	978	238.72	X60992, class C, 20 probes, 20 in all_X60992 2450-3021, H.sapiens CD6 mRNA for T cell glycoprotein CD6
486.14	384.25	462.38	248.3	276.9	352.6	270.26	776.85	602.67	283.75	279.52	642.9	111.16	Y08837, class A, 20 probes, 17 in Y08837 275-473: 3 not in GB record, H.sapiens mRNA for RAD51-like protein. /gb=Y08837 /ntype=RNA
240.56	1215.05	242.75	1678.95	1402.69	1121.4	656.9	1747.05	552.75	1224.2	1075.09	1721.4	86.41	U63336, class A, 20 probes, 20 in U63336 1666-2062, Human MHC Class I region proline rich protein mRNA, complete cds
642.55	1034.05	556.42	859.05	937.27	1088.25	675.64	634.05	382.76	1523.05	165.11	1256.4	1169.41	M64572, class A, 20 probes, 20 in M64572 3431-3923, Human protein tyrosine phosphatase mRNA, complete cds
285.11	737.7	351.29	255.6	229.95	814.75	1111.22	377.1	156.13	281	177.91	769.95	947.59	X85545, class C, 20 probes, 20 in all_X85545 1122-1591, H.sapiens mRNA for protein kinase, PKX1
652.38	1303.9	637.1	1361.9	203.08	1222.95	257.17	2773.05	1681	781.4	773.01	1732.75	185.9	X12433, class C, 20 probes, 20 in all_X12433 1247-1734, Human pHS1-2 mRNA with ORF homologous to membrane receptor proteins
1387.06	2676.5	1148.77	2394.5	1335.16	2705.65	773.79	1981.35	383.61	3201.45	1179.67	2523.55	334.25	S40719, class A, 20 probes, 20 in S40719 2440-2964, glial fibrillary acidic protein [human, glioma cell line U-251 MG, mRNA, 3033 nt]

174.3	227.8	288.36	104.3	14.85	230.85	257.6	361.7	421.86	99.6	58.97	68.75	29.91	U06233, class A, 20 probes, 20 in U06233 2545-3055, Human POU domain protein (Brn-3b) mRNA, complete cds
228.04	6076.1	46.39	5799.5	31.11	6437.1	514.35	6004.5	1013.71	4916.15	655.84	6246.1	758.58	U51336, class A, 20 probes, 20 in U51336 2520-3024, Human inositol 1,3,4-trisphosphate 5/6-kinase mRNA, complete cds
1186.6	2589.4	547.72	1649.4	1233.76	583.2	246.78	2105.4	87.96	1554.75	195.8	1090.2	197.99	AC002045, class A, 20 probes, 20 in AC002045mRNA#1 643-838, A-589H1.1 from Homo sapiens Chromosome 16 BAC clone CIT987-SKA-589H1 ~complete genomic sequence, complete sequence. /gb=AC002045 /ntype=DNA /annot=mRNA, A-589H1.1 from Homo sapiens Chromosome 16 BAC clone CIT987-SKA-589H1 ~complete genomic sequence, complete sequence. /gb=AC002045 /ntype=DNA /annot=mRNA
129.05	2186.45	319.12	1672.65	410.05	2624.4	446.75	1606.4	195.02	1509.2	178.9	1909.15	159.88	U51477, class A, 20 probes, 20 in U51477 2883-3444, Human diacylglycerol kinase zeta mRNA, complete cds
204.71	427.5	200.39	772.35	702.79	577.55	286.59	401.35	17.89	397.7	25.03	386.7	46.1	M89957, class A, 20 probes, 20 in M89957 675-1095, Human immunoglobulin superfamily member B cell receptor complex cell surface glycoprotein (IGB) mRNA, complete cds
489.04	2098.7	112.57	1142.15	608.04	983.9	1130.66	846.55	977.01	920.55	728.25	729.45	814.94	D28589, class A, 20 probes, 20 in D28589mRNA 281-743, Human mRNA (KIAA00167), partial sequence. /gb=D28589 /ntype=RNA
312.47	1596.8	491.16	1945.85	210.93	1672.6	836.93	1961.35	430.98	1400.5	337.01	1751.7	1670.47	X74295, class C, 20 probes, 20 in all_X74295 178-695, H.sapiens mRNA for alpha 7B integrin
274.85	250.3	299.53	444.55	460.82	1864.15	2493.47	472.4	526.79	651.5	716.44	183.7	13.86	X52009, class A, 20 probes, 19 in X52009cds 813-1315: 1 in reverseSequence, 1629, H.sapiens alpha-1 strychnine binding subunit of inhibitory glycine receptor mRNA.
373.56	871.1	313.81	473.4	347.19	1808.05	1284.88	313.6	152.88	392.75	124.8	743.05	406.8	M82809, class A, 20 probes, 20 in M82809 1465-1915, Human annexin IV (ANX4) mRNA, complete cds
207.96	888.8	311.69	1566.85	707.04	1315.35	542.14	584.4	330.78	844.55	343.44	850.6	327.25	U16997, class A, 20 probes, 20 in U16997 1218-1764, Human orphan receptor ROR gamma mRNA, complete cds
111.93	142.65	126.22	166.2	137.04	864.35	1010.1	371.9	346.91	420.35	452.34	146.35	36.27	D90279, class A, 20 probes, 20 in D90279 5154-5590, Human mRNA for collagen alpha 1(V) chain, complete cds
979.27	1874.8	1390.31	3576.35	3131.56	5343.85	4183.31	3618.3	3622.65	20703.2	2875.24	5404.9	2574.58	AB000450, class A, 20 probes, 20 in AB000450 1298-1730, Human mRNA for VRK2, complete cds
906.65	6220.35	1527.7	4683.4	1589.15	3453.2	1927.29	3908.25	873.77	3794.75	960.04	3865.8	1204.91	U02493, class C, 20 probes, 20 in all_U02493 2130-2485, Human 54 kDa protein mRNA, complete cds
33.59	105.55	2.05	132.6	14.14	772.6	999.71	112.9	18.67	112.1	93.34	95.25	74.88	D90070, class A, 20 probes, 20 in D90070 1329-1828, Human ATL-derived PMA-responsive (APR) peptide mRNA
201.31	849.75	174.3	1032.6	894.35	1262.6	718.42	1639.8	117.95	724.95	312.19	1149.3	331.77	U03858, class A, 20 probes, 20 in U03858 806-1040, Human flt3 ligand mRNA, complete cds
280.09	963.65	295.36	797.35	266.65	159.3	106.07	1027.95	238.51	442.45	464.64	445.05	118.58	M13981, class A, 20 probes, 20 in M13981 1253-1300, Human inhibin A-subunit mRNA, complete cds
299.04	368	40.87	393.45	165.39	486.15	103.03	314.7	71.42	423.8	109.18	459.7	90.37	M26682, class A, 20 probes, 20 in M26682 721-1171, Human T-cell translocation gene 1 (Ttg-1) mRNA, complete cds

120	1334.65	384.74	758.25	736.88	1611.45	613.27	1443.8	121.34	1482.2	310.28	1161.65	223.38	M69023, class A, 20 probes, 20 in M69023 524-1088, Human globin gene
661.92	1025.6	630.46	1854.15	505.79	949.75	146.02	1792	1127.55	1708.6	472.35	374.75	187.31	X17094, class C, 20 probes, 20 in all_X17094 3675-4180, Human fur mRNA for furin
10.89	47.15	15.77	89.1	74.67	147.7	127.14	633.7	812.47	662.8	864.79	145.8	13.01	L16782, class A, 20 probes, 20 in L16782 1994-2522, Human putative M phase phosphoprotein 1 (MPP1) mRNA, partial cds
327.39	520.25	269.9	669	695.23	1414.95	1747.9	710.85	564.91	464.2	484.37	371.6	34.08	U35340, class A, 20 probes, 20 in U35340 442-868, Human beta B1-crystallin mRNA, complete cds
274.64	834.4	282.28	763.35	397.18	431.55	369.18	1225.15	1410.32	1041	976.51	785.1	670.34	U17579, class C, 20 probes, 20 in all_U17579 1465-1982, growth hormone-releasing hormone receptor form b gene extracted from Human growth hormone-releasing hormone receptor gene, alternatively spliced forms a, b, and c, partial cds
49.57	270.55	307.66	303.2	35.92	78.75	54.24	78.1	1.27	78.55	25.53	628.4	826.47	D14660, class A, 20 probes, 20 in D14660 739-1249, Human mRNA for KIAA0104 gene, complete cds
891.94	2502.5	346.48	1702.2	384.24	1289.55	386.29	3965.95	35.71	3248.15	885.51	1999.95	157.9	M92843, class A, 20 probes, 20 in M92843 1144-1583, H.sapiens zinc finger transcriptional regulator mRNA, complete cds
586.69	559	584.49	324.15	260.71	513.45	216.45	493.85	215.6	1143.65	1457.42	335.9	14.99	L09708, class A, 20 probes, 20 in L09708mRNA 2255-2795, Human complement component 2 (C2) gene allele b
116.74	229	203.22	88.5	20.36	899.6	580.82	392.75	126.5	437.25	291.12	348.15	388.27	M90696, class A, 20 probes, 20 in M90696 1168-1738, Human cathepsin S (CTSS) mRNA, complete cds
14.5	177.4	185.54	1160.5	1296.98	1066.3	1179.88	402.6	290.06	348.5	215.81	1196.5	1090.92	X80923, class A, 20 probes, 20 in X80923mRNA 31-361, H.sapiens nov gene. /gb=X80923 /ntype=DNA /annot=mRNA
257.95	370.05	366.78	359.5	408.99	1408.35	1897.38	750	166.74	316.3	255.69	435.75	373.28	M94633, class A, 20 probes, 20 in M94633exon 1275-1611, Human recombination acitivating protein (RAG2) gene, last exon. /gb=M94633 /ntype=DNA /annot=exon
1309.49	1412.3	919.52	1350.85	1669.55	845.3	859.42	279.1	100.69	328.85	125.65	317.1	51.9	M81780, class C, 20 probes, 11 in reverse_M81780 4000-4487: 9 in M81780cds#3 175-296, SMPD1 gene (acid sphingomyelinase) extracted from Homo sapiens acid sphingomyelinase (SMPD1) gene, complete cds, ORF 1-3, complete cds, SMPD1 gene (acid sphingomyelinase) extracted from Homo sapiens acid sphingomyelinase (SMPD1) gene, complete cds, ORF 1-3, complete cds, SMPD1 gene (acid sphingomyelinase) extracted from Homo sapiens acid sphingomyelinase (SMPD1) gene, complete cds, ORF 1-3, complete cds
122.97	1513.1	62.65	1679.25	994.55	2214.75	440.32	1567.6	454.25	2641.35	1626.13	1925.1	166.88	Hepatocyte Growth Factor Receptor
319.47	1491.05	515.83	1690.3	48.08	641.5	345.49	1756.3	316.5	1374.05	261.98	1641.95	128.2	X69433, class C, 20 probes, 20 in all_X69433 1312-1733, H.sapiens mRNA for mitochondrial isocitrate dehydrogenase (NADP+)
506.29	1083.5	491.58	503.95	110.8	1725.35	1672.94	1424.45	7.42	710.55	302.15	1582.3	524.25	M20778, class A, 20 probes, 20 in M20778 401-974, Homo sapien, alpha-3 (VI) collagen
639.44	2154.35	690.07	1158.2	685.61	2238.25	198.91	2979.05	1571.54	3338.3	1660.99	2226.3	195.3	U19261, class A, 20 probes, 20 in U19261 1792-2320, Human Epstein-Barr virus-induced protein mRNA, complete cds
356.88	304.25	413.59	43.65	47.59	101.7	69.72	348.7	464.85	86.5	49.07	95.85	88.18	U31973, class A, 20 probes, 20 in U31973 2396-2914, Human phosphodiesterase A subunit (PDE6C) mRNA, complete cds

172.39	215.85	105.01	120.6	130.53	304.3	294.16	599.5	454.81	131.9	3.82	264.25	255.48	M81882, class A, 20 probes, 20 in M81882mRNA 1835-2261, Human glutamate decarboxylase (GAD65) mRNA, complete cds
812.32	1118.9	156.13	1428.75	1609.16	1439.2	28.57	1277.55	493.07	1340.6	1014.13	1995.45	724.86	X17644, class C, 20 probes, 20 in all_X17644 1976-2559, Human GST1-Hs mRNA for GTP-binding protein
135.84	255.05	142.91	935.25	652.73	1063.95	1339.9	407.45	347.26	402.3	361.33	744.25	751.87	D17427, class A, 20 probes, 20 in D17427 3421-3506, Human mRNA for desmocollin type 4
133.71	171.95	157.33	230.5	3.68	354.8	281.71	217.15	117.31	327.9	172.11	300.9	69.16	L04569, class A, 20 probes, 20 in L04569 8323-8890, Homo sapiens (clone hHT-1) L-type voltage-dependent calcium channel a1 subunit (hHT) mRNA, complete cds
36.2	46.6	42.43	56.05	35.85	85.7	91.36	75.3	14	645.25	836.44	140.75	165.11	U57452, class A, 20 probes, 20 in U57452 43-103, Human SNF1-like protein kinase mRNA, partial cds. /gb=U57452 /ntype=RNA
31.04	32.95	2.05	206.3	31.68	307.4	229.39	455.7	0.85	255.15	183.35	154.5	45.68	U54617, class A, 20 probes, 20 in U54617 1233-1737, Human pyruvate dehydrogenase kinase isoform 4 mRNA, complete cds
458.77	638.7	518.87	582.2	88.53	209.1	22.2	632.85	311.62	261.2	52.47	376.55	109.39	X00734, class C, 20 probes, 20 in all_X00734 8016-8239, Human beta-tubulin gene (5-beta) with ten Alu family members
146.44	417.45	290.83	545.5	297.69	468.95	591.64	403.95	226.34	488.2	162.78	257.1	271.1	X99586, class A, 20 probes, 19 in X99586cds 12-223: 1 in reverseSequence, 329, H.sapiens mRNA for SMT3C protein
106.99	131.6	70.99	93.8	63.64	238.9	281.99	60.15	14.78	65.9	24.47	70	40.45	M35999, class C, 20 probes, 20 in all_M35999 3904-4463, Human platelet glycoprotein IIIa (GPIIIa) mRNA, complete cds
90.44	292.7	258.24	614.95	678.19	1704.65	2151.37	299.3	164.33	827.85	339.76	339.4	285.25	M57464, class A, 20 probes, 20 in M57464 3899-4439, Human ret proto-oncogene mRNA, complete cds
651.1	1085.1	667.79	669.85	270.33	278.3	57.56	523.15	57.35	405.3	188.09	466.6	457.07	U90914, class A, 20 probes, 20 in U90914 1369-1795, Human clone 23587 mRNA sequence
40.87	18007.85	785.81	16884.95	5966.07	12831.4	9637.02	13392.5	2894.33	10862.3	8779.86	14094.85	8307.87	L08666, class A, 20 probes, 20 in L08666 953-1421, Homo sapiens porin (por) mRNA, complete cds and truncated cds
9554.85	19654.95	5363.33	13122.6	4050.45	21106.05	8010.6	25605.05	3376.22	25431.05	7152.03	24561.65	4241.72	X64177, class B, 20 probes, 12 in X64177cds 8-147: 8 in reverseSequence, 2-277, H.sapiens mRNA for metallothionein
193.04	172.1	169.56	178.3	215.81	81.7	72.83	323.1	281	1014.1	1385.65	78.35	48.72	U92459, class A, 20 probes, 20 in U92459 2702-3194, Human metabotropic glutamate receptor 8 mRNA, complete cds
45.33	1022.65	20.44	1494.5	843.15	705.35	833.89	843.35	600.83	531.95	486.14	906.9	720.97	S78271, class A, 20 probes, 20 in S78271 4580-5111, SB1.8/DXS423E=mitosis-specific chromosome segregation protein SMC1 homolog [human, HT1080 and M426 fibroblast cell lines, mRNA, 5190 nt]
42.78	35.7	39.46	261.75	353.62	104.95	112.5	86.4	86.27	861.95	1197.2	200.15	228.89	M10950, class C, 20 probes, 20 in all_M10950 57-304, Human alpha-fetoprotein (AFP) gene
569.65	497.5	565.26	254.3	237.73	247.55	63.99	215.9	143.54	110.25	49.14	250.1	138.73	Z35309, class A, 20 probes, 20 in Z35309cds 3385-3727, H.sapiens mRNA for adenylyl cyclase
386.93	432.75	215.74	497	188.94	692.5	389.05	1290.4	576.72	1800.75	1322.78	969.35	390.68	Y00451, class A, 20 probes, 17 in Y00451cds 1461-1890: 3 in reverseSequence, 2009-2037, Human mRNA for 5-aminolevulinate synthase
300.31	586.45	360.55	358	353.55	161.5	99.14	1029	173.67	1034.75	201.03	570.1	360.06	M73720, class A, 20 probes, 20 in M73720mRNA 1032-1584, Human mast cell carboxypeptidase A (MC-CPA) gene

960.96	4413.95	1258.16	4155	444.49	3466.6	3189.19	4996.1	1413.22	4326.75	678.61	2907.7	39.6	X93510, class A, 20 probes, 17 in X93510cds 548-956: 3 in reverseSequence, 1021-1069, H.sapiens mRNA for 37 kDa LIM domain protein
205.49	585.1	149.48	406.2	166.88	1627.55	1654.42	505.2	322.86	657.6	676.56	562.35	715.66	D25539, class A, 20 probes, 20 in D25539 3997-4375, Human mRNA for KIAA0040 gene, complete cds
353.91	1361.6	447.17	1965.15	744.8	1968.6	75.52	3053.3	1021.34	1790.6	473.48	2004.5	813.88	Fibroblast Growth Factor Receptor K-Sam, Alt. Splice 1
55.72	186.6	134.35	363.05	95.11	276	71.7	280.85	124.66	379.55	323.22	241.55	86.2	M20203, class A, 20 probes, 20 in M20203cds 242-364, Human neutrophil elastase gene
347.61	471.25	328.31	885.15	690.91	372.3	300.94	455.5	301.65	374.6	359.35	405.35	374.27	D87440, class A, 20 probes, 20 in D87440 3807-4245, Human mRNA for KIAA0252 gene, partial cds
508.13	3051.2	487.06	2785.55	406.09	2169.95	288.99	4383.2	851.5	3878.65	384.74	2660.65	206.4	Z47038, class A, 20 probes, 20 in Z47038cds 267-698, Human partial cDNA sequence, clone x101, putative microtubule-associated; protein 1A (MAP1A). /gb=Z47038 /ntype=DNA /annot=CDS
66.61	584.7	240.7	1143.95	944.06	293.15	95.25	225.45	84.5	482.75	137.96	553.4	10.75	D49817, class A, 20 probes, 20 in D49817 1233-1725, Human mRNA for fructose 6-phosphate,2-kinase/fructose 2,6-bisphosphatase, complete cds
782.34	1991.45	224.22	3520.35	1896.39	3153.55	2140.48	1119.3	239	1456.55	778.88	1556.75	19.73	M90656, class A, 20 probes, 20 in M90656 2044-2590, Human gamma-glutamylcysteine synthetase (GCS) mRNA, complete cds
151.96	182.15	148.99	81.05	89.31	302.75	368.61	59.05	22.42	96.35	16.19	149.05	84.22	M30773, class A, 20 probes, 20 in M30773 1983-2499, Human calcineurin B mRNA, complete cds
61.45	67.05	63.99	74.2	42	164.75	100.76	171.3	120.21	54.1	15.84	59.95	22.42	X15217, class C, 20 probes, 20 in all_X15217 2292-2875, Human sno oncogene mRNA for snoA protein, ski-related
8023.47	9351.85	7963.37	20995.3	26846.44	47136.2	48196.11	14628.05	12887.38	18467.55	16802.34	8478.6	1640.91	X03453 Bacteriophage P1 cre recombinase protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)
482.03	430.4	593.55	447.35	470.3	2422.3	2450.83	457.45	611.58	130.15	93.55	135.3	12.45	U80226, class A, 20 probes, 20 in U80226 1440-1476, Human gamma-aminobutyric acid transaminase mRNA, partial cds. /gb=U80226 /ntype=RNA
7378.24	27354.35	8511.23	24041.65	12891.33	22384.75	21498.95	27018.1	11114.02	30484.65	2624.43	32032.9	2300.64	Nucleoside Diphosphate Kinase Nm23-H2s
1761.12	4510.9	1601.17	3721.25	382.62	2416.05	3002.45	4949.3	720.4	4414.4	13.86	3652.95	189.43	U05875, class A, 20 probes, 20 in U05875 1655-2105, Human clone pSK1 interferon gamma receptor accessory factor-1 (AF-1) mRNA, complete cds
1012.86	1216.1	687.02	777.8	294.72	2444.1	530.05	1760.45	1207.95	3386.95	3406.49	1201.65	12.8	D13305, class A, 20 probes, 20 in D13305 1572-1992, Human mRNA for brain cholecystokinin receptor
24.4	396.8	136.33	480.6	10.47	120.55	42.5	658.65	353.62	986.65	482.03	689.45	90.16	L10123, class A, 20 probes, 20 in L10123 3572-4100, Homo sapiens surfactant protein A mRNA, complete cds
3.11	1029.65	255.62	491.25	356.03	2334.95	1773.78	1372.8	477.01	1269.8	862.1	1346.25	824.42	M68520, class A, 20 probes, 20 in M68520 1708-2170, Human cdc2-related protein kinase mRNA, complete cds
253.99	1391.05	168.08	2316.45	543.55	1092.8	1298.67	766	662.84	1838.05	327.32	1834.2	922.63	U81802, class A, 20 probes, 20 in U81802 2557-3043, Human PtdIns 4-kinase (PI4Kb) mRNA, complete cds.
31.82	103.95	67.25	75.8	51.05	307.85	35.57	169.3	94.33	185.35	163.7	181.6	64.35	M24748, class C, 20 probes, 20 in all_M24748 1170-1531, THRA1 gene (thyroid receptor alpha-1) extracted from Human thyroid hormone receptor alpha 1 (TR-alpha-1) gene, complete cds
1277.95	5354.25	1324.2	2912.6	74.25	6579.15	4159.56	8276.95	1182.78	8556.7	741.05	5356.8	1348.31	X02419, class A, 20 probes, 20 in X02419mRNA 1754-2210, H.sapiens uPA gene
37.69	106.3	37.48	209.05	37.69	125.9	16.83	445.6	562.29	126.95	98.22	81.5	4.53	U03911, class A, 20 probes, 20 in U03911 2485-3013, Human mutator gene (hMSH2) mRNA, complete cds

99.91	148.2	74.81	119.1	46.95	412.65	361.97	336.8	12.87	212.2	10.61	235.8	25.88	X04143, class B, 20 probes, 15 in X04143cds 49-265: 5 in reverseSequence, 1430-1592, Human gene for bone gla protein (BGP)
502.33	603.4	219.91	542.7	372.36	640.35	636.04	1366.15	739.56	1588.1	1049.21	820.6	340.83	U43083, class A, 20 probes, 20 in U43083 1408-1582, Human G alpha-q (Gaq) mRNA, complete cds
123.89	199.35	114.2	215.15	137.39	1101.9	1010.88	725.4	650.54	288.8	82.45	347.15	99.91	M57732, class A, 20 probes, 20 in M57732mRNA 2643-3165, Human hepatic nuclear factor 1 (TCF1) mRNA, complete cds, clones HCL10, HCL12, HCL17, and HCL20
42.78	182.05	199.19	96.3	0.57	372.25	331.14	420.6	113	732.6	996.6	108.4	58.83	D87683, class A, 20 probes, 20 in D87683 6419-6941, Human mRNA for KIAA0243 gene, partial cds
167.66	621.1	168.43	945.9	78.49	258.15	216.45	1077.55	393.79	764.15	318.83	414.05	280.51	M80359, class A, 20 probes, 20 in M80359 2409-2835, Human protein p78 mRNA, complete cds
185.9	255.25	161.43	328.15	177.41	523.55	199.19	483.4	551.26	669.6	450.29	538.25	541.01	L19297, class A, 20 probes, 20 in L19297 602-1034, Human nuclear-encoded mitochondrial carbonic anhydrase (CA5) mRNA, complete cds
259.65	1422.4	214.54	1298.25	310.35	659.05	79.13	1071.55	240.49	1624.5	144.82	1517.35	20.15	Ras-Like Protein Tc21
204.64	269.35	198.06	815.8	951.62	1022.35	177.84	1024.9	868.33	777.05	315.86	445.85	120.7	U62800, class A, 20 probes, 20 in U62800 181-535, Human cystatin M (CST6) mRNA, complete cds
513.08	1963.85	147.86	483.35	267.5	1890.85	741.54	1490	91.22	2030.85	6.58	1999.35	614.69	M95610, class A, 20 probes, 20 in M95610 1971-2493, Human alpha 2 type IX collagen (COL9A2) mRNA, partial cds
27.51	98.85	83.23	100.25	31.75	473.75	326.19	164.45	92.56	267.6	159.52	127.4	11.88	D87011, class B, 20 probes, 10 in D87011cds 1308-1590: 10 in fullSequence, 24060-24222, Human (lamuda) DNA for immunoglobin light chain
2118.85	3891.85	1276.54	2115.1	559.32	1700.65	892.86	1507.9	638.66	2751.75	63	1629.75	1413.72	U70321, class A, 20 probes, 20 in U70321 1127-1643, Human herpesvirus entry mediator mRNA, complete cds
92.7	247.9	167.3	210.4	79.62	153.45	102.46	323.1	353.41	157.2	63.36	153.65	73.19	U43672, class A, 20 probes, 20 in U43672 2939-3443, Human putative transmembrane receptor IL-1Rrp mRNA, complete cds
237.38	364.7	170.41	162.45	20.86	855.75	879.43	219.1	99.7	909.9	1025.02	268.05	21.85	U43843, class A, 20 probes, 20 in U43843 813-1374, Human h-neuro-d4 protein mRNA, complete cds
55.65	60.85	47.87	46.2	49.21	523.9	678.26	79.1	71.13	69.1	45.54	193.05	145.88	Z49208, class C, 20 probes, 20 in reverse_Z49208 20545-20696, Human DNA from cosmid L161A8, Huntington Disease Region, chromosome 4p16.3
25.24	29.25	13.79	226.95	76.72	84.3	82.73	57.1	58.12	558.75	644.24	174.15	119.29	U53468, class A, 20 probes, 20 in U53468 862-1390, Human NADH:ubiquinone oxidoreductase subunit B13 (B13) mRNA, complete cds
55.93	541.45	206.4	1028.35	406.09	507.6	306.88	451.45	314.73	1799.2	475.32	334	118.37	M32373, class A, 20 probes, 20 in M32373mRNA 2201-2753, Human arylsulfatase B (ASB) mRNA, complete cds
93.41	121	104.09	264.95	345.56	1689.75	2179.94	858.3	909.48	181.8	189.79	221.75	90.72	X68090, class A, 20 probes, 20 in X68090cds 5-61, H.sapiens Fc-gamma-RIIA gene for IgG Fc receptor class IIA (5 flank). /gb=X68090 /ntype=DNA /annot=CDS
32.39	49.15	16.48	279.15	155.21	959.65	1150.53	519.9	233.63	453.5	568.51	45.6	2.69	X66533, class C, 20 probes, 20 in all_X66533 1884-2365, H.sapiens soluble guanylate cyclase small subunit mRNA
354.97	555.1	258.24	251.25	76.44	980.15	963.43	592.25	509.33	628.9	625.65	1045.5	7.92	U26403, class A, 20 probes, 20 in U26403 991-1495, Human receptor tyrosine kinase ligand LERK-7 precursor (EPLG7) mRNA, complete cds
63.43	101.7	75.94	230.65	2.05	104.05	11.53	113.95	60.88	185.8	75.09	601.6	335.59	L06633, class A, 20 probes, 20 in L06633 1153-1669, Human transcription factor mRNA, complete cds

307.24	230.6	0.42	153.45	82.24	85.7	40.45	284.8	333.33	69.85	36.27	351.9	433.88	U89355, class A, 20 probes, 20 in U89355 3288-3618, Homo sapiens clone cRT16 CREB-binding protein mRNA, partial cds.
787.65	1140.2	820.67	766.25	386.01	725.05	314.17	1195.9	1007.91	1380.15	501.27	1417.45	1073.46	L04953, class A, 20 probes, 20 in L04953 2585-3065, Human x11 protein (x11) mRNA, 3 end
321.38	541.95	352.78	254.2	20.08	1333.35	650.75	337.25	400.86	292.1	121.62	581.1	72.97	J05401, class A, 20 probes, 20 in J05401 1158-1554, Human sarcomeric mitochondrial creatine kinase (MtCK) gene, complete cds
31.82	81.1	25.03	499.3	458.77	630.5	764.38	331.5	175.93	184.45	155.07	293.4	343.51	Y10517, class A, 20 probes, 20 in Y10517mRNA 252-606, H.sapiens mRNA for CD108 protein. /gb=Y10517 /ntype=RNA
615.4	884.65	593.33	544.5	215.53	109.55	0.78	529.05	0.64	336.9	335.17	165.75	99.49	U59423, class A, 20 probes, 20 in U59423 1481-1943, Human Smad1 mRNA, complete cds.
238.37	530.25	93.97	243.25	136.4	493.6	69.86	587.2	702.72	418.2	398.95	399.4	299.95	X53800, class C, 20 probes, 20 in all_X53800 377-961, Human mRNA for macrophage inflammatory protein-2beta (MIP2beta)
880.35	1188.15	1007.7	1087.1	1133.78	1462.7	962.94	620.2	559.75	283.35	171.19	724.15	273.16	M26901, class A, 20 probes, 16 in M26901cds 808-1187: 4 in reverseSequence, 218-293, Human renin gene
41.44	77.85	13.65	76.05	49	187.4	139.02	142.1	47.23	73.45	39.53	140.45	45.89	U67988, class A, 20 probes, 20 in U67988 1623-2163, Human guanylate kinase associated protein (GKAP) mRNA, complete cds
949.57	5130.15	299.6	4035.9	590.15	2459.4	2507.12	3443.35	2205.68	3562.85	650.18	3093.4	417.9	U52522, class A, 20 probes, 20 in U52522 1047-1581, Human arfaptin 2, putative target protein of ADP-ribosylation factor, mRNA, complete cds
896.26	1092.45	700.95	627.45	290.27	971.15	751.3	327.75	105.15	1498.1	1433.87	777.65	441.02	M32598, class B, 20 probes, 12 in M32598cds 2214-2448: 8 in reverseSequence, 146-368, Human muscle glycogen phosphorylase (PYGM) gene
166.38	403.05	217.86	146.35	143.05	345.95	11.1	303.6	123.74	604.75	466.34	738.75	313.6	U01120, class A, 20 probes, 20 in U01120 2484-2982, Human glucose-6-phosphatase mRNA, complete cds
1933.23	31300.45	2304.53	27879.5	5430.86	41083.7	30488.04	42618.55	2908.54	41293.95	401.28	38678.25	11707.21	J03592, class A, 20 probes, 20 in J03592 707-1085, Human ADP/ATP translocase mRNA, 3 end, clone pHAT8
49.64	83.7	76.79	589.25	741.26	275.3	133.5	130.5	101.54	522.25	603.09	475.35	266.23	M61826, class A, 20 probes, 20 in M61826exon 72-568, Human alpha-spectrin gene
22.06	281.5	47.38	243.15	206.4	348.8	358.64	512.55	420.66	745.1	837.78	215	210.01	X56807, class C, 20 probes, 20 in all_X56807 2617-3194, Human DSC2 mRNA for desmocollins type 2a and 2b
6573.34	11232.75	4505.61	14747.65	9039.72	10046.35	5938.21	13610.95	2269.74	8042.9	5860.93	13650.8	4001.23	M21142, class B, 20 probes, 12 in M21142cds#1 889-1109: 8 in reverseSequence, 1884-2122, guanine nucleotide-binding protein G-s-alpha-3 gene extracted from Human guanine nucleotide-binding protein alpha-subunit gene (G-s-alpha)
1811.18	2472.6	1357.93	2665.2	58.97	2117.55	601.54	1961.65	367.34	2371.35	142.62	1790.95	1271.73	Ras-Related C3 Botulinum Toxin Substrate
60.81	156.85	34.58	229.55	125.09	468.1	130.67	376.6	415.07	248.15	256.61	183.7	143.54	L05568, class A, 20 probes, 20 in L05568 1937-2459, Human Na+/Cl- dependent serotonin transporter mRNA, complete cds
14.85	54.8	2.69	508.35	660.93	690.65	575.66	263	161.36	105.55	15.63	118.8	37.05	Oncogene Ret/Ptc, Fusion Activated
36.13	1403.4	61.24	863.35	815.65	626.15	718.77	441.35	261.98	976.75	562.08	1048.8	164.19	M95809, class A, 20 probes, 20 in M95809 1310-1850, Human basic transcription factor 62kD subunit (BTF2), complete cds

233.2	2346.2	640.64	994.95	1193.1	1046.75	1093.82	659.35	570.14	871.8	846.41	1658.45	966.83	X01060, class C, 20 probes, 20 in all_X01060 4427-4986, Human mRNA for transferrin receptor
781.64	519.1	474.47	465.6	388.48	487.2	70.85	732.05	544.12	609.45	484.16	230.4	67.74	U31201, class C, 20 probes, 20 not in GB record, Human laminin gamma2 chain gene (LAMC2), Human laminin gamma2 chain gene (LAMC2)
668.71	1122.15	909.69	1535.8	1242.81	479.85	394.64	485.35	307.52	398.75	152.38	1039.55	471.43	U73514, class A, 20 probes, 20 in U73514 376-892, Human short-chain alcohol dehydrogenase (XH98G2) mRNA, complete cds.
585.91	1179.7	538.25	1098.9	629.33	3007.45	3173.28	1009.7	739.35	1142.35	474.26	637.2	513.93	U57341, class C, 6 probes, 6 in all_U57341 2-129, Human neurofilament triplet L protein mRNA, partial cds. /gb=U57341 /ntype=RNA, Human neurofilament triplet L protein mRNA, partial cds. /gb=U57341 /ntype=RNA
27.29	5714.6	945.4	5034.45	533.65	2339.2	2069.56	4397.05	1050.12	3066.15	709.72	4331.45	129.61	X04654, class B, 20 probes, 10 in X04654cds 1467-1819: 10 in reverseSequence, 2524-2634, Human mRNA for U1 RNA-associated 70K protein
98.92	298.9	56.57	502.7	101.12	368	403.19	209.4	19.09	152.05	89.59	158.65	22.56	U48730, class A, 20 probes, 20 in U48730 2629-2690, Human transcription factor Stat5b (stat5b) mRNA, complete cds.
168.01	649.4	109.46	638.2	26.87	872.45	282.49	673.15	150.68	647.85	15.2	634.25	94.68	Homo sapiens /REF=M27830 /DEF=Human 28S ribosomal RNA gene, complete cds /LEN=1955 (_5, _M, _3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
99.42	153.95	22.42	748.6	673.02	148.8	101.26	700.75	146.44	143.6	22.2	653.95	588.24	U49020, class A, 20 probes, 20 in U49020mRNA#1 4711-5228, MEF2A gene (myocyte-specific enhancer factor 2A, C9 form) extracted from Human myocyte-specific enhancer factor 2A (MEF2A) gene, first coding
869.95	1445.85	990.02	1453.6	615.18	625.25	112.36	1157.9	748.97	925.6	509.4	1143.35	100.9	Atpase, Ca2+ Transporting, Plasma Membrane 1, Alt. Splice 6
891.8	1432.6	914.85	1632.3	948.23	1318.05	604.08	853.85	275.84	2664.25	1168.35	1880.65	1078.27	S71043, class A, 20 probes, 20 in S71043mRNA 442-970, Ig alpha 2=immunoglobulin A heavy chain allotype 2 {constant region, germ line} [human, peripheral blood neutrophils, Genomic, 1799 nt]
984.58	1032.85	800.23	339.2	126.29	320.75	225.07	634.35	483.87	1648.75	1398.59	1213.9	499.08	M34516, class C, 11 probes, 11 in all_M34516 426-469, Human omega light chain protein 14.1 (Ig lambda chain related) gene, Human omega light chain protein 14.1 (Ig lambda chain related) gene
123.18	204.6	121.62	146.65	121.13	697.85	876.18	264.5	240.27	319.95	138.1	153.75	6.15	K03494, class A, 19 probes, 18 in K03494cds 557-1062: 1 in reverseSequence, 115, Human green cone photoreceptor pigment gene 1
110.03	1557.9	126.85	1446.35	123.53	1827.4	994.48	1754.4	388.06	805.15	811.69	1363.9	128.83	X98263, class C, 20 probes, 20 in all_X98263 520-1019, H.sapiens mRNA for M-phase phosphoprotein, mpp6
760.56	2696.3	969.87	2033.35	1415.98	3935.7	1313.24	3553.8	2599.75	6131	4623.91	1806.2	1268.13	D11086, class A, 20 probes, 20 in D11086 976-1408, Human mRNA for interleukin 2 receptor gamma chain
66.11	107	10.47	365.4	279.87	189.65	210.93	310.9	394.71	92.55	10.82	141.05	145.73	M81883, class A, 20 probes, 20 in M81883mRNA 3226-3538, Human glutamate decarboxylase (GAD67) mRNA, complete cds
97.23	116.65	23.55	170.9	47.66	1503.95	1571.26	702.65	447.25	740.1	67.18	678.1	117.95	S78693, class A, 20 probes, 20 in S78693 2-40, alpha CREB-1=cyclic AMP response element-binding protein-1 alpha isoform {alternatively spliced, internal fragment} [human, placenta, mRNA Partial, 64 nt]. /gb=S78693 /ntype=RNA

18.53	409.9	161.64	927.5	749.11	849.85	731.08	502.6	101.4	439.95	150.97	726.1	758.87	D26361, class A, 20 probes, 20 in D26361 5991-6543, Human mRNA for KIAA0042 gene, complete cds
98.22	62.5	50.06	366.35	376.39	750.15	489.11	248.25	184.77	159	142.13	95.35	27.51	X89399, class C, 20 probes, 20 not in GB record, H.sapiens mRNA for Ins(1,3,4,5)P4-binding protein
34.01	100.55	42.64	86.35	112.92	953.55	948.87	71.3	51.05	59	61.24	145.15	135.41	U79751, class A, 20 probes, 20 in U79751 2171-2615, Human basic-leucine zipper nuclear factor (JEM-1) mRNA, complete cds. /gb=U79751 /ntype=RNA
68.45	2272.85	195.23	1914.95	153.94	2307.05	341.18	1851.65	1079.12	827.25	615.25	2213.3	758.44	Y08302, class B, 20 probes, 12 in Y08302cds 807-1122: 8 in reverseSequence, 1385-1445, H.sapiens mRNA for MAP kinase phosphatase 4
104.37	870.3	364.3	1038.95	195.8	509.05	349.81	599.95	21	1376.55	809.28	730	869.03	U19247, class A, 20 probes, 20 in U19247mRNA 1469-1939, Human interferon-gamma receptor alpha chain gene
32.24	1823.8	85.7	1332	1078.76	1660.2	799.6	1583.5	179.61	1540.5	513.36	2000.5	760.28	X81198, class C, 20 probes, 20 in all_X81198 3084-3673, H.sapiens mRNA (clone p5) for archain
370.1	640.4	320.89	642.9	336.87	590.4	53.88	616.25	128.76	613.55	178.97	467.4	6.36	M65199, class A, 20 probes, 20 in M65199 735-1101, Human endothelin 2 (ET2) mRNA, complete cds
368.61	894.6	32.24	869.1	976.51	637.3	71.84	806.3	92.91	817	696.22	588.05	656.27	Guanine Nucleotide-Binding Protein Ral, Ras-Oncogene Related
449.37	461.55	332.55	768.85	537.61	542.15	83.93	1137.7	1412.66	658.75	250.25	647.45	441.16	U83303, class C, 20 probes, 20 in all_U83303 1160-2035, GCP-2 gene (granulocyte chemotactic protein-2) extracted from Human line-1 reverse transcriptase gene, partial cds, and granulocyte chemotactic protein-2 (GCP-2) gene, complete cds
114.9	397.55	335.1	286.85	179.82	326.25	15.63	271.5	23.76	1628.25	1690.48	351.55	90.58	M11591, class C, 20 probes, 18 in all_M11591 5495-6174: 2 in M11591cds 567-598, Human MHC class II HLA-SX-alpha gene
176.64	507.1	280.44	1228.25	824.7	411.4	143.26	93	15.41	1754.2	1836.36	293.75	179.82	U10550, class A, 20 probes, 20 in U10550 1591-2107, Human Gem GTPase (gem) mRNA, complete cds
256.89	499.2	351.01	900.15	716.94	393.35	315.3	835.85	739.99	213.6	118.23	582.2	355.25	U58032, class A, 20 probes, 20 in U58032 1007-1322, Human myotubularin related protein 1 (MTMR1) gene, partial cds. /gb=U58032 /ntype=DNA /annot=CDS
268.84	178.3	67.03	218.55	184.06	590.1	38.75	261.15	52.26	229.7	60.95	790.5	470.23	L40402, class A, 20 probes, 20 in L40402mRNA 593-1157, Homo sapiens (clone Zap2) mRNA fragment
245.15	225.4	301.23	215.85	287.86	157	66.89	96.9	71.28	427.3	532.88	517.5	295.85	M27093, class A, 20 probes, 20 in M27093 2049-2509, Homo sapiens nuclear-encoded mitochondrial branched chain alpha-keto acid dehydrogenase transacylase subunit (E2b) mRNA, complete cds
162.99	274.3	105.64	377.5	72.83	376.15	228.61	177.55	79.83	882.15	1105.56	206.95	71.35	X73874, class C, 20 probes, 20 in all_X73874 3675-4156, H.sapiens PHKA 1 mRNA
233.42	406.95	453.04	948.5	881.9	431.6	371.8	307.85	369.04	768.65	848.74	615.45	12.8	U21943, class A, 20 probes, 20 in U21943 2246-2678, Human organic anion transporting polypeptide (OATP) mRNA, complete cds
2796.32	3308.6	2362.02	5076.1	5390.56	14360.35	13634.93	4296.75	3578.17	7033.8	5119.03	2894.7	631.3	J04423 E coli bioD gene dethiobiotin synthetase (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)
148.78	411.15	170.48	190.35	106.56	789.1	1095.03	531.2	133.5	279.45	249.54	83	9.05	U18271, class A, 20 probes, 20 in U18271exon#3 51-512, Human thymopoietin (TMPO) gene, Human thymopoietin (TMPO) gene
357.94	3600.05	390.96	3362.8	536.13	1702.95	1347.96	3895	391.17	2543.45	1185.04	3466.8	967.04	U44755, class A, 20 probes, 20 in U44755 965-1487, Human PSE-binding factor PTF delta subunit mRNA, complete cds

1660.15	7502.9	3361.16	5372.25	1240.76	4879.4	957.56	4176.6	1752.07	3793.55	2329.85	4630.55	68.66	M77349, class A, 20 probes, 20 in M77349 2102-2642, Human transforming growth factor-beta induced gene product (BIGH3) mRNA, complete cds
1210.78	1974.45	1379.49	3803.5	3825.73	8445.25	6374.21	3312.45	2114.89	6968.5	7063.01	2629.25	341.18	J04423 E coli bioB gene biotin synthetase (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
321.1	744.95	374.84	539.15	81.39	1567.1	226.98	1256.1	1480.54	1235.4	872.15	1065.2	257.25	L35269, class A, 20 probes, 20 in L35269exon 1602-2040, Homo sapiens zinc finger protein 35 (ZNF35) gene
236.53	541.85	555.01	848.3	253.29	1232.1	1491.71	610	57.84	832.45	172.04	356.8	328.38	M20137, class A, 20 probes, 20 in M20137mRNA 400-820, Human interleukin 3 (IL-3) mRNA, complete cds, clone pcD-SR-alpha
774.71	956.05	616.24	1375.2	1289.9	1462.25	83.79	723.75	13.08	1166.65	941.94	910.1	697.77	X03663, class C, 20 probes, 20 in all_X03663 3391-3824, Human mRNA for c-fms proto-oncogene
70.43	533.35	447.67	794.4	891.66	835.9	934.23	466.05	190.71	1138.6	695.09	433.85	350.51	Z30425, class A, 20 probes, 18 in Z30425cds 649-1009: 2 in reverseSequence, 1299-1311, H.sapiens mRNA for orphan nuclear hormone receptor.
339.34	894.1	914.43	647.2	176.78	1561.95	1518.37	740.65	612.57	511.2	15.84	911.1	254.28	U32674, class A, 20 probes, 18 in U32674cds 593-1060: 2 in reverseSequence, 1181-1191, Human orphan receptor GPR9 (GPR9) gene, partial cds
757.59	7258.2	127.42	6766.35	1683.41	4066.65	3620.74	7428.1	2832.53	6270.75	3163.67	7857.6	493.56	Y00486, class A, 20 probes, 20 in Y00486mRNA 252-786, adenine phosphoribosyltransferase (aprt) gene extracted from Human APRT gene for adenine phosphoribosyltransferase
7721.39	11699.05	3958.03	15583.75	5909.93	14456.25	9751.36	13538.65	5740.36	13637.55	4468.14	15911.9	1539.37	D38047, class A, 20 probes, 20 in D38047 327-825, Human mRNA for 26S proteasome subunit p31, complete cds
98.08	141.4	84.99	101.65	60.74	417.6	83.44	450.55	188.16	144.6	104.23	278.85	243.74	U80456, class A, 20 probes, 20 in U80456 3416-3788, Human transcription factor SIM2 long form mRNA, complete cds
85.77	150.75	68.8	683.8	793.23	515.3	98.43	224.7	76.79	129.1	89.38	161.2	26.73	X65550, class A, 20 probes, 20 in X65550exon#15 2051-2549, H.sapiens mki67a mRNA (long type) for antigen of monoclonal antibody Ki-67
19.37	175.95	103.17	204.8	176.21	804.3	1004.37	94.15	62.15	122.25	89.59	176.05	53.25	D50915, class A, 20 probes, 20 in D50915 7282-7750, Human mRNA for KIAA0125 gene, complete cds
38.18	80.8	57.98	73.05	19.45	91.85	20.29	317.55	177.13	242.55	313.18	92.05	61.59	X59710, class A, 20 probes, 19 in X59710cds 75-585: 1 in reverseSequence, 714, H.sapiens mRNA for CAAT-box DNA binding protein subunit B (NF-YB)
142.84	60	25.6	228.4	236.03	287.25	159.17	320.25	104.3	385.1	472.35	96.45	30.48	U41815, class A, 20 probes, 20 in U41815 3037-3559, Human nucleoporin 98 (NUP98) mRNA, complete cds
6.15	154.6	7.07	314.8	283.97	413.75	317.84	131.3	22.91	245.5	171.97	974.65	5.59	U00946, class A, 20 probes, 20 in U00946 1382-1916, Human clone A9A2BRB5 (CAC)n/(GTG)n repeat-containing mRNA
4012.97	5908.55	1714.1	6926.85	5486.37	4314.9	2645	2549.95	524.04	3217.15	984.36	6777.15	882.26	L40397, class A, 20 probes, 20 in L40397mRNA 979-1267, Homo sapiens (clone S31i125) mRNA, 3 end of cds
474.19	608.65	468.74	1225.75	826.68	716.75	335.24	993.95	451.49	1009.7	266.44	864.7	826.89	X52896, class C, 20 probes, 20 in all_X52896 1629-2195, H.sapiens RNA for dermal fibroblast elastin
403.76	3556.95	588.67	3584.35	990.44	2983	3436.96	4487.35	69.65	3612.3	256.82	4562.25	84.78	U09196, class A, 20 probes, 20 in U09196mRNA 725-1169, Human 1.1 kb mRNA upregulated in retinoic acid treated HL-60 neutrophilic cells

15214.82	56473.05	13712.71	55929.7	21204.86	54593.8	41171.57	54067.45	25079.45	40746.1	34065.72	67766.55	18628.09	M84711, class A, 20 probes, 20 in M84711 345-831, Human v-fos transformation effector protein (Fte-1), mRNA complete cds
94.33	840.65	691.34	1224.9	731.43	803.7	502.89	575.05	518.95	820.4	933.1	547.3	318.76	X75962, class C, 20 probes, 20 in all_X75962 913-1340, H.sapiens mRNA for OX40 homologue
150.47	399.95	359.14	690.5	524.81	723.9	749.82	317.85	242.61	247.5	116.39	260.25	5.02	L36870, class A, 20 probes, 20 in L36870mRNA 3077-3533, Homo sapiens MAP kinase kinase 4 (MKK4) mRNA, complete cds
461.39	293.4	197.14	376.15	214.61	1178.7	126.15	356.4	379.29	171.85	0.78	462.95	309.36	M29550, class A, 20 probes, 20 in M29550 2005-2317, Human calcineurin A1 mRNA, complete cds
297.55	212.05	167.66	300.8	95.46	148.25	97.37	429.9	142.84	811.4	1110.44	450.75	512.16	U69140, class A, 18 probes, 18 in U69140 297-846, Human zyginII mRNA, partial cds
200.04	102.6	104.23	293.7	278.18	733.05	983.94	332.65	424.33	638.25	625.86	103.8	80.47	X84908, class C, 20 probes, 20 in all_X84908 3722-3975, H.sapiens mRNA for phosphorylase-kinase, beta subunit
1255.19	1832.1	1562.71	2211.95	344.43	2344.2	2007.62	1039.95	424.48	3449.1	538.67	2203.55	1016.18	D79999, class A, 20 probes, 20 in D79999 4526-4922, Human mRNA for KIAA0177 gene, partial cds
585.06	1336.2	582.09	1104.7	15.7	1543.3	1206.75	1550.45	463.79	1033.9	1151.59	1106.95	290.55	M21539, class A, 20 probes, 20 in M21539 521-557, Human small proline rich protein (sprII) mRNA, clone 1292
451.77	4700.55	258.73	4434.75	96.66	4533.05	478.5	4655.9	519.3	3880.75	137.39	5120.75	770.82	U62739, class A, 20 probes, 20 in U62739 986-1430, Human branched-chain amino acid aminotransferase (ECA40) mRNA, complete cds
45281.07	105984.55	27992.45	67570.55	2971.61	67665.1	22248.97	95733.75	30056.35	103874.41	63196.26	101639.2	18459.45	Homo sapiens /REF=X00351 /DEF=Human mRNA for beta-actin /LEN=1761 (_5, _M, _3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
328.59	732.65	676.06	1098.55	459.55	1966	1546.16	1094.1	39.03	468.35	456.15	939.25	4.17	L19183, class A, 20 probes, 20 in L19183mRNA 1533-1959, Human MAC30 mRNA, 3 end
2980.1	14887.8	202.37	14539.6	6535.65	5656.45	7110.6	9924.05	3137.36	10395.65	5117.54	6309.1	1111.01	S37730, class A, 20 probes, 16 in S37730cds 625-916: 4 in reverseSequence, 496-635, insulin-like growth factor binding protein-2 [human, placenta, Genomic, 4575 nt 4 segments]
304.06	401.85	335.95	194.55	88.74	333.3	218.5	279.35	306.25	344.5	10.18	775	191.06	M23197, class A, 20 probes, 20 in M23197 848-1388, Human differentiation antigen (CD33) mRNA, complete cds
5.59	66.5	40.45	75.3	23.33	145.6	133.64	179.9	164.9	120.25	17.32	75	23.48	U63312, class A, 20 probes, 17 in U63312exon#1 4-199: 3 not in GB record, Human cosmid LL12NC01-242E1, ETV6 gene, exons 1B and 3 and partial cds. /gb=U63312 /ntype=DNA /annot=exon
2889.95	5352.35	1578.9	5143.9	2814.14	4172.7	1491.85	3438.45	2256.73	3351.8	3029.53	3843.3	905.52	L10678, class A, 20 probes, 20 in L10678 1128-1650, Human profilin II mRNA, complete cds
88.25	2093.6	142.69	2560.95	405.81	2188.3	1899.29	3581.95	1097.64	3754.9	2313.79	1882.85	1647.63	M22490, class A, 20 probes, 20 in M22490 1282-1630, Human bone morphogenetic protein-2B (BMP-2B) mRNA
696.92	642.8	753.35	247.1	196.86	1061.45	1276.68	592.1	653.65	1851.7	1862.66	976.3	758.73	U21049, class B, 20 probes, 12 in U21049cds 61-319: 8 in reverseSequence, 592-760, Human DD96 mRNA, complete cds
39.32	68.6	79.48	390.35	395.63	941.45	1219.41	215.3	220.05	336.75	39.67	353.55	238.08	U83117, class A, 20 probes, 20 in U83117 1201-1477, Human sentrin mRNA, complete cds.
185.76	2180.45	244.02	1751.6	1441.93	1918.65	733.2	2089.75	294.79	1885.85	527.29	2194.15	551.19	X81003, class A, 20 probes, 20 in X81003mRNA 1032-1560, H.sapiens HCG V mRNA

42.5	238.9	165.04	148.05	14.64	1128.4	1384.23	211.85	116.74	303.35	192.26	201.7	52.61	L29339, class A, 20 probes, 20 in L29339mRNA 1862-2324, Homo sapiens Na+/glucose co-transporter (SGLT1) gene
464.29	627.5	796.63	799.45	678.47	1184.45	1575.65	1775.65	1899.64	695	361.33	649.85	449.08	M20543, class C, 20 probes, 20 in all_M20543 2890-3542, Human skeletal alpha-actin gene, complete cds
630.88	1440.15	356.31	2602.7	452.41	804.35	987.62	295	30.55	1582.6	927.87	1024.25	638.73	X72727, class C, 20 probes, 20 in all_X72727 2460-2794, H.sapiens tunp mRNA for transformation upregulated nuclear protein
1258.93	10730.55	1373.84	7616.1	1754.33	7887.4	2011.86	9744.35	1708.44	9637	1778.51	9078.65	1088.17	D28124, class A, 20 probes, 20 in D28124 1346-1886, Human mRNA for unknown product, complete cds
793.8	3136.1	895.76	3499	777.11	1380.7	1442.92	3792.45	655.84	3142.65	832.34	3324.25	1175.57	U01337, class A, 20 probes, 20 in U01337exon#16 412-553, Human Ser/Thr protein kinase (A-RAF-1) gene, complete cds
84.15	568.2	87.96	658.55	83.65	509.65	317.84	435.75	343.44	541.55	87.47	924.25	42.36	X92110, class A, 20 probes, 20 in X92110mRNA 714-1242, H.sapiens mRNA for hcgVIII protein
228.68	317.2	228.4	324.75	305.12	218.8	36.35	342.5	142.41	360.8	223.87	380.7	298.68	U45255, class A, 20 probes, 20 in U45255mRNA 1576-1952, Human paired-box protein PAX2 (PAX2) gene
1117.72	3048.95	1112.21	2630.2	243.95	732	694.8	2128.65	338.49	824	899.16	1967.5	320.89	L27624, class A, 20 probes, 20 in L27624 373-917, Homo sapiens tissue factor pathway inhibitor-2 mRNA, complete cds
121.69	266.5	48.93	219.45	190	230.05	93.97	295.05	286.59	168.95	196.65	173.6	62.37	Adrenergic, Beta, Receptor Kinase 2
105.71	140.5	119.5	902.75	1158.45	397.5	443.78	240.05	77.99	399.85	396.9	305.25	67.81	X71348, class C, 20 probes, 20 in all_X71348 2835-3436, H.sapiens vHNF1-C mRNA
1097.57	2960.75	986.2	2714.65	950.85	1488.9	733.98	1956.1	1605.13	1462.65	1366.63	3139.35	17.75	U76388, class A, 20 probes, 20 in U76388 1630-1842, Human steroidogenic factor 1 mRNA, complete cds
155.63	181.6	129.12	275.9	314.8	226	118.37	163.65	54.52	113.8	64.77	433.95	336.23	U77735, class A, 20 probes, 20 in U77735 1516-2026, Human pim-2 protooncogene homolog pim-2h mRNA, complete cds
51.97	127.15	43.63	115.85	48.01	575.1	631.45	740.9	561.3	313.3	42.43	232.9	107.2	U08015, class A, 20 probes, 20 in U08015mRNA 2388-2700, Human NF-ATc mRNA, complete cds
129.12	183.85	113.35	145.9	9.33	418.9	383.53	199.25	30.48	678.35	774.21	395.25	299.32	L35592, class A, 20 probes, 20 in L35592mRNA#1 1633-2107, Homo sapiens germline mRNA sequence
386.36	2863.8	22.63	1958.15	1651.73	1775.6	1298.53	3243.15	801.22	2256.9	294.58	2755.85	906.44	Nuclear Mitotic Apparatus Protein 1, Alt. Splice Form 2 - Also Represents: HG2238-HT2323, Z11584
147.57	667.9	597.65	543.6	490.02	546.3	243.39	325.55	5.02	324.25	138.24	333.6	131.52	U03494, class A, 20 probes, 20 in U03494 2213-2393, Human transcription factor LSF mRNA, complete cds
82.87	211.3	39.74	798.7	694.38	487.6	196.58	341.05	382.76	228.2	280.01	314.7	208.31	D90042, class A, 20 probes, 20 in D90042 675-1215, Human liver arylamine N-acetyltransferase (EC 2.3.1.5) gene
378.02	714.3	591.57	788.4	118.51	433.4	472.06	574.1	166.17	1156.5	626.92	1230.95	771.81	U19345, class A, 20 probes, 20 in U19345 2258-2756, Human AR1 protein (AR) mRNA, complete cds
695.3	1739.75	468.6	1429.4	635.97	2352.15	1762.04	2006.7	614.76	1862.75	124.24	1660.85	25.67	X55019, class B, 20 probes, 15 in X55019cds 1128-1521: 5 in reverseSequence, 1538-1701, H.sapiens mRNA for acetylcholine receptor delta subunit.
62.58	120	36.49	281.3	273.37	214.4	113.84	124.4	2.26	249	154.71	98.05	18.03	D83657, class A, 20 probes, 18 in D83657exon#1-3 13-167: 1 in reverseSequence, 2025: 1 not in GB record, Human DNA for CAAF1 (calcium-binding protein in amniotic fluid 1), complete cds
248.62	29.95	7.99	98.8	69.86	196.95	74.03	670.15	98.78	494.6	239.43	389.3	355.53	U94320, class A, 20 probes, 20 in U94320 828-1392, Human neuropeptide Y5 receptor (NPYY5) mRNA, complete cds.

55.79	88.95	56.21	179.65	183.49	227.1	160.23	124.15	95.95	169.75	198.91	162.85	19.16	X65663, class A, 20 probes, 20 in X65663cds 83-137, H.sapiens Sox-6 mRNA. /gb=X65663 /ntype=RNA
58.41	155.95	143.47	382.95	524.32	80	43.27	137.6	96.31	641.05	556.56	105.7	19.8	U43885, class A, 20 probes, 20 in U43885 1914-2442, Human Grb2-associated binder-1 mRNA, complete cds
173.95	1324.05	61.16	1889.7	832.83	808.6	227.97	1601.8	944.41	1167	338.28	629.9	32.95	L14778, class A, 19 probes, 19 in L14778 1665-2225, Human calmodulin-dependent protein phosphatase catalytic subunit (PPP3CA) mRNA, complete cds and alternative exon
758.87	969.9	1190.06	637.65	66.4	1119.35	137.67	482.05	211.92	1010.15	791.46	563.85	451.06	D79989, class A, 20 probes, 20 in D79989 3469-3919, Human mRNA for KIAA0167 gene, complete cds
33.45	110.35	49.99	314.2	343.51	155.95	141.63	194.85	7.14	83.4	0.28	85.2	11.17	X77166, class C, 20 probes, 20 in all_X77166 798-1183, H.sapiens gene for kunitz-type protease inhibitor, HKIB9
671.61	2808.7	248.34	2620.85	1986.33	1259	394.57	3724.4	116.81	2950.55	1248.68	2795.55	674.37	U52112, class A, 20 probes, 20 in U52112mRNA#1 3929-4463, Homo sapiens Xq28 genomic DNA in the region of the L1CAM locus containing the genes for neural cell adhesion molecule L1 (L1CAM), arginine-vasopressin receptor (AVPR2), C1 p115 (C1), ARD1 N-acetyltransferase related protein (TE2), renin-binding protein (RbP), host cell factor 1 (HCF1), and interleukin-1 receptor-associated kinase (IRAK) genes, complete cds, and Xq28lu2 gene., Homo sapiens Xq28 genomic DNA in the region of the L1CAM locus containing the genes for neural cell adhesion molecule L1 (L1CAM), arginine-vasopressin receptor (AVPR2), C1 p115 (C1), ARD1 N-acetyltransferase related protein (TE2), renin-binding protein (RbP), host cell factor 1 (HCF1), and interleukin-1 receptor-associated kinase (IRAK) genes, complete cds, and Xq28lu2 gene.
360.77	273.2	110.03	387.25	240.2	279.35	177.41	563.05	568.44	934.75	517.53	1048.5	342.38	U46194, class A, 20 probes, 20 in U46194 1466-1997, Human renal cell carcinoma antigen RAGE-4 mRNA, complete putative cds
54.52	126.25	86.9	459.1	592.84	440.15	20.29	48.8	11.88	192.5	198.7	52.9	1.41	Z48510, class A, 20 probes, 19 in Z48510exon#5-7 47-6: 1 in reverseSequence, 471, H.sapiens XG mRNA (clone FB1). /gb=Z48510 /ntype=RNA
119.78	123.8	120.07	107.55	117.31	1833.55	2245.42	508.45	342.17	195.05	118.44	387.3	321.59	U66497, class A, 20 probes, 20 in U66497 3549-4047, Human leptin receptor splice variant form 13.2 mRNA, complete cds.
54.16	1006.65	3.89	1260.8	289.77	479.05	394.49	580.5	294.3	502.8	414.79	615.2	7.35	M92439, class A, 20 probes, 20 in M92439 4255-4633, Human leucine-rich protein mRNA, complete cds
60.74	4408.6	68.45	4666.55	1299.31	2911.25	1496.31	3839.65	2120.68	4238.8	1224.14	4693.35	765.02	L41351, class A, 20 probes, 20 in L41351mRNA 1269-1695, Homo sapiens prostasin mRNA, complete cds
590.58	593.65	589.66	271.5	138.31	617	348.75	1330.95	586.4	1088.7	1357.08	419.05	34.29	S62028, class A, 20 probes, 20 in S62028 496-1041, recoverin [human, retina, mRNA, 1108 nt]
274.99	1428.9	102.39	979.1	491.16	961.45	589.8	1131.7	242.96	1245.85	1022.55	956.3	601.75	U41163, class B, 20 probes, 12 in U41163exon#9 36-266: 6 in reverseSequence, 2625-2862: 2 not in GB record, Human creatine transporter (SLC6A10) gene, partial cds
26.02	60.65	40.23	76.65	84.64	434.8	548.43	114.45	110.52	193.6	205.34	146.05	131.88	AF001620, class A, 20 probes, 20 in AF001620 1478-2000, Homo sapiens trabecular meshwork-induced glucocorticoid response protein (TIGR) mRNA, complete cds.

666.87	622.8	749.39	165	79.62	263.1	65.9	593.2	144.11	547.5	228.4	346.85	31.04	V00536, class A, 20 probes, 20 in V00536mRNA 811-1135, IFNG gene extracted from Human immune interferon (IFN-gamma) gene
296.35	552.5	180.17	899.5	1014.84	534.4	361.9	645.7	104.09	1018.9	1101.25	685.75	212.34	J05096, class A, 20 probes, 20 in J05096mRNA 5006-5456, Human Na,K-ATPase subunit alpha 2 (ATP1A2) gene, complete cds
224.36	251.55	307.52	129.9	0.85	729.95	266.93	216.55	206.83	599.7	463.01	398.5	371.23	D26561, class C, 20 probes, 20 in all_D26561 2433-3022, ORF for L1 protein gene extracted from Human papillomavirus 5b genome integrated into human carcinoma DNA, ORF for L1 protein gene extracted from Human papillomavirus 5b genome integrated into human carcinoma DNA, ORF for L1 protein gene extracted from Human papillomavirus 5b genome integrated into human carcinoma DNA
124.73	116.45	138.52	441.8	129.54	164.7	76.79	192.5	154.01	302.85	213.62	479.4	461.18	X81836, class C, 20 probes, 20 in all_X81836 812-1414, H.sapiens mRNA for Dents Disease candidate gene
469.87	931.3	215.81	822.9	155.7	577.35	252.08	1282.75	52.4	1062.05	307.52	613.5	365.01	L05188, class B, 20 probes, 12 in L05188cds 2-194: 8 in reverseSequence, 281-342, Homo sapiens small proline-rich protein 2 (SPRR2B) gene, complete cds
806.95	812.3	645.45	346.55	119.29	1362.9	1270.81	779.9	134.63	405.2	28.99	719.85	874.76	L13042, class A, 20 probes, 17 in L13042exon#2-3 11-216: 3 not in GB record, Homo sapiens calbindin D-9k gene, 5 end cds
235.33	1474.1	338.28	2164.9	745.15	1193.35	410.19	853.35	606.34	1509.8	390.46	1426.1	20.36	U94836, class A, 20 probes, 20 in U94836 3450-3894, Human ERPROT 213-21 mRNA, complete cds.
8.56	140.75	63.71	1205.25	1022.41	713.15	7.71	986.95	404.54	1063.95	1287.29	240.05	42.21	Y10209, class A, 20 probes, 20 in Y10209mRNA 79-331, H.sapiens mRNA for CD30L protein. /gb=Y10209 /ntype=RNA
360.48	908.5	268.56	759.15	342.45	420.6	275.63	539.65	497.87	967.2	934.09	431.55	324.35	M32053, class C, 20 probes, 20 in all_M32053 2900-3489, Human H19 RNA gene, complete cds (spliced in silico)
361.83	421.25	459.27	73.55	2.33	1767.55	1980.82	549.95	114.9	293.15	94.4	415.35	428.29	M65292, class A, 20 probes, 20 in M65292 667-1202, Human factor H homologue mRNA, complete cds
141.14	550.1	240.56	335.8	115.12	896.55	1199.18	107.15	83.65	506.4	68.31	547.6	422	- Also Represents: M58460
425.04	4845.7	757.45	4139.45	15.2	3529.3	1434.58	4349.45	953.11	5792.4	652.66	5192.05	504.52	U32986, class A, 20 probes, 20 in U32986 3609-4126, Human xeroderma pigmentosum group E UV-damaged DNA binding factor mRNA, complete cds
891.24	6602.9	263.89	5218.4	4546.56	8742.5	5693.34	7975.05	1335.8	2646.95	3662.46	8752.1	1479.69	X79200, class C, 4 probes, 4 in all_X79200 380-600, H.sapiens mRNA for SYT-SSX, synovial sarcoma translocation junction. /gb=X79200 /ntype=RNA, H.sapiens mRNA for SYT-SSX, synovial sarcoma translocation junction. /gb=X79200 /ntype=RNA
316.01	542.85	214.89	653.4	420.45	954.7	1067.87	410.9	91.92	620.6	286.52	1213.2	786.44	D38496, class A, 20 probes, 20 in D38496 3615-4170, Human mRNA for LZTR-1, complete cds
472.91	2915.05	324.07	3422.05	711.84	2169.4	804.83	2820.15	1185.18	2315	1213.4	2314.25	298.75	X12451, class A, 20 probes, 20 in X12451mRNA 974-1496, Human mrRNA for pro-cathepsin L (major excreted protein MEP)
416.98	2143.7	56.43	1737.3	989.1	1485.2	1498.93	881	732.56	1292.85	417.83	1158.3	308.58	M23294, class A, 20 probes, 20 in M23294mRNA#1 1219-1651, Human beta-hexosaminidase beta-subunit (HEXB) gene

132.44	101	114.41	385.2	536.27	1121.55	1081.38	480.35	659.8	416	558.61	225.45	257.88	D00408, class C, 9 probes, 9 in all_D00408 1838-1946, Human fetal liver cytochrome P-450 (P-450 HFLa), complete cds, Human fetal liver cytochrome P-450 (P-450 HFLa), complete cds
126.29	265.8	72.12	184.2	162.78	138.45	9.26	134.7	80.89	159.6	105.64	106.65	89.59	D88460, class A, 20 probes, 20 in D88460 1233-1695, Human mRNA for N-WASP, complete cds
260.78	325.6	379.01	223.75	4.17	79.4	0	101.15	29.06	577.05	247.56	234.5	263.47	L27071, class A, 20 probes, 20 in L27071 1963-2527, Human tyrosine kinase (TXK) mRNA, complete cds
52.82	163.25	76.16	224.25	142.62	274.2	293.45	892.45	1056.63	259	157.26	184.3	93.9	X86401, class C, 20 probes, 20 in all_X86401 1686-2217, H.sapiens mRNA for L-arginine:glycine amidinotransferase
527.57	1410.25	9.97	1040.15	54.38	1062.15	590.93	1072.35	109.11	889.55	171.19	1135.3	1044.82	Elastin, Alt. Splice 2 - Also Represents: HG2994-HT3149, HG2994-HT3148, HG2994-HT4851, U77846_rna1
929.92	808.65	1033.58	342.1	192.76	1065.9	1087.53	217	63.64	155.05	39.95	423.6	204.35	D29833, class A, 20 probes, 20 in D29833 194-680, Human mRNA for salivary proline rich peptide P-B, complete cds
288.36	288.05	286.73	93.9	6.08	108.6	96.59	94.15	21	185.05	66.11	142.15	81.25	M68516, class A, 20 probes, 20 in M68516mRNA 1662-2172, PCI gene (plasminogen activator inhibitor 3) extracted from Human protein C inhibitor gene, complete cds
169.21	151.25	69.93	227.05	154.93	269.8	134.77	306.35	18.31	325.7	106.07	713.1	126.57	D89501, class A, 20 probes, 20 in D89501exon#3 206-441, Human PBI gene, complete cds
254.56	254.95	231.72	516.5	359.78	220.85	19.45	185.45	122.12	172.15	163.27	111.65	44.76	Glucocorticoid Receptor, Beta
520.01	526.1	635.97	842.8	1087.11	1125.85	1238.78	994.45	852.42	844.25	814.37	763.05	75.45	U24153, class A, 20 probes, 20 in U24153 1796-1994, Human p21-activated protein kinase (Pak2) gene, complete cds
13.72	118.75	43.63	223.05	108.54	337.9	0.99	170.65	146.44	730	885.01	239.35	36.13	U34844, class A, 20 probes, 20 in U34844exon 40-259, Human mercurial-insensitive water-channel gene, 5 region and partial exon 1 /gb=U34844 /ntype=DNA /annot=exon
4.38	225.75	265.38	914.8	650.68	884.55	1091.84	1153.35	557.27	321.85	319.97	554.75	614.97	M82827, class A, 20 probes, 20 in M82827mRNA 2078-2228, Human fusion protein mRNA, complete cds.
106.7	282.75	47.02	284.1	60.95	374.5	3.11	357.35	295.64	337.05	108.4	364.85	241.19	U48231, class A, 20 probes, 20 in U48231exon#2 1478-2015, Human bradykinin B1 receptor (BDKRB1) gene, first
288.36	531.4	106.21	512.8	200.96	1408.85	1095.8	436.85	5.59	403	342.1	915.3	288.08	M28212, class A, 20 probes, 20 in M28212 175-691, Homo sapiens GTP-binding protein (RAB6) mRNA, complete cds
143.61	4039.4	786.87	2692.55	773.08	5835.25	38.4	1626.55	611.29	1024	209.87	3804.1	1200.1	L48546, class A, 20 probes, 16 in L48546exon#1-4 54-264: 4 not in GB record, Homo sapiens tuberin (TSC2) gene
641.2	2164	251.16	2685.45	427.02	1646.05	170.91	2030.85	594.89	1626	140.57	1396.2	4.53	D78132, class A, 20 probes, 20 in D78132 373-907, Human mRNA for Ras homologue enriched in brain (RHEB) gene, Ras-related GTP binding protein gene, complete cds
6.93	69.3	50.2	86.9	77.36	5575.75	7092.63	1278.05	1702.08	844.25	1120.13	152.25	96.66	M96132, class C, 6 probes, 6 in all_M96132 32-66, Human MHC class II HLA-DR-beta-1*09012 (HLA-DRB1*09012) gene, 3 end cds
831.91	1682.75	568.44	2373.6	880.77	1817.3	711.63	2048.1	735.67	1325	142.55	1997.85	728.39	Y00815, class C, 20 probes, 20 in all_Y00815 7107-7684, Human mRNA for LCA-homolog. LAR protein (leukocyte antigen related)

14154.86	125768.8	15325.98	104982.2	3427.21	117133.7	72611.23	133136.59	41706.43	111974.66	51893.22	138690.95	38239.41	M81757, class A, 20 probes, 20 in M81757 49-421, H.sapiens S19 ribosomal protein mRNA, complete cds
88.6	149.7	87.4	156	89.66	684	428.22	232.1	187.81	447.5	137.46	180.35	25.53	U20980, class A, 20 probes, 20 in U20980 1596-2118, Human chromatin assembly factor-I p60 subunit mRNA, complete cds
100.41	417.3	25.46	294.25	12.23	484.1	89.1	546.4	301.93	714.15	566.46	306.7	107.76	Z14093, class C, 20 probes, 20 in all_Z14093 1190-1743, H.sapiens mRNA for branched chain decarboxylase alpha subunit
25.31	40.65	32.6	244.5	320.18	674.5	891.94	28.5	14.42	620.95	859.49	525.65	671.4	X59739, class A, 20 probes, 20 in X59739mRNA 5061-5473, Human ZFX mRNA for put. transcription activator, isoform 2
164.76	320.45	10.54	198	108.61	307.5	329.09	143.65	68.09	199.4	59.4	297.3	2.55	Y10207, class A, 20 probes, 20 in Y10207mRNA 61-475, H.sapiens mRNA for CD171 protein. /gb=Y10207 /ntype=RNA
36.27	105.7	15.13	114.6	87.4	777.65	440.46	70.35	1.48	569.05	170.91	420.4	463.44	U27768, class A, 20 probes, 20 in U27768 235-709, Human RGP4 mRNA, complete cds
145.17	2055.45	14.21	2195.55	308.51	1652.6	1572.18	1926.45	975.17	2203.9	822.08	2770.1	1280	X57522, class C, 20 probes, 20 in all_X57522 2229-2788, H.sapiens RING4 cDNA
190.85	378.15	424.19	497.4	296.98	649.3	363.31	229.55	115.05	550	263.75	135.7	92.91	U58130, class A, 20 probes, 20 in U58130 2887-3301, Human bumetanide-sensitive Na-K-2Cl cotransporter (NKCC2) mRNA, complete cds
413.23	191.75	93.83	310.45	145.45	581.95	635.9	646.9	591.99	1036.7	896.05	342.8	17.11	U90918, class A, 20 probes, 20 in U90918 1794-2094, Human clone 23654 mRNA sequence
12706.14	18258.1	12717.32	74699.8	86197.73	436073.03	560306.69	117648.75	132832.48	48938.85	59894.28	31241.15	184.06	Homo sapiens /REF=M10098 /DEF=Human 18S rRNA gene, complete /LEN=1969 (_5, _M, _3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
771.67	14887.75	1146.29	11753	1259.22	17695.7	11366.6	16954.7	1540.36	19239.45	3301.13	23990.65	1602.23	U47025, class A, 20 probes, 20 in U47025 3462-3994, Human fetal brain glycogen phosphorylase B mRNA, complete cds
304.48	1120.5	813.6	1161.6	232.07	2347.85	755.4	1525.45	210.08	1657.9	118.23	1684.05	803.91	Collagen, Type Iv, Alpha 2, N-Terminus
203.93	241.55	193.25	328.15	245.44	260.7	205.34	202.05	49.14	198.15	62.72	283.7	80.04	Adp-Ribosylarginine Hydrolase - Also Represents: L13291
55.01	272.9	160.94	504.45	54.38	999.05	88.32	726.45	741.97	113.8	9.33	612.15	25.95	M15881, class A, 20 probes, 20 in M15881 1752-2310, Human uromodulin (Tamm-Horsfall glycoprotein) mRNA, complete cds
70.57	80.45	58.34	51.35	42.78	206.45	95.81	120.55	146.58	1120.45	1498.15	163.9	189.65	M14123, class A, 20 probes, 20 in M14123cds#4 3521-3935, pol from Human endogenous retrovirus HERV-K10. /gb=M14123 /ntype=DNA /annot=CDS, pol from Human endogenous retrovirus HERV-K10. /gb=M14123 /ntype=DNA /annot=CDS, pol from Human endogenous retrovirus HERV-K10. /gb=M14123 /ntype=DNA /annot=CDS, pol from Human endogenous retrovirus HERV-K10. /gb=M14123 /ntype=DNA /annot=CDS
134.35	342.95	298.05	282.45	1.63	903.6	403.05	369.2	143.26	136.5	13.72	650.75	679.03	Z78289, class A, 20 probes, 20 in Z78289 35-153, H.sapiens mRNA (clone 1D2).
865.78	1019.75	894.7	883.05	131.03	1819.6	702.72	982.3	661	570.8	459.34	561.6	268.42	U92015, class A, 20 probes, 20 in U92015 605-1031, Human clone 143789 defective mariner transposon Hsmar2 mRNA sequence

46.03	149.25	55.23	221.4	135.91	266.85	31.75	49.6	0.42	349.85	18.03	441.15	420.09	U24186, class B, 20 probes, 12 in U24186cds 465-747: 8 in reverseSequence, 1223-1391, Human replication protein A complex subunit homolog Rpa4 gene, complete cds
3202.98	9681	3565.23	10316	2995.02	9276.8	5472.3	9629.2	1813.73	11028.8	702.44	9594	4336.26	Epstein-Barr Virus Small Rna-Associated Protein
114.48	117.9	115.4	594.2	773.29	185.15	205.27	431.2	508.27	142.45	83.51	152.15	29.49	Thyroid Hormone Receptor, Beta-2
25.1	648.8	111.16	1334.4	1105.49	675.25	742.39	1117.1	589.3	74.1	18.1	389.4	304.76	M90356, class B, 20 probes, 15 in M90356cds 222-618: 5 in reverseSequence, 1194-1281, Human BTF3 protein homologue gene, complete cds
39.88	83.45	84.36	301.8	394.28	55.45	34.44	127.9	62.23	889.35	868.54	254.2	69.3	U26712, class A, 20 probes, 20 in U26712 2910-3318, Human cbl-b truncated form 2 lacking leucine zipper mRNA, complete cds
1515.97	7707.65	3023.94	6873.6	2886.55	6928.1	7275.28	8742.75	191.56	7636.3	2683.75	7646.85	2989.86	D83735, class A, 20 probes, 20 in D83735 1551-2085, Human adult heart mRNA for neutral calponin, complete cds
558.19	617.45	613.42	172.2	4.95	201.1	11.88	274.5	94.47	601	558.33	274.8	13.15	U00928, class A, 20 probes, 20 in U00928 611-1151, Human clone CE29 4.1 (CAC)n/(GTG)n repeat-containing mRNA
69.72	122.95	121.41	138.95	145.31	245.25	72.2	157.15	54.66	172.15	170.2	339.2	225.71	M28585, class A, 20 probes, 20 in M28585 329-839, Human leukocyte interferon-alpha mRNA, complete cds, clone pIFN105
88.46	928.75	107.55	1788.85	1737.29	674.05	396.33	646	524.25	382.75	52.11	1039.8	446.89	U41518, class A, 20 probes, 20 in U41518 1761-2253, Human channel-like integral membrane protein (AQP-1) mRNA, clone AQP-1-2344, partial cds
471.07	2325.55	171.61	2715.6	1118.78	4000.2	410.83	2965.3	366.99	4708.05	1835.44	4071.05	1594.6	X69910, class C, 20 probes, 20 in all_X69910 2339-2892, H.sapiens p63 mRNA for transmembrane protein
625.79	1551.05	381.91	4407.1	1958.26	969.55	251.8	4029.9	363.03	2750.55	2027.77	1225.9	350.44	Sry-Related Hmg-Box 12 Protein (Gb:X73039)
205.41	99.8	88.67	623.25	321.52	201.75	5.3	315.7	350.3	202.6	163.2	286.15	318.27	U33202, class A, 19 probes, 19 in U33202 104-354, Human mdm2-D (mdm2) mRNA, complete cds. /gb=U33202 /ntype=RNA
557.98	2787.45	467.89	2156.25	508.48	2399.75	566.46	2424.95	298.19	2922.75	1081.38	2523.45	169.92	U59831, class A, 20 probes, 20 in U59831mRNA 1876-2385, Human transcription factor, forkhead related activator 4 (FREAC-4) gene, complete cds
149.84	139	141.28	47.2	44.83	143.75	145.59	38.05	24.54	674.4	525.8	88.55	19.87	U59914, class A, 20 probes, 20 in U59914 802-1240, Human chromosome 15 Mad homolog Smad6 mRNA, complete cds
6.51	58.2	31.54	143.15	57.35	215.7	157.97	338.9	291.19	316.8	65.2	235.95	122.26	AF012270, class A, 20 probes, 20 in AF012270 815-1325, Homo sapiens peropsin (Rrh) mRNA, complete cds. /gb=AF012270 /ntype=RNA
1270.95	1701.55	1200.6	1127.2	971.71	2067.45	1580.17	1240.35	961.45	1363.95	305.68	375.85	31.61	D79992, class A, 20 probes, 20 in D79992 6597-6897, Human mRNA for KIAA0170 gene, complete cds
165.46	68.85	63.29	116.7	3.96	290	152.31	155.75	91.85	493.65	632.51	260.6	277.04	M31166, class A, 20 probes, 20 in M31166mRNA 1286-1784, Human tumor necrosis factor-inducible (TSG-14) mRNA, complete cds
923.55	4239.6	998.72	3911.1	1323	4185.65	2995.66	3704.55	501.83	3423.25	2243.58	5570.8	1215.38	D26308, class A, 20 probes, 20 in D26308 241-691, Human mRNA for NADPH-flavin reductase, complete cds
347.76	646.3	421.44	542.75	500.14	1181.85	527.29	1867.4	1016.82	3211.4	2535.26	1378.4	12.02	D13897, class C, 20 probes, 20 in all_D13897 1402-1545, peptide YY precursor gene extracted from Human DNA for peptide YY, complete cds
222.24	26.8	0.85	162.35	93.27	597.5	675.15	322.2	86.13	756.15	918.04	549.25	582.3	M14113, class A, 20 probes, 20 in M14113mRNA 8440-8986, Human coagulation factor VIII:C mRNA, complete cds

1075.93	3534.05	1014.34	3823.9	2246.76	957.35	616.24	7274.35	1910.11	5005	4826.57	3780.55	799.53	X78992, class A, 20 probes, 20 in X78992cds 1349-1373, H.sapiens ERF-2 mRNA
390.82	595.55	445.12	231.85	82.94	622.65	525.45	172.8	22.91	256.15	45.75	346.35	73.33	J04102, class A, 20 probes, 20 in J04102 1830-2184, Human erythroblastosis virus oncogene homolog 2 (ets-2) mRNA, complete cds
90.3	269.75	27.37	288.5	121.2	1925.95	2433.51	249.5	26.3	319.25	179.96	535.4	599.34	Z83805, class A, 20 probes, 20 in Z83805 199-463, H.sapiens mRNA for axonemal dynein heavy chain (partial, ID hdhc8).
49.14	313.4	209.3	247.75	28.07	377.6	151.75	164.65	13.93	1531.7	1279.01	587.9	328.52	Phosphoglucomutase 1, Alt. Splice
74.39	341.1	34.51	318.65	325.48	756.75	592.34	384.3	135.06	340.4	180.31	263.8	24.61	M95549, class A, 20 probes, 20 in M95549 1774-2194, Homo sapiens sodium/glucose cotransporter-like protein mRNA, complete cds
239.64	258.7	346.06	212.35	122.68	1330.3	1451.27	481.5	366.14	228.05	132.02	348.7	450.85	U72935, class A, 20 probes, 20 in U72935mRNA#1 7752-7898, ATRX gene (putative DNA dependent ATPase and helicase) extracted from Human putative DNA dependent ATPase and helicase (ATRX) gene
54.31	116.65	32.6	37.4	13.44	86.45	43.49	359.45	432.68	278.05	56.5	133.7	47.23	M29335, class B, 15 probes, 15 in M29335 2-180, Human MHC class II DO-alpha mRNA, partial cds, Human MHC class II DO-alpha mRNA, partial cds
46.17	116.45	44.9	120.6	47.23	156.4	82.17	99.5	18.38	152.6	75.52	93.45	33.16	U28251, class A, 20 probes, 20 in U28251exon 1192-1642, Human Krueppel-type zinc finger protein (ZNF169) gene, partial cds
1077.77	962.25	895.13	185.65	173.17	163.85	225.5	296.2	356.52	197.5	132.94	385.05	225.35	AC000066, class B, 20 probes, 11 in AC000066cds 2985-3237: 9 in reverseSequence, 48519-48663, Human BAC clone RG293F11 from 7q21-7q22
337.57	532.45	246.14	254.65	130.04	443.4	182.86	616.7	461.46	263.85	69.08	935.55	517.25	M13928, class A, 20 probes, 20 in M13928mRNA 542-1020, Human delta-aminolevulinate dehydratase mRNA, complete cds.
36.84	59.65	51.12	124.9	14.28	80.8	12.59	58.55	3.46	486.05	308.93	274.5	242.4	L31860, class C, 20 probes, 20 in all_L31860 2084-2589, Human glycophorin A, MN-types (GYPA) mRNA, complete cds
219.91	1100.35	56.21	507.9	121.34	966.65	377.24	828.85	258.31	1340.9	22.06	1160.7	440.67	Serine Protease Met1
159.88	2849.3	185.54	2695.55	65.12	3535.4	1830.84	3187.85	203.86	5324.1	2083.56	3925.95	1012.65	D86965, class A, 20 probes, 20 in D86965 6166-6490, Human mRNA for KIAA0210 gene, complete cds
16.55	100.15	71.63	464.55	551.33	424.8	107.48	310.55	26.09	194.05	155.92	151.2	43.84	M60450, class A, 20 probes, 20 in M60450 1932-2404, Human voltage-gated potassium channel (HK1) mRNA, complete cds
101.82	152.95	93.83	178.1	88.53	184	123.89	177.25	27.51	172.15	106.99	145.6	44.12	X52056, class C, 20 probes, 20 in all_X52056 778-1304, Human mRNA for spi-1 proto-oncogene
150.83	1725.9	211.14	1599.85	583.15	1018.45	164.54	2159.45	376.25	1215.2	735.39	1495.85	786.8	L02867, class A, 20 probes, 20 in L02867 2179-2689, Homo sapiens 62 kDa paraneoplastic antigen mRNA, 3 end
68.09	167.45	13.22	200.55	6.58	331.5	296.14	124.75	120	507	278.03	945.6	657.04	M63623, class A, 20 probes, 20 in M63623 2390-2900, Human oligodendrocyte-myelin glycoprotein (OMGP) mRNA, complete cds
311.2	1798.5	224.44	1697.15	527.43	490.7	220.33	671.3	580.68	996.45	518.24	886.2	192.76	S74728, class A, 20 probes, 20 in S74728 1245-1773, antiquitin=26g turgor protein homolog [human, kidney, mRNA, 1809 nt]
8.84	301.15	141.49	313.95	168.79	180.35	64.56	256.3	195.02	270.45	45.33	183.1	56.14	X86428, class B, 20 probes, 14 in X86428cds 626-920: 6 not in GB record, H.sapiens gene for phosphotyrosyl phosphatase activator (exon 1)

105.36	699.95	113.21	451.7	342.81	1298.8	856.87	702.35	115.33	1173.6	16.4	967.45	43.49	U85992, class A, 20 probes, 20 in U85992 1236-1596, Human clone IMAGE:35527 unknown protein mRNA, partial cds
391.6	784.75	779.16	666.5	156.84	789.15	890.88	851.7	180.74	1341.6	877.8	1234.35	307.38	U03486, class A, 20 probes, 16 in U03486cds 625-1045: 4 in reverseSequence, 1121-1211, Human connexin40 gene, complete cds
322.65	365.45	312.33	1141.2	743.31	343.3	63.64	358.9	125.44	921.4	353.69	312.9	60.25	M22995, class A, 20 probes, 20 in M22995 1008-1542, Human ras-related protein (Krev-1) mRNA, complete cds
76.58	245.5	42	299	139.44	300.6	158.67	222.95	17.89	230.7	50.91	322.6	22.49	U52111, class A, 20 probes, 20 in U52111mRNA#4 1182-1671, Homo sapiens Xq28 genomic DNA in the region of the ALD locus containing the genes for creatine transporter (SLC6A8), CDM, adrenoleukodystrophy (ALD), Na+-isocitrate dehydrogenase gamma subunit (IDH), and translocon-associated protein delta (TRAP) genes, complete cds, plexin related protein (PLEXR) and serine kinase (SK) genes, partial cds, Xq28lu1 gene and cytochrome C (CCp) pseudogene., Homo sapiens Xq28 genomic DNA in the region of the ALD locus containing the genes for creatine transporter (SLC6A8), CDM, adrenoleukodystrophy (ALD), Na+-isocitrate dehydrogenase gamma subunit (IDH), and translocon-associated protein delta (TRAP) genes, complete cds, plexin related protein (PLEXR) and serine kinase (SK) genes, partial cds, Xq28lu1 gene and cytochrome C (CCp) pseudogene.
550.41	805.75	961.03	814.45	761.91	2124	1621.68	1998.6	1213.11	1536.65	51.55	1758.95	323.93	X59373, class A, 20 probes, 20 in X59373mRNA 531-1071, Human HOX4D mRNA for a homeobox protein
248.76	1887.55	290.83	1927.45	798.82	1664.05	826.96	1934.5	2.55	2128.75	508.2	2730.45	319.26	AC002115, class C, 20 probes, 14 in AC002115 66940-67151: 6 in AC002115cds#2 675-1000, COX6B gene (COXG) extracted from Human DNA from overlapping chromosome 19 cosmids R31396, F25451, and R31076 containing COX6B and UPKA, genomic sequence, COX6B gene (COXG) extracted from Human DNA from overlapping chromosome 19 cosmids R31396, F25451, and R31076 containing COX6B and UPKA, genomic sequence, COX6B gene (COXG) extracted from Human DNA from overlapping chromosome 19 cosmids R31396, F25451, and R31076 containing COX6B and UPKA, genomic sequence, COX6B gene (COXG) extracted from Human DNA from overlapping chromosome 19 cosmids R31396, F25451, and R31076 containing COX6B and UPKA, genomic sequence
354.33	403.4	368.83	219.15	140.64	257.65	35.57	143.85	0.64	146.75	81.39	144.2	19.09	AC000115, class B, 20 probes, 11 in AC000115cds#1 546-684: 9 in reverseSequence, 35138-35366, WUGSC:H_GS188P18.1a gene extracted from Human BAC clone GS188P18
252.58	308.55	27.79	850.9	27.29	2110.75	1412.3	1778.25	608.89	629.9	420.87	715.4	254.13	X02875, class C, 20 probes, 20 in all_X02875 158-628, Human mRNA (3 -fragment) for (2 -5) oligo A synthetase E (1,8 kb RNA)

866.28	1737.6	755.33	2607.05	443	4015.6	2554.64	3220.95	634.49	1113.35	1416.41	2745.9	18.95	Z86000, class C, 20 probes, 20 in reverse_Z86000 20444-20634, Human DNA sequence from PAC 151B14 on chromosome 22q12-qter contains somatostatin receptor subtype 3 (SSTR3), tRNA, ESTs, CpG island and STS.
812.18	2287.6	836.93	1836.05	809.57	2456.4	512.94	2360.85	551.05	1253.8	1208.16	1248.25	166.81	X15393, class A, 20 probes, 20 in X15393mRNA 19-541, H.sapiens motilin gene exon 2 (and joined CDS)
27.58	37.5	12.73	105.75	78.56	219.25	236.1	79.55	82.38	151.7	127.56	62.25	11.81	X04145, class B, 20 probes, 13 in X04145cds 286-454: 7 in reverseSequence, 593-689, Human mRNA for T-cell receptor T3 gamma polypeptide
4424.72	11538.25	3296.6	14392.1	5188.33	9013.8	8690.34	10806.65	3176.25	6965.85	4546.91	12166.3	2949.34	M30448, class A, 20 probes, 20 in M30448mRNA 1936-2447, Human casein kinase II beta subunit mRNA, complete cds.
695.51	2237.85	915.07	1434.25	1062.71	2404.5	1000.7	2457.5	914.29	1154.8	1120.76	636.5	162.63	X60655, class A, 20 probes, 20 in X60655mRNA 990-1474, H.sapiens EVX1 mRNA.
278.03	1068.1	850.08	509.55	64.98	874.75	325.48	1267.1	1065.33	469.05	84.5	1175.35	363.1	M76732, class C, 20 probes, 20 in all_M76732 736-1273, Human HOX7 gene
69.86	113.5	87.82	152.4	101.68	253.25	159.45	181.5	155.7	247.15	142.91	152.1	39.6	X81851, class A, 19 probes, 19 in X81851cds 26-326, H. sapiens IL-4 gene splice variant. /gb=X81851 /ntype=RNA
177.06	236.95	148.28	252	138.17	1151.4	1439.53	880.1	724.78	490.05	261.28	852.9	638.8	U32499, class A, 20 probes, 20 in U32499 848-929, Human d3 dopamine receptor mRNA, complete cds.
1056.21	1657.95	936.7	1243.45	408.78	810.45	628.69	2445.35	804.05	2217.2	1554.36	1260.7	867.9	U77604, class A, 20 probes, 20 in U77604 13-493, Human microsomal glutathione S-transferase (GST-II) mRNA, complete cds
2036.82	3102.25	1114.33	4671.95	1329.29	1640.2	1642.47	3303.65	857.65	2358.1	96.59	2383.8	319.47	D21235, class A, 20 probes, 20 in D21235 1418-1673, Human mRNA for HHR23A protein, complete cds
181.51	94.1	12.02	961.8	800.73	143.25	85.77	354.7	72.27	309.45	166.1	701.6	605.42	X69398, class A, 20 probes, 20 in X69398cds 898-927, H.sapiens mRNA for OA3 antigenic surface determinant
312.33	704.85	369.75	796.55	154.22	1916.7	1896.6	982.85	355.89	1616.85	242.33	1618.8	188.51	U89335, class A, 20 probes, 20 in U89335exon#30 375-909, NOTCH4 gene (notch4) extracted from Human HLA class III region containing notch4 (NOTCH4) gene, complete cds, complete sequence.
39.32	1762.3	933.66	988.5	623.53	892.3	784.32	2197.25	810.27	2552	389.76	1324	610.09	Mucin 4, Tracheobronchial
219.77	44595.5	1294.71	48830.05	7083.3	35447.05	16830.77	33300.6	4811.86	29201.15	7651.95	37339.7	719.13	M14328, class A, 20 probes, 20 in M14328mRNA 1144-1704, Human alpha enolase mRNA, complete cds
98.08	519.1	113.7	916.55	390.39	382.05	2.9	403.35	221.68	891.45	377.38	677.5	882.89	X69978, class A, 20 probes, 18 in X69978cds 3218-3494: 2 in reverseSequence, 3769-3781, H.sapiens mRNA for XP-G factor
90.58	95.8	96.45	96.1	97.44	1279.45	1687.51	142.95	93.69	211.15	253.64	238.85	71.06	L22569, class A, 20 probes, 20 in L22569 1685-2243, Homo sapiens cathepsin B mRNA, 3 UTR with a stem-loop structure providing mRNA stability
1141.91	1385.8	1156.69	940.65	418.68	352.1	164.33	829.2	718.28	285.2	143.4	1044.7	1112.14	X84746, class A, 20 probes, 20 in X84746cds 544-1012, H.sapiens Histo-blood group AB0 gene, exon 1
114.34	199.75	30.76	284.1	307.59	330.05	381.77	881.4	1162.77	618.45	663.05	326.6	51.34	AJ001047, class A, 20 probes, 17 in AJ001047cds 485-791: 3 in reverseSequence, 959-1013, Homo sapiens mRNA for matrilin-3. /gb=AJ001047 /ntype=RNA
1407.07	3551.05	1054.93	3556.8	901.14	3278.3	1126.56	3373.35	51.27	3331.45	443.99	3541.05	1201.45	Collage, Type Vii, Alpha 1 - Also Represents: L02870
79.2	558.45	135.27	338.1	180.31	911.05	298.33	1095.95	1133.85	206.6	3.11	510.8	343.23	L40933, class B, 20 probes, 14 in L40933cds 1218-1470: 6 in reverseSequence, 1639-1819, Homo sapiens phosphoglucomutase-related protein (PGMRP) gene, complete cds

281.71	777.75	252.65	614.5	186.25	960.85	493.07	1127.9	738.08	838.8	227.55	753.4	568.94	U47334, class A, 20 probes, 20 in U47334 7-307, Human gamma aminobutyric acid receptor beta4 subunit-like mRNA, partial cds. /gb=U47334 /ntype=RNA
1155.62	1415.6	982.31	1333.05	396.33	2015.15	1917.74	2137.6	1240.55	1382.8	845.28	421.9	187.81	U34587, class A, 20 probes, 20 in U34587 1545-2061, Human corticotropin-releasing factor receptor 2 mRNA, complete cds
389.19	456.05	596.59	801.75	727.54	770.1	1000.13	982.9	1266	445.3	138.45	758.45	9.26	Y00083, class B, 20 probes, 14 in Y00083cds 902-1201: 6 in reverseSequence, 1416-1641, Human mRNA for glioblastoma-derived T-cell suppressor factor G-TsF (transforming growth factor-beta2, TGF-beta2)
171.97	148	179.89	452.15	573.53	744.85	999.07	86.15	11.67	1442.3	249.89	291.25	319.12	S42457, class A, 20 probes, 20 in S42457 2418-2814, CNCG=rod photoreceptor cGMP-gated channel [human, retina, mRNA, 2857 nt]
194.1	349.05	328.88	402.85	169.92	684.1	637.81	724.8	173.95	758.3	766.65	187.25	169.63	M86699, class A, 20 probes, 20 in M86699 3355-3787, Human kinase (TTK) mRNA, complete cds
464.29	67.8	38.18	794.55	249.11	882.1	201.81	1060.85	756.53	479	151.18	667.75	799.67	X95632, class C, 20 probes, 20 in all_X95632 1680-1784, H.sapiens mRNA for Arg protein tyrosine kinase-binding protein
1335.23	3725.3	670.62	4722.3	510.81	3160.55	1144.88	3298.65	323.93	2716.65	26.52	4634.5	161.08	X99728, class A, 20 probes, 20 in X99728exon 13-247, H.sapiens NDUFV3 gene, exon 3 /gb=X99728 /ntype=DNA /annot=exon
109.88	535.25	390.11	368.85	199.33	256.75	52.26	866.8	600.48	305.75	41.65	461.35	83.65	U77970, class A, 20 probes, 20 in U77970 2339-2855, Human neuronal PAS2 (NPAS2) mRNA, complete cds.
577.92	463.65	614.12	67.1	53.03	300.4	82.87	64.25	59.75	133.05	59.89	300.7	156.69	S77393, class A, 20 probes, 20 in S77393 43-205, transcript ch138 [human, RF1,RF48 stomach cancer cell lines, mRNA, 235 nt]. /gb=S77393 /ntype=RNA
301.51	2776.35	549.21	2154.1	2382.53	2039.1	2592.11	2056.2	1881.47	459.7	203.51	4173.6	132.09	AF009368, class A, 20 probes, 20 in AF009368 901-1345, Homo sapiens Luman mRNA, complete cds. /gb=AF009368 /ntype=RNA
94.33	156	122.33	364.4	136.9	190.6	148.07	189.2	46.39	175.65	62.3	140.15	29.91	M71243, class B, 16 probes, 14 in M71243mRNA 25-38: 2 not in GB record, Human glycophorin Sta (type A) exons 3 and 4, partial. /gb=M71243 /ntype=DNA /annot=exon
38.82	1092.6	90.09	2202.65	1401.98	1016.05	352.21	1226.05	698.55	1633.8	511.66	1363.85	11.1	L34075, class A, 20 probes, 20 in L34075 7342-7912, Human FKBP-rapamycin associated protein (FRAP) mRNA, complete cds
17.11	201.3	37.05	215.15	55.79	359.5	65.05	337	205.49	371.4	337.29	273.8	35.92	D50931, class A, 20 probes, 20 in D50931 2527-2971, Human mRNA for KIAA0141 gene, complete cds
324.49	780.25	459.97	771.3	379.43	681.05	349.95	464.4	414.93	2249.1	1634.83	706.9	604.01	D86970, class A, 20 probes, 20 in D86970 5998-6400, Human mRNA for KIAA0216 gene, complete cds
433.46	515.4	618.29	707.4	615.18	369.1	34.22	314.3	41.72	893.45	868.68	753.95	44.48	U00930, class A, 20 probes, 20 in U00930 2705-3191, Human clone C4E 1.63 (CAC)n/(GTG)n repeat-containing mRNA
56.36	143.35	31.75	192.95	112.08	267.5	224.44	331.2	238.29	243	140.43	313.35	199.47	U40317, class A, 20 probes, 20 in U40317 5400-5965, Human protein tyrosine phosphatase PTPsigma (PTPsigma) mRNA, complete cds
1349.09	6734.3	2200.23	5549.7	1689.14	9268.1	3134.32	4845.2	848.95	7251.95	25.1	6697.95	1952.96	D16562, class A, 20 probes, 20 in D16562 514-1042, Human mRNA for ATP synthase gamma-subunit (L-type), complete cds
961.24	6307.25	2745.77	5250.05	1034.71	9331.9	8150.54	8738.45	3506.75	7336.25	4314.84	8889.95	549.35	M19888, class A, 20 probes, 20 in M19888 58-580, Human small proline rich protein (sprl) mRNA, clone 128

90.51	1438.15	152.24	1939.7	939.6	1110.15	188.16	1831.15	220.26	581.1	133.36	982.4	55.44	U72066, class A, 20 probes, 20 in U72066 2672-3212, Human CtBP interacting protein CtIP (CtIP) mRNA, complete cds
222.95	3754.35	731.08	3547.85	221.25	2003.6	857.86	2901	206.33	3105.05	783.83	3061.15	585.84	Z22548, class B, 20 probes, 13 in Z22548cds 310-547: 7 in reverseSequence, 684-894, H.sapiens thiol-specific antioxidant protein mRNA
1322.64	1757.25	1274.84	892.8	683.07	382.6	8.63	877.1	406.16	1809.3	1046.66	1573.8	478	U50330, class A, 20 probes, 20 in U50330 3071-3515, Human procollagen C-proteinase (pCP-2) mRNA, complete cds
22.13	1404.25	72.62	752.45	548.36	1831.7	2154.27	1030	805.96	1051.6	402.49	509	26.45	M19722, class A, 20 probes, 20 in M19722 1843-2245, Human fgr proto-oncogene encoded p55-c-fgr protein, complete cds
38.75	159.1	105.5	462	542.63	1242.45	766.15	83.35	16.9	927.65	1099.2	250.6	167.87	M21642, class C, 8 probes, 8 in all_M21642 180-301, Human (dysfunctional) antithrombin III (ATIII) Utah gene, Human (dysfunctional) antithrombin III (ATIII) Utah gene
18.81	1419.95	299.18	1781.25	439.47	577.65	450.21	265	157.4	1065.4	165.18	1348.1	436.71	U41060, class A, 20 probes, 20 in U41060 2936-3416, Human breast cancer, estrogen regulated LIV-1 protein (LIV-1) mRNA, partial cds
789.84	673.5	654.22	643.8	766.22	750.75	410.62	507.6	543.91	842.85	412.6	740.1	49.21	D67029, class A, 20 probes, 20 in D67029 4839-5355, Human SEC14L mRNA, complete cds
270.47	1108.7	399.66	1654.05	748.05	1984	1907.35	1792.2	332.2	760.75	867.97	1916.85	444.84	M16938, class A, 20 probes, 20 in M16938 1154-1659, Human homeo box c8 protein, mRNA, complete cds
217.29	308.75	13.36	216.25	110.8	293.4	98.85	318.45	33.45	205.3	103.52	183.2	50.06	D16532, class A, 20 probes, 20 in D16532exon 123-561, Human gene for very low density lipoprotein receptor, 5 flanking and
496.81	556.7	631.59	974.05	676.63	171	105.64	1262	851.64	749.45	183.49	981.95	485.15	L14812, class A, 20 probes, 20 in L14812 3349-3936, Human retinoblastoma related protein (p107) mRNA, complete cds
351.79	424.25	6.86	991.1	321.59	1060.7	466.97	461.9	74.53	592.65	142.62	543.55	185.9	Mg81
26.16	64.5	6.65	61.7	34.51	88.7	29.84	67.8	6.22	244.45	66.26	185.15	82.52	Gal Beta 1,3(4)Glcnac Alpha2,3-Sialyltransferase
23.12	2016.25	888.76	2824.75	1048.57	1270.55	478.5	2590.4	675	1898.3	896.33	979.4	428.08	U50136, class A, 20 probes, 20 in U50136mRNA 106-640, Human leukotriene C4 synthase (LTC4S) gene, complete cds
207.11	1140.65	215.46	864.85	5.16	1761.65	489.25	807	2.83	916.15	140.78	785.65	547.51	U26727, class A, 20 probes, 20 in U26727 512-938, Human p16INK4/MTS1 mRNA, complete cds
410.26	857.75	993.84	642.3	266.72	769.95	310.07	777.2	760.71	1265.85	642.27	872.3	373.35	L37199, class A, 20 probes, 20 in L37199 977-1313, Homo sapiens (clone cD24-1) Huntington disease candidate region mRNA fragment
1948.64	7382.65	749.32	7181.35	2178.67	4494.7	1401.91	7509.75	3280.62	7100.05	1330.28	5997.45	987.05	U53347, class A, 20 probes, 20 in U53347 2278-2812, Human neutral amino acid transporter B mRNA, complete cds
821.73	1966.15	287.44	2503.9	1003.67	2274.75	1328.3	2446.8	913.72	2797.75	491.65	1879.65	916.91	D38305, class A, 20 probes, 20 in D38305 701-1181, Human mRNA for Tob, complete cds
9543.25	62447.45	7129.54	51618.6	24341.87	43566.95	36216.38	53156.5	33546	37722.55	36333.05	74783.75	9923.6	X62691, class A, 20 probes, 20 in X62691cds 13-343, H.sapiens mRNA for ribosomal protein (homologuous to yeast S24)
1.27	179.15	71.35	355.65	432.82	172.6	7.64	91.5	58.97	304.4	200.96	386.75	519.65	J03810, class A, 20 probes, 20 in J03810 2627-3113, Human liver glucose transporter-like protein (GLUT2), complete cds
561.87	5927.95	413.87	5213.55	2330.69	4086.7	1347.18	5469.55	1457.13	4428.95	2552.02	6063.75	361.12	X51521, class C, 20 probes, 20 in all_X51521 2653-3026, Human mRNA for ezrin

4.1	225.4	96.87	731	454.95	153.45	58.9	574.7	350.72	423.15	108.26	334.65	239.92	X79204, class C, 20 probes, 20 in all_X79204 10002-10585, H.sapiens SCA1 mRNA for ataxin
171.26	2316.85	927.94	1366	1407.14	743.55	627.42	2377.45	385.59	1744.25	1059.32	445.75	44.62	U31501, class A, 20 probes, 20 in U31501 2359-2773, Human fragile X mental retardation syndrome related protein (FXR2) mRNA, complete cds
54.8	133.2	60.25	1097.25	1307.37	529.15	280.65	556.5	679.67	2465.8	3358.33	371.5	207.18	U27333, class C, 9 probes, 9 in all_U27333 2701-2753, Human alpha (1,3) fucosyltransferase (FUT6) mRNA, major transcript I, complete cds, Human alpha (1,3) fucosyltransferase (FUT6) mRNA, major transcript I, complete cds
644.25	11032.6	1338.27	13247.25	1405.09	9333.85	3460.93	13907.5	1207.6	10875.7	4639.19	13316	1498.36	X05130, class C, 19 probes, 19 in all_X05130 1362-1876, Human mRNA for prolyl 4-hydoxylase beta subunit (EC 1.14.11.2) (procollagen-L-proline, 2-oxoglutarate:oxygen oxidoreductase, 4-hydroxylating)
92.91	118.65	73.61	273.95	283.34	497.1	560.74	450.05	380.07	407.4	105.78	182.7	101.68	U39231, class A, 20 probes, 20 in U39231 1739-1979, Human GIP receptor (GIPR) mRNA, complete cds
1295.99	1279.85	1188.86	2095.9	709.09	1447.15	1159.3	2189.75	201.6	2802.25	842.38	2193.85	289.98	U46461, class A, 20 probes, 20 in U46461 1591-2152, Human dishevelled homolog (DVL) mRNA, complete cds
213.76	1494	98.29	1943.4	362.18	2018.85	710.57	1439.35	727.83	1271.05	849.31	2476.25	934.87	D38555, class A, 20 probes, 20 in D38555 3911-4421, Human mRNA for KIAA0079 gene, complete cds
155.7	227.6	102.95	604.3	681.23	1027.6	745.71	212.5	46.53	964.25	715.52	323.8	346.48	D16350, class A, 20 probes, 20 in D16350 1280-1826, Human SA mRNA for SA gene product, complete cds
254.91	754.3	519.02	766.75	265.09	1192.7	7.35	2296.95	2352.76	1493.35	213.05	1188.85	707.32	X07496, class C, 20 probes, 20 in all_X07496 2066-2268, Human Tangier apoA-I gene
30.76	89.5	11.03	303.45	49.85	188.1	110.45	286.8	248.62	120.25	61.02	104.7	19.66	L19063, class A, 20 probes, 20 in L19063exon 79-451, Human glial-derived neurotrophic factor gene, complete cds. /gb=L19063 /ntype=DNA /annot=exon
118.3	287.6	97.86	428	196.58	350.85	246.57	164	17.11	230.3	76.79	824.45	621.76	U76764, class A, 20 probes, 20 in U76764 2544-3054, Human CD97 mRNA, complete cds
595.24	1618.05	231.86	1198	14.57	994.35	887.91	626.4	172.11	1715.85	1682.56	759.25	259.86	X63380, class A, 20 probes, 20 in X63380 1061-1613, H.sapiens mRNA for RSRFR2
679.46	1719.95	408.64	1061.95	1000.06	914.15	972.77	1756.6	352	1064.65	1343.86	1207.4	136.75	U61374, class A, 20 probes, 20 in U61374 1265-1715, Human novel protein with short consensus repeats of six cysteines mRNA, complete cds
57.77	412.95	163.84	450.55	138.95	573.05	144.46	407	110.31	359.4	129.26	1132.05	865.29	D38128, class A, 20 probes, 20 in D38128exon 604-1126, Human IP gene for prostacyclin receptor
154.5	3997.25	928.22	2851.8	692.54	4400.7	719.98	2949.3	502.75	3258.15	336.23	2891.65	621.48	X76180, class C, 20 probes, 20 in all_X76180 2760-3115, H.sapiens mRNA for lung amiloride sensitive Na+ channel protein
11.31	191.7	105.78	678.8	790.26	262.4	4.53	217.6	16.97	241	140.86	223.85	14.78	X97674, class B, 20 probes, 12 in X97674cds 4092-4326: 8 in reverseSequence, 4536-4758, H.sapiens mRNA for transcriptional intermediary factor 2
53.39	212.5	169.56	708.95	830.21	2373.5	2236.3	754.25	674.37	1332.3	263.04	663.75	345.14	D12485, class A, 20 probes, 20 in D12485 2904-3444, Human mRNA for nucleotide pyrophosphatase, complete cds
618.86	1668.5	726.06	1488.55	1176.27	1788.95	684.69	745.9	75.94	2484.55	1893	2630.7	123.32	U79261, class A, 20 probes, 20 in U79261 883-1422, Human clone 23959 mRNA, partial cds.
10.32	64.15	18.31	370.65	416.13	264.65	234.12	157.85	78.56	209.85	179.53	120.85	41.51	D87457, class A, 20 probes, 20 in D87457 1568-2060, Human mRNA for KIAA0281 gene, complete cds
380.14	3775.25	442.15	3357.6	638.38	3358.65	371.02	2957.55	154.22	3287.2	1487.05	4461.3	595.1	S78771, class A, 20 probes, 20 in S78771 1149-1661, NAT=CpG island-associated gene [human, mRNA, 1741 nt]

62.37	617.9	178.62	460.9	263.89	704.8	576.01	1044.7	327.39	812.05	480.34	401.25	100.06	U89717, class A, 20 probes, 20 in U89717 635-1151, Human 9-cis-retinol specific dehydrogenase mRNA, complete cds
821.02	1053.7	831.84	954.85	582.3	1464.5	1008.05	381.15	12.94	757.25	690.49	1006.55	1144.45	Fmlp-Related Receptor I
232.57	189.35	216.59	136.65	53.81	212.95	205.7	73.3	32.67	204.15	216.45	108.2	56.29	Z47556, class A, 20 probes, 20 in Z47556mRNA#2 1596-1866, semenogelin II gene extracted from H.sapiens genes for semenogelin I and semenogelin II
185.62	420.55	355.75	688.95	366.92	300.25	268.49	261.75	130.74	378.25	178.4	252.35	165.53	X99584, class C, 20 probes, 20 in all_X99584 197-630, H.sapiens mRNA for SMT3A protein
48.58	80.6	44.26	244.5	253.99	111.05	99.77	131.55	21.57	108.15	57.06	130.55	7.57	J05200, class A, 20 probes, 20 in J05200mRNA 14740-15238, Human ryanodine receptor mRNA, complete cds
16.05	77.45	17.75	84.15	43.06	164	39.17	216.9	202.8	101.55	18.03	67.3	11.74	L34035, class A, 20 probes, 20 in L34035 1405-1813, Homo sapiens NADP-dependent malic enzyme mRNA, complete cds. /gb=L34035 /ntype=RNA
544.61	4236.9	97.44	3839.1	943.7	4202.95	1288.98	4484.25	1052.53	4155.6	1770.6	5097.7	840.89	D86966, class A, 20 probes, 20 in D86966 4491-5031, Human mRNA for KIAA0211 gene, complete cds
158.6	210.1	274.5	69.65	24.96	377.85	299.88	472.25	563.63	952.85	1290.12	70.55	68.94	Mucin 1, Epithelial, Alt. Splice 6 - Also Represents: M21868
221.18	234.8	117.95	221.9	83.3	1513.2	1526.22	508.55	70.07	438.55	271.6	528.7	41.15	X89894, class C, 20 probes, 20 in all_X89894 2017-2486, H.sapiens mRNA for nuclear receptor
957.35	864.85	1009.25	168.9	14.28	694.4	599.91	385.6	193.75	618.1	8.63	271.1	116.67	L05424, class C, 20 probes, 20 not in GB record, CD44 gene (cell surface glycoprotein CD44) extracted from Human hyaluronate receptor (CD44) gene
38.18	140.6	8.77	792.7	961.52	1359.25	1878.15	366.6	204.21	126.7	130.96	150.45	101.33	U46024, class A, 20 probes, 20 in U46024 2801-3377, Human myotubularin (MTM1) mRNA, partial cds
81.67	647.3	236.17	617.35	611.72	796.35	657.68	664.7	375.76	550.8	72.27	689.1	277.19	U68494, class A, 20 probes, 20 in U68494 1290-1764, Human hbc647 mRNA sequence
45.18	456	22.06	831.4	405.31	371.7	165.32	597.2	53.74	699.4	426.24	713	11.31	L42025, class A, 20 probes, 20 in L42025mRNA 1988-2504, Homo sapiens cellular co-factor (RAB) gene, complete cds
1073.67	2657.7	446.89	3165.3	1120.91	2262.7	1378.29	1447.45	695.16	2134.9	239.43	1782.6	575.16	M23114, class A, 20 probes, 20 in M23114mRNA 3623-4085, Homo sapiens calcium-ATPase (HK1) mRNA, complete cds
78.49	279.1	313.25	509.9	440.81	340.9	344.08	440.85	128.76	160.65	20.15	531.5	361.19	U79247, class A, 20 probes, 20 in U79247 1157-1559, Human clone 23599 mRNA sequence
450.07	833.4	661.57	1032	652.66	282.3	58.41	1107.65	444.84	1654.6	398.24	1855.1	699.89	U57623, class A, 20 probes, 19 in U57623exon#1-4 51-240: 1 in reverseSequence, 6798, Human fatty acid binding protein FABP gene, complete cds
1048.64	5522.4	860.27	4859.05	2337.34	4784.95	1595.02	5410.9	2055.7	4474	2779.5	5900.25	400.15	X13973, class B, 20 probes, 15 in X13973cds 996-1356: 5 in reverseSequence, 1770-1896, Human mRNA for ribonuclease/angiogenin inhibitor (RAI)
1079.82	2006.05	432.96	1676.6	2096.57	2018.6	65.48	2372.95	290.55	1646.5	301.09	2131.3	14.85	U07158, class A, 20 probes, 20 in U07158 641-1169, Human syntaxin mRNA, complete cds
484.51	4952.15	62.15	3599.35	1476.37	2484.9	1626.35	6434.3	3636.79	5134.8	2417.32	4374.65	2216.14	AF015913, class A, 20 probes, 20 in AF015913 1437-1947, Homo sapiens SKB1Hs mRNA, complete cds. /gb=AF015913 /ntype=RNA
5031.49	3961.05	1305.11	8233.4	5047.89	8670.6	5020.6	4877.65	1949.99	5476.45	2491.63	9989.8	5776.5	M19645, class C, 20 probes, 20 in all_M19645 4941-5470, Human 78 kdalton glucose-regulated protein (GRP78) gene, complete cds
165.75	273.95	244.73	365.7	166.31	1255.65	343.3	1495.5	1544.18	1147.8	772.3	333.35	310.35	S77415, class B, 20 probes, 13 in S77415cds 660-948: 7 in reverseSequence, 1449-1617, melanocortin-4 receptor [human, Genomic, 1671 nt]

71.98	105.35	62.72	111.1	80.75	243.2	179.75	453.3	8.06	1262.05	1477.64	266.65	149.69	M37981, class A, 20 probes, 20 in M37981 1751-1829, Human alpha-3 neuronal nicotinic acetylcholine receptor subunit mRNA, complete cds
158.75	500.7	124.59	354.2	201.67	497.45	352.63	401.5	130.11	399.05	113.77	414.55	10.96	Crystallin, Beta B
1788.56	5622.75	1405.8	5453.05	2204.12	4520.4	671.75	5710.65	380.07	3925.15	2221.52	4333.4	265.17	L14856, class A, 20 probes, 16 in L14856cds 746-1100: 4 in reverseSequence, 1324-1393, Human somatostatin receptor gene, complete cds
103.03	480.95	95.81	273.2	169.56	1426.5	1467.81	387.1	362.6	784.1	257.1	446.05	285.6	Z21488, class B, 20 probes, 15 in Z21488cds 2749-3016: 5 in reverseSequence, 3179-3326, H.sapiens contactin mRNA
4522.16	10431.15	3941.77	9796.9	6412.33	3814.6	576.43	3697.75	1600.39	5239.9	728.89	5966.9	591.99	M60858, class A, 20 probes, 20 in M60858mRNA 2193-2481, Human nucleolin gene, complete cds
99.49	104.25	131.03	529.6	463.58	169.05	119.29	226.55	233.27	910.5	742.18	76.5	39.74	X00948, class A, 20 probes, 20 in X00948cds 65-503, Human mRNA for prepro-relaxin H2. /gb=X00948 /ntype=RNA
684.27	1453.65	428.01	1082.05	499.29	1604.05	200.75	2199.55	627.84	493.7	316.64	1148.1	609.53	U49441, class A, 20 probes, 20 in U49441 258-648, Human mitochondrial trifunctional protein beta subunit mRNA, partial cds. /gb=U49441 /ntype=RNA
2474.59	2526.1	2003.8	2190.15	1462.37	1974.55	1134.41	2043.4	1094.6	876.65	274.15	1715.15	1875.04	J03915, class A, 19 probes, 19 in J03915 1313-1721, Human chromogranin A mRNA, complete cds
310.49	379.85	340.19	558.8	551.54	172.7	103.1	392.5	5.37	487.6	563.99	157.2	96.31	X77753, class C, 20 probes, 20 in all_X77753 2390-2787, H.sapiens TROP-2 gene
267.85	260.7	293.59	145.55	71.91	231.4	12.16	374.05	296.35	216.6	51.9	74.95	38.25	U79300, class A, 20 probes, 20 in U79300 930-1404, Human clone 23629 mRNA sequence
1082.51	2367.85	506.22	1397.55	1004.02	1414	1075.23	2645.15	504.38	836.85	430.7	1518.4	458.77	L14754, class A, 20 probes, 20 in L14754 3369-3813, Human DNA-binding protein (SMBP2) mRNA, complete cds
0.21	126.45	25.39	195.6	98.43	174	128.13	132	74.39	128.3	98.71	263.15	136.54	X95289, class C, 20 probes, 20 in all_X95289 48-625, H.sapiens mRNA for HCGIX protein
355.6	698.05	535.07	393.15	193.25	776.4	597.79	1129.9	53.17	667.05	162.28	288.95	49.57	L10955, class A, 20 probes, 16 in L10955cds#1 184-560: 4 in reverseSequence, 394-467, carbonic anhydrase IV gene extracted from Human carbonic anhydrase IV gene, promoter region and
125.16	154.35	117.59	108.65	82.94	795.85	829.65	320.7	342.38	949.1	1212.97	184.75	44.05	U36621, class A, 20 probes, 20 in U36621exon 17-536, Human Y-chromosome RNA recognition motif protein (YRRM) gene
436.14	725.7	596.52	386.3	71.98	1143.85	272.87	1013	237.59	203.6	14.42	993.75	350.51	L38933, class A, 20 probes, 20 in L38933mRNA 883-1393, the longest open reading frame predicts a protein of 202 amino acids, with fair Kozak consensus at the initial ATG codon; an in-frame TGA codon is seen at nucleotide 8; ORF; putative gene extracted from Homo sapiens GT198 mRNA, complete ORF
298.61	227.8	290.76	86.2	16.12	429.5	504.03	82.05	60.03	51.5	54.73	332	405.31	X79200, class C, 16 probes, 16 in all_X79200 380-600, H.sapiens mRNA for SYT-SSX, synovial sarcoma translocation junction. /gb=X79200 /ntype=RNA, H.sapiens mRNA for SYT-SSX, synovial sarcoma translocation junction. /gb=X79200 /ntype=RNA
168.72	1206.75	630.67	2483.35	877.31	2876.75	490.8	1748.65	421.22	1704.7	545.6	2780.1	193.61	X60003, class A, 20 probes, 20 in X60003mRNA 543-965, Human delta CREB mRNA for cAMP-responsive element (CRE) binding protein

547.16	533.7	519.3	227.4	157.26	185.75	165.96	590.6	428.79	849.95	1112.07	487.35	210.65	M91432, class A, 20 probes, 20 in M91432mRNA 1664-2144, Human medium-chain acyl-CoA dehydrogenase (MCAD) gene
17.11	346.35	170.91	398.4	366.56	255.7	105.22	330.45	213.62	370.4	199.26	228.55	91.15	U01317, class C, 20 probes, 20 in all_U01317 19502-63478, epsilon-globin gene extracted from Human beta globin region on chromosome 11, epsilon-globin gene extracted from Human beta globin region on chromosome 11, epsilon-globin gene extracted from Human beta globin region on chromosome 11
68.17	191.75	16.62	223.8	77.07	243.65	240.63	208.1	102.11	443.05	442.3	157.5	24.89	X85740, class A, 20 probes, 20 in X85740mRNA 1112-1604, H.sapiens mRNA for C-C chemokine receptor-4
366.78	702.05	432.11	277.5	207.18	1297.2	1658.87	537.9	520.85	292.3	263.89	775.2	875.26	U66578, class A, 20 probes, 19 in U66578cds 598-1036: 1 in reverseSequence, 1277, Human putative G protein-coupled receptor (GPR23) gene, complete cds
10377.5	51551.9	15406.73	50325.05	16023.39	49797.05	46455.85	45198.9	13705.85	44280.75	5093.79	52114.25	16431.96	Ribosomal Protein S12
273.16	114.6	89.52	179.75	21.14	419.45	475.25	70.45	19.3	140.4	46.53	177.45	118.58	U59913, class A, 20 probes, 20 in U59913 1640-2168, Human chromosome 5 Mad homolog Smad5 mRNA, complete cds
110.03	242.05	88.46	611.8	642.9	1043.8	1097.43	833.65	824.7	1400.15	1623.87	229.5	96.17	X07732, class A, 20 probes, 20 in X07732mRNA#2 991-1543, Human hepatoma mRNA for serine protease hepsin
103.66	130.4	125.3	302.5	346.2	185.85	166.67	144.7	87.4	105.6	46.81	96.2	21.07	Y09912, class A, 20 probes, 20 in Y09912mRNA 757-1315, H.sapiens AP-2 beta gene
848.53	2875.35	248.69	2492.7	1253.98	1910.6	334.6	2924.2	322.3	1630.85	714.96	2681.3	295.43	Rhesus (Rh) Blood Group System Ce-Antigenl, Alt. Splice 3, Rhviii - Also Represents: X63097
53.1	51.25	45.33	67.6	20.22	753.4	910.61	265.9	230.66	190.9	229.95	85	11.17	X74142, class C, 20 probes, 20 in all_X74142 1952-2535, H.sapiens HBF-1 mRNA for transcription factor
48.65	258.85	132.44	97.8	0.14	313.5	325.98	349.95	46.17	332.6	273.93	515.65	499.85	X63097, class C, 20 probes, 20 in all_X63097 2195-2670, H.sapiens mRNA for rhesus polypeptide (RhXIII)
517.74	872.95	365.64	607.35	624.59	1808.6	1045.1	1562.15	15.34	806.8	725.07	819.7	546.03	X58199, class A, 20 probes, 20 in X58199mRNA 2491-2573, Human mRNA for beta adducin
261.13	1020.1	154.43	671.1	282.42	670.45	540.72	337.65	21.57	526.05	258.16	355.1	227.41	L13738, class A, 20 probes, 20 in L13738mRNA 4076-4490, Human activated p21cdc42Hs kinase (ack) mRNA, complete cds
8084.07	79972	529.62	73008.55	10275.6	74554.2	51945.05	87557.65	2346.39	91428.35	11758.69	93708.41	1761.12	X63527, class A, 20 probes, 17 in X63527cds 99-549: 3 in reverseSequence, 637-655, H.sapiens mRNA for ribosomal protein L19
387.28	916.4	381.98	923.95	299.04	2338.3	1729.44	887.1	154.86	692.15	8.7	890.3	144.96	Homo sapiens /REF=M11507 /DEF=Human transferrin receptor mRNA, complete cds /LEN=2826 (_5, _M, _3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
14.42	467.35	26.66	630.05	298.33	1143.9	1169.84	195.1	185.83	422.4	340.12	813.65	408.92	X85137, class C, 20 probes, 20 in all_X85137 3131-3726, H.sapiens mRNA for kinesin-related protein
125.23	141.7	91.78	276.15	163.55	282.35	148.85	133.9	3.25	129.2	140.43	478	555.08	M74091, class A, 20 probes, 20 in M74091 925-1453, Human cyclin mRNA
917.26	9312	1875.39	9388.15	422.92	8994.05	4565.44	9710.4	255.55	6429.55	1107.12	8840.35	555.86	J04444, class B, 20 probes, 13 in J04444cds 596-944: 7 in reverseSequence, 3590-3740, Human cytochrome c-1 gene, complete cds

46.39	47	43.84	83.85	102.18	148.35	136.4	111.05	35	117	88.25	77.7	0.85	L49218, class A, 20 probes, 20 in L49218exon 4-91, Homo sapiens retinoblastoma susceptibility protein (RB1) E413K 1 bp deletion mutant (resulting in premature stop at amino acid 416) gene, exon 13 (L11910 bases 73717-73901).
286.66	134.2	112.43	812.95	694.03	995.4	766.65	440.55	420.23	1048.7	764.95	460.3	258.8	X07820, class B, 20 probes, 11 in X07820cds 1252-1378: 9 in reverseSequence, 1430-1694, Human mRNA for metalloproteinase stromelysin-2
295.92	1304.35	723.16	1755.1	946.96	1365.2	1295.56	1454.7	109.6	1568.5	637.24	1426.65	129.75	M37245, class B, 20 probes, 12 in M37245cds 272-434: 8 in reverseSequence, 113-353, Human Ig superfamily cytotoxic T-lymphocyte-associated protein (CTLA-4) gene
171.61	329.25	39.24	292.45	116.46	509.85	328.73	162.65	9.55	4406.5	5293.54	279.05	63.57	M94547, class A, 20 probes, 20 in M94547mRNA 55-565, HUMMLC2At; Homo sapiens; ; 593 base-pairs
411.47	7092.75	881.41	7708.9	506.71	5154.9	3377	5859.5	1940.02	4172.45	3119.97	4972.45	1040.51	J05272, class A, 20 probes, 20 in J05272 2311-2809, Human IMP dehydrogenase type 1 mRNA complete cds
263.11	316.6	238.72	620.4	329.09	325.9	140.15	514.3	180.6	427.3	158.39	304.85	271.03	X14787, class C, 20 probes, 20 in all_X14787 5124-5701, Human mRNA for thrombospondin
41.08	45.9	38.18	87.75	99.49	128.45	112.64	115.2	54.59	96.8	74.81	63.2	3.96	X16699, class C, 20 probes, 20 in all_X16699 2053-2130, Human mRNA for cytochrome P-450HP
83.72	753.6	393.15	1188.05	499.71	522.85	388.41	378.35	399.44	1165.55	583.15	985.1	383.39	M60483, class A, 18 probes, 18 in M60483mRNA 1636-2107, protein phosphatase-2A catalytic subunit-alpha gene extracted from Human protein phosphatase 2A catalytic subunit-alpha gene, complete cds
36.2	720.6	219.2	522.4	521.7	259.5	197.99	445.6	148.07	1319.95	1327.45	612.2	189.65	U77665, class A, 20 probes, 20 in U77665 393-873, Human RNaseP protein p30 (RPP30) mRNA, complete cds.
797.47	1324.5	544.61	1418.5	1263.74	919.25	273.16	1567.55	1200.74	777.25	324.21	1801.75	956.93	L48728, class C, 20 probes, 19 in all_L48728 380-657: 1 not in GB record, TCRBV10S1 gene extracted from Homo sapiens T cell receptor beta (TCRBV10S1) gene, complete cds
201.03	313	139.58	1297.95	846.62	2394.2	3062.2	1694.8	880.49	1006.2	729.59	512.05	80.82	S75989, class A, 20 probes, 20 in S75989 1492-1948, gamma-aminobutyric acid transporter type 3 [human, fetal brain, mRNA, 1991 nt]
298.82	645.2	102.11	1249.9	946.25	1019.4	1007.91	581.95	139.51	455.05	70.07	481.2	409.13	Y08991, class B, 20 probes, 11 in Y08991cds 3846-4038: 9 in reverseSequence, 4743-4845, H.sapiens mRNA for adaptor protein p150
633.21	1369.8	832.41	1530.05	606.34	515.45	192.12	1054.7	582.37	866.95	792.6	460.45	140.22	X99656, class B, 20 probes, 15 in X99656cds 798-1068: 5 in reverseSequence, 1251-1329, H.sapiens mRNA for protein containing SH3 domain, SH3GL1
46.24	67	39.6	60.5	0.28	278.55	224.93	127.4	125.87	413.2	454.67	89.1	55.58	U90268, class A, 20 probes, 20 in U90268 1427-1703, Human Krit1 mRNA, complete cds.
565.69	539.5	590.29	306.2	335.59	388.1	377.31	180.8	171.83	456.4	131.38	224.95	241.05	U79245, class A, 20 probes, 20 in U79245 1139-1541, Human clone 23586 mRNA sequence
37.97	66.7	64.21	75.1	16.55	274.9	232.36	194.65	25.53	254.75	293.94	59.8	48.51	X58022, class A, 20 probes, 20 in X58022mRNA 803-1223, Human mRNA for corticotropin-releasing factor binding protein (CRF-BP)
43.49	136.15	27.22	214.05	153.8	531.2	366.42	148.6	32.67	180.6	91.64	191.65	104.44	X99657, class C, 20 probes, 20 in all_X99657 783-1318, H.sapiens mRNA for protein containing SH3 domain, SH3GL2
753.42	19288.15	9673.57	11456.2	2920.92	12705.1	11735.43	9421.55	5689.59	9986.9	6461.97	13465.95	2276.25	D88422, class C, 20 probes, 15 in all_D88422 230-483: 5 in D88422cds 13-188, Human DNA for cystatin A

35.14	68.9	7.92	234.4	208.46	80.55	37.55	172.2	71.28	553.9	580.82	112.75	80.12	J05213, class A, 20 probes, 20 in J05213 430-958, Human sialoprotein mRNA, complete cds
241.12	216.05	238.79	305.95	299.46	879.45	792.03	619.45	451.35	1048.35	1340.6	367.85	62.3	M19507, class A, 20 probes, 20 in M19507mRNA#4 2620-3184, Human myeloperoxidase mRNA, complete cds
430.56	1447	63.22	2683.5	1069.57	1310.2	685.19	1485.7	177.2	1431.7	429.21	1008.45	86.9	M37104, class A, 20 probes, 20 in M37104 13-421, Human mitochondrial ATPase coupling factor 6 subunit (ATP5A) mRNA, complete cds
83.16	85.1	20.22	741.65	865.57	480.8	429.92	77.5	60.53	332.95	374.7	353.5	101.82	X77307, class B, 20 probes, 13 in X77307cds 1244-1382: 7 in reverseSequence, 1491-1701, H.sapiens mRNA for 5-HT2B serotonin receptor
134.92	157.05	111.23	594.8	82.17	368.3	327.25	189.55	202.16	926.05	1120.98	60.85	19.59	X58234, class A, 20 probes, 20 in X58234mRNA 31-253, Human mRNA for anti-lectin antibody epitope (clone p36/8-5)
359.63	634.1	469.38	902.95	1021.27	828.55	191.7	722.1	474.19	705.3	411.39	628.1	98.15	X52520, class C, 20 probes, 20 in all_X52520 2414-2673, Human mRNA for tyrosine aminotransferase (TAT) (EC 2.6.1.5)
1.13	108.55	3.04	178.55	167.51	250.75	221.11	169.25	74.32	174.55	1.2	252.5	67.46	J04449, class A, 20 probes, 20 in J04449 2290-2776, Homo sapiens (clone NF 10) cytochrome P-450 nifedipine oxidase mRNA, complete cds
81.18	2354.8	242.96	2497.45	980.83	1429.65	951.84	1918.4	389.47	2219.1	506.43	1898.45	322.23	U14550, class A, 20 probes, 20 in U14550 1319-1877, Human sialyltransferase SThM (sthm) mRNA, complete cds
1290.96	4904.35	315.86	4123.15	1266.36	6434.4	2956.56	6974.95	1514.55	5528.2	257.67	6865.1	3763.65	M34182, class A, 20 probes, 20 in M34182mRNA#1 1112-1517, Human testis-specific protein kinase gamma-subunit mRNA, complete cds
198.63	487.85	472.14	968.05	624.45	705.95	71.63	162.15	160.16	1048.5	418.47	862.45	975.31	S78653, class C, 20 probes, 20 in all_S78653 1782-2347, mrg=mas-related [human, Genomic, 2416 nt]
278.88	2377.45	651.32	2210.9	544.76	1008.5	1099.69	1170.3	119.22	2974.25	1314.44	2090.65	287.44	U72761, class A, 20 probes, 20 in U72761 2753-3233, Human karyopherin beta 3 mRNA, complete cds. /gb=U72761 /ntype=RNA
311.27	486.1	392.59	216.6	45.25	423.95	288.43	702.75	161.29	553.2	670.9	146.25	78.42	U15642, class A, 20 probes, 20 in U15642 975-1472, Human transcription factor E2F-5 mRNA, complete cds
67.95	234.5	235.33	548.35	162.99	753.25	549.78	418.05	168.64	315.15	166.38	199.7	170.27	U13045, class A, 20 probes, 20 in U13045 2126-2690, Human nuclear respiratory factor-2 subunit beta 1 mRNA, complete cds
442.58	355.5	417.33	169.5	88.53	183.05	110.1	107.65	43.35	270.75	218	154.4	3.11	X83107, class C, 20 probes, 20 in all_X83107 1867-2348, H.sapiens Bmx mRNA for cytoplasmic tyrosine kinase
90.86	313.6	19.37	274.2	45.11	737.95	568.02	369.05	55.65	298.9	4.81	373.2	128.13	X90846, class C, 15 probes, 15 in all_X90846 2935-3407, H.sapiens mRNA for mixed lineage kinase 2, H.sapiens mRNA for mixed lineage kinase 2
15.98	107.5	124.31	265.7	23.33	963.6	1056.84	231.4	39.74	897.8	1219.9	183.75	99.21	X00038, class C, 20 probes, 20 in all_X00038 599-718, Human H4 histone gene
6459.28	15574.6	5316.03	13542.35	5749.27	12814.85	9687.58	16398.85	6997.03	9628.1	12608.14	15579.4	3360.74	Z21507, class A, 20 probes, 16 in Z21507cds 424-802: 4 in reverseSequence, 901-955, H.sapiens EF-1delta gene encoding human elongation factor-1-delta
37.26	117.4	1.27	354.6	54.87	209.45	188.59	681.95	681.01	727.35	904.74	165.7	144.25	M21535, class A, 20 probes, 20 in M21535 2603-3101, M17390 Human erg protein (ets-related gene) mRNA, complete cds.

35.28	98.6	35.92	200	191.77	120.7	26.3	154.7	74.39	82.15	22.56	112.7	35.07	X02544, class A, 20 probes, 16 in X02544cds 256-544: 4 in reverseSequence, 688-772, Human mRNA for alpha1-acid glycoprotein (orosomucoid)
23.97	4376.85	272.87	3378.2	1398.66	2436.8	2474.59	2739.2	30.41	4665	2065.6	4591.35	1809.13	L49380, class A, 20 probes, 20 in L49380mRNA 2916-3414, Homo sapiens clone B4 transcription factor ZFM1 mRNA, complete cds
229.88	599.95	198.63	370.3	278.88	930.15	711.28	471.3	326.12	759.2	575.02	921.55	7.28	D63135, class A, 20 probes, 20 in D63135mRNA 31-499, Human mRNA for ETS-like 30 kDa protein. /gb=D63135 /ntype=RNA
135.34	140.7	115.12	190.25	5.02	204.85	162	99.55	2.47	285	24.75	229.35	212.2	Y10571, class B, 20 probes, 13 in Y10571cds 696-978: 7 in reverseSequence, 1002-1194, H.sapiens mRNA for dinG gene
203.36	210.2	246.64	798.4	688.58	375.4	179.46	747.8	394.42	782.9	1018.38	563.75	83.37	Y10506, class A, 20 probes, 20 in Y10506mRNA 251-593, H.sapiens mRNA for CD110 protein. /gb=Y10506 /ntype=RNA
94.33	158.05	79.27	661.8	848.39	1181.15	1489.94	535.75	588.95	228.65	63.85	466.5	554.23	L10403, class A, 20 probes, 20 in L10403 834-1254, Homo sapiens DNA binding protein for surfactant protein B mRNA, complete cds
6281.02	44608.05	9998.7	24884.3	15743.03	26602.1	11746.74	50355	20853	52159.35	24772.28	62917	16496.24	Fibronectin, Alt. Splice 1 - Also Represents: HG3044-HT2527, X02761
571.63	514.25	570	255.5	69.01	646.35	105.57	164.65	32.88	332	55.01	908.95	388.13	M96759, class A, 20 probes, 20 in M96759mRNA 783-1299, Human rod outer segment membrane protein 1 (ROM1) gene exons 1-3, complete cds
252.3	262.2	242.25	168.8	33.66	746.1	939.89	220.05	170.48	129.25	75.59	220.85	22.84	X16660, class C, 19 probes, 19 in all_X16660 1795-2049, open reading frame p25 (AA 1-223) gene extracted from Human HTLV-I related endogenous retroviral sequence (HRES-1/1)
291.4	1673.85	598.57	1711.7	736.1	3778.3	2890.79	1407.15	269.62	2590.35	738.43	1463.35	105.71	Laminin, A Polypeptide
2090.35	15035.35	5156.29	17108.3	41.44	16557.05	8229.95	14863.25	1441.72	11265.7	4723.9	16133.9	1344.49	M21388, class C, 6 probes, 6 in all_M21388 2-95, Human unproductively rearranged Ig mu-chain mRNA V-region (VD), 5 end, clone mu-3A1A., Human unproductively rearranged Ig mu-chain mRNA V-region (VD), 5 end, clone mu-3A1A.
227.41	452.4	221.18	544.5	28.14	347.65	102.18	537.35	143.19	595.25	623.17	763.4	328.52	U32439, class A, 20 probes, 20 in U32439 1324-1822, Human regulator of G-protein signaling similarity (RGS7) mRNA, partial cds
611.93	444	550.98	604.15	449.93	574.15	658.81	748.2	459.34	304.05	105.57	1050.05	204.14	D14043, class A, 20 probes, 20 in D14043 1834-2362, Human mRNA for MGC-24, complete cds
452.41	1062.6	136.61	682.95	412.17	1640.35	1006	1040	714.18	2021.6	1284.67	1210.35	939.67	M63962, class A, 20 probes, 20 in M63962mRNA 2985-3507, Human gastric H,K-ATPase catalytic subunit gene, complete cds
1211.27	5134.5	1155.13	6286.2	2560.29	4444.85	1806.02	5994	1752.92	4006.8	2489.58	4589.85	559.53	X01630, class B, 20 probes, 15 in X01630cds 883-1213: 5 in reverseSequence, 1312-1468, Human mRNA for argininosuccinate synthetase
72.05	418.1	308.58	726.65	402.41	523.75	33.3	834.9	280.44	230.2	251.45	484.35	505.65	M24594, class A, 20 probes, 20 in M24594mRNA 1077-1593, Human interferon-inducible 56 Kd protein mRNA, complete cds
514.42	442.95	452.76	666.85	112.5	545.35	511.03	1392.05	1104.57	131.5	105.36	99.05	8.13	Z46632, class C, 4 probes, 4 in all_Z46632 2953-3206, H.sapiens HSPDE4C1 gene for 3 ,5 -cyclic AMP phosphodiesterase, H.sapiens HSPDE4C1 gene for 3 ,5 -cyclic AMP phosphodiesterase

1511.79	3488.2	832.69	3570.85	1344.56	2403	2351.55	1734.75	910.82	2144.4	26.3	2453.05	538.32	M96982, class A, 20 probes, 20 in M96982 483-801, Homo sapiens U2 snRNP auxiliary factor small subunit, complete cds
411.25	590.3	612.5	166.25	144.6	1115	1366.7	331.75	18.31	957.75	1002.61	563.5	658.88	M23178, class C, 20 probes, 20 in all_M23178 2889-3818, Human homologue-1 of gene encoding alpha subunit of murine cytokine (MIP1/SCI), complete cds
241.12	1314.4	283.69	1404.65	2.05	2138.5	743.03	2376.9	938.47	2743.45	793.87	1470.7	1065.33	X83573, class A, 20 probes, 20 in X83573 1377-1803, H.sapiens ARSE mRNA
149.98	2627.85	295.92	1121.1	1207.03	1167.1	1230.51	2057.6	461.74	1291	749.53	2266.5	132.79	U34877, class A, 20 probes, 20 in U34877 511-997, Human biliverdin-IXalpha reductase mRNA, complete cds
194.74	41	27.58	217.7	97.44	247.45	227.9	683.95	231.72	814.35	971.35	339.8	241.12	M18079, class B, 20 probes, 12 in M18079cds 85-343: 8 in reverseSequence, 4158-4278, Human, intestinal fatty acid binding protein gene, complete cds, and an Alu repetitive element
299.46	1682.5	95.88	1628.8	627.49	1874.05	958.48	659.65	672.81	1077.6	1329.36	1317.7	171.54	L41559, class A, 20 probes, 20 in L41559mRNA 61-475, Homo sapiens pterin-4a-carbinolamine dehydratase (PCBD) mRNA, complete cds
18.67	112.35	7.42	243.05	27.51	456.35	40.38	178.4	183.71	161.4	8.49	116.5	45.82	X00540, class C, 20 probes, 20 not in GB record, H.sapiens gene encoding prolactin, exon 2 and joined CDS
281.85	107.35	37.12	354.6	142.55	184.35	67.25	656.6	199.26	525.25	104.44	223.8	205.63	Y10262, class A, 19 probes, 19 in Y10262cds 1163-1693, H.sapiens EYA3 gene. /gb=Y10262 /ntype=DNA /annot=CDS
980.33	817.7	723.09	2015.45	221.11	779.65	929.77	2470.5	954.31	969.85	851.99	723.95	412.03	U91985, class A, 20 probes, 20 in U91985 1014-1320, Human DNA fragmentation factor-45 mRNA, complete cds
39.1	51.75	48.15	107.2	101.96	721.15	837.43	219.3	1.27	930.7	1195.86	193.75	17.32	X89426, class C, 20 probes, 20 in all_X89426 1465-1958, H.sapiens mRNA for ESM-1 protein
300.94	1346.05	203.29	1008.1	989.53	817.85	517.96	1607.75	244.16	1827.5	1536.12	1553.85	209.66	U57316, class C, 20 probes, 20 in all_U57316 1593-1996, Human GCN5 (hGCN5) gene, complete cds
368.97	2113.25	449.22	4383.65	4920.12	6301.05	6664.55	2484.05	1576.49	6102.8	471.36	6552.1	4738.04	AB000897, class A, 20 probes, 20 in AB000897 43-373, Homo sapiens mRNA for cadherin FIB3, partial cds. /gb=AB000897 /ntype=RNA
282.42	743.5	272.24	421.6	102.25	1078.7	705.83	259.45	229.46	296.8	22.77	369.7	246.78	D28118, class A, 20 probes, 20 in D28118 1807-2263, Human mRNA for DB1, complete cds
35.99	43	30.97	70.3	54.73	523.45	716.51	46.6	7.5	473.6	582.94	146.5	156.13	X66360, class A, 20 probes, 16 in X66360cds 1134-1518: 4 in reverseSequence, 1629-1689, H.sapiens mRNA PCTAIRE-2 for serine/threonine protein kinase
1115.6	2771.3	584.64	2250	727.61	3447.55	1009.39	2512.5	1380.7	1861.55	429	2493.55	1559.38	J04599, class A, 20 probes, 20 in J04599 1078-1630, Human hPGI mRNA encoding bone small proteoglycan I (biglycan), complete cds
515.2	851.9	656.34	1113.85	798.39	782.95	94.12	1233	941.02	1478.35	819.89	661.6	539.38	M27492, class A, 20 probes, 20 in M27492 4336-4864, Human interleukin 1 receptor mRNA, complete cds
97.86	540.25	221.96	388.5	195.3	284.5	138.88	216.25	107.98	396.95	84.78	227.6	168.57	U11732, class A, 20 probes, 20 in U11732 997-1531, Human ets-like gene (tel) mRNA, complete cds
840.33	2620.5	744.16	3259.95	600.69	3743.35	1836.85	2551.55	261.84	1607.5	1505.01	3062.65	1288.7	M18700, class A, 20 probes, 20 in M18700cds 288-784, Human elastase III A gene, exon 8.
202.09	327.25	108.26	489.3	444.49	398.4	247.63	538.55	40.94	313.85	63.14	441.7	160.65	X80343, class A, 20 probes, 16 in X80343cds 435-807: 4 in reverseSequence, 1006-1036, H.sapiens p35 mRNA for regulatory subunit of cdk5 kinase

1158.45	3610.55	955.51	3219.3	597.65	3759.5	1099.98	2911.9	802.71	4436.7	2631.57	3591.55	479.06	X74795, class B, 20 probes, 14 in X74795cds 1923-2181: 6 in reverseSequence, 2272-2488, H.sapiens P1-Cdc46 mRNA
244.8	2588.75	326.05	1971.5	1526.36	1429.05	725.28	2824.75	1408.06	2800.1	247.63	1491.2	523.12	M36429, class A, 20 probes, 20 in M36429 827-1412, Human transducin beta-2 subunit mRNA, complete cds
23.33	26.7	6.51	46.2	12.16	77.4	22.63	53.9	2.12	143.6	105.22	246.9	213.4	M63154, class A, 20 probes, 20 in M63154 977-1541, Human intrinsic factor mRNA, complete cds
292.81	289.6	306.32	528	120.21	723.85	536.62	1399.3	102.53	492.7	282.7	1150.6	48.79	X54925, class C, 20 probes, 20 in all_X54925 1537-1904, H.sapiens mRNA for type I interstitial collagenase
3536.24	18119.8	2564.11	14999.05	7910.47	9808.65	7133.08	15430.25	9367.54	13183.55	8050.48	13091.5	2446.59	X04106, class C, 20 probes, 20 in all_X04106 907-1478, Human mRNA for calcium dependent protease (small subunit)
1189.42	1285.75	989.31	977.8	958.55	1793	1141.41	1673.4	792.95	1150	807.8	1336.9	76.23	M22919, class C, 20 probes, 20 in all_M22919 3226-3665, MLC gene (non-muscle myosin light chain) extracted from Human nonmuscle/smooth muscle alkali myosin light chain gene, complete cds
74.88	106.2	93.76	94.3	26.59	339.7	295	191.75	153.94	128.9	71.13	279.2	273.65	M95767, class A, 20 probes, 20 in M95767 1065-1563, Homo sapiens di-N-acetylchitobiase mRNA, complete cds
174.58	632.9	411.39	732.95	542.85	1789.9	657.04	595.95	454.6	1468.95	101.05	1253.85	663.76	U43408, class A, 20 probes, 20 in U43408 2308-2668, Human tyrosine kinase (Tnk1) mRNA, complete cds
160.94	203.05	133.71	698.7	691.41	536.75	406.8	784	196.01	296.5	106.49	429.25	173.03	U85767, class A, 20 probes, 20 in U85767 13-505, Human myeloid progenitor inhibitory factor-1 MPIF-1 mRNA, complete cds
340.61	327.3	293.59	546.85	652.45	184.45	29.49	396	263.89	255.9	75.52	104.85	7.85	Trithorax Homolog Hrx - Also Represents: L04731
96.8	330.1	266.01	779.25	428.44	1538.1	1964.2	153.2	22.49	738.55	286.45	545.5	714.88	L47276, class A, 20 probes, 20 in L47276 383-934, Homo sapiens (cell line HL-60) alpha topoisomerase truncated-form mRNA, 3 UTR. /gb=L47276 /ntype=RNA
111.3	139	98.71	175.1	97.44	106.75	32.88	237.8	216.23	147.45	68.09	136.4	76.93	X03473, class B, 20 probes, 13 in X03473cds 270-552: 7 in reverseSequence, 1054-1150, Human gene for histone H1(0)
24.75	9971.25	718.77	9467.8	2432.59	6833.1	4659.83	9986.45	1386	9187.75	3428.55	9109.3	2722.64	D29012, class A, 20 probes, 20 in D29012 230-752, Human mRNA for proteasome subunit Y, complete cds
418.54	372.2	450.57	158.5	70.71	177.9	78.06	259.5	116.96	358	201.38	301.1	173.81	S69231, class A, 20 probes, 20 in S69231 1444-1981, TYRP2=tyrosinase-related-protein-2 [human, melanocytic cell line SK-MEL-19, mRNA, 2056 nt]
314.38	1440.4	162.78	1613.95	1265.08	2140.35	1476.93	1609.4	151.04	1388.2	35.92	1402.85	315.58	U43030, class A, 20 probes, 20 in U43030 1121-1484, Human cardiotrophin-1 (CTF1) mRNA, complete cds
7.78	139.75	30.33	125.5	11.88	562.45	312.05	193.45	92.42	222.3	180.17	225.6	99.28	M55284, class A, 20 probes, 20 in M55284 1800-2364, Human protein kinase C-L (PRKCL) mRNA, complete cds
364.3	3138.8	231.08	3009.15	692.61	2836.35	1078.13	2386.25	735.46	2873	271.39	2664.75	935.43	U07857, class A, 20 probes, 20 in U07857 173-677, Human 18 kDa Alu RNA binding protein mRNA, complete cds
447.53	213.05	155.35	1155.25	920.3	278.15	273.16	334.55	296.35	495.35	193.96	559.3	665.25	D49354, class A, 20 probes, 20 in D49354 769-1293, Human mRNA for enhancer protein in hsp70 gene, partial cds
69.79	150.7	40.73	341.55	117.59	639.35	457.14	160.4	77.22	1307.65	1058.89	753.35	884.66	Atpase, Na+/K+ Transporting, Alpha 1 Polypeptide
408.99	998.4	604.86	1879.5	532.03	1086	1045.95	977.25	106.42	1352.3	258.38	918.9	205.49	U60975, class A, 20 probes, 20 in U60975 6398-6824, Human hybrid receptor gp250 precursor mRNA, complete cds

66.75	197.45	56.92	184.1	45.25	234.85	8.98	167.25	31.89	227.6	141.7	182.8	33.23	M62840, class A, 20 probes, 20 in M62840mRNA 1755-2175, Human acyloxyacyl hydrolase mRNA, complete cds
48.08	132.55	142.48	270.75	294.79	1348.95	1148.41	797.4	340.4	766.05	176.42	445.15	168.36	L77561, class A, 20 probes, 20 in L77561mRNA 583-1093, Homo sapiens DGS-D mRNA, 3 end
13.29	119	18.95	135.55	78.28	240.75	179.53	129.65	1.63	144.2	72.41	248.9	47.66	M76729, class A, 20 probes, 20 in M76729 6573-7077, Human pro-alpha-1 (V) collagen mRNA, complete cds
428.93	1038.6	640.36	1141.35	964.42	653.8	370.52	1741.65	215.88	1495.9	487.2	2488.1	913.58	L42621, class A, 20 probes, 20 in L42621mRNA 1775-2231, Homo sapiens Ly-9 mRNA, complete cds
26.09	80	58.83	113.8	12.45	201.1	59.96	105.6	26.59	186.75	233.84	478.5	64.35	M64358, class A, 20 probes, 20 in M64358exon 16-189, Human rhom-3 gene, exon. /gb=M64358 /ntype=DNA /annot=exon
874.97	2000.25	879.15	2622.1	2226.4	1178.5	1438.4	1354.85	103.59	2251.3	700.46	2029.95	492.5	L24804, class A, 20 probes, 20 in L24804 223-721, Human (p23) mRNA, complete cds
280.44	767.2	168.57	565	321.31	1181.6	442.08	1564.3	281.99	487.55	98.08	863.8	292.74	M55422, class B, 11 probes, 11 in M55422 2463-2733, Human Krueppel-related zinc finger protein (H-plk) mRNA, complete cds
0.78	38.95	13.51	75.65	18.17	1200.75	1645.79	79.45	22.42	102.9	94.89	112.65	19.87	U18004, class A, 20 probes, 20 in U18004 31-205, HSU18004 Homo sapiens cDNA
60.17	160.45	4.03	317.65	94.54	277.35	77.29	304.45	60.88	361.05	166.95	281.1	21.92	Transition Protein 2 - Also Represents: X63759
187.31	3375.1	990.52	2255.15	1780.85	1551.1	1220.89	3754.1	895.48	2224.3	1694.37	3600.3	581.81	S52969, class C, 20 probes, 20 not in GB record, Description: alpha-1,3 fucosyltransferase gene extracted from alpha-(1.3/1,4) fucosyltransferase/FucT-III, alpha-1,3 fucosyltransferase/FucT-VI [human, Genomic, 1045 nt 4 segments]
19.94	72.45	69.23	492.9	678.82	108.2	60.53	163.2	17.11	71.1	44.55	275.45	355.75	X83857, class C, 18 probes, 18 in all_X83857 1431-1809, H.sapiens mRNA for prostaglandin E receptor (EP3a1)
53.32	124.55	61.16	574.5	693.67	520.55	560.95	150.55	84.64	180.7	91.92	452.4	523.68	Collagen, Type Ii, Alpha 1 - Also Represents: J00116, X06268
832.69	877.25	409.2	1594.4	13.58	569.6	450.29	1363.75	355.46	1011.8	619.57	994.7	158.53	X75091, class A, 20 probes, 17 in X75091cds 300-653: 3 in reverseSequence, 848-892, H.sapiens mRNA for HLA-DR associated protein II (PHAPII)
825.69	1615.55	109.39	1201.25	155.63	3307.95	2105.69	2183.75	72.62	3328.35	1698.4	2229.95	735.04	U62437, class A, 20 probes, 20 in U62437 1943-2393, Human nicotinic acetylcholine receptor beta2 subunit precursor, mRNA, complete cds
76.58	90.55	72.62	98.1	57.28	120.85	40.8	144.2	99.84	579.95	726.55	90.35	68.94	U13061, class C, 20 probes, 20 in all_U13061 518-1020, Human dehydroepiandrosterone sulfotransferase (STD) gene
125.58	233.4	130.25	355.55	417.83	65.3	36.77	272.15	201.88	83.2	54.02	166.25	183.92	D86980, class A, 20 probes, 20 in D86980 4616-5192, Human mRNA for KIAA0227 gene, partial cds
120.7	321.4	42.28	331.5	167.02	1245.1	988.39	475.85	217.29	411.3	252.72	684	456.79	Z71389, class A, 20 probes, 20 in Z71389mRNA 85-295, H.sapiens mRNA for skin-antimicrobial-peptide 1 (SAP1).
395.06	674.9	654.78	498.3	536.55	839.85	912.52	290.3	7.35	85.45	45.04	377.7	126.29	Z29331, class C, 20 probes, 20 in all_Z29331 1560-1981, H.sapiens (23k/3) mRNA for ubiquitin-conjugating enzyme UbcH2
909.62	1629.5	271.25	642.05	441.59	682.35	49.29	1039.4	249.61	687.75	240.49	837.1	49.07	U19718, class A, 20 probes, 20 in U19718 479-947, Human microfibril-associated glycoprotein (MFAP2) mRNA, complete cds
255.48	304.05	260.29	300.45	241.19	616	554.51	258.55	156.62	217.25	99.49	321.55	166.95	Pan-2

1017.95	1116.3	1017.24	407.45	62.86	798.9	536.84	362.3	81.46	576	167.16	540.1	65.76	U82468, class A, 20 probes, 20 in U82468 1566-2091, Human tubby related protein 1 (TULP1) mRNA, complete cds.
29.06	218.85	181.09	205.85	210.36	1389.5	591.14	420.1	404.75	726.85	344.71	1436.6	692.12	Phosphoprotein Tal2
5.8	165.9	53.03	332.15	217.15	466.8	428.08	773.8	792.24	292.9	4.53	1227.3	1474.6	M60614, class A, 20 probes, 20 in M60614 1996-2060, Human Wilms tumor (WIT-1) associated protein mRNA, complete cds
610.3	1222.1	324.99	925.85	555.86	1104.25	119.71	311.55	348.67	357.85	79.69	803.85	538.04	U18291, class A, 20 probes, 20 in U18291 1439-1973, Human CDC16Hs mRNA, complete cds
353.48	2863.45	52.96	2483.05	268.77	2841.3	664.68	2693.15	802.92	1913.8	866.21	2037.7	587.75	U57877, class A, 20 probes, 20 in U57877 766-1144, Human integral membrane protein CII-3 mRNA, nuclear gene encoding mitochondrial protein, complete cds
711.07	976.6	509.12	1229.7	1141.69	1002.55	473.27	1348.9	33.38	2229.45	350.65	1910.2	409.56	L47738, class A, 20 probes, 20 in L47738 2286-2856, Homo sapiens inducible protein mRNA, complete cds
225.28	251.35	210.36	637.25	179.39	509	113.42	507.9	17.82	717.85	255.9	460.05	30.76	Zinc Finger Protein (Gb:M88357)
18.88	189.35	24.68	211	11.46	220.45	197.35	216.85	36.56	192.05	15.06	284.9	190.07	Cpg-Enriched Dna, Clone E35
89.66	265.7	277.89	140.05	62.86	589.3	350.44	369.35	72.2	2765.3	3613.17	537.95	433.81	S77154, class A, 20 probes, 20 in S77154 1862-2362, TINUR= NGFI-B/nur77 beta-type transcription factor homolog [human, T lymphoid cell line, PEER, mRNA, 2469 nt]
66.54	2731.9	639.08	2033.4	675.43	622.75	437.49	974.35	732.92	766.8	661.85	1263.5	528.92	X69141, class C, 20 probes, 20 in all_X69141 1444-1997, H.sapiens mRNA for squalene synthase
11.38	174.6	151.75	456.8	375.62	1411.1	1370.37	627.85	632.22	156.3	44.41	183.9	54.87	Thyroid Peroxidase, Alt. Splice 2 - Also Represents: M25715
270.82	457.5	300.38	600.45	429.43	505.35	38.11	394	466.55	587.55	143.47	313.05	216.87	D50924, class A, 20 probes, 20 in D50924 3807-4083, Human mRNA for KIAA0134 gene, complete cds
888.76	1258.65	641.13	1238.2	937.48	852.05	305.82	933.15	224.08	433.55	113.35	664.5	173.81	U14383, class A, 20 probes, 20 in U14383 958-1372, Human mucin (MUC8) mRNA, partial cds
320.96	1073.05	541.29	525.45	143.75	962.95	955.51	646.95	415.43	342.1	15.13	553.05	245.44	D14838, class A, 20 probes, 20 in D14838 915-1317, Human mRNA for FGF-9, complete cds
640.99	740.15	349.95	229.3	147.93	1026.1	1258.23	832.45	373.85	364.7	30.26	851.45	46.74	U55853, class A, 20 probes, 20 in U55853 1917-2433, Homo sapiens 130 kD Golgi-localized phosphoprotein (GPP130) mRNA, complete cds
136.61	1061.1	193.75	598.2	604.86	747.5	126.15	459.05	479.21	1542.95	1059.74	809.55	16.76	U22897, class A, 20 probes, 20 in U22897 1798-2338, Homo sapiens nuclear domain 10 protein (ndp52) mRNA, complete cds
1708.44	3379.85	1250.66	2453.9	71.42	4060.8	1312.81	4113.7	783.47	3666.65	870.8	3463.65	663.05	D16105, class A, 20 probes, 20 in D16105 2703-2979, Human mRNA for leukocyte tyrosine kinase, complete cds
80.89	119.5	61.52	143.35	99.91	517.15	627.56	105.9	12.59	200.35	96.38	96.4	12.87	X77533, class A, 20 probes, 17 in X77533cds 982-1390: 3 in reverseSequence, 1532-1544, H.sapiens mRNA for activin type II receptor
2254.68	4472.2	991.93	2212.2	2595.51	1193.9	739.78	1792.65	748.19	1995.7	1151.88	4668.25	2741.52	Mucin 6, Gastric (Gb:L07517) - Also Represents: HG881-HT881
189.15	292.3	234.05	718.2	356.95	826.7	1064.05	374.25	90.01	256.7	85.28	495.45	150.54	X17059, class B, 20 probes, 11 in X17059cds 522-840: 9 in reverseSequence, 1331-1418, Human NAT1 gene for arylamine N-acetyltransferase
582.87	731	635.26	890	307.73	919.15	161.15	809.65	211.21	803.1	606.13	1050.8	176.78	U09716, class A, 20 probes, 20 in U09716 1621-2124, Human mannose-specific lectin (MR60) mRNA, complete cds
174.94	180.4	215.95	109.8	80.47	269.5	294.72	134.75	23.83	492.7	463.58	329.95	238.22	L24893, class A, 20 probes, 20 in L24893cds 218-679, Human myelin protein zero (PO) gene

78.35	150.9	50.06	222.4	45.54	394.15	215.88	297.7	47.94	553.85	101.47	318.1	324.7	D49372, class A, 20 probes, 20 in D49372 197-755, Human mRNA for eotaxin, complete cds
1.48	160.7	50.35	63.5	10.32	297.95	120.42	63.45	27.22	74.25	43.91	115.75	41.93	U77949, class A, 20 probes, 20 in U77949 2079-2622, Human Cdc6-related protein (HsCDC6) mRNA, complete cds
7113.14	17235.15	6704.29	19766.9	4370.2	11207	6943.08	12907.55	5123.06	12554.6	5261.58	12161	4975.06	M19483, class A, 20 probes, 18 in M19483cds 1127-1559: 2 in reverseSequence, 2162-2204, Human ATP synthase beta subunit gene
796.7	1230.75	1211.34	657.4	731.43	203.55	29.91	888.05	408.92	930.05	495.89	1029.1	357.23	U53209, class A, 20 probes, 20 in U53209 968-1436, Human transformer-2 alpha (htra-2 alpha) mRNA, complete cds
52.89	110.2	1.7	86.1	27.15	226.35	163.98	344.05	413.59	161.1	78.91	143	60.95	U90910, class A, 20 probes, 20 in U90910 1208-1724, Human clone 23564 mRNA sequence
336.58	2879.15	505.37	1766.05	112.5	2662.55	2177.96	2602.35	318.41	1245.25	396.19	1048.85	632.51	M59916, class A, 20 probes, 20 in M59916 1784-2300, Human acid sphingomyelinase (ASM) mRNA, complete cds
140.93	419.55	124.52	448.6	112.29	454.5	319.19	167.45	78.84	298.9	75.8	655.6	145.38	S79873, class A, 20 probes, 20 in S79873 3394-3969, h-lamp-2=lysosome-associated membrane protein-2 {alternatively spliced} [human, liver, mRNA, 4006 nt]
390.68	644.25	242.89	395.95	101.05	764	137.18	531.15	49.57	377.15	242.18	386.5	125.3	D63484, class A, 20 probes, 20 in D63484 2642-3188, Human mRNA for KIAA0150 gene, partial cds
322.23	975.6	269.55	2048.7	2366.12	1300.6	1315.78	1159.55	817.77	671.65	484.72	1356.5	1203.5	X89101, class A, 20 probes, 17 in X89101exon#3 8-96: 3 in reverseSequence, 183-188, H.sapiens mRNA for Fas (Apo-1, CD95). /gb=X89101 /ntype=RNA
119.43	524.9	100.27	780.5	837.07	306.25	59.18	433.65	12.23	229.25	149.84	800.1	898.03	U31556, class A, 20 probes, 20 in U31556 1445-1679, Human transcription factor E2F-5 mRNA, complete cds
305.89	1312.75	58.05	912.05	214.89	177.9	41.86	569.25	447.67	313.85	165.82	477.4	369.53	L13848, class A, 20 probes, 20 in L13848 3675-4161, Human RNA helicase A mRNA, complete cds
82.66	77.1	87.96	96.65	94.82	152	177.2	58.5	32.53	115.95	109.81	207.15	232.14	Y10510, class A, 20 probes, 20 in Y10510mRNA 13-268, H.sapiens mRNA for CD67S protein. /gb=Y10510 /ntype=RNA
49.43	111.8	34.51	196.5	117.38	178	64.06	129.75	10.96	126.55	48.58	228.75	36.42	U66619, class A, 20 probes, 20 in U66619 1165-1699, Human SWI/SNF complex 60 KDa subunit (BAF60c) mRNA, complete cds
6226.29	12918.3	8823.99	10770.75	6152.75	12854.3	4001.09	14075.3	4273.47	12496.45	10820.93	13026.2	7363.81	L12711, class A, 19 probes, 19 in L12711 1468-2000, Homo sapiens transketolase (tk) mRNA, complete cds
64.98	799.35	258.73	672.4	402.77	642.6	370.1	557.95	613.13	479.8	595.95	516.2	373.49	U28413, class A, 20 probes, 20 in U28413 1440-1926, Human Cockayne syndrome complementation group A CSA protein (CSA) mRNA, complete cds
510.25	2371.2	63.5	3365.6	38.18	1860.45	885.23	2203.05	269.62	2194.3	1222.73	2454.05	1353.47	X79865, class A, 20 probes, 19 in X79865cds 267-411: 1 in reverseSequence, 922, H.sapiens Mrp17 mRNA
771.74	594.55	809.43	303.75	404.96	2041	2830.12	81.95	51.83	130	120.92	56.45	44.48	X04571, class C, 20 probes, 20 in all_X04571 4306-4835, Human mRNA for kidney epidermal growth factor (EGF) precursor
332.91	913.65	268.91	828.25	277.68	519.4	338.42	1465.4	125.16	1769.1	758.58	901	109.18	S38953, class B, 19 probes, 12 in S38953cds 611-820: 7 in reverseSequence, 4446-4527, XA [human, Genomic, 6873 nt]
380.21	549.75	608.04	398	199.26	640.95	762.61	79.25	21.85	409.25	185.05	214.8	30.26	U18259, class A, 20 probes, 20 in U18259 6077-6647, Human clone CIITA-8 MHC class II transactivator CIITA mRNA, complete cds

58.27	142.85	166.1	162.2	66.89	1373.5	1531.03	158.55	81.11	963.3	1316.77	43.45	3.61	M60724, class A, 20 probes, 20 in M60724 1847-2321, Human p70 ribosomal S6 kinase alpha-I mRNA, complete cds
178.97	133.45	177.55	321.35	449.22	115.75	56.36	59.45	52.96	33.25	26.94	270.9	359.78	M21305, class A, 16 probes, 16 in M21305cds 39-119, Human alpha satellite and satellite 3 junction DNA sequence. /gb=M21305 /ntype=DNA /annot=CDS
51.41	412.75	0.21	451.65	270.47	1061.45	863.02	513.5	308.58	550.1	426.53	686.85	322.23	AF003743, class A, 20 probes, 20 in AF003743 28-235, Homo sapiens delayed rectifier potassium channel (KVLQT1-Iso5) mRNA, 5 UTR and partial cds. /gb=AF003743 /ntype=RNA
699.54	1953	124.31	1083.5	342.1	2883.25	3441.14	2499.95	1181.93	1102.6	727.19	2417.65	1780.85	U61836, class A, 20 probes, 20 in U61836mRNA 540-972, Human putative cyclin G1 interacting protein mRNA, partial sequence
933.88	1632.05	1579.18	1001.7	371.51	2957.1	3333.58	2694.55	892.3	4814.7	2209	4262.1	1369.95	L27943, class A, 20 probes, 20 in L27943mRNA 291-867, Homo sapiens cytidine deaminase (CDA) mRNA, complete cds
1274.28	1443.45	956.08	1220.25	982.52	651.8	297.13	813.8	270.96	909.75	286.73	497.65	4.74	D79985, class A, 20 probes, 20 in D79985 3997-4393, Human mRNA for KIAA0163 gene, complete cds
605.28	749.75	425.32	367.55	343.87	375.4	408.99	361.55	417.26	451.25	164.54	544.5	609.81	X52011, class C, 20 probes, 20 in all_X52011 699-1144, H.sapiens MYF6 gene encoding a muscle determination factor
461.6	252.05	88.32	275.1	0.71	428.25	169.63	244.35	163.41	361.9	247.35	737.3	766.36	Protein Kinase Pitslre, Alpha, Alt. Splice 1-Feb - Also Represents: U07705, U04818, M37712, U04819
276.05	683.95	305.4	974.35	648.06	1505.5	925.89	734.3	324.28	250.7	83.58	629.4	189.22	L32164, class A, 20 probes, 20 in L32164 630-1158, Homo sapiens zinc finger protein mRNA, 3 end
34.93	185.65	1.06	174.55	16.9	468.4	311.13	181.05	37.55	411.7	117.24	418.75	156.77	D86062, class A, 19 probes, 19 in D86062 286-862, Human mRNA for KNP-Ib, complete cds
2.76	421.8	107.2	201.2	149.06	1388.7	1828.15	234.55	52.26	121.4	56.57	433.05	70.5	M62843, class A, 20 probes, 20 in M62843 906-1404, Human brain protein recognized by the sera of patients with paraneoplastic sensory neuronopathy mRNA, complete cds
288.43	790.05	86.76	965.65	376.39	769.05	229.46	1257.15	229.17	1649.05	1264.8	815.65	19.73	Spliceosomal Protein Sap 62
515.98	1239.15	504.38	976.65	135.98	1171.55	452.9	1351.35	576.93	2302.85	2116.58	1515.6	109.6	Neural Cell Adhesion Molecule
14	21.95	12.23	47.35	44.19	870.3	691.27	299.45	361.4	274.75	269.62	317.15	317.14	U44105, class A, 20 probes, 20 in U44105 314-574, Human Rab9 expressed pseudogene mRNA, complete cds
224.79	302.7	79.62	412.6	247.49	605.3	25.46	271.9	45.96	675.1	455.09	427.95	101.33	X75346, class B, 20 probes, 15 in X75346cds 788-1157: 5 in reverseSequence, 1205-1305, H.sapiens mRNA for MAP kinase activated protein kinase
54.45	222.55	25.53	760.05	256.47	647.45	145.88	409.75	241.76	652.95	480.48	336	21.07	U62317, class A, 20 probes, 20 in U62317mRNA#6 1634-2156, hypothetical protein 384D8_2 gene extracted from Chromosome 22q13 BAC Clone CIT987SK-384D8 complete sequence., hypothetical protein 384D8_2 gene extracted from Chromosome 22q13 BAC Clone CIT987SK-384D8 complete sequence., hypothetical protein 384D8_2 gene extracted from Chromosome 22q13 BAC Clone CIT987SK-384D8 complete sequence.
106.07	358.8	69.16	788.8	505.58	214.1	72.12	152.8	108.89	342.85	210.08	599.85	640.14	Z68747, class A, 20 probes, 19 in Z68747cds 656-1106: 1 in reverseSequence, 1177, H.sapiens mRNA for imogen 38.

360.62	1835.65	27.08	1673.7	299.95	695.15	346.84	1712.45	1.77	1002.2	388.91	764.75	580.46	X07767, class C, 20 probes, 20 in all_X07767 1948-2516, Human mRNA for cAMP-dependent protein kinase catalytic subunit type alpha (EC 2.7.1.37)
116.81	337.95	110.66	741.45	526.44	2623.75	2671.1	1339.25	659.94	1172.65	1245.43	732.4	185.26	U10685, class B, 20 probes, 12 in U10685exon#2 992-1142: 8 in reverseSequence, 3043-3271, Human MAGE-10 antigen (MAGE10) gene, complete cds
21.64	242.2	47.66	563	487.62	1235.95	698.98	813.7	629.04	270.25	7.42	264.5	250.6	M54968, class C, 20 probes, 20 in all_M54968 5180-5775, Human K-ras oncogene protein mRNA, complete cds
518.1	713.25	249.11	1051.2	1045.25	225.3	73.4	455.7	325.69	671.95	223.23	172.55	34.01	D14533, class A, 20 probes, 20 in D14533 812-1322, Human mRNA for XPAC protein
405.67	326.35	391.67	85.65	13.08	372.25	81.81	102	76.51	212.65	95.53	70.2	5.52	D10922, class A, 20 probes, 20 in D10922 1288-1808, Human mRNA for FMLP-related receptor (HM63)
14.92	881.1	258.24	1549.7	559.89	1028.55	212.77	1215.4	750.24	583.95	415	1083.6	724.93	Z68204, class A, 20 probes, 20 in Z68204cds 43-373, H.sapiens mRNA for succinyl CoA synthetase. /gb=Z68204 /ntype=RNA
202.16	4854.35	845.91	3696.65	405.24	4515.2	3365.26	4830.75	1789.62	5450.65	1664.88	5285.95	786.37	L19605, class A, 20 probes, 20 in L19605 1483-1915, Homo sapiens 56K autoantigen annexin XI gene mRNA, complete cds
239.57	202.5	163.2	232.35	69.65	1157.9	1008.05	539.45	13.65	827.45	739.56	259.35	159.31	U04520, class A, 20 probes, 20 in U04520mRNA 6221-6641, Human type IV collagen a5 chain (COL4A5) gene
442.37	5366.55	205.56	5931	674.44	4223.35	176.28	4858.2	545.6	5496.05	936.14	5348.9	819.96	M64571, class A, 20 probes, 20 in M64571mRNA 4553-4931, Human microtubule-associated protein 4 mRNA, complete cds
1076.29	1061.3	936.92	370.25	28.35	1097.8	141.28	337.05	98.92	292.6	176.35	402.15	122.97	Serine/Threonine Protein Kinase Cdk3 - Also Represents: X66357
39.39	4984.65	471.71	5400.3	2283.81	4409.4	2149.04	5334.65	625.29	5525.85	1167.08	5084.6	1175.07	U57342, class A, 20 probes, 20 in U57342 967-1459, Human myelodysplasia/myeloid leukemia factor 2 (MLF2) mRNA, complete cds
35.78	38.25	10.68	544.75	620.77	711.6	593.4	258.4	307.73	328.3	155.14	54.35	6.15	M31423, class B, 20 probes, 13 in M31423cds 322-640: 7 in reverseSequence, 1200-1320, Human cerebellar-degeneration-related antigen (CDR34) gene, complete cds
33.02	475.75	425.32	830	656.05	1317.45	1203.14	829.1	700.32	1244.9	739.63	452.3	418.47	M13232, class A, 20 probes, 20 in M13232mRNA 1850-2405, Human factor VII serine protease precursor mRNA, complete cds, clone lambda-HVII2463
32.74	6083.3	519.44	5388.05	1718.76	4428.85	60.6	7009.05	705.34	4511	978.49	6758.4	63.22	M64098, class A, 20 probes, 20 in M64098 3873-4305, Human high density lipoprotein binding protein (HBP) mRNA, complete cds
879.99	4245.9	969.16	4035	527.08	1996.15	2376.52	1504.85	1249.67	1606.6	162.92	1898.7	192.47	M91670, class A, 20 probes, 20 in M91670 301-787, Human ubiquitin carrier protein (E2-EPF) mRNA, complete cds
136.47	150.55	159.45	85.35	29.34	144.2	70.14	679.8	907.36	219.95	73.33	112.7	59.4	Immunoglobulin Heavy Chain, Vdjrc Regions (Gb:L23566)
825.97	869.85	680.45	706.7	700.18	821.6	725.77	1103.85	158.6	427.95	197.35	1664.95	1511.58	U79241, class A, 20 probes, 20 in U79241 849-1347, Human clone 23759 mRNA, partial cds.
209.8	6799.5	1055.29	6537.9	699.33	6241.2	3446.86	6361.45	452.76	7050.15	563.07	5565.85	572.54	M75126, class A, 20 probes, 20 in M75126 3159-3537, Human hexokinase 1 (HK1) mRNA, complete cds
629.82	1228.85	441.59	493.4	7.35	908.1	360.34	1599.75	1257.73	1237.4	511.8	907.8	218.21	M34276, class B, 20 probes, 11 in M34276cds 2066-2228: 9 in reverseSequence, 213-354, Human plasminogen gene

1619.27	2125.55	1346.83	2302.9	2045.66	3460.45	1696.28	3865.15	2222.08	1752.35	740.41	1984.5	1834.66	M98045, class A, 20 probes, 20 in M98045 1593-2097, Homo sapiens folylpolyglutamate synthetase mRNA, complete cds
519.02	760.35	572.26	1341.25	813.81	2045.1	491.44	1873.9	759.15	1030.2	881.9	1529.85	638.02	M11726, class B, 20 probes, 10 in M11726exon#1 49-163: 9 in reverseSequence, 1559-2534: 1 not in GB record, Human pancreatic polypeptide gene, complete cds
299.6	393.4	136.75	625.45	315.16	1607.9	1955.86	970.95	407.65	298.25	292.11	905.1	696.36	M27161, class C, 20 probes, 20 in all_M27161 6940-7265, Human MHC class I CD8 alpha-chain (Leu-2/T8) gene, complete cds
159.38	329.15	407.36	358.25	446.54	426.3	323.15	87.15	24.68	886.6	48.65	462.2	252.86	L32140, class A, 20 probes, 20 in L32140 1681-2227, Human afamin mRNA, complete cds
1333.82	2770.1	940.31	3315.05	1082.51	1993.15	2126.06	2240.2	610.09	1399.55	1250.52	4146.65	403.12	D31887, class A, 20 probes, 20 in D31887 4050-4512, Human mRNA for KIAA0062 gene, partial cds
378.51	317.95	427.16	288.05	292.67	105.6	23.9	49.45	39.81	323.3	89.1	135.9	115.82	D13626, class A, 20 probes, 20 in D13626 1857-2373, Human mRNA for KIAA0001 gene, complete cds
10.61	51.3	15.98	139.85	119.57	267.6	289.63	514.7	663.69	121.7	135.48	64.4	2.97	S60415, class A, 20 probes, 20 in S60415 2937-3333, myasthenic syndrome antigen B [human, fetal brain, mRNA, 3477 nt]
35.71	112.65	141.92	194.2	3.54	585	731.57	199.5	144.25	173.05	221.4	208.7	178.62	U65918, class A, 20 probes, 20 in U65918 1248-1820, Human putative RNA binding protein (DAZH) mRNA, complete cds
4.24	238.1	27.29	718.65	342.73	735.2	321.45	857.5	857.44	2034.8	1356.51	1548.25	231.29	X98833, class A, 20 probes, 20 in X98833mRNA 3475-3937, H.sapiens mRNA for zinc finger protein, Hsal1
745.86	2470	142.69	2749.9	134.92	1813.65	1025.66	1936.85	716.94	1824.4	102.53	2001.05	417.83	X54304, class C, 20 probes, 20 in all_X54304 391-878, Human mRNA for myosin regulatory light chain
0.78	270.8	52.89	243.15	112.78	351.1	15.56	369.5	145.24	303.5	30.83	277.5	42.57	D79987, class A, 20 probes, 20 in D79987 6109-6523, Human mRNA for KIAA0165 gene, complete cds
3284.58	5055.25	2227.32	5368.6	2631.85	2665.5	65.2	2129.25	679.03	2203.1	1013.57	2758.75	1161.28	D43950, class A, 20 probes, 20 in D43950 1355-1739, Human mRNA for KIAA0098 gene, partial cds
9320.66	17817.2	4891.34	13098.4	4514.74	11868.8	8873.77	15337.65	5869.62	13037.95	5683.37	12522.65	5290.64	D90209, class A, 20 probes, 20 in D90209 1420-1972, Human mRNA for DNA binding protein TAXREB67
314.1	1664.9	1015.83	1024.35	920.16	895.25	105.15	905.15	373.85	2209.3	755.05	2334.5	525.38	M59807, class A, 20 probes, 20 in M59807mRNA 369-933, Human NK4 mRNA, complete cds
316.36	650.25	224.65	430.05	26.23	473.1	16.55	820	128.27	395.4	225.71	686.05	314.73	U07559, class A, 20 probes, 20 in U07559 1832-2366, Human ISL-1 (Islet-1) mRNA, complete cds
195066.72	197274.45	135485.13	123422.16	148958.06	276929.34	156525.23	252335.5	32989.08	255398.09	62064.74	367157.25	82447.46	J04617, class B, 18 probes, 11 in J04617cds 1069-1364: 7 in reverseSequence, 3823-4030, Human elongation factor EF-1-alpha gene, complete cds
951.06	1425.35	667.44	1622.65	1508.19	2937.95	2501.39	1277	863.8	2473.2	1725.91	1373.4	626.78	U78793, class A, 20 probes, 20 in U78793 6-29, Human folate receptor alpha (hFR) mRNA, partial cds. /gb=U78793 /ntype=RNA
35.85	47.75	44.76	31.05	17.61	133.2	135.91	56.85	6.72	70.4	32.1	68.4	1.84	S77835, class A, 20 probes, 20 in S77835 88-380, IL-2=interleukin-2 [human, brain, mRNA, 418 nt]. /gb=S77835 /ntype=RNA
196.01	678.5	64.91	642.45	192.12	592.85	321.52	788.55	216.73	664.55	100.9	532.9	11.74	S67070, class A, 20 probes, 20 in S67070 55-421, heat shock protein HSP72 homolog [human, thyroid associated ophthalmopathy patient, mRNA Partial, 450 nt]
123.67	187.7	106.07	245.4	219.49	511.4	246.64	152.6	58.55	510.7	42.99	1072.6	1200.38	U36601, class A, 20 probes, 20 in U36601 2620-3166, Human heparan N-deacetylase/N-sulfotransferase-2 mRNA, complete cds

432.4	586.15	632.79	400	182.29	125.45	117.87	193.7	58.27	878.75	259.01	408.85	501.41	X77744, class C, 20 probes, 20 in all_X77744 1467-1750, H.sapiens F11 mRNA
442.3	1364.2	248.76	1854.9	1137.73	1833.15	312.47	872.65	308.09	1790.4	745.71	1033.1	31.96	D14889, class A, 20 probes, 20 in D14889 760-1240, Human mRNA for small GTP-binding protein, S10, complete cds
470.93	1194.25	389.4	1174.25	539.45	3063.6	1973.39	1370.65	369.46	2790.7	385.8	1497.2	384.95	Translocation-Associated Notch (Drosophila) Homolog 1
79.76	1089.6	323.57	1213.85	64.42	928.25	14.35	1275.75	549.35	857.5	39.32	1221.75	50.98	M23161, class A, 20 probes, 20 in M23161 3287-3791, Human transposon-like element mRNA
298.33	655.8	319.75	373.05	13.79	974.9	1071.55	428.85	182.79	1036.4	1041.43	1382.15	888.48	X62466, class C, 20 probes, 20 in all_X62466 25-410, H.sapiens mRNA for CAMPATH-1 (CDw52) antigen
423.91	2737.75	398.31	1723.05	221.96	1159.35	20.29	3000.75	592.34	1409.25	863.73	1089.5	527.5	X79882, class A, 20 probes, 19 in X79882cds 2116-2656: 1 in reverseSequence, 2773, H.sapiens lrp mRNA
5163.5	27650.65	1410.04	20043.35	3066.23	7183.85	4062.12	8033.45	2498.7	13378.25	3650.44	9162.55	2182.91	M58026, class C, 20 probes, 20 in all_M58026 831-1240, Human NB-1 mRNA, complete cds
77.85	1424.65	413.87	1410.45	66.11	1267.7	1504.3	1118.2	99.14	514.7	596.37	322.05	305.82	M93143, class A, 20 probes, 20 in M93143 461-499, Human plasminogen-like protein (PLGL) mRNA, complete cds.
212.91	339.85	191.84	956.25	1067.38	298.95	34.86	410.55	67.1	543.45	193.82	583.85	22.84	X58529, class A, 20 probes, 20 in X58529mRNA 1754-2276, Human rearranged immunoglobulin mRNA for mu heavy chain enhancer and constant region
772.51	2460.1	781.49	1873.5	1582.36	1049.1	15.13	2400.75	2203.56	3093.45	1540.57	2238.8	143.12	L20316, class A, 20 probes, 20 in L20316 1565-2003, Human glucagon receptor mRNA, complete cds
6.65	25.9	11.74	85.45	0.49	248.75	162.14	150.6	65.34	676.85	919.88	407.7	261.91	U50361, class A, 20 probes, 20 in U50361 16-319, Human calcium, calmodulin-dependent protein kinase II delta mRNA, partial cds. /gb=U50361 /ntype=RNA
93.13	623.1	146.8	207.95	23.26	2385.7	3054.7	955	181.02	1001.9	30.55	280.2	40.16	Z84483, class A, 20 probes, 20 in Z84483cds 1166-1676, Human DNA sequence from PAC 46H23, BRCA2 gene region chromosome 13q12-13 contains Klotho, ESTs.
99.35	486.45	288.99	523.35	428.44	1150.8	1139.57	487.6	152.03	444.2	73.68	426.35	115.89	Z26256, class A, 20 probes, 20 in Z26256exon 64-364, H.sapiens isoform 1 gene for L-type calcium channel, exon 1 /gb=Z26256 /ntype=DNA /annot=exon
1210.64	2513.6	919.52	2373.05	194.95	3609.65	2917.59	2050.5	574.31	1458.55	546.24	2225.2	692.82	U87459, class A, 20 probes, 20 in U87459 331-703, Human autoimmunogenic cancer/testis antigen NY-ESO-1 mRNA, complete cds
2013.77	15223.15	4638.13	7278.95	545.67	14075.35	9760.83	22233.6	3521.39	21793.6	5797.99	20931.6	1480.26	S72493, class A, 20 probes, 20 in S72493 617-819, keratin=keratin 16 homolog [human, tracheobronchial epithelial cells, mRNA Partial, 976 nt]
2300.22	2695.4	2701.15	3334.85	3555.83	6357.75	4907.82	15571.95	20405.48	20587.45	15647.78	7016.6	1999.27	AB000115, class A, 20 probes, 20 in AB000115 1469-1973, Human mRNA, complete cds
153.16	232.4	144.25	157.95	67.95	1143.65	1393.35	163.5	115.4	119.8	36.06	132.35	14.64	X52599, class A, 20 probes, 20 in X52599cds 295-649, Human mRNA for beta nerve growth factor
411.11	750.2	481.12	673.1	10.47	400.5	422.43	985.7	332.62	1245.15	1125.5	703.7	353.69	U72209, class A, 20 probes, 20 in U72209 432-990, Human YY1-associated factor 2 (YAF2) mRNA, complete cds
1295.7	1065.2	1002.96	2048.4	2.26	1250.7	605.28	1604.25	474.68	1277.75	758.37	1466.95	1356.73	U35113, class A, 20 probes, 20 in U35113 2039-2555, Human metastasis-associated mta1 mRNA, complete cds
794.15	1368.4	1044.54	1721.25	208.1	2098.25	590.36	874.25	97.23	1404.65	226.06	1127.15	150.4	D79994, class A, 20 probes, 20 in D79994 4227-4749, Human mRNA for KIAA0172 gene, partial cds

2123.16	3177.6	1570.48	2457.85	1990.43	2341.15	267.64	2056.8	666.8	2959.75	509.33	2671.5	1.13	M86667, class A, 20 probes, 20 in M86667 1037-1517, H.sapiens NAP (nucleosome assembly protein) mRNA, complete cds
155.35	176.1	176.92	245.05	105.15	232.4	128.69	585.15	694.87	254.1	184.84	269.7	276.76	Mage-4a Antigen
324.7	1615.4	1160.79	1487.05	95.81	1469.85	478.78	1923.9	424.97	2718.5	1316.35	1450.7	384.1	M16405, class C, 20 probes, 20 in all_M16405 1994-2553, Human m4 muscarinic acetylcholine receptor gene
211.78	84.8	55.86	410	245.51	172.6	54.59	673.45	125.65	397.95	85.21	662.7	360.34	Z69915, class A, 20 probes, 20 in Z69915mRNA 31-244, H.sapiens mRNA (clone ICRFp507L1876).
435.93	523.3	366.14	424.25	273.16	809.7	592.98	417.7	475.32	171.55	107.83	398.3	266.01	U18467, class A, 20 probes, 20 in U18467 1436-1946, Human pregnancy-specific beta 1-glycoprotein 7 (PSG7) mRNA, complete cds
179.39	1147.75	693.18	1815.95	995.68	449.85	28.92	1981.95	1989.44	2517.1	636.11	476	201.67	X96401, class A, 20 probes, 20 in X96401 1673-2186, H.sapiens mRNA for ROX protein
14.07	105.6	53.32	202.55	19.73	202.2	71.28	1016.25	1182.78	902.3	744.58	274	55.72	D38500, class A, 20 probes, 20 in D38500 574-958, Human PMS6 mRNA (yeast mismatch repair gene PMS1 homologue), partial cds (C-terminal region)
39.39	731.55	287.01	512.45	319.68	634.6	157.68	355.7	42.28	1115.35	724.43	1159.65	676.91	U60061, class A, 20 probes, 20 in U60061 905-1325, Human FEZ2 mRNA, partial cds
205.63	270.8	170.84	623.25	494.48	770.2	289.49	420.95	352.07	449.85	462.66	894.55	217.01	Amyloid Beta (A4) Precursor Protein, Alt. Splice 4 - Also Represents: HG3638-HT4808
10.39	230.25	15.91	348.85	211.64	89.3	32.39	472.85	508.62	222.75	167.37	883.05	248.27	L38929, class A, 20 probes, 20 in L38929mRNA 5668-6190, Homo sapiens protein tyrosine phosphatase delta mRNA, complete cds
3.96	1558.75	481.89	1478.7	1307.72	579	261.77	721.05	155.78	621.35	158.04	1142.15	16.76	M22324, class A, 20 probes, 20 in M22324 2954-3416, Human aminopeptidase N/CD13 mRNA encoding aminopeptidase N, complete cds
82.17	101.5	102.25	201.95	233.56	144.95	134	63.95	33.16	316.45	156.62	81.2	6.79	Homo sapiens /REF=M97935 /DEF=transcription factor ISGF-3 mRNA, complete cds /LEN=3970 (_5, _MA, MB, _3 represent transcript regions 5 prime, MiddleA, MiddleB, and 3 prime respectively)
217.44	200.6	234.76	60.6	37.34	81.2	73.68	131.1	54.02	329.95	350.51	349.7	414.36	U84551, class C, 20 probes, 20 not in GB record, Human dystrobrevin (DTN) gene
541.36	478.45	605.07	1102.45	1239.49	1311.9	1039.73	478.7	335.59	324.2	112.01	206.75	137.11	U82979, class A, 20 probes, 20 in U82979 785-1371, Human immunoglobulin-like transcript-3 mRNA, complete cds.
164.83	6145.6	1380.56	6491.45	27.22	4793.45	1108.81	6102.45	1387.41	5962.7	211.85	5560.4	1064.76	S68616, class A, 20 probes, 20 in S68616 4005-4491, Na+/H+ exchanger NHE-1 isoform [human, heart, mRNA, 4516 nt]
119.43	309.15	32.03	546.4	643.33	268.9	117.1	249.45	141.07	243.85	115.75	222.85	23.41	U68162, class A, 20 probes, 20 in U68162mRNA 3059-3585, MPL gene (thrombopoietin receptor) extracted from Human thrombopoietin receptor (MPL) gene
361.69	721.8	202.66	976.05	66.82	637.45	626.43	176.95	184.91	306.8	191.34	1270.8	659.16	J04058, class A, 20 probes, 20 in J04058 767-1217, Human electron transfer flavoprotein alpha-subunit mRNA, complete cds
206.4	396.15	247.7	151.45	138.1	257.65	288.71	166.2	23.76	405.65	276.55	240.5	155.28	M30269, class A, 20 probes, 20 in M30269 4417-4849, Human nidogen mRNA, complete cds
2968.36	1052.2	249.18	2823.7	1215.8	2063.35	1716.93	5290.05	715.52	4012.6	2118.77	3667.7	440.95	U60644, class A, 20 probes, 20 in U60644 1537-1957, Human HU-K4 mRNA, complete cds

478	134.55	57.63	176.05	39.24	948.95	1205.12	229.65	208.24	140.8	27.44	587.3	695.37	D29675, class C, 4 probes, 4 in all_D29675 1092-1149, Human inducible nitric oxide synthase gene, promoter and exon 1 /gb=D29675 /ntype=DNA /annot=exon, Human inducible nitric oxide synthase gene, promoter and exon 1 /gb=D29675 /ntype=DNA /annot=exon
59.61	228.45	108.12	289.4	209.45	379.8	285.25	529.7	16.12	1002.65	1172.03	187	23.76	Y10260, class C, 20 probes, 20 in all_Y10260 1483-2048, H.sapiens EYA1 gene
366.49	337.2	313.67	666.45	602.24	152.7	32.1	290.05	181.66	770.35	427.87	540.75	68.09	J02986, class A, 20 probes, 20 in J02986exon#3 1617-1983, FGF4 gene (transforming protein) extracted from Human transforming protein (hst) gene, complete cds
1207.38	1840.3	557.91	1916.05	2333.95	1925.4	834.1	1691.7	577	2421.1	493.28	1251.35	156.34	Stimulatory Gdp/Gtp Exchange Protein For C-Ki-Ras P21 And Smg P21
740.13	1683.65	181.09	2153.45	1200.74	1933.85	1148.84	1884.45	1100.33	2637.4	1252.29	2198.15	581.03	J05412, class C, 20 probes, 18 in all_J05412 3866-4062: 2 in J05412cds 411-439, Human regenerating protein (reg) gene, complete cds
143.83	268.45	95.11	223.3	133.36	237.5	36.77	248.6	39.03	1150.75	1440.73	283.95	181.94	M67439, class A, 20 probes, 20 in M67439cds 941-1355, Human D5 dopamine receptor (DRD5) gene, complete cds.
773.72	1711.45	551.61	1599.8	1207.17	1097.7	1094.88	1305.6	234.19	2473.65	1242.88	1322.1	46.81	D86479, class A, 20 probes, 20 in D86479 2250-2814, Human mRNA for AEBP1 gene, complete cds
77.99	159.95	175.72	115.7	67.46	297.05	288	340.8	204.07	598.95	794.15	262.3	332.34	X78925, class C, 20 probes, 20 in all_X78925 1966-2447, H.sapiens HZF2 mRNA for zinc finger protein
67.1	2654.85	121.41	1768.5	1109.31	767.5	884.17	1445.9	962.51	1546.4	301.51	1096.45	1135.97	U37022, class A, 20 probes, 20 in U37022mRNA 860-1322, Human cyclin-dependent kinase 4 (CDK4) gene, complete cds
129.26	199.2	93.06	102.55	69.65	1358.1	1112	335.05	57.77	434.6	524.67	393.65	219.84	K02054, class A, 20 probes, 20 in K02054mRNA 238-676, Human gastrin-releasing peptide mRNA, complete cds
349.17	1084.3	182.29	1506.95	479.63	1360.3	611.93	1251.35	522.91	785.7	294.72	1639.5	322.02	D38293, class A, 20 probes, 20 in D38293 2926-3400, Human mRNA for clathrin-like protein, complete cds
161.86	406.7	296.84	84.9	2.69	1331.95	1269.75	445.05	260.57	529.35	281.07	579.65	271.46	X81892, class B, 20 probes, 14 in X81892cds 2760-2994: 6 in reverseSequence, 3126-3204, H.sapiens mRNA for HE6 Tm7 receptor
597.08	1216.9	559.04	533.1	405.6	466.1	417.33	812.1	241.55	1437.55	227.19	818.1	119.22	X76057, class C, 20 probes, 20 in all_X76057 1206-1765, H.sapiens PMI1 mRNA for phosphomannose isomerase
396.47	1473.85	18.46	1108.8	59.68	612.9	555.22	614.6	647.85	1895.25	2575.21	1823.85	5.44	Guanine Nucleotide Exchange Factor 1
641.2	1331.2	389.76	1960.15	407.51	2255.1	278.32	1752.85	267.22	2954.3	1123.31	1679.2	235.18	X93499, class B, 20 probes, 14 in X93499cds 285-591: 6 in reverseSequence, 1241-1451, H.sapiens mRNA for RAB7 protein
594.11	1685.35	997.23	1759.95	370.88	1894.15	634.06	1459.25	42.78	1731.3	184.27	1964.75	81.11	M33308, class A, 20 probes, 20 in M33308 4519-5071, Human vinculin mRNA, complete cds
164.97	184	142.84	198.1	179.18	388.7	264.32	168.75	37.83	171.6	130.96	146.65	35.14	U13021, class A, 19 probes, 19 in U13021 844-1392, Human positive regulator of programmed cell death ICH-1L (Ich-1) mRNA, complete cds
390.04	660.4	42.71	699.95	0.35	378.9	195.16	943.95	173.03	384.3	328.52	588.2	67.18	U33448, class B, 19 probes, 10 in U33448cds 773-1108: 9 in reverseSequence, 1666-1883, Human putative G-protein-coupled receptor (GPR16) gene, complete cds
32.17	102.65	38.54	587.3	726.48	214.3	252.44	232	2.55	329.85	290.27	803.3	1036.05	X59766, class C, 20 probes, 20 in all_X59766 583-1166, H.sapiens mRNA for Zn-alpha2-glycoprotein

315.44	2107.75	105.71	2795.3	946.82	754.6	457.5	1396.25	15.06	880.05	595.88	1027.45	85.91	M16342, class A, 20 probes, 20 in M16342mRNA#2 1287-1581, Human nuclear ribonucleoprotein particle (hnRNP) C protein mRNA, complete cds
66.89	155.4	114.98	145.85	89.45	1094	1395.26	212.6	190.35	278.4	116.39	272.7	15.84	M77235, class A, 20 probes, 20 in M77235 7902-8418, Human cardiac tetrodotoxin-insensitive voltage-dependent sodium channel alpha subunit (HH1) mRNA, complete cds
124.03	137.7	84.15	269.95	86.2	300.55	167.23	50.2	9.9	179.35	115.75	167.1	109.18	X86098, class C, 20 probes, 20 in all_X86098 2367-2704, H.sapiens mRNA for BS69 protein
376.61	421.95	97.65	800.45	1043.19	855.05	424.48	736.8	455.24	1350.35	108.97	443.65	343.87	D14827, class A, 20 probes, 20 in D14827 1564-1966, Human mRNA for Tax helper protein 1, complete cds
91.29	312.1	158.25	367.75	178.26	617.9	329.65	368.65	50.7	1229.25	1057.76	783.35	646.22	Adducin, Alpha Subunit, Alt. Splice 3 - Also Represents: Z68280_cds2, HG651-HT4201
240.77	573.25	515.83	510.05	307.1	1031.85	200.18	1022.55	125.37	696.05	99.35	1259.6	375.05	U43747, class A, 20 probes, 20 in U43747 993-1425, Human frataxin (FRDA) mRNA, complete cds
131.45	165.95	89.17	180.8	164.61	174.9	155	80.25	24.82	689.85	785.38	173.35	125.09	U82279, class A, 20 probes, 20 in U82279 1505-2069, Human immunoglobulin-like transcript 2 mRNA, complete cds
2221.38	4345.35	1829.21	2998.65	663.34	2398.65	1220.25	3188.65	3173.71	1972.45	131.31	840.15	110.95	Tumor Necrosis Factor Receptor 2 Associated Protein Trap3 - Also Represents: U12597
9.05	53.35	26.52	359.2	495.12	88.1	34.93	70.35	65.27	332.55	397.46	177.75	143.75	D14497, class A, 20 probes, 20 in D14497 2222-2726, Human mRNA for proto-oncogene protein, complete cds
98	594.3	344.36	811.65	307.38	374.9	18.1	834.65	703.78	1068.75	10.82	677.45	276.27	L37360, class A, 20 probes, 20 in L37360 146-698, Homo sapiens (clone hEHK1-L) EHK1 receptor tyrosine kinase ligand (EFL-2) mRNA, complete cds
142.76	203.55	257.46	771.4	850.79	578	532.45	253	239	253.45	231.86	792.6	783.05	D13631, class A, 20 probes, 20 in D13631 2795-3373, Human mRNA for KIAA0006 gene, complete cds
8.63	62.1	44.41	144.85	94.82	851.05	411.32	345.25	341.18	652.4	787.86	667.15	365.64	M76559, class A, 20 probes, 20 in M76559 3065-3521, Human neuronal DHP-sensitive, voltage-dependent, calcium channel alpha-2b subunit mRNA, complete cds
222.03	144.35	28.92	282.7	247.63	178.15	5.73	187.3	140.86	88.6	7.35	161.3	23.05	X51953, class A, 20 probes, 19 in X51953exon#1-2 37-64: 1 not in GB record, Human UCP gene for uncoupling protein exons 3 and 4 /gb=X51953 /ntype=DNA /annot=exon
1376.95	1213.7	600.76	1548.45	885.37	1015.8	1077.77	1441.35	636.75	2430.55	1748.46	1843.2	431.76	L35240, class B, 20 probes, 14 in L35240cds 1065-1323: 6 in reverseSequence, 1431-1533, Human enigma gene, complete cds
34.65	1362.65	410.62	1172.35	288.71	1506.75	1135.4	1106.3	911.18	1062	818.26	1378.2	164.9	D63876, class A, 20 probes, 20 in D63876 3171-3717, Human mRNA for KIAA0154 gene, partial cds
1141.69	2097.65	544.4	1111.3	752.64	1151.6	989.81	2698	294.86	2741.2	1658.02	2115.8	327.11	L35475, class B, 20 probes, 14 in L35475cds 642-888: 6 in reverseSequence, 1441-1675, Human olfactory receptor-like gene, complete cds
569.86	623.3	527.64	1150.65	577.21	266	134.92	340.5	10.75	162.35	10.96	328.1	90.09	U28043, class A, 20 probes, 20 in U28043 1989-2499, Human plasma membrane Na+/H+ exchanger isoform 3 (NHE3) mRNA, complete cds
734.54	815.95	848.17	782.8	277.75	2601.55	1909.54	1485.3	383.53	1041.45	167.94	2065.8	170.13	M74542, class A, 20 probes, 20 in M74542 1131-1611, Human aldehyde dehydrogenase type III (ALDHIII) mRNA, complete cds
109.96	134.8	104.09	167.75	136.54	205.2	169	122.9	34.79	111.8	44.26	110.05	10.68	Z83803, class A, 20 probes, 20 in Z83803 7-259, H.sapiens mRNA for axonemal dynein heavy chain (partial, ID hdhc4).

366.92	300.1	179.18	553.6	422.28	798	759.57	579.85	261.84	617.6	144.39	1050.35	418.54	U08191, class A, 20 probes, 20 in U08191 4687-5220, Human R kappa B mRNA, complete cds
132.37	382.15	190.57	658.35	181.8	1123.35	314.59	1474.7	1151.03	855.45	593.19	1031.6	982.45	Z37976, class C, 20 probes, 20 in all_Z37976 6411-6916, H.sapiens mRNA for latent transforming growth factor-beta binding protein (LTBP-2)
414.15	3654.7	214.11	2651.25	2526.42	1948.15	2171.88	1936.2	539.24	2478.85	1741.53	3275.25	978.42	D45248, class A, 20 probes, 20 in D45248 389-773, Human mRNA for proteasome activator hPA28 subunit beta, complete cds
963.36	1390.1	983.02	791.2	179.61	2557.45	2677.04	995.55	565.61	2712.4	1512.78	597.8	39.03	U43431, class A, 20 probes, 20 in U43431 3166-3712, Human DNA topoisomerase III mRNA, complete cds
154.5	231.45	9.12	261.75	157.19	2563.05	2351.48	350.7	318.2	476.4	379.72	315.65	3.04	M29610, class B, 14 probes, 14 in M29610 179-451, Human glycophorin E mRNA, complete cds, Human glycophorin E mRNA, complete cds
10.25	152.85	138.95	196.15	53.1	273.2	236.17	110.95	26.52	220.9	105.5	431.45	241.19	L34657, class A, 20 probes, 20 in L34657mRNA 2757-3219, Homo sapiens platelet/endothelial cell adhesion molecule-1 (PECAM-1) gene
1598.06	97369	25391.92	90150	4331.03	137924.91	122700.71	136512.09	20997.53	152889.75	21901.99	158275.75	5398.41	Homo sapiens /REF=X00351 /DEF=Human mRNA for beta-actin /LEN=1761 (_5, _M, _3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
37.05	587.6	161.64	789.1	823.21	1036.4	1306.45	581.6	616.17	692	757.59	906.8	676.98	X66894, class C, 20 probes, 20 in all_X66894 3881-4417, H.sapiens FACC mRNA from complementation group C (FA(C))
674.37	988.55	518.1	425.7	254.98	707	16.12	1464.25	1438.18	1501.6	1180.3	560.6	762.54	D89377, class C, 4 probes, 4 in all_D89377 1587-2173, Human mRNA for MSX-2, complete cds, Human mRNA for MSX-2, complete cds
1186.88	1187	1045.39	1813.55	2041.06	976.45	482.74	833.65	534.78	1819.55	1705.33	539.4	133.64	L08044, class A, 19 probes, 19 in L08044 8-358, Human intestinal trefoil factor mRNA, complete cds
159.74	995.05	265.24	840.15	117.59	1161.8	571.63	927.8	371.8	870.8	749.82	1646.8	1035.91	X69878, class C, 20 probes, 20 in all_X69878 3909-4372, H.sapiens Flt4 mRNA for transmembrane tyrosine kinase
75.66	449.15	220.26	319.7	123.89	680.05	376.25	371.55	38.54	987.35	975.03	475.15	215.6	U84573, class A, 20 probes, 20 in U84573 2882-3422, Homo sapiens lysyl hydroxylase isoform 2 (PLOD2) mRNA, complete cds
234.9	2112.2	334.89	939.5	955.73	1891.7	1737.5	1763.65	179.25	1756.2	237.73	2063.2	144.25	U66052, class A, 20 probes, 20 in U66052mRNA 7-271, Human clone W2-6 mRNA from chromosome X. /gb=U66052 /ntype=RNA
323.22	2195.25	719.76	2353	99.14	2766.05	1480.33	1573.1	134.77	1277.5	299.67	2339.45	521.63	Cytosolic Acetoacetyl-Coenzyme A Thiolase
36.06	536.75	163.13	1275.5	649.27	485.95	382.05	913	173.52	1226.15	1014.77	711.9	771.88	U35835, class A, 20 probes, 20 in U35835 2404-2859, Human DNA-PK mRNA, partial cds
221.4	56.75	7.57	375.8	282.98	1471.55	1096.23	191.1	180.45	144.65	28.21	183	71.56	M35252, class A, 20 probes, 20 in M35252 602-998, Human CO-029
69.23	192.35	121.55	152.7	1.98	376.3	233.91	219.9	44.97	319.85	94.68	485.85	461.95	U84487, class A, 20 probes, 20 in U84487 2776-3238, Human CX3C chemokine precursor, mRNA, alternatively spliced, complete cds
98.99	180.65	11.1	110.25	10.96	391.1	68.73	584.35	725.28	405.25	185.05	148.7	3.96	M30607, class A, 18 probes, 17 in M30607mRNA 2131-2301: 1 in reverseSequence, 2607, Human zinc finger protein Y-linked (ZFY) mRNA, complete cds
143.19	169.4	179.75	483.95	362.68	271.7	3.25	604.6	304.62	343.95	338.35	265.8	119.78	S81221, class A, 20 probes, 20 in S81221 2246-2546, lanosterol synthase [human, fetal liver, mRNA Partial, 2637 nt]

329.37	686.8	636.96	526.9	552.11	350.15	302.71	177.65	63.43	410.4	382.26	615.55	624.45	M97639, class A, 20 probes, 20 in M97639 3533-4037, Human transmembrane receptor (ror2) mRNA, complete cds
181.37	92.3	29.42	461.05	203.58	1182.45	857.08	245.1	16.83	151.25	160.02	278.9	141.14	U37143, class A, 20 probes, 20 in U37143 1256-1832, Human cytochrome P450 monooxygenase CYP2J2 mRNA, complete cds
395.98	1297.55	645.8	1063.85	578.77	540.2	487.2	722.4	414.08	1843.55	1181.79	977.65	227.9	Dystrophin-Associated Glycoprotein, 50 Kda, Alt. Splice 2 - Also Represents: L34355
111.23	751.85	172.32	836.6	12.59	916.6	566.96	1360.2	631.02	1545.6	748.68	1167.55	953.39	Y11710, class A, 20 probes, 20 in Y11710mRNA 2222-2642, H.sapiens mRNA for extracellular matrix protein collagen type XIV, C-terminus
115.47	644.35	110.38	498.2	413.52	1677.85	1139.64	178.7	53.46	499.7	291.89	125.15	58.05	D86985, class A, 20 probes, 20 in D86985 5502-5946, Human mRNA for KIAA0232 gene, complete cds
582.87	1239	826.18	1195.3	538.96	930.2	314.38	1799.4	1675.7	1840.55	760.49	719	549	X14085, class C, 19 probes, 19 in all_X14085 1251-1422, H.sapiens mRNA for beta-1,4-galactosyltransferase (EC 2.4.1.22)
3824.32	7992.8	4623.77	8503	3713.02	5903.05	1035.84	6714.75	1723.15	5720.15	5286.54	5414.85	732.63	Transformation-Related Protein - Also Represents: D16431
244.87	1077.7	144.11	1161.35	600.69	813	794.93	904.3	240.7	510.55	198.48	1119	44.55	U30999, class A, 20 probes, 20 in U30999 25-379, Human (memc) mRNA, 3 UTR. /gb=U30999 /ntype=RNA
104.3	129.55	150.83	192.15	139.65	107.45	39.95	62.15	56.64	151.05	102.46	214.6	139.3	U17033, class A, 20 probes, 20 in U17033 5092-5578, Human 180 kDa transmembrane PLA2 receptor mRNA, complete cds
71.56	117.25	98.64	206.1	151.6	239.4	43.27	822.2	840.33	128.05	53.81	1062.6	638.09	S69370, class A, 20 probes, 20 in S69370 234-761, PAX3B=transcription factor {alternatively spliced} [human, adult cerebellum, mRNA, 841 nt]
29.56	861.45	218.57	1479.8	937.06	3173.95	2228.02	1701.4	723.23	2263.3	2270.8	1155.75	217.29	U68488, class A, 20 probes, 20 in U68488 1013-1397, Human 5-hydroxytryptamine7 receptor isoform d mRNA, complete cds
92.35	266.2	93.76	272.05	113.49	349.55	221.25	1146	1456.92	1031.15	1085.2	286.65	106.84	S80905, class A, 40 probes, 40 in S80905cds 17-1067, PRB2 (PRB2L CON1+)=Con1 {exon 3} [human, peripheral blood leukocytes, subject R.S., Genomic Mutant, 1179 nt]
2094.95	6033.4	1201.66	5818.3	15.42	4913.7	2208.72	4312.4	1908.2	4091.35	805.61	5994.35	2418.52	Z49254, class A, 20 probes, 16 in Z49254cds 75-435: 4 in reverseSequence, 549-651, H.sapiens L23-related mRNA
180.24	782.75	197.49	1306.65	781	511.5	404.18	1167.25	25.67	1372.25	44.34	845.25	730.37	Utrophin
286.38	550.7	516.33	316.35	17.32	414.85	176.28	1177.1	255.41	631.4	576.29	1191.35	200.46	M26041, class A, 20 probes, 20 in M26041 1273-1405, Human MHC class II DQ alpha mRNA, complete cds
240.27	37.5	34.65	296.95	268.63	318.55	130.18	177.8	168.57	71.9	54.73	54.3	8.34	U69611, class A, 20 probes, 20 in U69611 2905-2985, Human TNF-alpha converting enzyme mRNA, complete cds
268.42	250.3	299.67	581.2	798.04	142.5	139.87	112.95	49	166	11.17	166.8	179.61	X56465, class A, 20 probes, 19 in X56465cds 1607-2069: 1 in reverseSequence, 3400, Human znf6 mRNA for zinc finger transcription factor
238.65	169.2	18.81	1052.65	923.41	195.95	7.71	201.35	59.04	124.35	36.27	342.7	336.58	L40392, class A, 20 probes, 20 in L40392mRNA 1818-2280, Homo sapiens (clone S164) mRNA, 3 end of cds
1718.2	3162.3	1546.02	2542	989.1	3479.3	473.05	3290.35	210.22	2355	1153.57	3533.5	1114.68	U75276, class A, 20 probes, 20 in U75276 2760-3225, Human TFIIB related factor hBRF (HBRF) mRNA, complete cds.
9.4	107.8	12.73	73.35	56.78	683.8	612.5	57.55	11.24	61.4	45.96	355.35	202.3	L78267, class A, 20 probes, 20 in L78267mRNA 2573-3113, Homo sapiens PAR-5 mRNA, probable 5 end

112.08	247.35	241.05	101.5	66.61	284	101.12	147.8	6.65	686.8	246.64	483.65	524.04	AB002365, class A, 20 probes, 20 in AB002365 5053-5617, Human mRNA for KIAA0367 gene, partial cds. /gb=AB002365 /ntype=RNA
1.27	568	348.46	404.8	234.19	123.45	84.92	412.7	363.17	364.15	417.26	536.5	290.76	Retinoblastoma 1
1412.45	4010.1	966.9	3953.35	1524.59	3786.3	326.12	4291.3	858.14	2884.15	1687.65	3949.6	1401.63	U14588, class A, 20 probes, 20 in U14588 3012-3570, Human paxillin mRNA, complete cds
221.04	117.3	24.61	511.25	596.16	2040.85	2558.67	533.85	421.36	126.75	17.75	148.25	74.32	L36861, class A, 20 probes, 20 in L36861exon#4 247-757, Homo sapiens guanylate cyclase activating protein (GCAP) gene exons 1-4, complete cds
1217.78	3778.4	616.46	2302.75	835.16	1657.65	665.6	2378.95	759.22	1605.65	1906.29	2848.4	1203.78	M14016, class A, 20 probes, 20 in M14016mRNA 644-947, Human uroporphyrinogen decarboxylase mRNA, complete cds
98.64	443.05	42.07	630.1	134.63	377.45	363.24	1327.85	1326.74	1386.75	22.27	379.8	340.83	U38480, class A, 20 probes, 20 in U38480 1008-1521, Human retinoid X receptor-gamma mRNA, complete cds
395.77	1892.9	57.84	913.8	763.53	1396.7	707.67	1425.75	117.03	1237.9	1218.63	1501.25	133.01	L19711, class A, 20 probes, 20 in L19711 4952-5414, Human dystroglycan (DAG1) mRNA, complete cds
4.74	190.25	74.6	544.95	713.12	545.65	597.86	148.95	111.37	214.3	39.03	103.9	113.56	Statherin - Also Represents: M32639
287.72	659.7	122.61	851.65	668.57	1071.1	56.57	810.1	555.08	518.65	346.27	534.9	128.98	X68561, class B, 20 probes, 11 in X68561cds 2234-2324: 9 in reverseSequence, 2547-2943, H.sapiens SPR-1 mRNA for GT box binding protein
1371.57	1855.9	1130.81	1601.3	864.79	2079.9	344.36	1829.45	271.46	1359.6	925.18	2267.7	585.77	S79639, class A, 20 probes, 20 in S79639 2588-3068, EXT1=putative tumour suppressor/hereditary multiple exostoses candidate gene [human, placenta, mRNA, 3183 nt]. /gb=S79639 /ntype=RNA
149.77	103.8	36.63	797.5	942.01	145.55	115.47	202.55	237.94	613.15	497.87	58.9	37.76	Y08136, class B, 20 probes, 12 in Y08136cds 292-496: 8 in reverseSequence, 520-820, H.sapiens mRNA for ASM-like phosphodiesterase 3a
357.94	1160.5	739.07	1684.85	1514.69	1188.75	476.52	748.1	746.28	1156.45	195.66	995.5	469.38	U82671, class B, 20 probes, 12 in U82671mRNA#2 1536-1776: 8 in reverseSequence, 106561-106657, HSP1-A gene extracted from Homo sapiens cosmids Qc14E2, Qc12H12, Qc11F9, Qc10G9, LA1733 and Qc17B8 from Xq28, complete sequence.
1233.97	6267.7	598.78	6228.95	654.99	4220.55	1777.88	6226.85	196.79	4877.35	2484.42	5207.7	466.97	U36341, class A, 20 probes, 20 in U36341mRNA#1 3376-3862, SLC6A8 gene (creatine transporter) extracted from Human Xq28 cosmid, creatine transporter (SLC6A8) gene, complete cds, and CDM gene, partial cds
161.15	495.7	186.39	1019.25	1136.96	293.65	76.44	431.55	404.96	219.45	93.13	418.55	183.78	L76937, class A, 20 probes, 20 in L76937mRNA 4642-5098, unnamed protein product gene extracted from Homo sapiens Werner syndrome gene, complete cds
346.91	1151.2	318.48	1871.85	1733.76	474.95	359.28	1484.7	90.09	1175.75	145.45	1369.4	578.13	M91463, class A, 20 probes, 20 in M91463mRNA 2634-3168, Human glucose transporter (GLUT4) gene, complete cds
1552.52	2963.1	1654.35	2580.3	794.79	1557.55	613.13	2339.45	2303.68	1748.05	1953.67	2089.9	571.91	X55079, class C, 20 probes, 4 in X55079mRNA 3257-3366: 16 not in GB record, GAA gene extracted from Human lysosomal alpha-glucosidase gene exon 1
2575.49	5597.3	5.66	1803.75	338.35	2454.85	3318.1	3917.85	955.37	6677.25	2799.51	4266.55	1936.13	Myosin, Light Chain, Alkali, Smooth Muscle (Gb:U02629), Non-Muscle, Alt. Splice 2 - Also Represents: HG2815-HT1357, M22919_rna2
88.11	155.3	106.49	715.4	872	345.6	177.48	160.45	19.3	185.1	162.35	216.1	125.44	S76617, class A, 20 probes, 20 in S76617 2203-2569, blk=protein tyrosine kinase [human, B lymphocytes, mRNA, 2608 nt]

478.78	499.7	383.68	319.7	1.27	416.25	204.71	612.45	767	488.2	332.62	460.2	511.1	Z34975, class C, 20 probes, 20 in all_Z34975 2303-2862, H.sapiens LDLC mRNA
251.16	226.05	262.69	298.1	320.89	762.95	232.28	103.25	7	182.85	27.65	208.95	81.67	M90391, class A, 20 probes, 20 in M90391 1736-2866, Human putative IL-16 protein precursor, mRNA, complete cds
1569.28	5325.25	921.15	5740.2	3543.45	5765.1	3402.17	4511.5	1849.93	6483.55	951.55	7758.25	2102.72	D16217, class A, 20 probes, 20 in D16217 1904-2414, Human mRNA for calpastatin, complete cds
4856.48	6510.85	4384.84	8010.3	2598.19	4081.9	2124.29	4363.2	1224.43	4609.65	868.54	5417.2	174.8	M11353, class A, 20 probes, 20 in M11353 707-1226, Human H3.3 histone class C mRNA, complete cds
1375.46	4876.1	616.17	4983.65	1476.65	2447.3	2209.99	3246.2	130.81	3358.55	2363.93	3362.75	1054.79	D25328, class A, 20 probes, 20 in D25328 2086-2536, Human mRNA for platelet-type phosphofructokinase, complete cds
153.8	249.7	174.09	232.85	128.06	210.05	53.39	239.65	14.35	417.3	413.8	299.55	12.8	M55682, class B, 20 probes, 13 in M55682cds 1132-1467: 7 in reverseSequence, 439-571, Human cartilage matrix protein (CMP) gene
144.11	155.45	160.44	281.25	126.64	311.15	245.01	822.6	512.09	291.35	206.4	192.4	103.38	AB000381, class A, 20 probes, 19 in AB000381exon#2-4 45-395: 1 not in GB record, Human DNA for GPI-anchored molecule-like protein, complete cds
60.17	110.8	72.83	176.55	124.8	355.3	313.81	138.85	45.47	208.1	117.66	127.4	55.72	Guanine Nucleotide-Binding Protein Rap2b, Ras-Oncogene Related
18.1	186.5	175.08	741.75	731.5	441.5	251.45	788.95	797.97	630.2	570.78	813.35	616.38	D13720, class A, 20 probes, 20 in D13720 3754-4319, Human mRNA for LYK, complete cds
2299.87	18848.5	6625.87	19254.15	5246.1	13135.55	8493.41	15514.7	4416.31	10968.1	7669.99	16162.85	3354.16	X77584, class B, 20 probes, 14 in X77584cds 5-215: 6 in reverseSequence, 43-481, H.sapiens mRNA for ATL-derived factor/thiredoxin
63.92	529.65	395.48	372.6	287.93	506.25	402.41	540.6	475.46	863.85	735.18	461.35	344.43	U76369, class A, 20 probes, 20 in U76369 13-325, Human cationic amino acid transporter-2B (ATRC2) mRNA, partial cds. /gb=U76369 /ntype=RNA
71.63	758.5	3.68	1107	762.83	726.65	11.38	653.25	94.82	1164.2	419.17	1216.75	244.73	M29877, class A, 20 probes, 20 in M29877mRNA 1434-1932, Human alpha-L-fucosidase, complete cds
141.63	1078.5	300.24	928.9	140.15	572.3	4.1	727.1	828.59	667.25	319.26	985.6	310	L07493, class A, 20 probes, 20 in L07493 193-631, Homo sapiens replication protein A 14kDa subunit (RPA) mRNA, complete cds
209.94	299.3	241.55	139.65	38.96	289.1	154.01	334.7	300.66	320.85	156.06	480.15	271.32	D87002, class B, 20 probes, 14 in D87002cds#2 4-201: 6 not in GB record, POM121-like 1 gene extracted from Human (lambda) DNA for immunoglobin light chain
33.87	365.4	44.83	200.75	121.55	728.25	419.95	726.5	339.69	75.75	40.66	393.05	276.13	L19593, class A, 20 probes, 18 in L19593mRNA 2261-2747: 2 in reverseSequence, 2765-2771, Homo sapiens interleukin 8 receptor beta (IL8RB) mRNA, complete cds
348.67	5778.95	1251.37	5326.4	970.43	2871.9	528.35	4101.9	1956.85	4422.05	399.44	2685.8	373.07	M60047, class A, 20 probes, 20 in M60047 641-1097, Human heparin binding protein (HBp17) mRNA, complete cds
678.54	2009.4	242.68	2334.15	719.76	2059.65	1390.24	1227.7	416.2	2513.6	628.9	1681.65	475.67	U66618, class A, 20 probes, 20 in U66618 1518-2016, Human SWI/SNF complex 60 KDa subunit (BAF60b) mRNA, complete cds
514.21	2391	417.62	2514.15	142.91	2027.75	479.77	2738.95	967.53	1776.45	1286.02	2183.3	228.11	U06863, class A, 20 probes, 20 in U06863 1416-1938, Human follistatin-related protein precursor mRNA, complete cds
27.72	45.45	34.72	24.45	9.69	59.4	10.61	67.6	9.33	65.45	14.35	263.85	339.48	L41349, class A, 20 probes, 20 in L41349mRNA 3238-3646, Homo sapiens phospholipase C beta 4 (PLCB4) mRNA, complete cds

380.14	399.85	140.5	448.4	391.03	1910.6	1029.26	1254.35	776.47	1262	88.25	1912.35	146.44	M93056, class A, 20 probes, 20 in M93056 859-1273, Human mononcyte/neutrophil elastase inhibitor mRNA sequence
39.74	539	103.38	334.1	99.14	1794.25	1459.26	849.5	456.79	728.75	174.3	292.95	66.4	M89955, class A, 20 probes, 16 in M89955cds 635-1085: 4 in reverseSequence, 1391-1439, Human 5-HT1D-type serotonin receptor gene, complete cds
69.37	3476.05	197.35	3294.35	481.04	1751.95	2083.07	2497.25	1343.29	3426.35	983.8	3260.6	889.68	U49785, class A, 20 probes, 20 in U49785 311-641, Human D-dopachrome tautomerase mRNA, complete cds.
136.83	133.1	121.91	566.55	692.33	133	84.71	84.8	12.3	1019.25	905.87	547.2	598.5	X77748, class C, 20 probes, 20 in all_X77748 2815-3296, H.sapiens mRNA for metabotropic glutamate receptor type 3
193.18	2338.45	354.33	2240.95	309.64	1846.75	845.35	2732.8	111.44	2150.85	96.66	2725.8	483.24	M22960, class A, 20 probes, 20 in M22960mRNA 1352-1760, Human protective protein mRNA, complete cds
87.12	238.9	134.63	558.05	270.89	1195.7	1449	296.45	231.15	266.3	66.19	554.95	609.03	U90902, class A, 20 probes, 20 in U90902 939-1407, Human clone 23612 mRNA sequence
539.38	1093.6	80.47	512.1	369.39	629.85	267.07	296.45	4.03	816.1	521.84	631.9	270.26	U95090, class B, 20 probes, 15 in U95090mRNA 2166-2418: 5 in fullSequence, 36716-36854, Human chromosome 19 cosmid F19541, complete sequence.
32.81	175.05	97.65	489.55	405.24	342.5	153.44	723.6	301.23	581.6	484.93	84.45	85.21	X78669, class C, 20 probes, 20 in all_X78669 1114-1643, H.sapiens ERC-55 mRNA
219.98	573.5	446.89	184.05	155.35	271.05	6.86	713.9	694.52	326.85	151.53	562.35	315.44	U03056, class A, 20 probes, 20 in U03056 1988-2468, Human tumor suppressor (LUCA-1) mRNA, complete cds
152.24	100.85	26.8	146.95	73.33	156.25	3.89	76.15	44.05	69.15	55.37	181.85	144.74	L03427, class A, 20 probes, 20 in L03427 4325-4844, Human zinc finger protein basonuclin mRNA, complete cds
86.69	107.7	133.5	58.2	1.13	812	1039.16	57.25	14.07	183.15	32.74	362.75	127.35	X60382, class A, 20 probes, 20 in X60382mRNA 2717-3149, H.sapiens COL10A1 gene for collagen (alpha-1 type X)
296.7	617.8	9.9	712.75	352.21	483.6	307.45	403.35	200.89	519.15	311.06	596.45	172.6	U58681, class A, 20 probes, 17 in U58681cds 807-1116: 2 in reverseSequence, 1191-1434: 1 not in GB record, Human neurogenic basic-helix-loop-helix protein (NeuroD2) gene, complete cds.
225.35	116	108.61	250.75	190	159.3	144.53	39.7	32.24	271.65	142.2	85.4	74.25	M15353, class A, 20 probes, 20 in M15353mRNA 1277-1769, Homo sapiens cap-binding protein mRNA, complete cds
1.34	113.55	39.67	126.85	74.46	133.7	42.14	150.7	94.89	352.6	120.49	266.65	120.42	M82962, class A, 20 probes, 20 in M82962mRNA 2313-2835, Human N-benzoyl-L-tyrosyl-p-amino-benzoic acid hydrolase alpha subunit (PPH alpha) mRNA, complete cds
557.34	2628.4	388.2	2899.35	1131.16	1977.7	279.17	1874.6	599.63	1261.2	288.22	1472.5	520.15	X66785, class C, 20 probes, 20 in all_X66785 2930-3511, H.sapiens mRNA for transacylase (DBT)
187.17	833.95	200.32	650.9	201.67	560.85	562.08	1152.6	140.57	1016.25	473.13	1753.65	1179.95	X75304, class C, 20 probes, 20 in all_X75304 9705-10252, H.sapiens giantin mRNA
495.75	892.9	974.68	997.3	166.45	364.9	248.62	267.45	221.25	472.15	507.63	897.5	52.47	D29640, class A, 20 probes, 20 in D29640 5767-6325, Human mRNA for KIAA0051 gene, complete cds
62.51	292.4	2.69	110.2	64.63	319.25	100.34	553.85	273.44	418.25	477.08	331.35	61.31	L04510, class A, 20 probes, 20 in L04510 2769-3285, Human nucleotide binding protein mRNA, complete cds
54.87	182.7	62.93	307.7	262.34	427.2	245.22	355.3	165.89	421.35	89.45	376.7	131.95	X53961, class A, 20 probes, 16 in X53961cds 1772-2060: 4 in reverseSequence, 2450-2564, Human mRNA for lactoferrin

70	821.05	380.07	631.4	622.68	791.05	88.03	989.05	939.96	676.05	454.88	396.05	152.24	M96789, class A, 20 probes, 20 in M96789 1026-1548, Homo sapiens connexin 37 (GJA4) mRNA, complete cds
2831.18	5085.35	2168.2	5461.25	2796.11	3152.75	940.95	1956.5	925.04	3691.1	1831.41	3132.25	712.98	D21260, class A, 20 probes, 20 in D21260 5600-6002, Human mRNA for KIAA0034 gene, complete cds
17.04	968	288.78	1187.2	30.69	1377.2	1034.64	165.95	113.49	666.7	482.67	954.85	330.29	L34600, class A, 20 probes, 20 in L34600 1958-2426, Human nuclear-encoded mitochondrial initiation factor 2 mRNA, complete cds
293.52	252.25	298.19	703.9	916.98	412.7	497.1	265.65	165.39	787.2	883.46	180.5	75.66	U73167, class C, 20 probes, 20 in all_U73167 4971-35099, H_LUCA14.2a gene extracted from Human cosmid LUCA14, H_LUCA14.2a gene extracted from Human cosmid LUCA14, H_LUCA14.2a gene extracted from Human cosmid LUCA14, H_LUCA14.2a gene extracted from Human cosmid LUCA14, H_LUCA14.2a gene extracted from Human cosmid LUCA14
108.4	1465.75	694.45	1040.75	717.5	1606.3	535	1281.45	1109.24	1598.6	571.06	761.85	471.43	X78549, class C, 20 probes, 20 in all_X78549 1912-2186, H.sapiens brk mRNA for tyrosine kinase
95.6	211.5	85.84	346.4	338.85	164.8	70.29	186.9	39.6	247.05	101.05	247.8	35.78	X58964, class A, 20 probes, 20 in X58964cds 2500-2914, H.sapiens gene for MHC class II regulatory factor RFX
13.01	516.4	121.62	464.95	536.06	1286.15	32.17	917.95	233.42	338.55	236.53	940.45	459.97	L43575, class A, 20 probes, 20 in L43575mRNA 1021-1106, Homo sapiens (clone 48A8) mRNA
1782.62	3182.55	719.06	2404.05	678.19	1550.4	313.25	3136.3	602.17	3025.5	1359.2	2983.9	448.45	X16316, class B, 20 probes, 11 in X16316cds 2109-2355: 9 in reverseSequence, 2483-2693, Human mRNA for vav oncogene
278.53	3206.6	591	2089.7	969.44	1618.6	425.54	3560.85	966.97	2894.45	2548.06	2360.9	1994.04	M76424, class C, 20 probes, 20 in M76424gene 6565-7070, H.sapiens carbonic anhydrase VII (CA VII) gene
252.01	232.85	96.94	351.95	168.36	602.55	93.83	379.25	359.42	325.25	356.17	233	131.24	L35035, class A, 20 probes, 20 in L35035mRNA 629-1079, Homo sapiens ribose 5-phosphate isomerase (RPI) mRNA
417.62	5411.15	419.39	4491.4	2573.44	3674.15	1536.9	4589.85	878.3	3482.3	3335.14	4725.25	1698.54	X90872, class A, 20 probes, 19 in X90872cds 288-600: 1 in reverseSequence, 799, H.sapiens mRNA for gp25L2 protein
2262.32	1785.7	1292.59	3954.4	278.74	1947.8	1019.79	2289.55	10.54	2180.9	689.43	2147.75	539.17	D90084, class A, 20 probes, 20 in D90084exon#10-8 36-41, Human pyruvate dehydrogenase (EC 1.2.4.1) alpha subunit gene, exons 11-Jan
1345.55	3242.35	1923.54	3457.35	1618.36	3123.1	740.91	1870.1	255.69	2437.65	946.46	3669.5	1430.05	M76482, class A, 20 probes, 20 in M76482 2855-3251, Human 130-kD pemphigus vulgaris antigen mRNA, complete cds
249.04	276	353.98	257.85	119.85	76.9	7.64	65	41.58	204.5	72.97	412.8	240.13	S78234, class A, 20 probes, 20 in S78234 2755-3259, nuc2 homolog [human, fibroblasts, mRNA, 3320 nt]
770.75	945.95	563.07	767.2	708.24	1446.55	916.91	332.05	334.11	1742	545.74	958.65	132.87	D26528, class A, 20 probes, 20 in D26528 1017-1515, Human mRNA for RNA helicase, complete cds
269.62	293	210.86	158.75	39.53	279.25	103.45	273.7	124.73	406.95	17.04	375.8	124.17	X79780, class A, 20 probes, 19 in X79780cds 94-622: 1 in reverseSequence, 646, H.sapiens YPT3 mRNA
767.42	616.5	732.99	103.6	34.22	829.7	953.18	149.8	49.36	1236.9	1354.53	168	79.76	L14269, class A, 20 probes, 20 in L14269 1369-1729, Homo sapiens synaptic vesicle amine transporter (SVAT) mRNA, complete cds
71.84	480.7	117.8	514.15	211.35	631.75	127.92	375	15.13	427	89.94	463.85	221.11	Y07829, class A, 20 probes, 20 in Y07829exon#1 7-283, exon from H.sapiens gene encoding RING finger protein. /gb=Y07829 /ntype=DNA /annot=exon, exon from H.sapiens gene encoding RING finger protein. /gb=Y07829 /ntype=DNA /annot=exon

166.74	628	260.64	594.6	43.13	1077.6	256.54	523.9	435.15	820.1	79.76	1106.8	293.03	AC002486, class A, 20 probes, 20 in AC002486cds 812-1358, Human BAC clone RG367O17 from 7p15-p21, complete sequence. /gb=AC002486 /ntype=DNA /annot=CDS
424.69	347.6	406.59	126.85	114.48	205.45	226.63	581	534.01	279.5	332.76	408.2	516.05	X95425, class A, 20 probes, 16 in X95425cds 2672-3061: 4 in reverseSequence, 3810-3857, H.sapiens mRNA for EHK-1 receptor tyrosine kinase
134.63	482.45	432.82	833.05	164.97	205.75	158.6	172.3	40.31	313.9	169.71	1185.55	899.65	X63717, class C, 20 probes, 20 in all_X63717 1962-2473, H.sapiens mRNA for APO-1 cell surface antigen
774.28	882.45	895.83	613.9	384.1	290.15	186.61	558.2	517.74	1537.85	425.47	1401.2	112.85	L13977, class A, 20 probes, 20 in L13977 1675-2017, Human prolylcarboxypeptidase mRNA, complete cds
52.26	181.35	12.8	121.5	8.34	216.2	62.93	274.55	127.21	254.65	104.86	282.4	116.11	M96684, class A, 20 probes, 20 in M96684 609-867, H.sapiens Pur (pur-alpha) mRNA, complete cds
95.6	1083.35	226.91	1056.75	134.99	596.6	179.89	245	154.86	608.55	381.63	1017.65	183.78	M11058, class A, 20 probes, 20 in M11058mRNA 2351-2879, Human 3-hydroxy-3-methylglutaryl coenzyme A reductase mRNA, complete cds
5.73	530.55	50.13	396.4	298.82	371.75	251.38	202.05	29.49	434.95	158.89	145	145.24	X79536, class C, 20 probes, 20 in all_X79536 1005-1156, H.sapiens mRNA for hnRNPcore protein A1.
446.26	1535.05	312.19	974.25	969.09	871.85	605.64	2225.35	573.82	1742	480.83	2771.65	257.17	U44975, class A, 20 probes, 20 in U44975 791-1301, Human DNA-binding protein CPBP (CPBP) mRNA, partial cds
1870.44	3119.35	1176.7	1281.65	1291.95	2873.85	512.86	3058.55	267.92	1576.6	24.61	2884.5	175.08	X03656, class A, 20 probes, 20 in X03656mRNA 971-1391, G-CSF protein gene extracted from Human gene for granulocyte colony-stimulating factor (G-CSF)
1964.2	4711	121.91	6237.95	1255.61	5849.05	5139.75	6413.9	2292.86	4882.3	1677.96	5919.8	1901.13	D32050, class A, 20 probes, 20 in D32050 2761-3307, Human mRNA for alanyl-tRNA synthetase, complete cds
503.74	2254.55	461.53	2350.85	754.98	1551.45	1268.48	1173.75	246.14	1742.7	1208.87	2544.3	1182	U36759, class A, 20 probes, 20 in U36759 486-1016, Human pre-T cell receptor alpha-type chain precursor, mRNA, complete cds
456.3	880.15	510.88	674.65	425.32	240.15	83.09	399.2	110.17	238.3	110.87	379.2	94.75	U40271, class A, 20 probes, 20 in U40271 3598-3999, Human transmembrane receptor precursor (PTK7) mRNA, complete cds
3155.25	11156.3	2599.18	9966.05	4535.17	6548.15	1065.96	11284.5	2399.5	8544.8	5982.69	7618.45	47.87	X69550, class C, 20 probes, 20 in all_X69550 1266-1801, H.sapiens mRNA for rho GDP-dissociation Inhibitor 1
3215.57	14622.05	4916.72	11812.9	6874.49	9088.2	5173.19	11695.2	5744.54	11051.4	10304.1	9849.9	1307.3	K03515, class A, 20 probes, 20 in K03515mRNA 1392-1938, Human neuroleukin mRNA, complete cds
63.5	114.3	65.62	158.45	19.59	251.15	211.07	262.8	38.75	343	25.88	281.95	149.55	U70370, class A, 20 probes, 20 in U70370 1511-2012, Human hindlimb expressed homeobox protein backfoot (Bft) mRNA, complete cds
11.53	34.75	1.48	76.4	58.83	695.7	893.78	115.75	40.66	114.45	77.99	149.9	184.41	S78569, class A, 20 probes, 20 in S78569 5723-6161, laminin alpha 4 chain [human, fetal lung, mRNA, 6204 nt]
169.07	823.75	355.46	1515.65	1327.45	1256.25	22.98	2271.45	1365.21	2389.2	1646.14	755.6	764.38	J03779, class A, 20 probes, 20 in J03779mRNA 4985-5459, Human common acute lymphoblastic leukemia antigen (CALLA) mRNA, complete cds
60.6	91.55	52.68	157.05	102.88	825.85	1127.76	59.55	38.96	327.75	148.85	62.55	16.48	D86981, class A, 20 probes, 20 in D86981 5936-6410, Human mRNA for KIAA0228 gene, partial cds
80.68	824	574.74	474.95	223.23	244.65	127.35	363.4	308.16	790.35	605.07	250.85	166.67	L40387, class A, 20 probes, 20 in L40387cds 31-433, Homo sapiens thyroid receptor interactor (TRIP14) gene, 3 end of cds. /gb=L40387 /ntype=DNA /annot=CDS
675.99	292.35	32.03	534.75	6.58	465.95	124.95	1076.3	740.91	1832.7	1613.48	2273.2	889.54	X53683, class A, 20 probes, 20 in X53683mRNA 377-638, Human LAG-1 mRNA

387.21	1713.5	901.42	3846.1	2679.51	4030.5	2657.31	2149.45	1130.03	3090.4	16.4	2205.9	688.3	X80818, class C, 20 probes, 20 in all_X80818 3601-3860, H.sapiens mRNA for metabotropic glutamate receptor type 4
246.64	137.7	93.48	465.35	363.81	593.15	396.19	240.75	214.18	126.85	21	98.55	48.3	X06956, class C, 20 probes, 20 in all_X06956 5299-5359, Human HALPHA44 gene for alpha-tubulin, exons 3-Jan
198.48	254.35	173.88	242.35	199.76	266.85	113.07	162.45	54.66	399.9	290.62	129.8	9.9	J03600, class A, 20 probes, 20 in J03600 2058-2364, Human lipoxygenase mRNA, complete cds
117.87	1091.8	258.66	1013.9	176.35	994.85	1146.57	609.3	328.95	1983	929.56	1546.65	295.08	X90999, class B, 20 probes, 15 in X90999cds 390-726: 5 in reverseSequence, 828-942, H.sapiens mRNA for Glyoxalase II
58.97	41.15	29.2	161.25	187.31	533.95	681.58	95.85	13.65	551.55	357.16	244.45	157.76	U93091, class A, 20 probes, 20 in U93091 4410-4806, Human Toll protein homolog mRNA, complete cds and LINE-1 reverse transcriptase homolog, pseudogene.
689.36	802.55	824.27	351.9	42.99	581.45	240.91	653.8	194.03	624	234.19	745.7	225.99	U90912, class A, 20 probes, 20 in U90912 1074-1584, Human clone 23865 mRNA sequence
412.6	1317.5	41.29	1217.1	518.03	1758.6	978.92	1571.1	597.79	1650.3	790.97	886.3	1135.75	Transcription Factor Iia
226.98	363.55	240.77	281.45	312.05	472.75	103.45	472.2	602.45	448.15	547.94	550	710.36	M65134, class A, 20 probes, 20 in M65134mRNA 3588-4102, Human complement component C5 mRNA, 3 end
111.72	306.8	123.18	339.95	166.52	469.15	373.14	652.55	418.96	477.05	467.61	684.4	305.89	U32645, class A, 20 probes, 20 in U32645 3566-4112, Human myeloid elf-1 like factor (MEF) mRNA, complete cds
700.39	1090.1	409.7	598.8	399.37	827.95	901.91	485	256.26	1823.3	1213.54	668.95	495.47	Serum Amyloid A1
211.78	627.95	12.8	354.55	153.37	1539.75	1228.88	657.35	398.6	358.85	230.59	287.9	10.61	L22454, class A, 20 probes, 20 in L22454 2803-2944, Homo sapiens nuclear respiratory factor-1 (NRF-1) mRNA, complete cds
553.45	671.55	459.69	558.95	258.73	493.45	208.53	399	13.44	1071.1	880.49	861.5	451.7	L13698, class A, 20 probes, 20 in L13698 2227-2791, Human gas1 gene, complete cds
2396.38	15779.1	6175.87	11970.8	1258.37	10185.75	7197.71	9841.25	1568.58	10801.8	51.62	13168.7	105.22	X04347, class C, 20 probes, 20 in all_X04347 618-917, Human liver mRNA fragment DNA binding protein UPI homologue (C-terminus)
1160.01	3866	1134.48	4951.95	293.66	4067.95	1062.85	3270.85	1619.63	4493.6	53.88	4406.1	9.76	U24105, class A, 20 probes, 20 in U24105 4121-4355, Human coatomer protein (HEPCOP) mRNA, complete cds
106.14	191.5	98.85	329.75	121.27	171.85	19.16	347.35	244.73	239.05	185.62	264.85	20.44	Acetyl-Coenzyme A Carboxylase
45.4	181.6	52.47	189.4	80.89	498.6	466.55	568.6	212.13	305.3	60.39	283.95	226.91	D38462, class A, 20 probes, 20 in D38462exon 57-549, Human gene for a1 chain of type XIX collagen, exon +3 . /gb=D38462 /ntype=DNA /annot=exon
104.02	393.15	246.71	350	1.27	278.1	19.23	435.55	304.55	219.4	12.45	286.25	70.22	X60787, class A, 20 probes, 20 in X60787mRNA 2434-3000, Human mRNA for transcription factor ILF
712.06	1937.35	869.25	1670.6	336.02	1204.3	1333.6	2530.25	212.06	2780.2	30.69	1824.75	598.14	D87469, class A, 20 probes, 20 in D87469 8413-8899, Human mRNA for KIAA0279 gene, partial cds
49.5	793.55	120.99	1177.5	535.14	609.6	649.41	1335.6	147.22	849.1	446.89	1171.35	540.87	X61072, class A, 20 probes, 20 in X61072mRNA 43-325, Human mRNA for T cell receptor, clone IGRA17.
389.05	488.8	292.6	217.65	86.48	485.6	151.32	632.5	793.66	638.6	680.8	179.95	23.69	X56687, class A, 20 probes, 19 in X56687cds 1698-2154: 1 in reverseSequence, 2348, H.sapiens mRNA for autoantigen NOR-90
6.93	72.85	44.05	192.8	67.18	315.4	369.11	83.6	21.07	356.75	214.47	63.85	38.82	X67337, class A, 20 probes, 20 in X67337 2825-3365, H.sapiens HPBRII-4 mRNA
411.96	8822.85	232.85	10099.1	2118.21	10756.3	6082.39	8571.1	895.34	8552.6	1065.19	8505.95	985.35	L76200, class A, 20 probes, 20 in L76200 268-796, Human guanylate kinase (GUK1) mRNA, complete cds

75.8	86.9	50.2	169.3	107.06	672.95	916.2	365.15	166.95	231.05	95.25	507	459.48	M98833, class A, 20 probes, 20 in M98833 2383-2899, Human ERGB transcription factor (FLI-1 homolog) mRNA, complete cds
87884.47	90329.85	84631.28	281053.56	358100.03	881057.88	939231.75	131998.16	92079.1	264267.19	291518.78	117816.59	16454.66	E. coli /GEN=bioC /DB_XREF=gb:J04423.1 /NOTE=SIF corresponding to nucleotides 4218-4566 of gb:J04423.1 /DEF=E.coli 7,8-diamino-pelargonic acid (bioA), biotin synthetase (bioB), 7-keto-8-amino-pelargonic acid synthetase (bioF), bioC protein, and dethiobiotin synthetase (bioD), complete cds.
39.24	133.3	15.84	126.9	28.14	177.1	73.26	171.85	96.8	174.6	18.95	169.7	52.61	Cd4 Antigen
193.39	431.8	281.43	832	902.55	2837.25	2691.74	543.35	172.18	3564.9	1040.15	1657	1142.97	D11428, class A, 20 probes, 20 in D11428 1253-1757, Human mRNA for PMP-22(PAS-II/SR13/Gas-3) of peripheral myelin, complete cds
90.44	428.6	130.81	483.9	451.56	758.7	343.94	1204.95	267.22	1209.35	465.63	325.75	137.82	U10991, class A, 20 probes, 20 in U10991 6267-6789, Human G2 protein mRNA, partial cds
60.74	116.8	56.71	338.6	369.53	156.8	48.08	340.45	322.65	315.8	329.65	302.2	215.1	Cea Family, Bi-Like Domain
546.95	4504.4	435.58	3157.7	248.76	1720	178.19	2597.75	182.5	2664.5	55.58	1702.1	345.49	U73379, class A, 20 probes, 20 in U73379 193-661, Human cyclin-selective ubiquitin carrier protein mRNA, complete cds
11.03	868.95	89.02	1008.45	70.5	172.5	92.21	903.7	634.42	509.95	193.25	305.95	183.92	M34057, class A, 20 probes, 20 in M34057 4720-5044, Human transforming growth factor-beta 1 binding protein mRNA, complete cds
1564.97	2987.75	455.87	1978.25	1434.65	1435.45	521.92	3185.65	658.95	2782.8	36.06	3138.2	639.65	D21089, class A, 20 probes, 20 in D21089 2957-3515, Human mRNA for XP-C repair complementing protein (p125), complete cds
16.05	125.3	73.26	370.95	480.9	968.5	738.93	462.85	447.25	145.45	54.94	237.2	192.33	M84424, class A, 20 probes, 20 in M84424exon 1073-1217, Human cathepsin E (CTSE) gene
1311.61	10241.4	2512.92	9792.9	1952.89	7239.2	1794.07	8054.2	5635.78	7605.9	2717.69	7004.95	1299.73	X54232, class A, 20 probes, 20 in X54232mRNA 3259-3643, Human mRNA for heparan sulfate proteaglycan (glypican)
322.09	420.1	239.14	593.3	360.06	925.85	1128.05	413.85	303.84	887.65	834.46	633.25	533.09	J05252, class A, 20 probes, 20 in J05252 1611-2178, Human Kex2-like endoprotease mRNA, complete cds.
59.75	129.15	93.69	337.15	303.42	266.45	159.31	86.9	10.75	707.85	797.83	661.15	839.27	M93119, class A, 20 probes, 20 in M93119 2345-2777, Human zinc-finger DNA-binding motifs (IA-1) mRNA, complete cds
41.58	116.9	29.42	130.55	85.49	317.9	188.23	150.15	49	105.05	1.34	379.2	357.65	U68723, class A, 20 probes, 20 in U68723 2126-2576, Human checkpoint suppressor 1 mRNA, complete cds.
701.59	2536.2	944.69	2549.9	1167.15	2149.95	2306.65	2831.8	894.35	1143.75	867.97	3504.3	1154.14	Mucin 3, Intestinal (Gb:M55405)
146.02	1969.65	155.35	2834.05	1564.47	582.2	55.72	555.4	27.29	2458.55	2584.55	516.85	41.79	X05299, class C, 20 probes, 20 in all_X05299 2010-2572, Human mRNA (~95%) for major centromere autoantigen CENP-B
180.24	554.35	498.16	588.25	702.65	612	482.67	338.45	265.8	245.1	74.67	453.55	457.57	M16707, class B, 13 probes, 13 in M16707mRNA 6-357, Human histone H4 gene, complete cds, clone FO108, Human histone H4 gene, complete cds, clone FO108
131.1	358.6	100.55	941.15	1023.96	2585.05	3228.15	366.65	90.3	393.4	14.42	290.05	36.42	M58297, class A, 20 probes, 20 in M58297 2215-2653, Human zinc finger protein 42 (MZF-1) mRNA, complete cds
215.03	252.85	222.1	1270.75	1610.86	386.25	416.56	464	160.94	640.8	782.77	224.5	138.59	M33600, class A, 20 probes, 20 in M33600 581-1109, Human MHC class II HLA-DR-beta-1 (HLA-DRB1) mRNA, complete cds

411.47	905.1	473.34	868.4	331.77	2127.7	91.92	1923.85	674.08	2088.35	131.17	1461.4	97.86	M21984, class A, 20 probes, 20 in M21984 441-951, Human (clone PWHTnT16) skeletal muscle Troponin T mRNA, complete cds
8095.6	30692.1	13021.65	25616.15	5705.43	20792.45	19783.36	24043.75	6398.68	28705.95	9498.35	24611.45	9450.84	M38591, class A, 20 probes, 20 in M38591 120-600, Homo sapiens cellular ligand of annexin II (p11) mRNA, complete cds
238.93	875.85	205.27	834.2	696.92	671.15	669.7	1383.95	812.68	772.3	193.04	478.15	283.34	D32202, class A, 20 probes, 20 in D32202 2017-2263, Human mRNA for alpha 1C adrenergic receptor isoform 2, complete cds
396.97	801.5	376.32	1328.6	798.89	732.95	122.82	1548	1116.52	2240.3	1281.28	556.9	198.41	S76638, class A, 20 probes, 20 in S76638 2553-3003, p50-NF-kappa B homolog [human, peripheral blood T cells, mRNA, 3113 nt]
12274.95	61156.35	4905.84	75670.9	21768.14	59661.5	55457.82	45829.7	13438.56	63967.45	5545.62	98207.91	15303.63	Ribosomal Protein L5
948.09	3430.55	697	3455.5	2477.7	2906.15	2934.71	1752.05	1854.67	1758.3	1258.93	2069.45	87.61	D00763, class A, 20 probes, 20 in D00763 609-1035, Human mRNA for proteasome subunit HC9
78.21	159.95	79.55	180.05	125.09	390.5	290.06	198.25	129.75	298.75	293.94	370.25	236.81	Y08409, class A, 20 probes, 19 in Y08409cds 4-385: 1 in reverseSequence, 431, H.sapiens spot14 gene
264.25	967.1	115.26	526.85	441.45	757.3	132.23	1578.1	257.95	516.65	206.55	1476.8	407.86	U90907, class A, 20 probes, 20 in U90907 1150-1612, Human clone 23907 mRNA sequence
866.77	709.9	538.67	1308.95	139.65	967.15	12.8	572.25	566.18	877.45	586.69	776.25	84.92	U20240, class A, 20 probes, 20 in U20240 448-898, Human C/EBP gamma mRNA, complete cds
75.8	115.7	114.13	439.85	515.83	183	80.89	56.2	0.28	292.35	326.05	140.35	8.56	U03905, class A, 20 probes, 20 in U03905 1438-1858, Human monocyte chemoattractant protein 1 receptor (MCP-1RB) alternatively spliced mRNA, complete cds
32.6	620.25	2.47	384	46.1	426.85	180.52	1313.45	160.02	1003.95	29.49	819.2	324.84	M26692, class A, 19 probes, 19 in M26692exon#1 37-195, Homo sapiens lymphocyte-specific protein tyrosine kinase (LCK) gene, exon 1, and downstream promoter region. /gb=M26692 /ntype=DNA /annot=exon
200.11	460.25	343.3	1031.2	1005.08	647.25	586.55	285.4	262.05	809.6	666.38	346.05	150.97	X81882, class C, 20 probes, 20 in all_X81882 2153-2682, H.sapiens mRNA for for vasopressin activated calcium mobilizing receptor-like protein
277.75	696.85	236.81	506.95	311.9	284.55	265.24	390.95	344.15	315.05	133.86	768.85	543.13	X77548, class C, 20 probes, 20 in all_X77548 2835-3418, H. sapiens cDNA for RFG
388.56	312.8	391.88	282.2	364.58	1177.15	1448.08	229.9	253.29	135.2	119.93	97.1	25.03	L09753, class A, 20 probes, 20 in L09753 1377-1827, Homo sapiens CD30 ligand mRNA, complete cds
115.47	15940.9	3398.36	12673.9	1196.14	4634.9	2703.13	9128.05	1199.89	10012.15	2872.2	5873.65	576.93	X13461, class B, 20 probes, 11 in X13461cds 88-422: 9 in reverseSequence, 1502-1737, H.sapiens intronless calmodulin-like gene (CLP gene) for calmodulin-like protein
12.45	425.05	159.74	473.2	36.63	721.55	39.39	655.65	137.67	1235.4	972.98	708.4	455.52	U09002, class A, 20 probes, 20 in U09002 5527-6082, Human N-methyl-D-aspartate receptor modulatory subunit 2A (hNR2A) mRNA, complete cds
1638.93	9988.15	1396.18	13028.65	3380.18	14882.15	5163.22	25162	18156.95	21966.55	16486.13	11678.8	1656.47	D26600, class A, 20 probes, 20 in D26600 354-822, Human mRNA for proteasome subunit HsN3, complete cds
833.89	4738	354.97	3763.6	778.38	2240.65	345.42	3967.55	524.18	3511.3	663.27	4468.2	317.07	M54915, class A, 20 probes, 20 in M54915 2208-2236, Human h-pim-1 protein (h-pim-1) mRNA, complete cds
1768.4	6020.85	491.51	3942.35	566.89	8013.05	3326.87	6636.25	155.63	3582.55	1856.23	4981.7	1027	Luteinizing Hormone, Beta Subunit
87.12	62.1	41.3	451.3	326.68	310.35	118.72	837.75	456.15	765	793.8	256.5	43.27	X63575, class C, 20 probes, 20 in all_X63575 3903-4466, H.sapiens mRNA for plasma membrane calcium ATPase

271.53	362.9	54.87	540.55	454.6	239.35	82.24	761.8	600.62	362.35	349.1	245.15	139.65	Z33642, class A, 20 probes, 20 in Z33642mRNA 2763-3291, H.sapiens V7 mRNA for leukocyte surface protein
46.88	2612.9	146.37	2618.4	850.08	2012.4	323.71	2141	186.11	2863.9	1342.37	2399.35	130.6	M38690, class A, 20 probes, 20 in M38690 584-1106, Human CD9 antigen mRNA, complete cds
59.04	219.85	171.19	662.25	127.21	319.15	104.86	546.2	642.76	357.2	40.02	611.2	91.5	Dna Excision Repair Protein Ercc6 - Also Represents: L04791
489.18	2245.65	2086.32	633.1	131.95	4019.7	3227.8	2621.8	354.97	1054.55	811.12	612.55	279.24	U22970, class A, 20 probes, 20 in U22970mRNA#1 205-749, 16-Jun gene (interferon-inducible peptide precursor) extracted from Human interferon-inducible peptide (6-16) gene, complete cds
312.26	270.55	343.3	216.05	95.95	375.8	325.13	130.65	111.65	716.1	859.84	97.1	90.93	Z29090, class B, 20 probes, 11 in Z29090cds 2967-3183: 9 in reverseSequence, 3201-3393, H.sapiens mRNA for phosphatidylinositol 3-kinase
1098.49	2475.6	1731.42	2834.6	631.73	1389.25	1520.77	3827.15	191.84	4144.9	1435	1742.25	269.34	X04828, class C, 20 probes, 20 in all_X04828 1116-1702, Human mRNA for G(i) protein alpha-subunit (adenylate cyclase inhibiting GTP-binding protein)
1156.12	2195.45	828.8	1311.15	637.17	2605.2	1243.38	1600.2	814.59	1504	82.45	1107.85	1019.29	M35296, class A, 20 probes, 20 in M35296 3284-3758, Human tyrosine kinase arg gene mRNA
4277.08	2512.4	1598.06	5716.55	3270.16	3948.25	858.64	3062.1	1519.01	1918.3	566.39	3092.8	422.85	M27396, class A, 20 probes, 20 in M27396mRNA 1389-1842, Human asparagine synthetase mRNA, complete cds
57.06	93.65	112.64	226.25	59.47	536.6	591	97	52.61	170.4	203.51	268.65	242.04	Y07596, class B, 20 probes, 11 in Y07596cds 1035-1149: 9 in reverseSequence, 1173-1509, H.sapiens mRNA for GPI8 protein
378.94	870.35	387	873.9	204.92	512.75	543.69	730.7	212.27	440.45	250.1	263.5	25.6	U90651, class A, 20 probes, 20 in U90651 1122-1576, Human embryonic ectoderm development protein homolog (eed) mRNA, partial cds.
574.03	733.8	702.72	927.1	817.7	586.15	463.79	1389.7	772.87	972.7	645.59	366	298.12	Z35491, class A, 20 probes, 20 in Z35491mRNA 797-1253, H.sapiens mRNA for novel glucocorticoid receptor-associated protein
240.91	766.75	668	2066.55	1115.89	2554.65	38.96	1288.15	414.72	2821.75	1502.96	1257.85	546.95	U24056, class A, 20 probes, 20 in U24056 1237-1787, Human inward rectifier K+ channel protein (hirk2) mRNA, complete cds
8.27	40.2	30.69	72.8	47.66	711.8	955.16	92.8	13.86	531.65	490.94	80.25	46.88	L36642, class A, 20 probes, 20 in L36642mRNA 3940-4474, Homo sapiens receptor protein-tyrosine kinase (HEK11) mRNA, complete cds
157.4	1195.2	304.34	1289.1	654.92	1777.75	1041.92	1066.1	197.57	1584.65	1104.15	1153.1	913.72	S67325, class A, 20 probes, 20 in S67325 1328-1712, propionyl CoA carboxylase beta subunit [human, liver, placenta, HL1008, mRNA, 1791 nt]
68834.23	68999.85	66478.57	130956.3	156955.5	671039.75	713247.81	100529.05	76252.2	183476.55	181284.56	77972.65	8147.07	E. coli /GEN=bioD /DB_XREF=gb:J04423.1 /NOTE=SIF corresponding to nucleotides 4980-5256 of gb:J04423.1, not 100% identical /DEF=E.coli 7,8-diamino-pelargonic acid (bioA), biotin synthetase (bioB), 7-keto-8-amino-pelargonic acid synthetase (bioF), bioC protein, and dethiobiotin synthetase (bioD), complete cds.
3099.46	10309.8	5102.91	8470.35	3501.8	9787.55	7015.28	11568.8	2027.42	7847.2	7075.73	10300.8	2021.05	M31520, class A, 16 probes, 16 in M31520mRNA 2-106, Human ribosomal protein S24 mRNA, Human ribosomal protein S24 mRNA, Human ribosomal protein S24 mRNA
75.94	544.35	125.09	675.2	119.5	733.9	814.59	923.9	606.56	1163.1	571.77	1033.75	52.96	U90716, class A, 20 probes, 20 in U90716 1825-2299, Human cell surface protein HCAR mRNA, complete cds

208.46	193.4	260.07	80.7	48.93	360	118.37	140.55	98.78	211.35	18.46	720.15	569.72	M91556, class A, 20 probes, 20 in M91556 4785-5343, Human voltage-gated sodium channel mRNA, complete cds.
21.85	26.55	14.92	124.2	113.99	448.65	191.13	380.15	430.13	220.85	61.31	72.05	31.89	M83941, class A, 20 probes, 20 in M83941 2764-3124, Human receptor tyrosine kinase (HEK) mRNA, complete cds
189.65	234.5	261.21	220.8	131.8	1265.75	1185.61	179	41.15	696.95	723.3	436.6	322.58	U50950, class A, 20 probes, 20 in U50950 1546-2074, Human infant brain unknown product mRNA, complete cds
416.98	2936.6	370.52	1857.85	1054.51	2555.05	309.08	2492.45	1163.26	2855.15	656.69	3032.55	105.43	X54938, class C, 20 probes, 20 in all_X54938 1175-1752, Human mRNA for inositol 1,4,5-triphosphate 3-kinase
245.86	569.65	493.63	834.2	650.96	691.6	616.31	1223.9	557.77	250.4	86.69	1295.95	744.65	D55638, class A, 20 probes, 20 in D55638 227-659, Human B-cell PABL (pseudoautosomal boundary-like sequence) mRNA, clone Bc4. /gb=D55638 /ntype=RNA
9.48	44.25	0.49	115.45	141.92	662.8	905.8	126.3	88.67	75	65.48	464.9	592.98	Clathrin, Light Polypeptide B, Alt. Splice 1 - Also Represents: X81637
371.09	959.8	190.07	353.75	394.35	450.3	446.61	510.05	288	652.45	314.87	398.45	211.35	D49396, class A, 20 probes, 20 in D49396 948-1446, Human mRNA for Apo1_Human (MER5(Aop1-Mouse)-like protein), complete cds
1720.18	1756.45	1243.31	247.2	52.33	708.45	837.14	1195	1156.54	397.8	131.95	971.7	77.36	U65404, class A, 20 probes, 20 in U65404 1140-1530, Human erythroid-specific transcription factor EKLF mRNA, complete cds.
1598.06	1358.6	1643.46	2630.35	3269.45	6420.5	7952.26	2347.75	2167.21	10084.3	4581.63	3366.3	1301.78	D12625, class A, 20 probes, 20 in D12625 2036-2219, Human mRNA for NF1 protein isoform (neurofibromin isoform), complete cds
873.06	617.6	402.06	1875.55	186.89	1779.25	998.36	441.85	384.74	1540.75	801.65	738.95	517.53	AF005887, class A, 20 probes, 20 in AF005887 1969-2413, Homo sapiens ATF family member ATF6 (ATF6) mRNA, complete cds. /gb=AF005887 /ntype=RNA
151.04	657.75	28.07	340.55	233.27	677.25	609.6	466.5	320.6	266.85	43.91	688.5	529.91	U87223, class A, 20 probes, 20 in U87223 4830-5196, Human contactin associated protein (Caspr) mRNA, complete cds
313.96	428.35	333.26	346.95	207.25	170.4	26.16	231.8	111.58	418.3	232.78	653.85	459.97	M36067, class A, 20 probes, 20 in M36067mRNA 2602-3040, Human DNA ligase I mRNA, complete cds
254.56	1982.55	77.71	1504.75	279.66	1711.05	655.84	2139.55	419.67	932.8	730.3	1454.35	156.2	U79260, class A, 20 probes, 20 in U79260 1035-1341, Human clone 23745 mRNA, complete cds.
38.96	171.7	72.97	187.75	91.43	1031.7	786.44	99.35	84.5	338.45	189.01	287.95	358.86	L78833, class A, 20 probes, 20 in L78833exon#24 1038-1476, BRCA1 gene extracted from Human BRCA1, Rho7 and vatI genes, complete cds, and ipf35 gene, partial cds, BRCA1 gene extracted from Human BRCA1, Rho7 and vatI genes, complete cds, and ipf35 gene, partial cds, BRCA1 gene extracted from Human BRCA1, Rho7 and vatI genes, complete cds, and ipf35 gene, partial cds
115.26	3088.6	95.6	2791	541.64	2709.35	2003.87	2777.4	173.24	2241.95	1224.5	2439.35	285.32	U07424, class A, 20 probes, 20 in U07424 1266-1764, Human putative tRNA synthetase-like protein mRNA, complete cds
77.99	126.15	111.37	242.55	164.97	721.05	415.28	404.1	439.25	585.4	425.11	308.35	34.29	M25322, class A, 20 probes, 20 in M25322mRNA 2577-3039, Human granule membrane protein-140 mRNA, complete cds
343.87	701.3	281.99	600.45	103.59	758.3	110.87	963.3	382.26	673.35	242.18	525	402.2	D17516, class A, 20 probes, 20 in D17516 1075-1615, Human mRNA for PACAP receptor, complete cds

225.14	266.95	325.48	298.9	275.91	861.1	1153.72	256.55	31.04	775.95	1067.38	116.4	117.24	M75110, class A, 20 probes, 20 in M75110 814-1312, Human H,K-ATPase beta subunit mRNA, complete cds
277.68	540.25	222.81	398.85	421.51	688.75	586.97	780.9	729.45	1247.7	397.11	752.95	119.57	U78798, class A, 20 probes, 20 in U78798 1699-2203, Human TNF receptor associated factor 6 (TRAF6) mRNA, complete cds.
864.51	619.8	622.11	1871.25	313.18	2085.85	1357.43	695.85	77.99	486.9	245.08	774.15	829.08	L42373, class A, 20 probes, 20 in L42373mRNA 2651-3071, Homo sapiens protein phosphatase 2A B56-alpha mRNA, complete cds
80.96	283.6	6.36	775	665.81	348.1	142.55	280.75	62.3	336.45	58.76	358.65	183.78	M77829, class A, 20 probes, 20 in M77829 737-1269, Human channel-like integral membrane protein (CHIP28) mRNA, complete cds
382.12	2380.95	224.93	2200.95	606.91	3391.7	267.85	2092	742.18	2759.25	93.55	3214.1	127.99	D-Amino-Acid Oxidase
227.9	296.75	120.7	705	707.39	149.65	47.87	351.65	365.64	371.9	140.15	493.55	282.91	L29218, class A, 20 probes, 20 in L29218mRNA 1408-1894, Homo sapiens clk2 mRNA, complete cds
1918.66	70817.9	5081.27	60840.7	10916.32	65704.15	64208.9	64409.95	32689.62	59186.7	35206.98	79947.25	27965.16	D14530, class A, 20 probes, 20 in D14530 55-403, Human homolog of yeast ribosomal protein S28, complete cds
296.14	810.95	11.38	514.55	148	830.2	19.94	408.05	149.55	1165.85	839.12	1035.35	898.94	M15517, class B, 20 probes, 14 in M15517cds#3 155-419: 6 in reverseSequence, 803-923, TTR gene extracted from Human mutant prealbumin gene directly linked to familial amyloidotic polyneuropathy (FAP), TTR gene extracted from Human mutant prealbumin gene directly linked to familial amyloidotic polyneuropathy (FAP)
250.17	584.9	112.71	426.4	351.15	574.6	492.43	375.2	183.14	974.65	362.53	549.5	338.28	U88892, class A, 20 probes, 20 in U88892 31-241, Human tenascin-C mRNA, splice variant TNCfn-ad2, partial cds. /gb=U88892 /ntype=RNA
84.92	1007.65	2.62	912.1	740.62	1211.2	211.99	683.25	730.37	1528.45	255.76	1103.8	463.15	M80783, class A, 20 probes, 20 in M80783 3001-3451, Human B12 protein mRNA, complete cds
204.14	711.3	204.5	939.15	586.4	402	160.65	951.75	90.72	739.5	393.43	889.6	189.36	U59289, class A, 20 probes, 20 in U59289 3350-3824, Human H-cadherin mRNA, complete cds
6.36	106.2	21.92	210.25	152.38	146.55	92.84	95.1	5.66	259.55	285.88	708.1	200.25	D17408, class A, 20 probes, 20 in D17408 906-1481, Human mRNA for calponin, complete cds
41.22	75.4	59.82	104.05	100.62	316.6	293.17	300.75	255.19	218.9	252.15	128.85	14.21	X95095, class C, 20 probes, 20 in all_X95095 487-1058, H.sapiens mRNA for PDGFRalpha protein. /gb=X95095 /ntype=RNA
64.77	144.45	78.7	897	931.83	254.35	50.7	185.4	70.29	200.9	86.27	188.8	26.02	S80335, class A, 20 probes, 20 in S80335 2197-2719, integrin beta 7 subunit [human, mRNA, 2798 nt]
413.37	295.4	8.63	480.45	61.73	2318.65	2335.22	494.75	217.72	349.8	301.65	1564.85	381.2	U43519, class A, 20 probes, 20 in U43519 2976-3474, Human dystrophin-related protein 2 (DRP2) mRNA, complete cds
385.16	479.15	628.69	196.35	85.21	134	132.65	201.8	217.65	550.9	219.34	274.95	370.59	X62534, class C, 20 probes, 20 in all_X62534 726-1137, H.sapiens HMG-2 mRNA
216.66	174	50.77	394.5	98.71	829.85	169.07	531.85	245.86	341.75	118.58	809.8	798.89	U43292, class A, 20 probes, 20 in U43292 706-1252, Human MDS1B (MDS1) mRNA, complete cds
283.83	2794.9	173.24	2832.9	619.57	2085.3	449.3	3089.6	1052.32	2760.25	423.77	2608.7	535.42	Homeotic Protein 7, Notch Group
1129.82	1218.3	1147.78	302.5	12.59	717.85	264.67	1797.95	1013.92	1637.6	856.31	827	407.86	D49410, class A, 20 probes, 20 in D49410exon 48-534, Human gene for interleukin 3 receptor alpha subunit
78.7	1942.25	69.79	2275.4	48.08	1117.05	1426.16	1291.7	492.71	1544.65	144.18	1353.6	662.98	U24266, class A, 20 probes, 20 in U24266 2533-3103, Human pyrroline-5-carboxylate dehydrogenase (P5CDh) mRNA, long form, complete cds
7952.76	10434.15	5553.83	15285.95	10927.7	8222.4	6332.42	8484.25	4509.86	7821.15	4299.42	11274.85	60.6	Transcription Factor Btf3b

14.99	54.9	9.19	63.5	49.5	108	16.97	148.8	36.91	143.95	46.88	43.2	10.75	M11437, class C, 20 probes, 20 in all_M11437 1562-2440, KNG gene (kininogen) extracted from Human kininogen gene, KNG gene (kininogen) extracted from Human kininogen gene
76.58	249.55	78.14	480.2	335.73	385.2	272.8	517.8	619.43	300.3	52.89	719.85	488.68	D38498, class A, 20 probes, 20 in D38498 37-604, Human PMS5 mRNA (yeast mismatch repair gene PMS1 homologue), partial cds (C-terminal region)
211148.66	271313.69	163438.39	200505.5	138715.11	397120.66	260968.58	348283.81	71175.25	337818.06	9527.2	470455.13	119028.72	V00594, class B, 12 probes, 12 in V00594mRNA 15-316, Human mRNA for metallothionein from cadmium-treated cells, Human mRNA for metallothionein from cadmium-treated cells
258.45	395.2	174.8	688.05	49.85	885.2	67.03	239.75	132.02	519.2	634.7	298.65	177.84	X95648, class C, 20 probes, 20 in all_X95648 1063-1658, H.sapiens mRNA for eIF-2B alpha subunit
48.22	137.25	124.66	254.45	26.23	777.35	381.77	113.4	22.63	294.4	7.92	834.35	925.11	M63256, class A, 20 probes, 20 in M63256 1975-2497, Human major Yo paraneoplastic antigen (CDR2) mRNA, 3 end
628.97	2291.1	47.38	2512.45	1470	1295.65	506.92	612.45	456.72	841.15	770.53	1674.55	528.28	L77886, class A, 20 probes, 20 in L77886 5390-5696, Human protein tyrosine phosphatase mRNA, complete cds
151.82	174.2	150.61	529.2	596.09	245.2	184.27	134.2	48.22	252.3	196.15	129.25	37.41	X76498, class A, 20 probes, 16 in X76498exon#3 63-369: 4 in reverseSequence, 2899-3043, H.sapiens gene for uterine bombesin receptor
1026.72	2827.05	508.2	1163.8	289.77	3228.65	1050.41	3687.55	1180.09	5019.2	1275.34	3108.55	1496.17	D38535, class A, 20 probes, 20 in D38535 2583-2973, Human mRNA for PK-120
2.4	100.65	104.3	124.9	71.13	836.65	1141.62	286.75	309.08	129.75	18.46	136.45	26.66	M73482, class A, 20 probes, 20 in M73482mRNA 757-1279, Human neuromedin B receptor (NMB-R) mRNA, complete cds
202.94	3268.55	136.54	3328.6	332.34	2054.65	193.54	2608.95	1353.05	2462.15	564.06	2627.75	55.08	X73478, class C, 20 probes, 20 in all_X73478 2042-2637, H.sapiens hPTPA mRNA
730.72	730.75	747.48	627.7	291.33	543.7	710.36	199.4	85.14	391.55	233.56	367.95	309.78	Z79693, class C, 20 probes, 20 in all_Z79693 2421-3018, H.sapiens mRNA for protein-tyrosine phosphatase NC-PTPCOM1.
194.1	313.35	261.7	348.25	303.28	154.5	114.13	686.95	221.96	272.9	225.99	429.75	289.7	U80669, class A, 20 probes, 20 in U80669 863-1403, Human androgen regulated homeobox protein (NKX3.1) mRNA, complete cds
770.46	1302.3	589.59	632.7	420.87	712.3	621.69	1371.2	280.3	906	224.44	802.6	10.75	AF000545, class A, 20 probes, 20 in AF000545cds 461-983, Homo sapiens putative purinergic receptor P2Y10 gene, complete cds. /gb=AF000545 /ntype=DNA /annot=CDS
2849.71	16484.4	890.67	12843.45	1615.95	13010.25	5696.1	15031.65	458.56	16498.05	2798.52	16915.5	920.37	U15008, class A, 20 probes, 20 in U15008 25-433, Human SnRNP core protein Sm D2 mRNA, complete cds
136.33	285.35	132.3	990	1184.26	1152.1	763.53	357.6	121.34	277.05	11.24	240.4	42.14	U47928, class A, 20 probes, 20 in U47928 2047-2491, Human protein A alternatively spliced form 2 (A-2) mRNA, complete cds
385.87	717.55	640.29	1585.7	898.59	2009.5	315.79	935.5	370.67	1351.65	964.99	1112.05	269.62	X63337, class C, 20 probes, 20 in all_X63337 548-1101, H.sapiens HB2A gene for high sulfur keratin
110.17	108.55	101.75	111.85	52.96	120.4	109.74	226.05	133.29	581.45	718.21	108.6	25.46	X52001, class C, 20 probes, 20 in all_X52001 1770-2281, H.sapiens endothelin 3 mRNA
11.03	140.6	143.68	243.25	96.94	328.35	14.64	455.4	537.26	298.9	280.44	370.5	372.36	Z21707, class A, 20 probes, 16 in Z21707cds 552-912: 4 in reverseSequence, 954-984, H.sapiens p18 mRNA
1146.86	4318.45	961.59	4368.8	1070.14	4745.6	1736.37	2050.55	2056.2	5348.35	2455.43	3404.2	234.05	X59932, class A, 20 probes, 20 in X59932mRNA 1557-2063, Human mRNA for C-SRC-kinase

84.36	388.25	87.47	1574.7	2079.18	2614.1	3371.77	1720.25	1840.1	555.15	268.06	856.15	710.43	J04423 E coli bioC protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)
200.32	360.9	451.56	155.65	124.95	288	101.54	418.65	425.61	945.55	763.32	352.65	261.28	X04707, class A, 20 probes, 18 in X04707cds 815-1343: 2 in reverseSequence, 1649-1673, Human c-erb-A mRNA for thyroid hormone receptor
267.07	883.55	385.59	1238.05	1156.76	2049.05	149.27	645.35	248.83	1731.85	183.49	1217.55	522.06	L05779, class A, 20 probes, 20 in L05779 1535-2069, Human cytosolic epoxide hydrolase mRNA, complete cds
797.12	5241.7	646.3	5119.55	134.56	2982.85	1996.37	3586	101.82	3916.45	1715.09	5189.9	1780.35	K01396, class A, 20 probes, 20 in K01396mRNA 769-1201, Human alpha-1-antitrypsin mRNA, complete cds
168.43	1157	44.12	466.5	207.18	396.3	84	689.95	507.35	426.55	358.72	846.7	451.7	J03133, class A, 20 probes, 20 in J03133 2096-2612, Human transcription factor SP1 mRNA, 3 end
331.21	261.85	45.61	623.1	272.8	806.4	689	880.85	208.38	608	744.58	1536.95	1130.59	K02882, class A, 20 probes, 20 in K02882cds 660-1098, IGHD gene (immunoglobulin delta-chain) extracted from Human germline IgD chain gene, C-region, C-delta-1 domain
27.15	135.1	26.87	161.5	141.14	138.6	136.61	400.45	418.11	246.9	73.82	101.15	49.71	U00921, class B, 20 probes, 15 in U00921exon#3 273-313: 5 not in GB record, Homo sapiens Lst-1 gene, complete cds
22.84	154.8	10.47	307.25	37.26	713.35	332.69	165.3	41.15	194.25	82.1	274.4	54.45	X79568, class B, 20 probes, 10 in X79568cds 1038-1314: 10 in reverseSequence, 1399-1627, H.sapiens BDP1 mRNA for protein-tyrosine-phosphatase
2009.39	4224.95	1626.7	5376.1	1220.61	3776.2	2776.53	4498.25	3162.11	5091.15	560.81	4619.8	1339.83	Male Enhanced Antigen
190.57	293.65	138.66	395.65	90.58	429.05	178.69	547.7	442.22	208.1	162.78	504.55	506.64	M74297, class A, 20 probes, 20 in M74297 1161-1551, Human homeobox 1.4 protein mRNA, complete cds
4.31	381	281.71	189.9	88.81	221.75	245.72	119.3	71.98	417	542.92	511.55	655.7	U84540, class A, 20 probes, 20 in U84540mRNA 1083-1341, Human dystrobrevin isoform DTN-3 (DTN) gene, exon 11B and complete cds. /gb=U84540 /ntype=DNA /annot=mRNA
30318.76	39384.7	26210.89	39089.8	25429.68	31188.3	25715.63	36637.05	8815	26545.5	10124.64	37979.05	10465.96	U06155, class B, 14 probes, 14 in U06155cds 43-495, Human chromosome 1q subtelomeric sequence D1S553. /gb=U06155 /ntype=DNA /annot=CDS, Human chromosome 1q subtelomeric sequence D1S553. /gb=U06155 /ntype=DNA /annot=CDS
223.87	678.55	202.3	242.1	94.47	107	18.95	432.65	67.25	635.25	647.21	228.6	40.73	D50640, class A, 20 probes, 20 in D50640exon 700-1132, Human DNA for phosphodieaterase 3B
1138.02	1110.95	1269.47	2221.4	588.03	1628.25	1409.48	1462.6	603.73	1237.85	992.28	2448	237.02	U32519, class A, 20 probes, 20 in U32519 1294-1708, Human GAP SH3 binding protein mRNA, complete cds
283.2	675.65	336.65	270.75	7	234.85	123.11	204.05	73.04	533.4	254.28	778.55	8.56	S76965, class A, 20 probes, 20 in S76965 1718-2066, protein kinase inhibitor [human, neuroblastoma cell line SH-SY-5Y, mRNA, 2147 nt]
19.52	276.85	102.46	270.7	210.72	525.05	168.36	218.15	219.27	391.6	321.59	560.7	663.69	L17328, class A, 20 probes, 20 in L17328 1400-1868, Human pre-T/NK cell associated protein (3Cl) mRNA, complete cds
880.49	1191.05	237.38	1353.15	614.41	618.8	390.61	1935.6	1271.1	1735.05	1284.18	1789.75	423.2	Tyrosine Kinase, Receptor Axl, Alt. Splice 2
338.7	568.65	263.4	467.8	322.44	579.6	246.21	528.75	54.52	476.6	282.14	594.2	53.74	M92432, class A, 20 probes, 20 in M92432 3098-3566, Homo sapiens retinal guanylyl cyclase (retGC) mRNA, complete cds
16.97	185.2	167.73	406.45	24.82	821.85	262.27	779.3	129.54	1012.85	541.86	710.95	297.9	X89986, class B, 20 probes, 12 in X89986cds 271-387: 8 in reverseSequence, 794-1043, H.sapiens mRNA for NBK apoptotic inducer protein

70.64	222.8	68.02	114	9.19	854.15	635.62	330.5	211.99	822.5	820.39	429.2	260.64	D82346, class A, 20 probes, 20 in D82346 944-1316, Human mRNA for HNSPC, complete cds
135.84	450.8	248.19	1438.05	1307.23	480.85	508.34	1785.65	765.58	743.35	257.46	1317.45	1715.94	X90761, class A, 20 probes, 20 in X90761mRNA 1365-1683, H.sapiens hHa2 gene
534.93	3614.15	1028.77	3711.15	782.84	3259.5	1148.77	2623.4	47.94	1832.95	28.92	1130.4	1047.23	X14850, class C, 20 probes, 20 in all_X14850 1122-1555, Human H2A.X mRNA encoding histone H2A.X
39.88	165.4	1.41	139.5	140.86	225.2	250.74	161	93.2	480.85	534.36	440.85	360.55	X02158, class A, 20 probes, 20 in X02158mRNA 949-1219, Human gene for erythropoietin
81.18	202.8	91.36	215.85	169.78	158.3	142.69	63.35	53.95	319.3	227.12	474.9	38.47	X82125, class C, 20 probes, 20 in all_X82125 2007-2368, H.sapiens HOK-2 mRNA for zinc finger protein
1092.9	4067	50.77	2782.15	2737.14	1434.95	504.24	1911.45	469.02	907.55	50.28	3201.85	603.66	D63486, class A, 20 probes, 20 in D63486 5877-6267, Human mRNA for KIAA0152 gene, complete cds
153.51	337.25	113.63	447.2	301.23	191.9	23.33	316.3	150.33	281.55	47.59	326.2	128.13	X97675, class A, 20 probes, 20 in X97675mRNA 3636-4212, plakophilin 2a gene extracted from H.sapiens mRNA for plakophilin 2a and b
121.76	186.6	165.75	128	118.37	1258.7	1500.62	378.7	207.75	261.75	245.01	216.75	162.71	S85963, class C, 20 probes, 20 in all_S85963 4324-4544, hIRS-1=rat insulin receptor substrate-1 homolog [human, cell line FOCUS, Genomic, 6152 nt]
113.7	175.2	177.91	239.05	109.25	267.9	217.93	512.55	467.89	870.5	949.08	285.75	172.75	U07223, class A, 20 probes, 20 in U07223mRNA 1930-2410, Human beta2-chimaerin mRNA, complete cds
1448.15	21648.05	477.37	17420.15	3893.26	18659.85	641.98	28087.7	4102.63	14112.05	10163.17	23472.25	1121.68	Z48199, class A, 20 probes, 20 in Z48199exon#4 1510-2026, H.sapiens syndecan-1 gene (exons 2-5)
143.12	206.85	72.76	769.7	795.21	214.45	178.4	389.25	323.78	269.6	248.62	161.8	12.16	M62958, class A, 20 probes, 20 in M62958 2486-2942, Human retinal degradation slow (RDS) mRNA
115.89	880.05	277.68	580.4	72.97	244	133.93	872.9	785.17	1002	247.2	920.6	254.56	U71087, class A, 20 probes, 20 in U71087 1092-1126, Human MAP kinase kinase MEK5b mRNA, complete cds.
158.89	731.9	280.16	339.7	339.98	320.25	7.42	505.25	437.63	392.1	265.73	976.1	292.88	U64520, class A, 20 probes, 20 in U64520 308-650, Human synaptobrevin-3 mRNA, complete cds
60.25	218	10.32	252.5	173.81	773.2	1021.91	705.25	789.06	252.7	182.86	521.15	272.59	U19252, class A, 20 probes, 20 in U19252 4495-5045, Human putative transmembrane protein mRNA, complete cds
423.49	1120.75	279.52	978.4	1145.65	1136.7	660.01	718.05	411.47	1644.65	80.4	1460.65	1034.71	D83783, class A, 20 probes, 20 in D83783 6249-6579, Human mRNA for KIAA0192 gene, partial cds
89.73	153.3	74.1	101.4	55.01	266.15	238.22	152.3	23.19	341.65	357.87	218.1	146.09	Z11933, class C, 20 probes, 20 in all_Z11933 1730-1942, H.sapiens mRNA for N-Oct 3, N-Oct5a, and N-Oct 5b proteins
774.14	817.3	174.66	1296.5	139.58	1487.55	384.88	1746.7	820.39	1376.6	42.99	1131.75	146.87	Dna-Binding Protein Ap-2, Alt. Splice 3
1689.99	3767.55	1376.24	5845.25	382.19	3305.2	681.65	4647.55	800.8	4960.15	1564.62	4502.7	251.73	X16135, class C, 20 probes, 20 in all_X16135 1552-2003, Human mRNA for novel heterogeneous nuclear RNP protein, L protein
92.98	433.85	172.32	185.9	81.32	770.1	0.57	383.15	301.16	649.5	347.19	1322.8	661.14	X17093, class C, 20 probes, 20 in all_X17093 3834-4023, Human HLA-F gene for human leukocyte antigen F
79.13	240.45	79.69	257.3	113	291.3	75.52	528.95	111.51	253.3	121.91	259.7	1.7	Z48475, class B, 20 probes, 14 in Z48475cds 1531-1795: 6 in reverseSequence, 1918-2128, H.sapiens GCKR mRNA for glucokinase regulator
16.33	258	160.23	209.9	188.23	1083.35	1319.67	117.6	15.98	176.1	83.86	188	50.91	M13560, class A, 20 probes, 18 in M13560exon 3-562: 2 not in GB record, Human Ia-associated invariant gamma-chain gene
254.91	230.65	274.71	214.45	252.37	329.95	241.19	75.75	9.26	89.5	64.21	432.4	149.2	X77922, class C, 20 probes, 20 in all_X77922 1492-2000, H.sapiens GD3 synthase mRNA

268.13	435.85	243.03	938.15	365.79	185.6	27.44	296.4	87.54	305.8	142.84	1370.05	505.09	U26398, class A, 20 probes, 20 in U26398 2628-3168, Human inositol polyphosphate 4-phosphatase mRNA, complete cds
316.01	1718.8	227.69	1732.8	376.75	1304.2	745.29	1404.95	160.16	1711.9	186.11	1419.75	135.41	U69645, class A, 20 probes, 20 in U69645 551-1037, Human zinc finger protein mRNA, complete cds
7182.65	7511.95	3075.28	9690.85	8542.2	4100.05	1667.29	9929.35	6102.83	6218.8	5974.91	7102.35	1755.82	Epican, Alt. Splice 11 - Also Represents: HG2981-HT3126, HG2981-HT3931, HG2981-HT3125, HG2981-HT3932, HG2981-HT3934, L05424_cds2
111.02	1182.25	207.11	1732.2	745.43	1598.45	162.85	1958.3	1267.7	1811.75	870.52	2252.6	727.47	L35545, class A, 20 probes, 20 in L35545mRNA 689-1223, Homo sapiens endothelial cell protein C/APC receptor (EPCR) mRNA, complete cds
2507.12	3919.85	2552.87	3387.85	653.86	4474.6	1239.28	4986.6	518.59	4029.2	1119.21	3832.7	585.34	U86529, class A, 20 probes, 20 in U86529 584-1130, Human glutathione transferase Zeta 1 (GSTZ1) mRNA, complete cds. /gb=U86529 /ntype=RNA
69.58	261.7	25.6	324.95	278.39	73.35	59.04	461.75	13.22	154.05	83.37	211.85	95.81	U57796, class A, 20 probes, 20 in U57796 3322-3784, Human zinc finger protein (LD5-1) mRNA, complete cds
44.41	128.6	17.11	195.5	152.59	148.65	124.95	112.9	21.5	176.1	21.5	157.25	45.61	X99897, class B, 20 probes, 10 in X99897cds 6561-6760: 10 in reverseSequence, 7057-7274, H.sapiens mRNA for P/Q-type calcium channel alpha1 subunit
420.3	247.1	61.24	1861.2	2424.39	426.75	353.62	350.05	166.38	1727.4	2093.88	395.7	325.98	Mucin 1, Epithelial, Alt. Splice 9 - Also Represents: J05582, J05581
167.58	59.9	66.89	86.55	53.67	646.75	799.81	569.15	660.08	1342.5	1812.31	79.15	24.82	M86873, class A, 20 probes, 20 in M86873mRNA 155-367, H.sapiens type A plasminogen related gene
1135.54	939.35	659.66	2065	944.27	2925.55	2195.78	2794.85	375.69	2992.9	1444.34	2124.35	960.89	M60165, class A, 20 probes, 20 in M60165mRNA 571-1069, HLA-DQB1 gene extracted from Human guanine nucleotide-binding regulatory protein (Go-alpha) gene
216.02	639.7	118.79	321.9	262.48	695.25	363.38	950.55	968.1	1151.55	638.45	784.4	772.3	U28049, class A, 20 probes, 20 in U28049 1684-2221, Human TBX2 (TXB2) mRNA, complete cds
110.73	541.3	372.22	698.8	354.4	644.75	204.28	613.35	451.49	846.65	381.77	822.9	371.94	D87075, class A, 20 probes, 20 in D87075 5013-5469, Human mRNA for KIAA0238 gene, partial cds
1014.77	1574.2	1204.49	2292.95	723.58	1676.4	1247.05	972.4	280.3	1387.7	1100.97	1886.25	352.35	M32886, class A, 20 probes, 20 in M32886 351-843, Human sorcin CP-22 mRNA, complete cds
19666.55	79121.95	11982.99	83750.85	20829.04	94382.1	31502.31	86541	30314.09	88814.4	34911.98	110204.7	11096.06	M14199, class A, 20 probes, 20 in M14199 2-381, Human laminin receptor (2H5 epitope) mRNA, 5 end
278.67	299.9	374.34	83.55	61.45	179.25	42.92	378.65	190.14	46.55	25.39	220.9	143.68	M74525, class A, 20 probes, 20 in M74525 2002-2536, Human HHR6B (yeast RAD 6 homologue) mRNA, complete cds
496.95	1284.75	488.12	728.15	389.83	501.4	407.43	1093.3	599.06	1133	129.82	731.1	415.5	D10656, class A, 20 probes, 20 in D10656 988-1528, Human mRNA for CRK-II, complete cds
62.23	1273.25	18.6	1870.35	663.34	1957.25	323.36	970.7	748.83	2361.1	748.26	1514.55	831.35	X79483, class C, 20 probes, 20 in all_X79483 1063-1556, H.sapiens ERK6 mRNA for extracellular signal regulated kinase
195.44	424.95	342.31	492.4	195.87	633.6	308.86	190.4	64.63	295.05	278.53	188.25	17.89	D26018, class A, 20 probes, 20 in D26018 2865-3381, Human mRNA for KIAA0039 gene, partial cds
2861.17	12478.2	1181.01	5546.35	217.15	1719.75	837.29	4665	558.05	2197.9	2316.34	2839.5	476.73	M26311, class A, 19 probes, 19 in M26311 27-504, Human cystic fibrosis antigen mRNA, complete cds.
145.31	3533.6	210.43	2874.25	46.88	3322.8	1415.49	4295.55	343.44	2245	1531.59	3953.05	42.36	Y09616, class C, 20 probes, 20 in all_Y09616 1443-1948, H.sapiens mRNA for putative carboxylesterase
2125.42	3085.8	1948.79	4343.35	2149.11	3260.5	81.32	2741.8	404.89	2561.15	521.07	3706.7	556.92	Z29083, class C, 20 probes, 20 in all_Z29083 1644-2023, H.sapiens 5T4 gene for 5T4 Oncofetal antigen
351.86	409.65	386.72	393.35	174.87	755.65	668.15	97	65.76	512.15	79.97	96	56.14	M77140, class A, 20 probes, 20 in M77140 91-409, H.sapiens pro-galanin mRNA, 3 end

731.5	660.35	844.64	777.6	800.87	908.5	804.55	251.55	110.66	1325.25	525.17	441.3	458.77	U23850, class A, 20 probes, 20 in U23850 8177-8697, Human inositol 1,4,5 trisphosphate receptor type 1 mRNA, partial cds
991.79	1483.25	185.9	1553.1	122.9	826.45	809.85	1417.05	772.94	689.15	637.17	488.4	191.2	M31211, class A, 20 probes, 20 in M31211mRNA 150-714, Human myosin light chain 1 slow a (MLC1sa) mRNA, complete cds
210.58	2856.8	210.29	1597.8	1518.58	2809.7	1142.4	2585.3	258.24	2217.2	106.91	1784.55	503.81	U31903, class A, 20 probes, 20 in U31903 2052-2510, Human CREB-RP (creb-rp) mRNA, complete cds
933.95	2515.45	637.6	1978.25	187.45	1797.5	903.26	2728.05	774.78	3031.45	651.32	2300.05	562.5	Transcription Factor Mef2, Alt. Splice 2 - Also Represents: HG4668-HT5082, L16794
342.88	958.95	329.44	1149	643.61	451.7	374.34	1079.35	211.21	697	106.21	472.95	410.76	M21154, class A, 20 probes, 20 in M21154mRNA 1234-1756, Human S-adenosylmethionine decarboxylase mRNA, complete cds
1368.25	1881	604.01	2117.6	35.07	1396.15	1807.72	2637.6	34.37	1523.7	530.9	2010.95	36.7	X61123, class A, 20 probes, 20 in X61123mRNA 1212-1608, Human BTG1 mRNA
84.22	111.2	17.68	429.65	237.8	211.8	112.43	184.35	153.94	354.8	379.57	95.95	22.56	D88795, class A, 20 probes, 20 in D88795 13-379, Homo sapiens mRNA for cadherin, partial cds. /gb=D88795 /ntype=RNA
29529.28	135876.66	9323.41	114851.09	28614.77	120345.25	85862.07	193909.44	42916.93	137133.84	51009.06	188777.66	14715.1	V01516, class B, 20 probes, 11 in V01516cds 713-1044: 9 in reverseSequence, 1070-1293, Human messenger fragment encoding cytoskeletal keratin (type II). mRNA from cultured epidermal cells from human foreskin
327.04	328	266.72	109.9	42	346.2	80.47	173.3	97.44	1209.1	1429.63	563.7	514.92	X52282, class A, 20 probes, 20 in X52282cds 1092-1597, Human mRNA for atrial natriuretic peptide clearance receptor (ANP-C receptor)
358.57	488.3	182.57	915	93.34	758.45	33.3	280.75	4.03	1455.65	1039.38	535.3	470.23	L33881, class A, 20 probes, 20 in L33881 1822-2206, Human protein kinase C iota isoform, complete cds
254.84	473.7	36.91	374.1	299.39	436.2	265.59	439.5	163.62	768.85	506.22	1033.05	330.86	U18062, class A, 20 probes, 20 in U18062 1678-2152, Human TFIID subunit TAFII55 (TAFII55) mRNA, complete cds
184.06	2572.2	3.54	2132.1	30.69	2577.6	200.25	3980.2	2545.73	4993.05	2789.32	2530.1	150.9	M96739, class A, 20 probes, 20 in M96739 1964-2510, Human NSCL-1 mRNA sequence
434.02	603.15	493.35	417	6.93	111.1	102.39	253.65	216.59	87.3	52.18	121.95	2.9	M21064, class C, 20 probes, 20 in all_M21064 1360-1426, Human migration inhibitory factor-related protein 14 (MRP14) gene, complete cds
104.51	219	164.05	190.4	79.62	113.65	17.47	171	91.64	1191.55	1528.13	106.75	46.46	J03060, class A, 20 probes, 20 in J03060exon#11 168-666, Human glucocerebrosidase (GCB) gene
118.86	240.35	221.68	757.05	980.26	795.75	969.37	579.75	289.28	273.3	102.25	846.65	204.71	U22961, class A, 20 probes, 20 in U22961 2627-3194, Human mRNA clone with similarity to L-glycerol-3-phosphate:NAD oxidoreductase and albumin gene sequences
11.03	28.5	17.82	132.1	100.69	106.7	15.56	46.95	21.28	76.2	50.2	127.1	134.63	X83228, class C, 20 probes, 20 in all_X83228 3359-3561, H.sapiens mRNA for LI-cadherin
900.01	1023.4	63.78	1709.65	1348.38	1812.25	2117.01	1667.1	1209.86	668.05	242.47	1106.7	153.3	AF009674, class A, 20 probes, 20 in AF009674 2870-3308, Homo sapiens axin (AXIN) mRNA, partial cds.
264.53	1838.75	418.25	1913.7	619.99	1122.3	1181.15	847.25	198.06	2222.95	1917.74	1939.7	1059.1	Ras Inhibitor Inf
178.76	1200.7	376.32	688.55	340.05	1035.3	766.22	608.4	151.75	863.4	325.41	826.45	115.89	U33429, class A, 20 probes, 20 in U33429 2481-2994, human K+ channel beta 2 subunit mRNA, complete cds
662.77	700.35	613.27	485.1	327.67	460.45	125.23	265.95	50.13	414.7	169	303.2	83.16	D85418, class A, 20 probes, 20 in D85418 875-1403, Human mRNA for phosphatidylinositol-glycan-class C (PIG-C), complete cds

381.48	1141.95	73.33	1414.3	728.74	1185.7	689.43	875.1	159.95	621.85	407.51	1358.35	182.93	Z24680, class A, 20 probes, 20 in Z24680mRNA 3558-4044, H.sapiens garp gene mRNA, complete CDS
16.62	934.9	206.19	1004.55	138.81	1049.3	549.56	1561.85	15.63	2280.6	880.63	1662.75	223.09	X97302, class A, 20 probes, 20 in X97302mRNA 40-235, H.sapiens mRNA for Ptg-1 protein. /gb=X97302 /ntype=RNA
56.21	139.15	48.01	175.6	83.72	242.05	152.95	511.6	624.8	257.7	81.6	249.8	96.73	AF014958, class A, 20 probes, 20 in AF014958 1175-1619, Homo sapiens chemokine receptor X (CKRX) mRNA, complete cds. /gb=AF014958 /ntype=RNA
221.61	353.85	211.92	657.95	715.95	959.65	1049.7	313.35	201.31	424.2	211.99	454.45	413.73	X86570, class A, 20 probes, 20 in X86570 1208-1532, H.sapiens mRNA for acidic hair keratin 1
168.08	351.75	107.13	961.9	769.76	958.9	748.54	650.25	324.35	285.5	283.97	1171.9	830.14	X76942, class A, 20 probes, 16 in X76942cds 24-420: 4 in reverseSequence, 487-527, H.sapiens mRNA for 72.1 protein.
127.14	828.55	141.07	538.7	513.22	283.7	58.41	174.1	80.04	456.7	196.29	1037.35	230.16	U68019, class A, 20 probes, 20 in U68019 1774-2218, Human mad protein homolog (hMAD-3) mRNA, complete cds
189.58	254.95	172.75	191.15	146.44	201.15	32.6	214.8	141.42	328.5	157.4	285.2	95.18	V01510, class A, 20 probes, 20 in V01510mRNA 506-1022, POMC gene (proopiomelanocortin) extracted from H.sapiens gene coding for ACTH and beta-LPH precursors. Gene codes for the common precursor of the pituitary hormones corticotropin (ACTH) and beta-lipotropin (beta-LPH)
4.31	82.6	4.1	135.05	52.82	138.3	128.13	142.9	73.54	309.4	194.45	440.75	506.5	U15306, class A, 20 probes, 20 in U15306 3076-3466, Human cysteine-rich sequence-specific DNA-binding protein NFX1 mRNA, complete cds
151.25	317.15	30.19	585.8	631.87	346.2	204.35	541	537.4	139.65	5.73	300	10.32	D26561, class A, 20 probes, 18 in D26561cds#2 153-435: 2 in reverseSequence, 3711-3717, ORF for L1 protein gene extracted from Human papillomavirus 5b genome integrated into human carcinoma DNA, ORF for L1 protein gene extracted from Human papillomavirus 5b genome integrated into human carcinoma DNA, ORF for L1 protein gene extracted from Human papillomavirus 5b genome integrated into human carcinoma DNA
254.77	526.85	210.79	426.25	351.79	553	507	312	38.61	346.9	245.08	409.05	490.1	D79993, class A, 20 probes, 20 in D79993 2741-3167, Human mRNA for KIAA0171 gene, complete cds
293.31	2692.1	151.18	2638.9	1717.28	1410.2	1399.51	2317.85	1289.13	1367.95	540.87	2605.4	327.67	X85373, class C, 20 probes, 20 in all_X85373 43-404, H.sapiens mRNA for Sm protein G
557.41	547.7	682.5	140.9	100.55	141.35	11.67	688.5	561.02	158.2	0.85	253.95	151.11	D25215, class A, 20 probes, 20 in D25215 4320-4839, Human mRNA for KIAA0032 gene, complete cds
504.94	934.4	873.98	991.5	967.46	473.5	412.53	793.7	158.82	789.15	275.42	1118.65	1123.24	X90978, class A, 20 probes, 20 in X90978 1187-1751, H.sapiens mRNA for an acute myeloid leukaemia protein (1793bp)
131.88	799	16.69	714.4	218.78	419.2	455.24	523.5	493.84	433.7	162.21	483.45	8.56	D76444, class A, 20 probes, 20 in D76444mRNA 2828-3362, Human hkf-1 mRNA, complete cds
373.21	787.55	455.87	1183.6	911.46	1059.4	767.78	827.85	0.49	520	310.42	649.75	551.33	M64269, class A, 20 probes, 18 in M64269cds 389-718: 2 in reverseSequence, 7859-7876, Human mast cell chymase gene, complete cds
556.99	480.75	132.44	1103.2	701.45	363.9	17.11	770.1	391.74	972.2	895.62	1150.75	862.03	M24766, class A, 20 probes, 20 in M24766 1513-2055, Human (clone pHAIV2-12) alpha-2 collagen type IV (COL4A2) mRNA, 3 end

178089.88	191404.84	126186.11	206333	227980.14	389687.38	159691.16	310789.81	63962.91	304302.84	18474.36	406499.34	84413.48	L06499, class A, 20 probes, 20 in L06499mRNA 4-301, Homo sapiens ribosomal protein L37a (RPL37A) mRNA, complete cds
15.77	99.9	61.38	160.7	32.53	264.7	30.12	380.05	148.99	297.95	252.08	104.35	7.85	U18422, class C, 20 probes, 20 in all_U18422 1719-2254, Human DP2 (Humdp2) mRNA, complete cds
157.47	2466.5	523.97	5846.55	4201.56	2034.9	1087.95	1950.95	419.1	2505.5	1114.4	1715.25	471	X91247, class C, 20 probes, 20 in all_X91247 3261-3700, H.sapiens mRNA for thioredoxin reductase
242.75	1104.65	186.32	509.05	153.51	1108.75	267.92	699.45	254.35	520.2	462.45	723.8	884.59	S59184, class A, 20 probes, 20 in S59184 2487-2979, RYK=related to receptor tyrosine kinase [human, hepatoma, mRNA, 3068 nt]
13.72	668.85	57.06	1091.5	776.12	869.2	620.27	631.2	118.94	883	323.15	1104.8	827.74	L13689, class A, 20 probes, 20 in L13689mRNA 2674-3076, Human prot-oncogene (BMI-1) mRNA, complete cds
135.91	913.35	90.01	717.3	451.56	1389	855.74	956.8	17.54	632.2	13.58	705.4	630.88	U01160, class A, 20 probes, 20 in U01160 1056-1635, Human transmembrane 4 superfamily protein (SAS) mRNA, complete cds
649.83	723.3	1.98	1396.15	1025.66	727.55	350.8	2773.9	371.8	1991.35	769.54	3442	1239.28	X04445, class A, 20 probes, 20 in X04445mRNA 521-1075, H.sapiens inhA gene exon 1 (and joined CDS)
254.06	1705.75	1054.51	2226	555.93	1670.75	196.93	1892.45	241.48	1373.65	496.18	1768.2	1522.83	X99142, class C, 20 probes, 20 in all_X99142 1281-1702, H.sapiens mRNA for hair keratin, hHb6
265.66	373.45	72.05	1058.7	692.54	574.1	485.92	653.25	227.62	833.3	493.28	661.2	198.98	U88726, class A, 20 probes, 20 in U88726 55-433, Human symplekin mRNA, partial cds. /gb=U88726 /ntype=RNA
2912.22	7918.7	3409.95	7729.3	3715	4565.9	4867.86	5491.55	1947.02	5077.45	560.24	8839.45	1003.74	D89052, class A, 20 probes, 20 in D89052 428-938, Human mRNA for proton-ATPase-like protein, complete cds
299.25	886.8	248.05	1416.9	1751.79	1623.45	1686.94	1510.25	1219.83	1835.25	1976.15	791.25	201.74	Latent Membrane Protein Lmp1
224.93	576.05	466.05	353.15	33.3	405.1	129.82	852.75	802.35	404.9	244.66	371.3	3.96	M60721, class A, 20 probes, 20 in M60721mRNA 1630-2182, Human homeobox gene, complete cds
508.62	1275.35	678.89	851.15	730.23	365.3	132.37	728	692.96	1056.25	83.23	1421.55	270.19	U78575, class A, 20 probes, 20 in U78575 3124-3634, Human 68 kDa type I phosphatidylinositol-4-phosphate 5-kinase alpha mRNA, clone PIP5KIa1, complete cds
176.71	1176.35	210.22	1284.7	519.02	759.9	562.86	1174.15	526.02	1317.1	1537.96	964.85	400.72	U10117, class A, 20 probes, 20 in U10117mRNA 474-954, Human endothelial-monocyte activating polypeptide II mRNA, complete cds
81.03	372.7	168.29	457.25	2.9	799.9	361.19	940.95	666.59	1625.65	1534.07	90.7	67.88	D25278, class A, 20 probes, 20 in D25278 2024-2510, Human mRNA for KIAA0036 gene, complete cds
26.73	226.2	75.09	162.55	64.84	133.8	69.72	456.55	591.49	181.95	93.27	44.4	2.26	U94319, class A, 20 probes, 20 in U94319 2044-2404, Human autoantigen DFS70 mRNA, partial cds.
145.52	318.25	160.3	181.55	157.76	270.85	103.17	151.45	43.49	870.85	262.27	791.25	219.7	U17195, class A, 20 probes, 20 in U17195 1567-2143, Human A-kinase anchor protein (AKAP100) mRNA, complete cds
2983.71	34087.35	6270.27	33920.15	5798.21	26430.9	10830.75	37930.45	5680.68	31827.1	8127.06	34510.85	790.19	L42611, class A, 20 probes, 20 in L42611 1374-1954, Homo sapiens keratin 6 isoform K6e (KRT6E) mRNA, complete cds
3624.49	69342.7	20791.63	50617.5	4551.36	58541.65	48762.16	48740	3865.61	83213.05	14130.61	83499.35	7560.88	X12447, class B, 20 probes, 13 in X12447mRNA#13 1-241: 7 not in GB record, Human aldolase A gene (EC 4.1.2.13)
220.83	515.7	109.46	471.2	169.28	2120.25	1981.38	1130.3	808.79	547.35	68.66	570.1	13.44	D13789, class A, 20 probes, 20 in D13789 1594-2062, Human mRNA for N-acetylglucosaminyltransferase III, complete cds

27.65	196.15	147.01	103.8	113.42	680.75	322.23	312.75	139.23	963.4	899.86	344	358.79	D50370, class A, 20 probes, 20 in D50370 2047-2581, Human mRNA for nucleosome assembly protein, complete cds
489.81	855.15	590.08	1717.6	1379.28	1543.3	1665.66	1041.8	183.71	1160.2	1253.56	495.45	72.76	U15197, class A, 20 probes, 20 in U15197 74-363, Human histo-blood group ABO protein mRNA, partial 3 UTR sequence
3.11	169	130.81	466	470.65	667.05	527.57	639.7	866.35	605.45	558.26	457.1	565.26	L01664, class A, 20 probes, 20 in L01664 43-529, Human eosinophil Charcot-Leyden crystal (CLC) protein (lysophospholipase) mRNA, complete cds
377.03	5279.4	320.18	4902.3	1021.2	4659.8	148.07	4804.6	531.18	4258.5	570.78	5233.75	467.04	M88468, class A, 20 probes, 20 in M88468 1378-1906, Homo sapiens mevalonate kinase mRNA, complete cds
159.03	124.25	9.97	366.3	464	417.15	56.07	203.15	232.43	897.8	1166.16	130.7	81.6	M61176, class A, 20 probes, 20 in M61176 1093-1549, Human brain-derived neurotrophic factor (BDNF) mRNA, complete cds
61.24	401.45	119.71	802.85	491.09	480.9	60.53	809.3	636.54	195.9	85.98	627.2	71.13	U21551, class A, 20 probes, 20 in U21551 728-1076, Human ECA39 mRNA, complete cds. /gb=U21551 /ntype=RNA
422.92	2374.65	931.33	1810.05	1206.68	1111.6	983.3	1971.15	1115.04	2439.35	218.14	1815.05	389.55	U41654, class A, 20 probes, 20 in U41654 1159-1525, Human adenovirus protein E3-14.7k interacting protein 1 (FIP-1) mRNA, complete cds
915.35	967.75	742.82	608.55	377.52	241.55	264.39	166.65	97.79	605.4	589.02	746.4	530.19	D42138, class A, 20 probes, 20 in D42138 1371-1833, Human mRNA for PIG-B, complete cds
82.45	174.45	52.68	134.45	66.11	198.4	133.36	276.35	39.81	149.65	14.92	231.65	72.48	U77594, class A, 20 probes, 20 in U77594 245-599, Human tazarotene-induced gene 2 (TIG2) mRNA, complete cds
48.15	207.1	64.21	270.45	171.33	324.65	71.77	185.55	109.25	209.55	22.7	292.1	142.84	X99894, class A, 20 probes, 20 in X99894 936-1482, H.sapiens mRNA coding for insulin promoter factor 1
5453.91	28621.45	3735.01	18169.6	4067.56	19736.7	12667.53	24092.95	6193.05	19884.9	15263.75	25333.95	5181.47	Myosin, Light Chain, Alkali, Smooth Muscle (Gb:U02629), Smooth Muscle, Alt. Splice 4 - Also Represents: HG2815-HT1357, M22919_rna2
20.29	76.6	22.34	324.1	402.49	198	165.46	419.55	382.76	263.35	191.56	338.65	235.11	U20536, class A, 20 probes, 20 in U20536 935-1428, Human cysteine protease Mch2 isoform alpha (Mch2) mRNA, complete cds
109.25	416.8	343.94	624.1	556.78	329.9	340.26	396.1	1.56	676.6	303.63	228.1	247.49	U44754, class A, 20 probes, 20 in U44754 809-1253, Human PSE-binding factor PTF gamma subunit mRNA, complete cds
8.56	342.4	120.49	251.9	105.36	667.45	141.92	604.95	271.74	222.95	114.34	372.6	277.75	M37825, class A, 20 probes, 20 in M37825 624-1044, Human fibroblast growth factor-5 (FGF-5) mRNA, complete cds
74.53	60.45	4.74	230.25	97.51	76.65	64.56	41.5	18.95	186.35	47.02	316	233.77	D14664, class A, 20 probes, 20 in D14664 3255-3639, Human mRNA for KIAA0022 gene, complete cds
1769.75	7021.45	850.72	8675.6	4024.43	5300.6	3241.52	5440.4	2041.28	4876.65	1454.87	6538.2	542.78	U72511, class A, 20 probes, 20 in U72511 440-596, Human B-cell receptor associated protein (hBAP) mRNA, partial cds
1097.57	2264	2146.07	1753.7	1797.61	971.05	336.94	2205.7	1555.63	1459.9	1021.77	2142.45	826.4	X71490, class C, 20 probes, 20 in all_X71490 1059-1552, H.sapiens mRNA for vacuolar proton ATPase, subunit D
1059.67	1362.3	1162.34	991.65	58.62	629.15	361.97	530.15	414.29	568.95	401.14	2350.35	1594.17	X66142, class B, 20 probes, 14 in X66142cds 2216-2539: 6 in reverseSequence, 2600-2643, H.sapiens mRNA for rod cGMP phosphodiesterase
88.74	204.6	17.11	209.2	141.14	269.05	83.93	333.05	241.9	832.05	715.95	209.3	4.53	U91616, class A, 20 probes, 20 in U91616 1484-1988, Human I kappa B epsilon (IkBe) mRNA, complete cds

371.8	454.5	210.86	135.4	76.79	255.9	128.98	186.35	33.16	266.7	5.52	466.55	180.81	U51240, class A, 20 probes, 20 in U51240 1679-2171, Human lysosomal-associated multitransmembrane protein (LAPTm5) mRNA, complete cds
114.9	974.65	159.45	955.2	81.46	854.2	81.03	1315.35	942.79	1103.75	53.39	1160.6	425.54	Z46376, class A, 20 probes, 20 in Z46376mRNA 4703-5249, H.sapiens HK2 mRNA for hexokinase II
69.51	428.3	68.73	891.65	223.52	242.45	117.59	129.95	69.37	581.35	367.34	430.5	46.53	U24389, class B, 19 probes, 14 in U24389cds 1495-1696: 4 in reverseSequence, 207-339: 1 not in GB record, Human lysyl oxidase-like protein gene
103.59	231.75	52.68	196.05	17.32	358.85	318.13	710.05	777.04	224.15	130.46	284.95	199.05	S39329, class A, 20 probes, 20 in S39329 948-1368, glandular kallikrein-1 {alternatively spliced} [human, prostate, mRNA, 1541 nt]
386.01	474.65	464.78	128.05	74.74	257	41.44	301.55	118.44	529.4	573.75	108.5	29.42	U45878, class A, 20 probes, 20 in U45878 2591-2689, Human inhibitor of apoptosis protein 1 mRNA, complete cds
260	516.25	437.49	1299.55	391.95	749.55	329.44	885.15	849.87	932.15	549.78	1531.45	429.43	U25265, class A, 20 probes, 20 in U25265 1758-2052, Human MEK5 mRNA, complete cds
1246.28	2968.55	619.35	2141.7	530.9	2554.95	386.58	3237.05	258.59	3342.65	233.98	2609.3	1307.3	M17466, class C, 20 probes, 20 in all_M17466 3487-4040, Human blood coagulation factor XII (F12) gene
125.23	222.75	117.45	232.7	46.53	1095.95	1107.54	682.95	365.5	280.65	52.68	76.5	50.91	L19493, class A, 20 probes, 20 in L19493exon 1931-2064, Human FMR1 gene, 3 end
270.11	563.05	422.64	736.65	213.62	529.35	215.6	261	264.74	545.1	542.35	902.65	274.71	U60205, class A, 20 probes, 20 in U60205 1192-1726, Human methyl sterol oxidase (ERG25) mRNA, complete cds
5133.45	7288.5	4036.73	9768.9	4974.78	5306	2706.24	6797.2	2656.88	4175.75	2554.71	6202.6	596.94	U68105, class A, 20 probes, 20 in U68105mRNA 2540-2765, Human poly(A)-binding protein (PABP) gene, promoter region and
324.99	628.65	584.28	460.95	80.4	942.95	96.24	921.8	836.65	720.5	260.22	789.2	856.87	U60666, class A, 20 probes, 20 in U60666 1923-2415, Human testis specific leucine rich repeat protein (TSLRP), complete cds
179.89	1793.25	436.78	837.3	707.39	1179.05	357.02	1748.95	324.21	823.5	882.19	611.15	461.67	S81264, class A, 20 probes, 20 in S81264 50-243, Hs-TBX2=T-box gene {T-box region} [human, fetal kidney, mRNA Partial, 283 nt]. /gb=S81264 /ntype=RNA
559.68	743.8	407.86	1249.55	666.45	628.4	235.61	360.9	97.3	439	405.31	219.7	221.47	M74002, class A, 20 probes, 20 in M74002 2243-2681, Human arginine-rich nuclear protein mRNA, complete cds
115.19	2189.6	432.47	1934.75	192.69	1926.05	495.19	688.6	884.73	2436.15	1180.23	2006.65	440.88	M21259, class C, 20 probes, 20 not in GB record, Human Alu repeats in the region 5 to the small nuclear ribonucleoprotein E gene
665.03	3957.8	588.31	4128.75	1027.64	5532.8	2717.55	3867.4	2263.31	3955.4	136.75	4632.7	879.64	D87460, class A, 20 probes, 20 in D87460 2023-2503, Human mRNA for KIAA0270 gene, partial cds
224.93	189.65	47.45	1196.95	1276.96	567.35	260.57	1800.2	2007.76	452.95	226.2	1141.6	282.7	U22029, class A, 20 probes, 20 in U22029 1690-2227, Human cytochrome P450 (CYP2A7) mRNA, complete cds
115.68	208.65	117.45	219.3	158.53	459.75	312.61	283.25	53.25	246.9	63.5	271.35	86.05	Y07846, class A, 20 probes, 20 in Y07846exon#7 92-587, H.sapiens mRNA for GAR22 protein
662.98	691.2	725.49	154.35	11.24	178	132.65	604.2	285.11	635.2	630.6	521	597.22	X68742, class C, 20 probes, 20 in all_X68742 2942-3423, H.sapiens mRNA for integrin, alpha subunit
30.76	75.15	27.51	65.25	34.01	186.45	136.97	137.45	111.94	106.65	74.46	129.25	25.95	X56411, class A, 20 probes, 20 in X56411mRNA 1990-2470, H.sapiens ADH4 gene for class II alcohol dehydrogenase (pi subunit), exon 1

437.91	450.9	182.57	541.3	148.07	702	295.99	1251.3	420.02	631.3	461.88	547.7	73.96	X52611, class B, 20 probes, 14 in X52611cds 962-1273: 6 in reverseSequence, 1441-1555, Human mRNA for transcription factor AP-2
135.55	139.2	49.78	292.55	233.56	106	122.33	396	434.59	214.1	60.1	77.3	15.7	L13943, class A, 20 probes, 20 in L13943 1963-2019, Human glycerol kinase (GK) mRNA exons 1-4, complete cds
67.1	108.15	56.36	159.25	128.2	214.2	5.8	238.85	106.14	189.35	115.75	202.65	159.88	L20433, class A, 20 probes, 20 in L20433 3738-3780, Human octamer binding transcription factor 1 (OTF1) mRNA, complete cds
67.53	249.25	78.7	252.25	198.48	138.55	75.73	384.8	246.21	790.95	516.82	201.2	93.06	U77129, class A, 20 probes, 20 in U77129 2447-2975, Human SPS1/STE20 homolog KHS1 mRNA, complete cds
309.15	291.7	271.25	182.9	147.5	595.8	814.3	51.25	14.35	334.3	410.26	175.5	82.59	L05597, class A, 20 probes, 17 in L05597cds 729-1071: 3 in reverseSequence, 1307-1499, Human serotonin receptor gene, complete cds
176.07	808.2	204.5	507.95	332.13	510.65	78.28	357	141.42	340.75	275.7	1101.6	142.41	Z48579, class B, 20 probes, 10 in Z48579cds 1852-2050: 10 in reverseSequence, 2062-2392, H.sapiens mRNA for disintegrin-metalloprotease (partial)
253.92	128.9	40.59	214.15	124.95	250.65	26.09	136.05	49.57	268.7	105.92	501.4	127.56	M73077, class A, 20 probes, 20 in M73077 2770-3178, Human glucocorticoid receptor repression factor 1 (GRF-1) mRNA, complete cds
708.38	1042.95	854.4	689.9	290.9	424.2	438.41	1402.35	1041.36	1015.4	121.62	909.5	223.59	J03242, class A, 20 probes, 20 in J03242 1155-1324, Human insulin-lke growth factor II mRNA, complete cds
240.63	177.05	57.91	185.25	190	118.15	32.46	148.65	4.88	602	711.07	297.55	119.43	N-Methyl-D-Aspartate Receptor Subunit, Splice Variant Hnr1n
91.5	1638.35	26.52	1683.55	137.82	2315.05	286.87	1705	100.13	2497.65	1558.96	2064.6	50.35	D63476, class A, 20 probes, 20 in D63476 4587-4953, Human mRNA for KIAA0142 gene, complete cds
379.43	1412.45	299.88	1119.1	146.37	530.1	145.81	1049.15	3.89	499.15	445.83	1952.4	1118.08	S43646, class A, 20 probes, 20 in S43646 1904-2402, cytokeratin 2 [human, epidermis, mRNA, 2427 nt]
310.63	832.85	213.9	501.7	306.04	630.35	296.21	590.8	56.57	988.45	862.32	659.8	395.98	Lectin, Galactoside-Binding, Soluble, 2
810.27	659.8	696.78	683.45	849.73	419.7	503.04	722.65	901.91	968.75	920.16	813.35	85.63	J02923, class A, 20 probes, 20 in J02923 2574-3132, Human 65-kilodalton phosphoprotein (p65) mRNA, complete cds
103.24	4410.9	111.16	4732	756.32	5024.7	1801.14	5556.9	852.77	5396.9	1642.61	4813.8	1651.66	X74104, class C, 20 probes, 20 in all_X74104 650-1059, H.sapiens mRNA for TRAP beta subunit
831.06	2298.5	814.16	875.6	839.48	462.85	435.65	1282.4	507.28	1206.85	87.19	1055.05	858.78	X70040, class B, 20 probes, 10 in X70040cds 3944-4130: 10 in reverseSequence, 4236-4470, H.sapiens RON mRNA for tyrosine kinase
77.64	59.75	13.93	758.2	950.35	192.1	118.51	105.85	92.56	750.6	598.07	468.35	570.99	X58288, class A, 20 probes, 20 in X58288mRNA 4517-4955, H.sapiens hR-PTPu gene for protein tyrosine phosphatase
217.36	368.7	427.94	1014.75	459.27	474.1	493.84	296.1	132.37	139.7	19.8	303.1	32.95	U07231, class A, 20 probes, 20 in U07231 2329-2647, Human G-rich sequence factor-1 (GRSF-1) mRNA, complete cds
215445.61	258300.59	164633.98	635353.75	768498.88	975077.75	526453.56	563775.25	436554.66	1247886.88	1355694.25	429969.31	288520.78	Bacteriophage P1 /GEN=cre /DB_XREF=gb:X03453.1 /NOTE=SIF corresponding to nucleotides 1089-1495 of gb:X03453.1 /DEF=Bacteriophage P1 cre gene for recombinase protein.
50.2	175.7	28.28	156.95	107.55	1024.9	1213.96	229	66.33	199.2	163.77	265.65	20.01	Modulator Recognition Factor 2
170.2	430.5	198.27	747.6	757.74	1169.4	795.21	387.65	144.89	354.4	173.38	456.55	77.85	D83407, class C, 20 probes, 20 in all_D83407 2601-3184, ZAKI-4 mRNA in human skin fibroblast, complete cds

1287.43	5352.75	701.52	4540.8	300.8	4969.3	422.99	6599.8	1782.33	5477.25	2520.2	4569.2	240.42	X62078, class C, 20 probes, 20 in all_X62078 1997-2400, H.sapiens mRNA for GM2 activator protein
308.86	3239.9	151.75	1693.7	1029.69	2005.7	1533.86	3452.25	142.48	2983.4	469.66	3267.2	369.11	Protein Phosphatase Inhibitor Homolog
90.58	74.9	97.44	76.5	96.87	85.7	16.69	125.95	151.39	40.65	23.55	112.95	76.44	M60751, class C, 20 probes, 20 in all_M60751 910-1163, Human histone H2B.1 (H2B) gene, complete cds
594.89	641.5	646.3	253.35	106	373.4	62.08	593.55	301.72	1244.3	1250.31	877.5	284.82	L11573, class A, 20 probes, 20 in L11573 1101-1665, Human surfactant protein B mRNA, complete cds. /gb=L11573 /ntype=RNA
325.69	654.45	401.85	440.55	239.64	230.8	195.44	194.9	144.53	838.3	402.77	98.4	63.64	X95384, class A, 20 probes, 20 in X95384 435-807, H.sapiens mRNA for unknown 14kDa protein
323.01	1029.3	130.25	494.45	15.34	464.45	265.94	558.2	116.39	969.15	827.1	425.05	119.71	U18244, class A, 20 probes, 20 in U18244 1166-1640, Human excitatory amino acid transporter 4 mRNA, complete cds
132.87	1402.9	617.59	1738.4	3.39	628	570.35	816.05	44.48	755.5	924.47	1725.75	15.34	X97065, class A, 20 probes, 18 in X97065cds 1852-2260: 2 in reverseSequence, 2389-2395, H.sapiens mRNA for Sec23B isoform, 2450bp
116.32	2371.85	2606.04	2452.8	2647.83	3906.05	4739.81	3666.3	4637.21	1952.05	2153.07	5030.35	739.85	U58516, class A, 20 probes, 20 in U58516 1860-1893, Human breast epithelial antigen BA46 mRNA, complete cds
26445.87	26195.2	25911.08	67156	85202.98	207474.7	213783.69	38617.95	25817.53	69223.65	70912.41	28298.4	2362.16	E. coli /GEN=bioB /DB_XREF=gb:J04423.1 /NOTE=SIF corresponding to nucleotides 2032-2305 of gb:J04423.1 /DEF=E.coli 7,8-diamino-pelargonic acid (bioA), biotin synthetase (bioB), 7-keto-8-amino-pelargonic acid synthetase (bioF), bioC protein, and dethiobiotin synthetase (bioD), complete cds.
45.96	507.95	360.55	289	54.02	481.45	238.08	483.25	92.98	295.65	27.08	584.45	112.92	X52213, class A, 20 probes, 20 in X52213cds 810-1371, H.sapiens ltk mRNA
699.96	5393.9	1126.28	5001.75	2013.77	2940.15	1351.21	3406.35	593.76	2921.25	114.06	3543.1	176.07	J02621, class A, 20 probes, 20 in J02621mRNA 653-1214, Human non-histone chromosomal protein HMG-14 mRNA, complete cds
8.77	3189.15	401.14	3744.2	330.08	4496.1	496.25	3169.35	1121.54	3593.95	679.46	3269.9	93.2	L20859, class A, 20 probes, 20 in L20859 2655-3159, Human leukemia virus receptor 1 (GLVR1) mRNA, complete cds
229.46	274.3	227.41	573.95	737.72	231.75	1.77	231.6	156.41	241.8	185.26	237.15	158.04	M22403, class A, 20 probes, 20 in M22403exon#2 1749-2224, Human blood platelet membrane glycoprotein Ib-alpha (GPIB) gene, complete cds, clone N10
125.37	370.1	130.11	333.45	131.45	194.8	38.89	239.25	29.77	578.45	137.25	355.05	64.28	X77737, class C, 20 probes, 20 in all_X77737 992-1431, H.sapiens mRNA for red cell anion exchanger (EPB3, AE1, Band 3) 3 non-coding region
502.68	4240.9	690.7	2972.25	445.41	3400.6	951.62	4565.15	192.83	3119.4	959.4	2872.15	947.45	D85527, class A, 20 probes, 20 in D85527 37-349, Human mRNA for LIM domain, partial cds. /gb=D85527 /ntype=RNA
794.72	962.2	807.52	540.7	436.14	697.05	742.39	433.45	331.14	1041.3	881.2	975.75	706.89	Y08639, class A, 20 probes, 17 in Y08639cds 837-1353: 3 in reverseSequence, 1953-2001, H.sapiens mRNA for transcription factor RZRbeta
66.68	127.15	88.18	132.75	122.26	146.05	162.42	169.95	5.02	75.4	35.36	131.6	33.94	M13577, class A, 20 probes, 20 in M13577mRNA 1550-2096, Human myelin basic protein (MBP) mRNA, complete cds
664.04	791.6	386.36	565.45	711.84	337.65	205.13	308.95	80.4	912.8	1009.47	595.2	373.35	U21556, class A, 20 probes, 20 in U21556 709-1204, Human membrane protein-like protein mRNA, partial cds. /gb=U21556 /ntype=RNA

159.88	280.35	60.46	282.65	265.94	507.15	490.38	239.55	183.64	244.6	197.99	246	161.36	Prostaglandin Receptor Ep1 Subtype - Also Represents: L22647
1163.54	3099.55	726.84	2675.2	978.64	3056.3	2149.04	2634.65	1028.77	3279.15	572.69	3221.8	53.6	AB002356, class A, 20 probes, 20 in AB002356 5330-5807, Human mRNA for KIAA0358 gene, complete cds. /gb=AB002356 /ntype=RNA
37.55	204.6	43.7	202.75	45.89	971.15	846.62	382.2	242.82	283.7	111.02	893.25	855.1	X55668, class A, 20 probes, 20 in X55668mRNA 550-940, Human mRNA for proteinase 3
31.18	223.7	170.27	211.15	163.55	565.95	203.29	565.4	70	485.55	427.87	602.15	5.44	S77361, class A, 20 probes, 20 in S77361 25-184, transcript ch132 [human, RF1,RF48 stomach cancer cell lines, mRNA, 216 nt]. /gb=S77361 /ntype=RNA
2250.3	11290.15	1259.29	11474.7	4384.49	6983.05	5265.19	10451.25	4532.77	8144.35	5618.03	10563.95	1547.79	X57152, class A, 20 probes, 20 in X57152mRNA#1 536-962, Human gene for casein kinase II subunit beta (EC 2.7.1.37)
706.33	241.95	81.11	783.75	256.61	679.75	108.54	1155.15	956.08	632.05	440.6	573.05	106	Y10936, class A, 20 probes, 20 in Y10936 637-1027, H.sapiens mRNA for hypothetical protein downstream of DMPK and DMAHP
84.64	94	4.95	160.3	143.68	352.7	91.92	346.05	254.63	173.05	147.43	202.95	67.81	J02843, class A, 20 probes, 16 in J02843cds 1103-1451: 4 in reverseSequence, 14089-14119, Human cytochrome P450IIE1 (ethanol-inducible) gene, complete cds
246.85	555.55	290.97	752.65	353.62	208.9	97.58	786.9	441.66	1405.05	300.17	985.5	841.32	M57730, class A, 20 probes, 20 in M57730mRNA 975-1437, Human B61 mRNA, complete cds
29.34	109.65	1.2	162.65	182.5	68.7	41.72	82.9	25.17	161.7	131.1	104.85	10.25	D80011, class A, 20 probes, 20 in D80011 4259-4793, Human mRNA for KIAA0189 gene, complete cds
548.36	689.05	809.99	1185.5	1071.69	718.05	365.08	321.05	234.55	468.85	183.35	767.75	184.77	Potassium Channel (Gb:L02752)
1803.26	58951.35	2821.57	60634.85	7464.29	81776.95	40719.1	74076.85	5547.75	81139.7	30657.18	85116.8	7005.44	U14972, class A, 20 probes, 20 in U14972 103-499, Human ribosomal protein S10 mRNA, complete cds
64.56	762.55	113.91	532.55	605.21	1176.35	86.48	1100.65	706.75	374.95	194.81	853.65	65.12	D86968, class A, 20 probes, 20 in D86968 4437-4899, Human mRNA for KIAA0213 gene, partial cds
158.53	189	203.36	101.4	31.68	155.85	166.52	423.4	547.44	973.8	1156.12	317.9	140.57	X61118, class A, 20 probes, 20 in X61118mRNA 1457-1955, TTG-2a gene extracted from Human TTG-2 mRNA for a cysteine rich protein with LIM motif
16.62	791.65	352.35	1882.5	805.96	586.7	558.19	1771.1	1259.5	1975.45	2403.39	1451.35	864.16	X52008, class B, 20 probes, 15 in X52008cds 899-1325: 5 in reverseSequence, 1748-1814, H.sapiens alpha-2 strychnine binding subunit of inhibitory glycine receptor mRNA
180.38	115.5	2.12	245.25	54.09	328.05	241.9	1528.35	1421.07	317.6	107.62	654.15	405.95	M34309, class A, 20 probes, 20 in M34309 4410-4836, Human epidermal growth factor receptor (HER3) mRNA, complete cds
1421.21	6935.5	262.2	5488.55	947.88	7263.3	2288.34	7495.65	2083.77	7890.9	2417.88	5743.45	450.36	AF001359, class A, 20 probes, 20 in AF001359 2-27, Homo sapiens DNA mismatch repair protein (hMLH1) mRNA, alternatively spliced, partial cds. /gb=AF001359 /ntype=RNA
46.67	200.95	117.73	65.8	11.6	181.45	45.18	157.05	55.51	286.75	206.83	512.7	639.65	U32581, class A, 20 probes, 20 in U32581 2488-2884, Human lambda/iota-protein kinase C-interacting protein mRNA, complete cds
125.65	54.1	36.2	608.25	714.25	169.95	74.32	85.7	36.06	206.1	258.24	46.5	1.41	X87613, class B, 20 probes, 14 in X87613cds 1996-2236: 6 in reverseSequence, 2780-2912, H.sapiens mRNA for skeletal muscle abundant protein
711.14	667.1	589.44	295.5	67.6	1025.45	1104.57	451.1	114.83	251.25	63.14	489.6	353.13	U64197, class A, 20 probes, 20 in U64197 250-658, Human chemokine exodus mRNA, complete cds

718.77	2827.35	960.32	2613.95	651.32	1736.65	1100.75	2701.7	1395.4	2108.15	1327.73	3635.9	11.6	U09584, class A, 20 probes, 20 in U09584 1382-1835, Human PL6 protein (PL6) mRNA, complete cds
33.94	176.65	61.16	332.75	4.6	311.65	165.11	285.35	11.1	478.95	505.79	224.15	3.75	U77845, class A, 20 probes, 20 in U77845 1515-1905, Human hTRIP (hTRIP) mRNA, complete cds
2991.56	3762.45	3509.72	1345.95	432.68	1840.95	1317.69	2318.5	2464.83	3015.9	307.17	3408.85	583.43	Cpg-Enriched Dna, Clone E18
385.94	315.85	267.64	1286.7	722.66	430.15	193.39	406.35	148.56	229.25	71.77	421.75	295.5	X92521, class B, 20 probes, 11 in X92521cds 1127-1484: 9 in reverseSequence, 1618-1768, H.sapiens mRNA for MMP-19 protein
1232.91	3847.4	1565.53	4440.85	347.4	2569.4	935.36	3495.75	836.44	2795.9	1940.58	3865.2	721.53	M88338, class A, 20 probes, 20 in M88338 1465-1867, Human serum constituent protein (MSE55) mRNA, complete cds
81.88	181.45	39.1	110.05	10.11	1763.4	2253.41	136.1	29.98	371.8	409.98	101.25	22.7	M81780, class B, 20 probes, 11 in M81780cds#4 49-265: 9 in fullSequence, 3676-3940, SMPD1 gene (acid sphingomyelinase) extracted from Homo sapiens acid sphingomyelinase (SMPD1) gene, complete cds, ORF 1-3, complete cds, SMPD1 gene (acid sphingomyelinase) extracted from Homo sapiens acid sphingomyelinase (SMPD1) gene, complete cds, ORF 1-3, complete cds, SMPD1 gene (acid sphingomyelinase) extracted from Homo sapiens acid sphingomyelinase (SMPD1) gene, complete cds, ORF 1-3, complete cds
82.45	614.65	158.89	888.35	89.02	1173.85	152.81	658	28.99	741.4	358.22	871	323.71	X12517, class B, 20 probes, 13 in X12517cds 261-441: 7 in reverseSequence, 480-660, Human mRNA for U1 small nuclear RNP-specific C protein
288.5	828.85	314.31	333.15	167.37	975.4	556.63	290.65	169.63	2180.35	1289.55	627.05	32.31	U33053, class A, 20 probes, 20 in U33053 2381-2879, Human lipid-activated protein kinase PRK1 mRNA, complete cds
1048.5	2613.5	1160.79	2740.2	30.26	1193.95	571.41	641.8	613.49	2414.45	870.8	1914.4	148.21	D44466, class A, 20 probes, 20 in D44466 2575-3115, Human mRNA for proteasome subunit p112, complete cds
13.29	71.4	24.61	88.45	39.67	138.75	113.35	109.8	83.58	182.7	174.94	99.45	62.86	L10381, class C, 20 probes, 20 in all_L10381 2086-2579, Human 2-5A-dependent RNase gene, complete cds
27.72	103.2	86.13	395.85	456.3	187.75	108.54	638.95	772.8	283.7	234.34	435.95	317.42	X77777, class A, 20 probes, 20 in X77777 2210-2771, H.sapiens intestinal VIP receptor related protein mRNA
748.83	1301.25	61.59	1812.8	1035.77	1622.35	2070.9	1512.9	738.08	1708.95	270.19	1329	171.54	L38961, class A, 20 probes, 20 in L38961 1919-2429, Human putative transmembrane protein precursor (B5) mRNA, complete cds
428.51	509.9	366.14	844.05	213.48	196.15	6.58	256	94.75	560.4	327.96	1272.9	1170.54	D50487, class A, 20 probes, 20 in D50487 3756-4098, Human mRNA for RNA helicase (HRH1), complete cds
58.9	170.65	64.42	155.15	159.74	231	43.13	527.65	157.05	108.25	78.56	164.45	177.13	X95073, class A, 20 probes, 20 in X95073 657-1119, H.sapiens mRNA for translin associated protein X
127.28	187.2	113.56	383.5	131.95	310.15	199.62	351.4	121.91	301.9	200.82	160.8	4.24	U57627, class A, 20 probes, 20 in U57627 4598-5078, Human fetal brain oculocerebrorenal syndrome (OCRL1) mRNA, complete cds
991.79	1872.5	212.56	2238.4	1729.3	2576.9	1260.49	2975.5	324.14	2233.6	1123.88	2350.2	449.72	U94586, class A, 20 probes, 20 in U94586 145-445, Human NADH:ubiquinone oxidoreductase MLRQ subunit mRNA, complete cds.
75.59	5854.25	916.62	4632.8	2439.66	4589.9	2932.51	3016	1573.03	3909.4	842.02	5948.1	1662.69	D21853, class A, 20 probes, 20 in D21853 1111-1543, Human mRNA for KIAA0111 gene, complete cds

26092.8	118684.45	5040.19	75868.75	17888.6	56604.95	49329.96	114706.5	32832.52	98959.65	77836.69	98055.5	11751.97	X00351, class B, 20 probes, 14 in X00351cds 855-1065: 6 in reverseSequence, 1154-1376, Human mRNA for beta-actin
118.09	134.1	154.71	194.8	160.8	90	28.14	62.3	11.6	547.1	52.61	118.45	10.39	M21119, class A, 20 probes, 20 in M21119 137-591, Human lysozyme mRNA, complete cds
325.2	702.05	226.77	628.1	203.65	1071.85	444.28	821.5	817.13	1378.95	1346.97	954.8	493.28	D28114, class A, 20 probes, 20 in D28114 780-1278, Human mRNA for MOBP (myelin-associated oligodendrocytic basic protein), complete cds, clone hOPRP2
1201.37	2488.45	303.42	2618.55	682.15	3524.45	1776.32	1931.6	956.57	1745.8	788.99	2105.95	214.47	L11284, class A, 20 probes, 20 in L11284 1672-2122, Homosapiens ERK activator kinase (MEK1) mRNA
555.86	3018.95	466.34	3256.9	783.47	1438.65	928.22	1565	984.43	2054	432.18	1790.1	532.03	Ribosomal Protein L39 Homolog
39.74	4408.8	260.64	3353.6	606.13	685.9	174.94	2190.25	290.83	1206.75	1314.44	1818.5	558.33	X53002, class B, 20 probes, 14 in X53002cds 2049-2356: 6 in reverseSequence, 2724-2849, Human mRNA for integrin beta-5 subunit
747.69	3135.5	1168.14	2656.4	195.73	3448.15	3692.58	3507.85	808.86	1917.2	1814.29	3522.85	1564.62	M87503, class A, 20 probes, 20 in M87503 1085-1535, Human IFN-responsive transcription factor subunit mRNA, complete cds
157.54	135.15	22.7	817.5	539.81	373.5	149.48	284.55	102.04	523	381.55	762.7	437.42	U28758, class A, 20 probes, 20 in U28758 45-626, Human NMDA receptor subtype 2B subunit (GRIN2B) mRNA, partial cds.
244.09	1366.9	280.01	2161.55	259.86	622.5	230.66	1118.3	217.65	1043.75	415.28	1222.95	501.83	X81625, class C, 20 probes, 20 in all_X81625 3058-3617, H.sapiens mRNA for Cl1 protein
837.14	5231.25	2291.38	3790.05	2605.19	1631.2	2067.86	1938.2	518.73	1554.3	1843.14	2237.45	404.82	M37583, class A, 20 probes, 20 in M37583mRNA 368-824, Human histone (H2A.Z) mRNA, complete cds
115.19	1536.4	22.34	807.65	903.61	1311.4	271.95	1297	760.85	851.55	445.97	1876.15	7.14	L29277, class A, 20 probes, 20 in L29277 2204-2762, Homo sapiens DNA-binding protein (APRF) mRNA, complete cds
1661.06	3202.25	831.49	1822.5	493.42	2798.4	1035.35	2136.6	513.08	4841	3211.82	1635.5	41.15	X66436, class C, 20 probes, 20 in all_X66436 1427-1980, H.sapiens hsr1 mRNA (partial)
45466.05	109336.25	23879.49	77535	27801.03	96316.59	68482.58	111080.3	42943.3	102401.6	46983.43	130999.05	42143.49	D00017, class A, 20 probes, 20 in D00017 851-1319, Human lipocortin II mRNA
57.98	1384.4	167.58	2108.25	363.81	1574.1	1543.9	1299.1	697.77	1473	505.58	1201.55	19.73	U24183, class A, 20 probes, 20 in U24183 2457-3031, Human phosphofructokinase (PFKM) mRNA, complete cds
767.99	2533.8	317.07	3371.3	1412.23	3432.5	366.85	1550.35	131.73	3335.9	537.4	2966.55	804.62	D31883, class A, 20 probes, 20 in D31883 6153-6711, Human mRNA for KIAA0059 gene, complete cds
22.42	135.8	32.39	135.6	36.49	203.7	30.55	202.3	5.66	338.55	329.02	217.75	171.33	M18731, class C, 20 probes, 20 not in GB record, Human galactose-1-phosphate uridyltransferase (GALT) mRNA, complete cds
505.37	3099.3	248.9	3908	857.01	4713.25	1291.11	3831.2	48.22	2504.4	927.3	4283.1	1441.79	D83542, class A, 20 probes, 20 in D83542 2250-2808, Human mRNA for cadherin-15, complete cds
74.46	332.8	21.78	357.8	309.43	242.1	86.83	492.2	516.33	939	424.69	415.95	18.88	U25975, class A, 20 probes, 20 in U25975 1675-1795, Human serine kinase (hPAK65) mRNA, partial cds
793.16	3131.05	520.22	2350.2	623.95	1804.45	1679.59	3380.4	1049.49	4803.05	2486.68	3415.4	1364.86	Z50781, class B, 20 probes, 13 in Z50781cds 100-205: 7 in reverseSequence, 346-394, H.sapiens mRNA for leucine zipper protein
161.86	548.45	78.28	516.55	239.92	5902.35	7201.81	1056.8	644.32	760	271.53	765.1	421.72	X51417, class C, 20 probes, 20 in all_X51417 1050-1651, Human mRNA for steroid hormone receptor hERR2
217.01	906.2	440.81	476	304.06	805.1	208.6	1269.85	81.81	825.65	242.18	669.85	342.73	M86849, class A, 20 probes, 20 in M86849 1747-2179, Human connexin 26 (GJB2) mRNA
656.9	734.9	880.63	384.4	263.89	799	772.87	1683.35	388.13	695.95	279.94	534.25	250.39	Homeotic Protein Hpx-5

304.41	271	306.46	163.8	87.4	1037	1364.01	86.5	16.83	250.95	18.6	296.65	217.44	U73167, class C, 20 probes, 20 in all_U73167 4971-35099, H_LUCA14.2a gene extracted from Human cosmid LUCA14, H_LUCA14.2a gene extracted from Human cosmid LUCA14, H_LUCA14.2a gene extracted from Human cosmid LUCA14, H_LUCA14.2a gene extracted from Human cosmid LUCA14, H_LUCA14.2a gene extracted from Human cosmid LUCA14
59.18	3631.95	487.83	3447.95	1640.42	1495.4	57.13	1025.25	266.23	1779.75	1663.75	370.2	87.4	U06643, class A, 19 probes, 19 in U06643 71-463, Human keratinocyte lectin 14 (HKL-14) mRNA, complete cds.
656.55	629.7	24.04	1219.4	932.11	726.6	536.69	688.8	55.44	604.95	638.02	660.75	358.15	M90516, class C, 20 probes, 20 in all_M90516 2559-3058, Human glutamine:fructose-6-phosphate amidotransferase (GFAT) mRNA, complete cds
162.35	318.2	321.31	586.25	508.62	118.05	47.59	165.95	159.45	207.85	106.42	243.35	254.63	D50916, class A, 20 probes, 20 in D50916 5465-5999, Human mRNA for KIAA0126 gene, complete cds
1204.7	1659.9	1116.66	2095.7	1983.86	2694.15	2557.68	1523.1	817.27	1340.6	435.15	1973.4	463.44	Z74615, class C, 20 probes, 20 in all_Z74615 5320-5852, H.sapiens mRNA for prepro-alpha1(I) collagen.
3.61	135.55	142.62	320.85	50.7	673.3	781.64	379.7	156.27	886	986.13	542.05	565.47	X51408, class C, 20 probes, 20 in all_X51408 1626-2017, Human mRNA for n-chimaerin
327.32	406.2	171.54	859.6	1125.86	636.75	42.92	369.35	184.77	793.4	1030.54	405.8	289.63	AB002366, class A, 20 probes, 20 in AB002366 5112-5616, Human mRNA for KIAA0368 gene, partial cds. /gb=AB002366 /ntype=RNA
293.38	520.45	353.34	750.75	522.76	431.5	44.97	182.7	19.23	1098.35	723.72	483.45	468.46	D88378, class A, 20 probes, 20 in D88378 2857-3157, Human mRNA for proteasome inhibitor hPI31 subunit, complete cds
80.82	231.2	10.47	134.05	42.36	344.05	278.25	140.75	41.08	276.1	70	93.65	41.65	M74290, class A, 20 probes, 20 in M74290 1261-1741, Human substance P receptor protein mRNA
1090.36	3218	1407.99	3283.6	536.84	2063.3	1962.5	3995.55	537.05	3048.4	935.22	3057.3	1191.9	U55054, class A, 20 probes, 20 in U55054 3199-3697, Human K-Cl cotransporter (hKCC1) mRNA, complete cds
369.68	961.4	130.53	810.1	502.47	1543.1	1358.64	670.6	225.99	2387.6	2174.92	976.45	435.51	M80333, class C, 20 probes, 20 in all_M80333 1628-2079, Human m5 muscarinic acetylcholine receptor gene, complete cds
12.16	89.85	71.63	172.05	64.13	126.75	89.73	203.5	105.64	282.9	47.8	209.6	5.66	X14346, class C, 20 probes, 20 in all_X14346 1969-2534, Human mRNA for eosinophil peroxidase
621.9	991.65	91.43	2019.9	416.63	1654.3	2182.7	420.45	204.99	1450.7	545.46	1120.95	58.62	X12791, class C, 20 probes, 20 in all_X12791 311-870, Human mRNA for 19kD protein of signal recognition particle (SRP)
217.51	166.05	101.05	221.35	13.36	323.05	116.46	306.55	52.11	227.55	119.29	197.75	2.19	X82676, class A, 20 probes, 20 in X82676 3333-3873, H.sapiens mRNA for tyrosine phosphatase
294.79	2137.3	643.47	2560.7	846.55	2141.05	775.91	1740.85	629.4	3373.05	132.3	3973.55	1821.01	D14695, class A, 20 probes, 20 in D14695 1259-1817, Human mRNA for KIAA0025 gene, complete cds
48.65	56.8	56.99	41.6	35.5	78.8	63.92	55.6	42.14	198.7	212.84	137.2	135.62	M57230, class A, 20 probes, 20 in M57230 2652-3000, Human membrane glycoprotein gp130 mRNA, complete cds
46.74	669.3	244.52	950.85	553.03	1783.25	1075.86	1392.85	478.36	1765.7	1258.37	1174.6	1026.86	U40279, class A, 20 probes, 20 in U40279cds 70-574, Human beta-2 integrin alphaD subunit (ITGAD) gene, exons 25-30, and partial cds. /gb=U40279 /ntype=DNA /annot=exon
293.1	2154.95	347.83	1688.3	1642.33	777.6	460.33	929.45	1051.26	1145.05	16.19	1552.8	210.43	D63878, class A, 20 probes, 20 in D63878 2850-3408, Human mRNA for KIAA0158 gene, complete cds
874.62	1041.1	753.92	968.2	530.75	1136.25	1414.28	633.25	628.55	1252.45	1115.18	1399.95	30.33	X82634, class A, 20 probes, 20 in X82634 1048-1378, H.sapiens partial mRNA for hair keratin acidic 3-II

68.94	92.4	47.52	168.2	69.16	204.15	27.93	171	105.64	368.05	426.6	920.9	1116.24	X05232, class C, 20 probes, 20 in all_X05232 1530-1771, Human mRNA for stromelysin
489.04	3097.55	520.5	2392.95	1845.9	3123.6	555.36	2027.9	1646	2424.35	1434.08	2995.05	734.33	D31767, class A, 20 probes, 20 in D31767 1338-1812, Human mRNA for KIAA0058 gene, complete cds
917.26	3687.4	1563.7	4303.5	3028.82	1927.25	419.67	1447.3	237.59	1747.2	1426.94	2928.4	553.52	X15729, class A, 20 probes, 16 in X15729cds 1538-1820: 4 in reverseSequence, 2069-2215, Human mRNA for nuclear p68 protein
405.53	883.4	96.31	789.65	397.32	754.85	393.08	1161.3	536.41	1157.45	857.23	735.5	323.85	M99701, class A, 20 probes, 20 in M99701 627-1137, Homo sapiens (pp21) mRNA, complete cds
130.11	330.15	144.89	1142.85	444.13	553.15	274.71	82.95	0.64	339.9	240.13	396.65	338.77	U04285, class A, 20 probes, 20 in U04285UTR#1 689-1148, Human lysosomal acid lipase, cholesteryl ester hydrolase (LIPA) gene
50.2	671.55	6.72	499.55	554.3	957.15	641.84	767.95	190.71	591.35	352.21	976.4	1180.44	X04085, class A, 20 probes, 20 in X04085mRNA 1684-2236, Human gene for catalase (EC 1.11.1.6) 5 flank and exon 1 mapping to chromosome 11, band p13 (and joined CDS)
698.76	2382.55	154.5	2859.85	1175.14	2084.75	297.2	1168.7	1077.07	1506.25	1172.03	1823.1	256.68	U91327, class A, 20 probes, 20 in U91327mRNA 156-654, Human chromosome 12p15 BAC clone CIT987SK-99D8 complete sequence. /gb=U91327 /ntype=DNA /annot=mRNA
10.04	38.45	15.49	354.1	450.99	122.65	132.44	66	14.14	202.55	192.12	37.25	0.49	U49973, class A, 20 probes, 20 in U49973cds#2 37-367, ORF1; MER37; putative transposase similar to pogo element from Human Tigger1 transposable element, complete consensus sequence. /gb=U49973 /ntype=DNA /annot=CDS, ORF1; MER37; putative transposase similar to pogo element from Human Tigger1 transposable element, complete consensus sequence. /gb=U49973 /ntype=DNA /annot=CDS
242.82	518.2	178.9	270.8	15.27	267.2	129.26	337.3	53.46	424.3	250.74	364.3	12.3	AF015910, class A, 20 probes, 20 in AF015910 41-433, Homo sapiens unknown protein mRNA, partial cds. /gb=AF015910 /ntype=RNA
46.74	393.9	13.01	931.35	1018.87	315.5	206.76	996.25	583.58	518.05	260.71	698.55	163.13	U18235, class A, 20 probes, 20 in U18235 61-493, Human ATP-binding cassette protein (ABC2) mRNA HFBCD04 clone, partial cds
111.51	1631.1	103.52	1924.1	36.06	4574.15	2986.04	1613.45	427.59	2246	741.76	2251.1	914.71	M21812, class A, 20 probes, 20 in M21812 61-592, Human (clone PWHLC2-24) myosin light chain 2 mRNA, complete cds
55.65	128	72.41	152.9	122.47	123.25	46.03	81.5	8.34	697.1	888.41	406.05	284.33	M19154, class A, 20 probes, 20 in M19154mRNA 2143-2503, Human transforming growth factor-beta-2 mRNA, complete cds
2391.01	7700.25	1443.28	4864.05	616.24	3941.85	1898.79	2603.45	204	3382.65	1036.55	4813.95	1535.06	J05614, class A, 20 probes, 20 in J05614mRNA 4-41, Human proliferating cell nuclear antigen (PCNA) gene, promoter region. /gb=J05614 /ntype=DNA /annot=mRNA
142.76	769.95	490.66	1207.45	527.57	1455.85	895.83	1572.7	462.59	1293.55	1639.57	459.65	231.58	Z48054, class C, 20 probes, 20 in all_Z48054 2544-3067, H.sapiens mRNA for peroxisomal targeting signal 1 (SKL type) receptor
486.07	506.65	332.69	310.7	9.33	947.4	910.05	245.4	2.26	632.65	451.77	469	200.54	M28879, class C, 20 probes, 20 in all_M28879 3325-4444, Human granzyme B (CTLA-1) gene, complete cds
2.97	198.2	205.2	579	746	181.5	105.92	207.2	114.69	151.9	34.65	168.9	14	U26174, class A, 20 probes, 20 in U26174 499-991, Human pre-granzyme 3 mRNA, complete cds
14.71	251.55	273.44	314.75	72.9	290.35	165.11	1433.05	1144.88	490.1	90.79	839.1	899.72	M11722, class A, 20 probes, 20 in M11722 1473-2037, Human terminal transferase mRNA, complete cds

25.03	34.95	27.22	135	151.75	644.8	849.52	53.35	36.42	125.7	111.86	51	4.95	U60665, class A, 20 probes, 20 in U60665 1603-2149, Human testis specific basic protein (TSBP), complete cds
58.55	72.45	79.55	253.75	321.8	231.2	128.27	366.75	447.67	155.25	189.01	266.05	321.52	Y07707, class A, 20 probes, 20 in Y07707 1136-1634, H.sapiens mRNA for ITBA4 gene. /gb=Y07707 /ntype=RNA
71.21	84.4	75.8	511.2	682.22	373.1	250.32	907	868.04	755.35	966.54	435.3	501.9	V00535, class A, 20 probes, 20 in V00535mRNA#1 253-692, interferon beta 1 gene extracted from Gene for human fibroblast interferon beta 1
330.22	309.85	225.5	578.05	43.49	534.9	430.77	241.8	210.72	1075.3	474.75	247.55	186.75	X97335, class C, 20 probes, 20 in all_X97335 3247-3704, H.sapiens mRNA for kinase A anchor protein
177.77	545.5	13.72	363.05	232.43	256.15	25.67	328.35	81.25	240.8	57.28	339.3	101.26	U83601, class A, 20 probes, 20 in U83601mRNA 58-136, Human calpastatin gene, exons 14 and 15, partial cds. /gb=U83601 /ntype=DNA /annot=mRNA
59.04	61.2	57.28	571.4	585.06	621.75	692.47	1288.35	1712.68	248.8	2.69	117.4	14	U81523, class A, 20 probes, 20 in U81523 1378-1870, Human endometrial bleeding associated factor mRNA, complete cds.
491.37	494.5	54.31	1646.45	898.52	850.55	350.09	926.65	848.88	1161.85	1298.88	912.2	635.26	Placental Protein 14, Endometrial Alpha 2 Globulin, Alt. Splice 3 - Also Represents: M34046, HG721-HT4827
25.1	40.25	12.37	90.6	58.55	112.05	112.5	36.9	16.97	86.75	75.45	56.65	24.68	U13616, class A, 20 probes, 20 in U13616 14235-14709, Human ankyrin G (ANK-3) mRNA, complete cds
344.57	1117.7	68.17	1436.6	865.5	1219.6	884.73	2084.4	884.02	2388.05	574.52	1208.15	479.63	M13485, class C, 20 probes, 2 in all_M13485 188-243: 1 in M13485cds 47: 17 not in GB record, Human metallothionein I-B gene
481.96	857.45	623.6	906.25	390.25	2300.35	2682.27	692.05	252.51	449.8	472.77	808.55	254.77	Z29678, class C, 20 probes, 20 in all_Z29678 1247-1740, H.sapiens mitF mRNA
496.39	3109.75	255.19	3768.05	857.79	2882.1	2185.53	3532.95	1769.39	2616.15	2163.53	1841.75	1554.15	L23333, class A, 20 probes, 20 in L23333 725-1305, Human corticotropin releasing factor receptor mRNA, complete cds.
301.02	961.7	203.65	1385.35	307.95	687.5	439.11	910.7	1043.12	474.95	572.97	888.85	94.26	Z97074, class A, 20 probes, 20 in Z97074 852-1176, Homo sapiens mRNA for Rab9 effector p40, complete cds.
31.82	2945	5.94	2728.4	577.28	1340.8	986.27	2076.65	1488.81	2725.15	694.17	2721.1	19.8	D63478, class A, 20 probes, 20 in D63478 2816-3308, Human mRNA for KIAA0144 gene, complete cds
1.06	127.65	6.15	343.75	219.98	296.85	193.54	278.55	148.56	145.95	40.09	246	49.21	Z49995, class A, 20 probes, 20 in Z49995mRNA 2014-2590, H.sapiens mRNA (non-coding; clone h2A)
80.26	194.1	116.39	504.25	211.07	659.4	132.51	267.75	171.47	277.85	86.34	171.25	31.32	X64072, class B, 20 probes, 14 in X64072cds 1948-2281: 6 not in GB record, H. sapiens CD18 exon 2
141.42	237.3	295.57	33.95	21.14	151.2	96.87	346.4	12.3	476.95	644.53	183.9	148.35	L39061, class A, 20 probes, 20 in L39061mRNA 1198-1654, Homo sapiens transcription factor SL1 mRNA, partial cds
1248.82	4615.85	880.28	4466.7	1610.36	3908.8	910.19	5022.95	777.04	3351.25	3300.84	5371.35	328.45	D38024, class C, 20 probes, 20 in all_D38024 2639-3228, Human facioscapulohumeral muscular dystrophy (FSHD) gene region, D4Z4 tandem repeat unit
74.18	117	53.88	88.45	61.02	330.4	258.8	142.05	52.68	228.75	162.28	499.65	567.74	U76272, class A, 20 probes, 20 in U76272mRNA 161-689, Human diadenosine triphosphate (Ap3A) hydrolase (FHIT) gene, 5 of
21.92	65.2	28.85	623.75	851.99	501.05	430.27	477.3	170.41	143.6	89.8	162.4	32.81	L41913, class C, 20 probes, 20 in all_L41913 305-502, Homo sapiens retinoblastoma susceptibility protein (RB1) gene, exon 26, bases 174145-174668 in L11910

408.92	842.35	289.42	1294.35	942.79	389.2	44.12	1547.2	317.92	495.7	7.92	1398.5	756.89	S58733, class C, 20 probes, 20 in all_S58733 264-357, pp52=B lymphocyte signal transduction gene {group 3, inverted repeat} [human, tumor cell line TC-32, Genomic Mutant, 356 nt]
78.14	723.95	214.04	557.7	76.65	821.15	686.67	1450.6	135.91	853.4	905.1	900.6	967.75	Zinc Finger Protein Znf155
1.2	119.25	99.35	145.85	54.09	235.65	241.76	99.25	6.01	370.6	170.13	242.85	286.02	L12468, class A, 20 probes, 20 in L12468 3246-3780, Homo sapiens aminopeptidase A mRNA, complete cds
0.99	125.45	166.52	371.3	396.26	162.1	31.82	219.8	240.98	279.85	361.12	116.2	62.65	M14306, class C, 20 probes, 20 in all_M14306 171-361, Human beta-A3/A1-crystallin gene (Hu-beta-A3/A1)
681.23	613	714.04	406	231.79	1434.45	1764.02	853.75	377.52	818.65	193.96	916.35	51.27	M23263, class A, 20 probes, 20 in M23263 3498-3648, Human androgen receptor mRNA, complete cds
191.27	522.25	274.85	423.8	25.46	437.4	47.38	565.95	220.26	625.6	155.85	714.9	52.33	D28588, class A, 20 probes, 20 in D28588 2747-3263, Human mRNA for KIAA0048 gene, complete cds
16.69	550.9	23.05	308.55	134.56	356.05	387.57	292.2	174.51	414.85	149.13	179.1	97.16	S83308, class A, 20 probes, 20 in S83308 920-1400, SOX5=Sry-related HMG box gene {alternatively spliced} [human, testis, mRNA, 1473 nt]
2324.05	1611.6	142.41	712.65	547.94	4084.65	310.07	6038.9	520.57	5021.35	1553.87	5736.25	1564.9	X04500, class A, 20 probes, 20 in X04500exon#7 244-778, Human gene for prointerleukin 1 beta
791.61	1252.8	234.48	1505.05	327.74	2812.95	1089.02	1533.15	476.1	2283.05	66.11	1698.55	245.86	D87437, class A, 20 probes, 20 in D87437 4613-5039, Human mRNA for KIAA0250 gene, complete cds
26.23	174.1	34.37	628.35	761.06	228.1	149.34	239.45	42.64	336	133.64	590.15	571.13	Y08417, class A, 20 probes, 20 in Y08417 1043-1558, H.sapiens mRNA for nicotinic acetylcholine receptor beta3 subunit precursor
6166.11	17505.05	5619.59	15884.05	10686.43	10280.95	5498.67	11374.45	3467.3	9804.05	6351.87	13517.85	3005.7	M86400, class A, 20 probes, 20 in M86400 2239-2743, Human phospholipase A2 mRNA, complete cds
176.78	721.25	176	778.1	122.33	1054.35	384.45	500.45	190.99	631.25	471.71	668.25	267.36	J02943, class A, 20 probes, 20 in J02943mRNA 869-1373, Human corticosteroid binding globulin mRNA, complete cds
1018.09	615.3	72.69	639.4	129.12	753.35	157.47	2089.5	2336.71	593	141.14	611.7	11.46	U09607, class A, 20 probes, 20 in U09607 3463-3730, Human JAK family protein tyrosine kinase (JAK3) mRNA, complete cds
57.98	130.35	120.99	231.7	115.26	200.35	143.33	195.35	175.29	236	171.26	169.9	166.45	U02683, class A, 20 probes, 20 in U02683 2758-3318, Human alpha palindromic binding protein mRNA, complete cds
645.45	1532.8	78.06	372.2	360.91	858.5	325.27	2250.1	267.99	2552.95	183.92	870	12.02	X52541, class C, 20 probes, 20 in all_X52541 2549-3102, Human mRNA for early growth response protein 1 (hEGR1)
149.2	174.35	2.76	277.35	32.74	1006.35	983.51	388.05	342.59	727.65	135.41	1157.35	43.35	U32849, class A, 20 probes, 20 in U32849 867-1383, Human Hou mRNA, complete cds
129.26	226.85	26.38	402.7	61.09	422.25	19.3	914.7	127.28	227.65	48.72	850.1	566.68	X83543, class C, 20 probes, 20 in all_X83543 7066-7427, H.sapiens APXL mRNA
21.43	610.85	405.95	614.8	20.51	2207.8	2521.54	630.3	497.1	293.95	13.79	521.25	47.45	D50923, class A, 20 probes, 20 in D50923 5018-5528, Human mRNA for KIAA0133 gene, complete cds
62.15	526.55	135.55	437.2	461.74	1505.5	1327.1	424.05	386.86	552	90.65	103.75	7.14	Macrophage Scavenger Receptor, Alt. Splice 2
355.67	458.85	327.32	539.65	438.19	209.5	91.08	616.3	481.4	738	985.57	675.7	858.85	X58723, class C, 20 probes, 20 in all_X58723 1862-2049, Human MDR1 (multidrug resistance) gene for P-glycoprotein
151.04	229.65	260.43	936.9	448.87	1344.35	1055.64	1026.85	464.64	845.9	386.79	550.7	631.87	U10690, class A, 20 probes, 20 in U10690exon#3 934-1499, Human MAGE-5b antigen (MAGE5b) gene, complete cds
995.89	1768.4	861.26	1069.9	659.31	896.35	563.78	704.6	539.38	1328.25	254.49	2214.55	147.43	Surfacant Protein Sp-A2 Delta - Also Represents: M30838, M68519_rna1, HG3928-HT4198, M13686

491.16	642.2	637.1	622.65	78.98	467.7	24.61	315.95	299.18	166.7	132.94	635.85	537.61	M13150, class A, 20 probes, 20 in M13150mRNA 776-1337, Human mas proto-oncogene mRNA, complete cds.
367.55	330.9	258.8	305	347.05	676.5	146.51	635.2	242.54	371.2	191.63	696.4	751.8	U26710, class A, 20 probes, 20 in U26710 3398-3878, Human cbl-b mRNA, complete cds
241.34	905.85	93.97	594.75	475.67	1286.35	0.49	661.55	232.85	577.1	560.88	648.2	509.82	U18242, class A, 20 probes, 20 in U18242 761-1265, Human calcium modulating cyclophilin ligand (CAMLG) mRNA, complete cds
153.09	322.2	88.53	201.95	19.45	415.7	318.2	295.65	8.41	250.7	89.52	244.55	94.12	U79280, class A, 20 probes, 20 in U79280 795-1359, Human clone 23575 mRNA, partial cds.
2543.53	13078.6	805.96	10727.25	3815.62	12602	4709.33	19400.15	9145.22	12065.85	10733.25	19934.2	2837.62	U31201, class A, 20 probes, 20 in U31201mRNA 4592-5106, Human laminin gamma2 chain gene (LAMC2), Human laminin gamma2 chain gene (LAMC2)
6.29	11.9	4.67	35.3	8.63	337.2	238.58	66.3	32.1	428.5	347.61	31.5	10.18	X51730, class C, 20 probes, 20 in all_X51730 4462-5003, Human mRNA and promoter DNA for progesterone receptor
278.32	342.55	367.34	459.2	364.73	944.5	226.7	469.15	335.66	615.15	217.29	410.55	290.13	L37378, class A, 20 probes, 20 in L37378mRNA 3182-3680, Homo sapiens guanylyl cyclase (RetGC-2) mRNA, complete cds.
19.8	992.55	200.61	873.85	326.05	1009.55	784.82	1345	858.14	1889.15	969.09	1092.75	622.61	Z67743, class A, 20 probes, 19 in Z67743cds 1792-2320: 1 in reverseSequence, 2350, H.sapiens mRNA for CLC-7 chloride channel protein
13.01	102.55	94.54	645.25	492.22	226.95	180.52	76.5	50.77	211.4	80.75	101.75	64.42	X75958, class C, 20 probes, 20 in all_X75958 1683-2170, H.sapiens trkB mRNA for protein-tyrosine kinase
2592.18	3513.75	2649.88	1547.7	158.25	2696.3	1491.71	3423.05	340.19	2127.4	16.83	1982.4	290.9	U59752, class A, 20 probes, 20 in U59752 524-938, Human Sec7p-like protein mRNA, partial cds
223.02	687.9	132.09	601.7	531.89	318.6	100.41	660.6	795.21	567.15	311.76	768.05	707.6	Y08612, class A, 20 probes, 18 in Y08612cds 1849-2197: 2 in reverseSequence, 2257-2269, H.sapiens mRNA for Nup88 protein
467.54	18014.45	2016.6	17131.8	3256.51	43764.9	34148.03	34574.3	2726.74	36369	1159.94	47340	6643.98	L20941, class A, 20 probes, 20 in L20941 615-1143, Human ferritin heavy chain mRNA, complete cds
35.57	82.25	43.35	96.2	38.61	613.6	719.55	379.25	346.69	162.85	76.72	113.05	49.29	M37485, class C, 20 probes, 20 in all_M37485 1008-1230, IGH@ gene (Ig Dxp heavy-chain gene) extracted from Human Ig germline H-chain D-region Dxp1 and Dxp1 genes, 3 end
758.16	2991.45	127.92	3726.25	63.57	1863.9	680.94	2391.1	655.91	2189.7	500.49	2909.1	353.98	L38932, class A, 20 probes, 20 in L38932mRNA 1076-1466, Homo sapiens GT197 partial ORF mRNA, 3 end of cds
66.61	66.9	69.58	124.65	129.47	545.15	310.91	83.25	18.46	517.1	510.81	102.35	73.89	L18920, class A, 20 probes, 20 in L18920exon#4 970-1461, Human MAGE-2 gene exons 1-4, complete cds
1402.83	2863.3	468.1	1474.5	3.39	2350.3	824.2	2441.25	889.19	1206.7	1163.19	1986.3	667.23	D50920, class A, 20 probes, 20 in D50920 2969-3419, Human mRNA for KIAA0130 gene, complete cds
235.11	358.1	292.74	710.2	898.03	315.4	194.6	622.45	177.98	575.2	31.11	480.15	648.2	Bone Morphogenetic Protein 3
844.43	831.65	774.35	393.35	232	201.2	191.06	996.5	92.63	123.85	45.33	179.9	90.65	U30246, class A, 20 probes, 20 in U30246 3599-4019, Human bumetanide-sensitive Na-K-Cl cotransporter (NKCC1) mRNA, complete cds
117.66	5381.15	32.6	4681	8.91	3982.3	2060.37	6587.05	842.52	6266.8	174.09	4973.35	55.51	X55448, class A, 20 probes, 20 in X55448exon#13 150-670, G6PD gene (glucose-6-phosphate dehydrogenase) extracted from H.sapiens G6PD gene for glucose-6-phosphate dehydrogenase, G6PD gene (glucose-6-phosphate dehydrogenase) extracted from H.sapiens G6PD gene for glucose-6-phosphate dehydrogenase

131.95	43082.15	2779.28	29543	9057.19	34171.5	17537.1	58924.8	24733.88	47558.05	29479.5	72182.8	12054.62	X02761, class C, 20 probes, 20 in all_X02761 7082-7646, Human mRNA for fibronectin (FN precursor)
604.22	865.65	596.73	707	608.68	332.05	385.59	241.7	167.16	1593.95	1533.08	179.5	85.42	X68264, class A, 20 probes, 20 in X68264mRNA 3300-3552, MUC18 gene (melanoma associated glycoprotein) extracted from H.sapiens MGF gene exons 1&2
415.14	4519.2	447.88	4082.1	2165.87	2839.15	2957.47	4474.4	391.45	3506.45	1055.64	1516.05	308.37	U56816, class A, 20 probes, 20 in U56816 1376-1838, Human kinase Myt1 (Myt1) mRNA, complete cds.
113.28	1632.15	787.22	1490.3	919.1	1548.35	1385.29	1113.75	227.05	1984.45	1469.86	2201.6	284.68	D30758, class A, 20 probes, 20 in D30758 1965-2469, Human mRNA for KIAA0050 gene, complete cds
933.38	1115.65	813.95	903.9	226.56	422.1	419.74	579.2	292.46	1461.55	553.74	1163.9	390.46	U96113, class A, 20 probes, 20 in U96113 1463-2003, Homo sapiens Nedd-4-like ubiquitin-protein ligase WWP1 mRNA, partial cds. /gb=U96113 /ntype=RNA
428.01	395.95	501.83	415.75	490.38	455	392.44	110.7	82.45	175.55	78.98	293.2	316.5	D45399, class A, 20 probes, 20 in D45399mRNA 155-629, Human adult neural retina mRNA for human cone cGMP phosphodiesterase gamma subunit, complete cds
188.09	324.9	219.63	543.25	295.78	562.25	310.49	227.8	206.19	728.75	219.13	492.85	374.41	L07541, class A, 20 probes, 20 in L07541 950-1436, Human replication factor C, 38-kDa subunit mRNA, complete cds
245.93	274.8	285.25	1181.65	1457.98	575.4	403.9	1208.8	520.43	330.95	319.4	208.4	86.83	M17254, class C, 20 probes, 20 in all_M17254 1366-1889, Human erg2 gene encoding erg2 protein, complete cds
45.89	969.65	98.22	824.95	729.38	1269.1	834.39	999.65	841.39	577.95	383.89	856.4	132.94	D21163, class A, 20 probes, 20 in D21163 3183-3693, Human mRNA for KIAA0031 gene, complete cds
552.25	750.2	760.56	1225.3	101.54	394.15	120.84	996.4	115.4	604.05	356.59	912.8	723.37	D21852, class A, 20 probes, 20 in D21852 3671-4241, Human mRNA for KIAA0029 gene, partial cds
5969.89	18411.55	6371.67	14437.3	2575.99	14447.05	8604.57	20883.6	2118.63	18662.5	4786.83	23346.4	587.04	J03077, class A, 20 probes, 20 in J03077 2159-2692, Human co-beta glucosidase (proactivator) mRNA, complete cds
410.83	1344	420.45	915.8	406.02	396.5	311.13	383.6	460.19	1655.1	1336.15	1080.25	389.4	U58089, class A, 20 probes, 20 in U58089 1599-1941, Human Hs-cul-3 mRNA, partial cds
428.58	411.4	353.69	570.9	721.11	541.1	545.74	117.4	15.41	86.55	65.97	219.15	70.92	U49187, class A, 20 probes, 20 in U49187 1780-2206, Human placenta (Diff48) mRNA, complete cds
37.55	210.4	21.78	287.75	246.99	1453.1	1781.06	277.55	108.26	191.4	36.35	226.95	98.5	U16258, class A, 20 probes, 20 in U16258 1212-1776, Human I kappa BR mRNA, complete cds
208.88	291.6	268.84	251	155.28	308.2	96.87	300.8	261.91	1642.65	1470.57	829.2	70	X84194, class A, 20 probes, 16 in X84194cds 61-271: 4 in reverseSequence, 501-555, H.sapiens mRNA for acylphosphatase, erythrocyte (CT) isoenzyme
1163.69	3967.6	1266	4532.05	618.08	5006	2499.62	5814.2	653.37	6081.1	1429.35	4546.35	430.7	M62762, class A, 20 probes, 20 in M62762 681-1083, Human vacuolar H+ ATPase proton channel subunit mRNA, complete cds
68.45	662.05	381.77	1157.25	877.59	1088.15	503.39	1648.45	410.05	1284.6	1201.52	421.1	100.69	L13197, class A, 20 probes, 20 in L13197 1853-2099, Human (clone D21S418E) pregnancy-associated plasma protein A (PAPP-A) gene, 5 UTR
85.35	165.9	71.7	209.3	31.25	183.35	71.21	142.05	17.61	313.65	344.15	146.1	102.53	L00022, class A, 20 probes, 18 in L00022cds 1249-1679: 2 in reverseSequence, 1750-1816, Human Ig active epsilon1 5 UT, V-D-J region subgroup VH-I, gene
743.52	2701.65	603.23	2362.7	363.03	3167.7	2215.22	2410.55	803.91	2034	2421.13	2848.2	554.09	M94172, class A, 19 probes, 19 in M94172 6837-7328, Human N-type calcium channel alpha-1 subunit mRNA, complete cds
76.23	162.4	71.13	401.3	14	127.4	17.96	183.05	73.75	1069.2	916.55	126.75	34.15	Guanine Nucleotide-Binding Protein, Alpha 12

1565.46	1525.8	1447.59	425.1	212.98	1495.65	1318.4	573.35	18.31	487.2	27.15	604.7	53.17	M85220, class C, 20 probes, 20 in all_M85220 15-201, Human heavy chain disease IgA chain gene, CH3 region with a 369 bp deletion, 3 end
298.54	1509.3	240.7	2188.25	505.09	840.35	227.62	1423.3	1303.06	998.05	32.88	1121.45	521.21	X12458, class A, 20 probes, 20 in X12458mRNA 1566-2046, P3 protein (AA 1-1382) gene extracted from Human P3 gene
3.46	70.95	1.48	108.4	47.8	513.7	18.53	118.45	12.09	96.85	86.48	99.5	48.37	L40384, class A, 20 probes, 20 in L40384mRNA 22-487, Homo sapiens thyroid receptor interactor (TRIP13) mRNA, partial cds. /gb=L40384 /ntype=RNA
529.91	1187.5	615.89	713.8	227.12	710.6	397.39	1102.65	469.31	1294.6	488.61	862.2	142.98	X63187, class A, 20 probes, 20 in X63187mRNA 91-511, H.sapiens HE4 mRNA for extracellular proteinase inhibitor homologue
273.72	436.85	133.29	341.8	78.91	771.75	61.16	835.25	409.49	234.8	78.91	724.45	277.4	X54816, class C, 20 probes, 20 not in GB record, Human gene for alpha-1-microglobulin-bikunin, exons 5-Jan (encoding alpha 1-microglobulin, N-terminus.)
463.44	598.55	360.84	901.6	1134.34	377.45	453.47	572	487.34	568.6	293.73	479.35	508.2	D89859, class A, 20 probes, 20 in D89859 2331-2841, Human mRNA for zinc finger 5 protein, complete cds
289.42	754.2	221.32	850.4	21.5	495.3	54.59	1279.55	84.5	613.7	324.42	682.95	702.37	S80050, class A, 20 probes, 20 in S80050mRNA 653-1133, UDP-N-acetylglucosamine: alpha-6-D-mannoside beta-1,6-N-acetylglucosaminyltransferase V|GlcNAc transferase V {5 region} [human, peripheral blood, Genomic, 1972 nt]. /gb=S80050 /ntype=DNA /annot=mRNA
185.12	243.8	42.57	1085.7	1020.64	431.9	461.88	1171.25	644.39	763.75	650.61	846.9	484.93	U12622, class A, 20 probes, 20 in U12622 4-358, Human beaded intermediate filament protein CP115 mRNA, partial cds. /gb=U12622 /ntype=RNA
220.48	278.75	266.51	1020	1268.97	510.3	530.9	732.35	401.57	219.85	99.21	524.6	439.54	X05323, class A, 20 probes, 18 in X05323cds 426-792: 2 not in GB record, Human MRC OX-2 gene signal sequence
86.34	73.6	84.29	37.85	34.58	612.05	845.91	554.85	771.1	187.6	246.5	320.15	417.26	U33838, class C, 4 probes, 4 in all_U33838 62-95, Human NF-kappa-B p65delta3 mRNA, spliced transcript lacking exons 6 and 7, partial cds. /gb=U33838 /ntype=RNA, Human NF-kappa-B p65delta3 mRNA, spliced transcript lacking exons 6 and 7, partial cds. /gb=U33838 /ntype=RNA
630.74	2635	761.98	1928.8	699.47	1117.45	584.14	2907.9	1188.51	1695.35	2157.59	2316.65	737.87	Z49989, class A, 20 probes, 19 in Z49989cds 779-1079: 1 in reverseSequence, 1540, H.sapiens mRNA for smoothelin
1028.77	5740.55	1127.76	4486.85	753.14	4845.1	3128.52	4932.15	28.07	6040.6	732	6152.9	1099.13	X97074, class B, 20 probes, 14 in X97074cds 182-398: 6 in reverseSequence, 704-782, H.sapiens mRNS for clathrin-associated protein
97.86	143	164.9	693.7	707.25	798.05	826.82	66.05	36.98	560.6	399.23	50.1	1.13	U46116, class A, 20 probes, 20 in U46116mRNA 5907-6477, Human receptor tyrosine phosphatase gamma (PTPRG) gene
85.49	643.9	199.55	718.4	166.03	1342.75	708.73	443.35	236.24	959.9	400.22	710.05	17.47	J04177, class A, 20 probes, 20 in J04177 5773-6133, Human alpha-1 type XI collagen (COL11A1) mRNA, complete cds
60.25	958.15	415.43	1300.5	486.49	577.7	168.01	515.7	405.31	1285.95	1215.87	939	367.13	D86962, class A, 20 probes, 20 in D86962 5028-5328, Human mRNA for KIAA0207 gene, complete cds
1256.88	5260	1200.95	4830.6	1326.11	5684.3	1933.09	4984.55	840.54	4859.5	8.06	3784.55	519.79	X67683, class A, 20 probes, 19 in X67683cds 34-319: 1 in reverseSequence, 13, H.sapiens mRNA for keratin 4 /gb=X67683 /ntype=RNA

197.14	311.05	399.44	102.1	4.1	314.8	291.61	525.55	318.13	463.8	387.35	60.55	22.13	U79272, class A, 20 probes, 20 in U79272 699-1179, Human clone 23720 mRNA sequence
3101.94	4909.65	1757.37	12316.7	14621.55	16005.55	15596.58	10202.85	7230.52	8104.15	4602.77	7112.4	2605.69	M16653, class A, 20 probes, 20 in M16653mRNA 652-742, Human pancreatic elastase IIB mRNA, complete cds.
1465.83	3059.8	557.62	4104.8	218.78	3731.35	2447.79	4783.65	57.2	2803.85	253.92	4889.35	1196.5	M88458, class A, 20 probes, 20 in M88458 585-1095, Human ELP-1 mRNA sequence
1890.59	3741.7	1499.91	4519.35	2224.2	3522.15	1267.06	2547.7	1716.86	3516.4	2124.71	3912.55	1199.18	D13988, class A, 20 probes, 20 in D13988 897-1353, Human rab GDI mRNA, complete cds
1068.65	1244	1345.91	1618.15	1106.69	580.9	83.3	1513.7	1202.51	1752	663.55	1283.6	287.65	X14675, class A, 20 probes, 20 in X14675cds 31-163, Human bcr-abl mRNA 5 fragment (clone 3c). /gb=X14675 /ntype=RNA
319.26	1155.3	148.35	1494.95	957.21	684.5	234.62	816.35	196.22	1629.5	1111.57	538.75	181.09	AB004884, class A, 20 probes, 20 in AB004884 2491-2995, Human mRNA for PKU-alpha, partial cds. /gb=AB004884 /ntype=RNA
4718.31	21301.55	8491.43	13974.9	7529.41	20768.6	17765.21	14996.5	6399.88	18234.5	14618.73	18383.85	112.36	X57351, class C, 12 probes, 12 in all_X57351 294-891, Human 1-8D gene from interferon-inducible gene family, Human 1-8D gene from interferon-inducible gene family
119.22	740.9	5.66	1515.4	1296.41	2190	1948.79	1486.65	1086.89	2151.15	1644.09	1144.95	43.35	AF005037, class A, 20 probes, 20 in AF005037 574-1030, Homo sapiens secretory carrier membrane protein (SCAMP1) mRNA, complete cds. /gb=AF005037 /ntype=RNA
334.04	782.9	107.06	348.5	160.23	1618	1019.22	1127.15	525.73	1355.55	852.42	726.7	45.11	U66048, class A, 20 probes, 20 in U66048mRNA 2400-2838, Human clone 161455-2-3 B cell expressed mRNA from chromosome X
38.82	96.45	5.02	158.45	120.84	146.25	43.49	551.55	394.64	106.45	4.17	254.35	260.99	X51956, class A, 20 probes, 20 in X51956mRNA 1881-2397, Human ENO2 gene for neuron specific (gamma) enolase
276.13	192.15	166.67	606.9	211.85	392.1	355.96	522.05	362.53	153.35	50.28	1670.35	694.59	X98311, class C, 20 probes, 20 in all_X98311 1901-2274, H.sapiens mRNA for carcinoembryonic antigen, CGM2
372.29	377.3	321.73	736.45	514	217.2	132.65	309.65	63	140.85	61.59	931.45	1079.82	Z48804, class A, 20 probes, 20 in Z48804mRNA 1006-1528, H.sapiens mRNA (ocular albinism type 1 related)
68.87	589.7	373.07	489.8	528.92	309.1	252.44	113.8	26.3	191.5	120.77	213.6	44.69	L25851, class A, 20 probes, 20 in L25851 3332-3812, Homo sapiens integrin alpha E mRNA, complete cds
36.84	244.45	24.11	472.45	12.52	661.1	290.76	613.25	124.66	724.4	225.28	593.5	171.26	U69263, class A, 20 probes, 20 in U69263 474-1002, Human matrilin-2 precursor mRNA, partial cds
67.95	250.4	106.91	453.6	255.12	218.5	222.6	876.8	519.72	634	801.15	1274.85	574.95	M25164, class B, 20 probes, 15 in M25164cds 5-383: 5 in reverseSequence, 265-1170, Human thyrotropin beta subunit gene
18516.51	86794	11899.76	73080.91	3406.56	64913.05	46353.89	82016.85	30636.18	73029.65	35822.24	103117.55	8885.86	X73460, class A, 20 probes, 19 in X73460cds 725-1133: 1 in reverseSequence, 1211, H.sapiens mRNA for ribosomal protein L3
0.92	387.5	46.81	259.95	195.94	818.9	318.62	661.25	575.8	239.3	143.4	1089.3	187.38	X68149, class A, 20 probes, 20 in X68149exon#2 2249-2687, H.sapiens BLR1 gene for Burkitt lymphoma receptor 1
222.67	622.2	232.64	491.7	100.83	655.7	231.51	872.85	317.7	891.85	413.87	723.3	34.65	U03642, class C, 20 probes, 20 in all_U03642 1060-1565, Human G protein-coupled receptor APJ gene, complete cds
160.37	2635.1	610.52	1847.25	31.04	2567.35	271.46	4122.55	186.18	4085.05	1387.98	3642.05	600.12	L38490, class A, 20 probes, 20 in L38490mRNA 785-1319, Homo sapiens ADP-ribosylation factor mRNA, complete cds

799.67	3148.7	867.48	2309.6	684.76	1370.3	573.46	3064.3	733.41	1803.05	1257.87	1273.5	966.19	L38487, class A, 20 probes, 20 in L38487mRNA 1623-2115, Human estrogen receptor-related protein (hERRa1) mRNA, 3 end, partial cds
474.19	2714.9	33.94	2028.1	289.35	3124.8	817.42	2246.9	692.82	3121.45	741.12	2593.35	504.52	L13203, class A, 20 probes, 20 in L13203 1536-2064, Human HNF-3/fork-head homolog-3 HFH-3 mRNA, complete cds
277.75	1385.3	415.35	1831.35	827.95	2335.1	1863.37	741.2	658.74	1378.45	23.12	658.8	283.13	U52426, class A, 20 probes, 20 in U52426 3469-3997, Human GOK (GOK) mRNA, complete cds
740.2	613.9	627.2	750.3	643.89	1043.05	855.25	1040.5	288.78	1400.9	870.02	1446.15	603.09	M55905, class A, 20 probes, 20 in M55905 1340-1820, Human mitochondrial NAD(P)+ dependent malic enzyme mRNA, complete cds
85.98	145.3	17.96	132.25	112.08	594.15	130.46	597.8	440.39	273.25	87.04	890.5	743.45	X59770, class A, 20 probes, 20 in X59770mRNA 685-1213, H.sapiens IL-1R2 mRNA for type II interleukin-1 receptor, (cell line CB23)
36.13	56.45	34.01	749.05	1017.88	197.7	215.1	154.1	51.76	699.85	903.47	56.05	27.37	Lymphocyte Antigen Hla-G3 - Also Represents: J03027
73.4	263.15	52.54	346.45	168.22	142.85	37.26	171.4	1.98	188	51.9	141.1	96.59	U11036, class A, 20 probes, 20 in U11036 121-553, Human lbd1 mRNA, partial cds. /gb=U11036 /ntype=RNA
35.21	368.85	195.94	266.6	49.92	1333.6	548.15	679.45	283.05	1372.2	1224.57	968	645.31	M91036, class A, 20 probes, 20 in M91036mRNA#1 37-529, G-gamma globin gene extracted from H.sapiens G-gamma globin and A-gamma globin genes, complete cds
83.65	458.95	45.04	372.9	102.67	1546.05	1780.99	963.4	51.62	1154.35	391.67	1523.55	1755.11	U66828, class A, 20 probes, 20 in U66828 2341-2500, Human carnitine palmitoyltransferase I (CPTI) mRNA, complete cds
12.66	481.75	352.07	246.35	85.06	271.55	88.6	205.35	16.05	118.4	5.23	217.95	19.45	X67325, class B, 20 probes, 13 in X67325cds 31-337: 7 in reverseSequence, 43-541, H.sapiens p27 mRNA
66.96	95	64.49	145.5	139.02	726.35	263.82	173.7	127.7	575.2	719.98	138.75	117.87	U28488, class A, 20 probes, 20 in U28488 1344-1847, Human putative G protein-coupled receptor (AZ3B) mRNA, complete cds
120.35	237	297.55	96.9	96.73	1276.35	1308.22	114.3	36.2	356.1	379.43	493.05	225.92	X69116, class C, 20 probes, 20 in all_X69116 2-434, H.sapiens ZNF37A gene for zinc finger protein
184.63	1097.75	276.69	432.6	208.74	1180.05	1401.13	801.55	565.19	744.55	426.03	622.4	170.55	M27281, class A, 20 probes, 20 in M27281 642-1116, Human vascular permeability factor mRNA, complete cds
81.53	166.35	48.58	1015.9	995.04	816.85	34.01	802.5	533.16	1076.2	664.96	248.8	181.3	U34044, class A, 20 probes, 20 in U34044 1143-1647, Human selenium donor protein (selD) mRNA, complete cds
1466.12	7273.85	285.74	4465	168.72	7195.3	2149.89	6071.65	78.98	7371.45	818.62	6544.8	1221.17	Adp-Ribosylation-Like Factor
105.15	648.35	260.71	829.85	241.9	565.55	307.1	1362.15	77	361.2	9.33	946.35	982.52	M36200, class C, 20 probes, 14 in all_M36200 67-428: 6 in M36200cds 194-339, Human synaptobrevin 1 (SYB1) gene
149.91	242.05	127.77	299.2	149.34	225.9	19.23	316.6	50.77	421.6	335.88	179.95	15.91	S70585, class A, 20 probes, 20 in S70585mRNA 138-612, thyroid-stimulating hormone alpha subunit [human, Genomic, 1327 nt 4 segments]
375.12	315.1	385.8	286.3	12.59	327.45	203.58	879.35	704.21	99.3	11.03	417.9	2.12	S80267, class A, 20 probes, 20 in S80267 1304-1872, p72syk {G insertion nucleotide 92} [human, Jurkat E6-1 J.CaM1 cells, mRNA Partial Mutant, 1909 nt]
95.25	118.2	143.83	199.35	239.36	208.4	121.91	591.15	718.07	323.95	368.47	400.9	270.68	X87767, class A, 20 probes, 20 in X87767exon 4-148, H.sapiens CD89 gene, exon S1. /gb=X87767 /ntype=DNA /annot=exon
181.94	813.9	164.33	1079.5	818.97	720.45	168.22	1433.1	1280.85	2390	2332.46	1587.15	586.55	X14766, class A, 20 probes, 20 in X14766mRNA 1363-1711, Human mRNA for GABA-A receptor, alpha 1 subunit

111.79	574.75	129.9	1945.75	998.36	1395.5	1063.06	769.05	219.98	1235.85	1280.08	972.95	721.04	D87024, class B, 20 probes, 11 in D87024cds#5 85-283: 9 in reverseSequence, 2475-39492, Human (lambda) DNA for immunoglobin light chain
348.89	676.6	428.65	365.5	325.13	1026.8	181.16	435	37.34	422.95	320.81	606.3	145.24	L49054, class A, 20 probes, 20 in L49054 587-1067, Homo sapiens t(3;5)(q25.1;p34) fusion gene NPM-MLF1 mRNA, complete cds.
253.29	1735.85	146.72	1381.15	871.65	1255.7	480.55	997.05	388.7	1296.95	451.2	1308.85	240.35	M94856, class A, 20 probes, 20 in M94856 163-619, Human fatty acid binding protein homologue (PA-FABP) mRNA, complete cds
101.68	907.15	788.92	455.65	358.15	581.1	384.95	385.85	283.48	279.15	71.63	576.95	503.25	M17252, class A, 20 probes, 20 in M17252mRNA 934-1198, Human cytochrome P450c21 mRNA, 3 end.
219.42	731.75	280.37	723.8	204.92	443.9	114.41	1444.25	387.99	686.65	389.12	776.05	222.67	U31248, class A, 20 probes, 20 in U31248 1663-2209, Human zinc finger protein (ZNF174) mRNA, complete cds
329.44	464.65	576.08	208.7	111.58	208.1	138.03	207.25	187.45	104	90.79	162.25	144.18	M22489, class A, 20 probes, 20 in M22489 1036-1504, Human bone morphogenetic protein 2A (BMP-2A) mRNA
3.11	590.15	146.44	287.5	142.13	1145.7	384.95	830.4	103.38	689.5	571.2	813.4	667.08	D79995, class A, 20 probes, 20 in D79995 4440-4806, Human mRNA for KIAA0173 gene, complete cds
415.57	3008.4	273.51	3946.1	587.18	2300.1	183	2733.5	135.91	1851.65	850.44	3095.9	454.81	Z35402, class A, 20 probes, 20 in Z35402mRNA 3912-4402, H.sapiens gene encoding E-cadherin, exon 3 and joined CDS
105.92	328.5	149.06	587.95	190	727.45	649.05	346.7	127	730	326.4	331.05	166.67	Transcription Factor Itf-1 - Also Represents: M31523
123.39	98.85	82.52	108.8	112.01	109.2	122.75	151.5	40.31	121.05	2.62	233.75	247.7	U53445, class A, 20 probes, 20 in U53445 2442-2928, Human ovarian cancer downregulated myosin heavy chain homolog (Doc1) mRNA, complete cds
1970.28	8711.7	2344.77	9807.5	3475.01	11538.1	9482.44	9126.25	2288.83	8766.9	1119.63	9936.2	3435.97	U94855, class A, 20 probes, 20 in U94855 696-1176, Human translation initiation factor 3 47 kDa subunit mRNA, complete cds.
462.73	1127.95	620.49	760.35	171.61	731.05	101.61	1304.35	224.51	816.3	905.1	827.3	279.31	M28983, class A, 20 probes, 20 in M28983 1823-2369, Homo sapiens interleukin 1 alpha (IL 1) mRNA, complete cds
490.02	491.5	251.45	547.7	591.28	426.6	509.54	516.95	365.08	967.3	1027.71	706.6	464.57	S71018, class A, 20 probes, 20 in S71018 282-798, cyclophilin C [human, kidney, mRNA, 883 nt]
64.28	2624.5	168.86	4416.7	2681.49	4401.2	2061.5	3747.45	328.31	2100.85	658.95	3061.85	1538.73	L11672, class B, 12 probes, 12 in L11672 3266-3562, Human Kruppel related zinc finger protein (HTF10) mRNA, complete cds, Human Kruppel related zinc finger protein (HTF10) mRNA, complete cds
34.15	1316.9	17.96	1144.25	349.1	837.85	788.49	438.55	502.82	511.6	268.98	1283.55	331	L48513, class A, 20 probes, 20 in L48513mRNA 941-1445, Homo sapiens paraoxonase 2 (PON2) mRNA, complete cds
57.42	3676.2	75.94	3818.25	875.19	3659.65	2583.98	3836.35	1323.77	4500.3	1569.78	4732.05	161.72	X93921, class C, 20 probes, 20 in all_X93921 942-1471, H.sapiens mRNA for protein-tyrosine-phosphatase (tissue type: testis)
87.47	718.45	395.34	753	734.68	846.1	696.36	513.15	258.02	623.7	56.85	902.15	711.7	U60276, class A, 20 probes, 20 in U60276 645-1191, Human hASNA-I mRNA, complete cds
659.02	3567.6	650.4	3527.55	583.72	4488.9	1443.49	3942.9	1167.72	2999.4	880.63	3857	493.7	D16581, class A, 20 probes, 20 in D16581 42-552, Human mRNA for 8-oxo-dGTPase, complete cds
601.96	2743.8	388.48	1739.95	340.75	2308	2455.07	2853.25	1871.08	2669.55	1182.35	3736.3	1326.11	X59434, class A, 20 probes, 20 in X59434mRNA 631-1129, Human rohu mRNA for rhodanese
100.48	129.05	164.12	146.65	70.07	750.85	808.29	195.75	146.87	121.85	79.55	90.3	28.28	U22816, class A, 20 probes, 20 in U22816 3784-4288, Human LAR-interacting protein 1b mRNA, complete cds

75.02	150.5	18.24	413.4	177.91	1187.8	1249.74	808.7	846.41	143.5	128.13	597.35	160.3	U86136, class A, 20 probes, 20 in U86136 8094-8472, Human telomerase-associated protein TP-1 mRNA, complete cds
49.36	186.05	24.54	204.5	139.72	588.05	478.64	478.2	441.66	299.5	83.72	410.7	263.61	X83441, class A, 20 probes, 20 in X83441mRNA 2724-3102, H.sapiens mRNA for DNA ligase IV
107.98	566.8	133.22	591.75	499.29	757.05	496.6	561.45	282.91	375.9	181.87	194.45	0.07	J03798, class A, 20 probes, 20 in J03798 1026-1536, Human autoantigen small nuclear ribonucleoprotein Sm-D mRNA, complete cds
178.19	159.3	134.07	187.65	138.81	209	95.32	488.6	61.8	248.75	4.74	359.3	81.18	Y09615, class B, 20 probes, 14 in Y09615cds 891-1131: 6 in reverseSequence, 1268-1472, H.sapiens mRNA for mitochondrial transcription termination factor
128.62	229.9	84.29	576.3	438.12	1689.75	2349.79	409.7	216.52	86.3	60.39	218.35	121.69	Y13115, class A, 20 probes, 19 in Y13115cds 2477-2879: 1 in reverseSequence, 3061, Homo sapiens mRNA for serine/threonine protein kinase SAK
35.43	261.05	110.1	605.9	652.24	182.9	53.03	508.85	521.35	246.45	239.78	123.85	124.66	S73205, class A, 20 probes, 20 in S73205 2183-2573, insulin activator factor [human, pancreatic insulinoma, mRNA Partial, 2622 nt]. /gb=S73205 /ntype=RNA
60.81	790	113	319.45	38.96	514.65	402.27	392.45	421.51	162.1	122.05	113.6	50.49	D50919, class A, 20 probes, 20 in D50919 3864-4404, Human mRNA for KIAA0129 gene, complete cds
311.98	559.75	210.51	537.3	324.28	143.85	33.02	809.5	724.22	709	387.49	488.65	178.26	D83776, class A, 20 probes, 20 in D83776 4596-5166, Human mRNA for KIAA0191 gene, partial cds
48.79	242.9	33.38	450.75	274.43	137.4	47.66	90.15	48.86	250.65	91.29	633.75	263.96	D49958, class A, 20 probes, 20 in D49958 1830-2346, Human fetus brain mRNA for membrane glycoprotein M6, complete cds
4005.83	8795.8	4526.47	8644.45	3449.2	5453.9	3569.62	5889	2328.93	6619.3	2456.63	4139.95	518.95	Calmodulin Type I
303.49	473.5	301.51	320.15	36.42	261.1	180.88	530.95	28.5	803.85	212.63	420.45	281.64	U12140, class A, 20 probes, 20 in U12140 3433-3673, Human tyrosine kinase receptor p145TRK-B (TRK-B) mRNA, complete cds
24.04	105.3	30.12	144.3	3.11	75.2	23.62	139	10.61	362	328.1	447	348.04	X15088, class C, 20 probes, 20 in all_X15088 1146-1276, Human GNAT1 mRNA for transducin alpha-chain
307.59	249.85	303.84	269.25	339.34	200.6	207.89	80.3	48.93	100.8	96.03	64.45	44.62	S67970, class A, 20 probes, 20 in S67970 962-1538, ZNF75=KRAB zinc finger [human, lung fibroblast, mRNA, 1563 nt]
19.87	111.4	1.98	260.6	208.17	267.8	299.81	197.05	1.2	319.05	217.15	253.65	48.72	Y10812, class A, 20 probes, 20 in Y10812 767-1253, H.sapiens mRNA for fructose-bisphosphatase
711	669.2	674.72	829.7	662.7	608.1	75.8	69.7	51.9	131.15	35.43	720.5	160.09	U46767, class A, 20 probes, 20 in U46767 288-768, Human monocyte chemoattractant protein-4 precursor (MCP-4) mRNA, complete cds
385.23	628.5	217.36	1114.2	79.2	1323.5	579.83	739.5	401.92	1424.4	781.07	959.4	205.77	Carboxyl Methyltransferase, Aspartate, Alt. Splice 1 - Also Represents: D13892
415.21	398.45	445.55	163.45	79.41	541.9	84.85	159.15	2.19	172.8	26.87	120.7	0.28	D79988, class A, 20 probes, 20 in D79988 6383-6899, Human mRNA for KIAA0166 gene, complete cds
602.38	999.8	654.07	1332.6	63.64	305.4	128.27	838.45	732.21	589.4	416.63	375.5	197.99	U61397, class A, 20 probes, 20 in U61397 875-1173, Human ubiquitin-homology domain protein PIC1 mRNA, complete cds
0.42	141.1	59.68	169.65	62.15	397.8	223.73	69.1	47.23	118.35	133.43	592.05	59.33	Y07566, class C, 20 probes, 20 in all_Y07566 951-1066, H.sapiens mRNA for RIT protein
142.27	849.5	386.79	286.9	56.71	903.55	280.93	711.55	652.02	596.2	300.38	627	597.08	M33493, class A, 20 probes, 20 in M33493 504-792, Human tryptase-III mRNA, 3 end
98.22	218.6	103.38	168.8	51.19	177.5	111.86	821.2	863.66	159.2	59.26	331.35	132.3	X06825, class C, 20 probes, 20 in all_X06825 679-1154, Human mRNA for skeletal beta-tropomyosin

41.01	69.35	44.9	759.7	989.38	184.75	126.5	158.1	54.31	238.9	69.01	205.65	136.4	Z78291, class A, 20 probes, 20 in Z78291 28-223, H.sapiens mRNA (clone 1D8).
107.2	285.75	293.1	331.1	87.82	367.3	174.23	424.7	85.56	143.2	94.61	496.6	183.28	X79888, class C, 20 probes, 20 in all_X79888 965-1500, H.sapiens AUH mRNA
168.64	1556.7	924.19	1145.5	512.79	615.2	320.32	1445.9	1217.35	1112.85	695.72	517.5	161.22	M63391, class A, 20 probes, 20 in M63391mRNA 1637-2189, Human desmin gene, complete cds
1112.77	1673.35	794.15	1882.55	1536.19	2806.75	2504.93	1557.4	666.94	1904.8	911.74	1885.85	1227.04	X59812, class C, 20 probes, 20 in all_X59812 1586-2025, H.sapiens CYP 27 mRNA for vitamin D3 25-hydroxylase
356.45	722.5	33.23	536.25	465.06	663.65	95.53	1159.15	744.65	985.3	863.09	1690.1	283.41	M99063, class A, 20 probes, 20 in M99063 1942-2452, Homo sapiens cytokeratin 2 mRNA, complete cds
225.85	1414.35	146.58	1306.5	711.35	707.6	457.22	1349.75	441.45	669.25	576.5	914.5	205.34	M96954, class A, 19 probes, 19 in M96954 806-1313, Homo sapiens nucleolysin TIAR mRNA, complete cds
316.43	351.7	217.22	1135.1	1188.79	259.6	212.41	673.45	715.52	784	708.38	686.1	687.02	M37190, class A, 20 probes, 20 in M37190mRNA 1220-1796, Human ras inhibitor mRNA, 3 end
178.05	781.65	283.62	746.9	349.59	229.5	179.32	444.15	192.97	855.55	1022.41	903	327.39	U28687, class A, 20 probes, 20 in U28687 1710-1983, Human zinc finger containing protein ZNF157 (ZNF157) mRNA, complete cds
408.99	539.45	626	207.3	8.49	396.6	31.11	496	211.85	706.5	791.68	398.95	205.98	U45880, class A, 20 probes, 20 in U45880 1969-2515, Human X-linked inhibitor of apotosis protein XIAP mRNA, complete cds
516.47	996.35	497.87	1122.1	423.7	1128.3	374.48	697.85	143.05	1198.5	92.07	1075.1	31.68	X82103, class B, 20 probes, 13 in X82103cds 660-840: 7 in reverseSequence, 954-1128, H.sapiens mRNA for beta-COP
55.01	189.4	102.67	108.5	32.39	98.4	105.78	39.5	30.41	53.55	48.3	45.5	5.37	L10717, class A, 20 probes, 20 in L10717 6303-6332, Homo sapiens T cell-specific tyrosine kinase mRNA, complete cds
869.88	3029.55	659.52	4115.45	2011.65	2226.7	2046.51	3046.65	1445.4	2485.1	1270.67	2058.5	1166.02	J00277, class C, 20 probes, 20 in all_J00277 3607-3724, Human (genomic clones lambda-[SK2-T2, HS578T]; cDNA clones RS-[3,4, 6]) c-Ha-ras1 proto-oncogene, complete coding sequence
73.54	246.05	24.82	277.15	242.33	419.85	388.98	244.1	71.98	248.2	6.36	246	54.87	D87463, class A, 20 probes, 20 in D87463 2511-2997, Human mRNA for KIAA0273 gene, complete cds
107.69	214.9	47.09	821.55	674.08	593.4	97.02	203.95	42.07	356.55	267.64	263.3	63.92	X90828, class A, 20 probes, 20 in X90828exon#2 233-752, H.sapiens mRNA for transcription factor, Lbx1
502.61	1864.8	412.1	1792.95	841.67	2570.75	2125.07	1199.45	851.43	2727.5	459.9	2102.6	1492.7	Peptide Yy - Also Represents: D13897_rna2
466.27	1348.3	53.46	1087.45	52.4	1087.8	358.93	893.1	208.88	1221.6	104.23	1318.95	289.98	U77948, class A, 20 probes, 20 in U77948 2714-3236, Human Bruton tyrosine kinase-associated protein-135 mRNA, complete cds.
182.93	684.65	454.88	769.95	461.81	684.35	748.19	1103.3	1305.6	1023.25	538.74	998.85	1092.41	S69369, class A, 20 probes, 20 in S69369 779-1115, PAX3A=transcription factor [human, adult cerebellum, mRNA, 1248 nt]
229.39	278.5	135.62	436.2	327.39	546.8	145.66	678.45	16.19	340.8	299.95	90.1	26.3	X15306, class A, 20 probes, 20 in X15306mRNA 3269-3707, H.sapiens NF-H gene, exon 1 (and joined CDS)
39.1	90.25	75.73	357.9	175.22	184.45	96.94	419.1	31.96	192.95	28.5	804.05	175.01	Gtp Cyclohydrolase I, Alt. Splice 1 - Also Represents: U19523
378.37	550.65	554.44	595.8	207.32	295.55	258.16	911.2	202.94	329.05	273.72	536.9	85.42	X92689, class A, 20 probes, 20 in X92689cds 1457-1853, H.sapiens mRNA for UDP-GalNAc:polypeptide N-acetylgalactosaminyl transferase. /gb=X92689 /ntype=RNA

587.82	1271.1	933.1	740.05	112.92	677.5	178.05	796.5	160.65	439.3	155.85	703.45	7.85	AF007551, class A, 20 probes, 20 in AF007551 16-574, Homo sapiens Bet1p homolog (hbet1) mRNA, complete cds. /gb=AF007551 /ntype=RNA
472.21	4168.2	172.68	2631.8	1023.89	2737.75	1645.79	2849.85	186.32	1643.15	975.74	3425.9	1186.95	X59798, class C, 20 probes, 20 in all_X59798 3705-4192, Human PRAD1 mRNA for cyclin
0.99	405.1	45.82	168.1	102.67	1139.1	450.29	270.55	70.22	227.6	79.9	310.6	192.33	X66503, class C, 20 probes, 20 in all_X66503 1125-1690, Human adenylosuccinate synthetase mRNA
463.58	1163.15	77.43	1530.45	402.41	850.35	839.27	1764	553.1	1054.55	198.34	1397.2	237.45	D31891, class A, 20 probes, 20 in D31891 3786-4275, Human mRNA for KIAA0067 gene, complete cds
34.72	1312.3	376.46	1263.5	841.46	594.1	160.23	874	153.58	1298.05	43.2	1160.85	236.39	L40391, class A, 20 probes, 20 in L40391mRNA 889-1435, Homo sapiens (clone s153) mRNA fragment
21.14	297.8	183.99	247.6	95.32	628.95	69.23	561.85	45.33	654.05	447.67	396.55	249.82	J02854, class A, 20 probes, 20 in J02854 531-1089, Human 20-kDa myosin light chain (MLC-2) mRNA, complete cds
198.63	572.45	139.79	837.1	354.12	793.5	278.32	637	252.72	781.9	467.26	808.75	470.44	L76703, class A, 20 probes, 20 in L76703mRNA 2687-3245, Homo sapiens B56epsilon mRNA, complete cds
180.81	345.3	40.73	159.3	10.47	1351	1643.46	947.8	1031.53	241.2	116.53	310.65	257.88	U82306, class A, 20 probes, 20 in U82306 135-225, Human unknown protein mRNA, partial cds. /gb=U82306 /ntype=RNA
1406.29	1554.85	1431.54	2005.8	998.29	853.3	291.75	1994.8	237.73	2507	526.37	1024.9	1057.69	M80899, class A, 20 probes, 20 in M80899 3582-4002, Human novel protein AHNAK mRNA, partial sequence
3939.08	18860.9	4319.29	19994.15	3828.63	19277.95	16196.21	23235.95	7856.17	18064.95	13092.29	24923.5	9215.44	U03057, class A, 20 probes, 20 in U03057 2172-2724, Human actin bundling protein (HSN) mRNA, complete cds
464.64	992.2	482.67	1237.7	757.59	2393.35	1765.43	449.05	381.2	1350.15	760.35	1278.45	556.99	M29581, class A, 20 probes, 20 in M29581 1469-1853, Human zinc-finger protein 8 (ZFP8) mRNA, 3 end
64.49	92.55	56.36	270.65	341.18	115.15	100.76	324.8	345.07	123.25	6.29	132.4	78.49	AB005535, class A, 20 probes, 20 in AB005535 1056-1635, Homo sapiens mRNA for Clock, partial cds. /gb=AB005535 /ntype=RNA
119.5	732.75	93.97	494.1	268.56	522.15	128.62	647.55	26.09	682.85	104.72	420.6	86.83	D80006, class A, 20 probes, 20 in D80006 4068-4596, Human mRNA for KIAA0184 gene, partial cds
2379.27	3617.65	1558.11	3186.75	1059.88	3956.3	2311.53	2226.05	481.89	3470.35	67.53	3981	2249.17	D30755, class A, 20 probes, 20 in D30755 1189-1633, Human mRNA for KIAA0113 gene, partial cds
640.64	1022.25	759.5	967.4	824.63	1133.95	814.09	869.8	959.4	601.65	381.2	1146.45	569.43	V00563, class A, 20 probes, 20 in V00563mRNA 19-127, Human gene for immunoglobulin mu, part of exon 8.
162.07	289.2	230.94	310.7	99.28	2129.55	2707.44	225.95	51.12	1042.8	366.85	607.4	676.84	X76059, class B, 20 probes, 15 in X76059cds 1208-1424: 5 in reverseSequence, 1709-1807, H.sapiens mRNA for YRRM1
47.38	74.85	42.36	571.95	742.39	114.3	104.79	111.15	114.2	273.4	207.32	474.15	575.09	U90544, class A, 20 probes, 20 in U90544 1665-2145, Human sodium phosphate transporter (NPT3) mRNA, complete cds
244.8	1322.85	343.72	1676.25	251.09	2797.45	756.68	1907.25	377.1	2184.2	637.53	1923.8	142.69	U34962, class A, 20 probes, 20 in U34962 1074-1560, Human transcription factor HCSX (hCsx) mRNA, complete cds
68.94	124.55	0.92	82.05	60.17	276	221.47	79.4	36.63	178.65	37.26	123	64.49	S83549, class A, 20 probes, 20 in S83549 19-571, Na+/H+ exchanger isoform NHE-2 [human, various tissues, mRNA Partial, 595 nt]. /gb=S83549 /ntype=RNA
154.57	829.05	111.51	1082.65	635.19	2706.9	2562.27	1325.7	724.22	886.15	254.63	1050.35	757.24	U78628, class A, 20 probes, 20 in U78628 7-199, Human leukemia inhibitory factor receptor mRNA, 5 untranslated region. /gb=U78628 /ntype=RNA
851.99	935.05	548.08	253.55	216.3	530.75	171.33	609.65	300.17	471.9	401.35	266.5	97.58	U90336, class A, 20 probes, 20 in U90336 2606-3074, Human PEG3 mRNA, partial cds.

3.54	120.65	22.98	310	261.77	281.95	145.45	340.55	86.62	349.55	280.93	366.55	147.29	L32606, class A, 20 probes, 20 in L32606 1862-2196, Human homeobox-like mRNA
356.24	3185.55	65.27	2280.8	113.84	1752.45	856.24	3241.85	641.28	3384.15	448.94	3160.2	483.52	U96629, class A, 20 probes, 20 in U96629mRNA#2 3194-3722, 2A8.2 gene (unknown protein CIT987SK_2A8_1) extracted from Human chromosome 8 BAC clone CIT987SK-2A8 complete sequence., 2A8.2 gene (unknown protein CIT987SK_2A8_1) extracted from Human chromosome 8 BAC clone CIT987SK-2A8 complete sequence.
286.8	753.4	237.87	1327.05	1335.09	934.85	510.88	871.75	124.24	1071.05	976.59	1208.55	244.02	D63813, class A, 20 probes, 20 in D63813 1154-1676, Human mRNA for rod photoreceptor protein, complete cds
25.1	108.8	23.48	254.35	237.94	312.65	239.92	176.9	111.44	141	88.39	107.5	21.07	U94333, class A, 20 probes, 20 in U94333 2877-3429, Human Clq/MBL/SPA receptor C1qR(p) mRNA, complete cds.
722.03	4490.8	1353.83	4389.7	8.2	3151.55	437.49	4442.75	259.3	2571	2714.58	3989.15	6.72	X80198, class C, 20 probes, 20 in all_X80198 1398-1939, H.sapiens MLN64 mRNA
62.23	1378.6	83.3	1298.95	237.38	681.9	180.74	1863.3	295.71	888.4	719.55	1625.15	228.75	U21931, class A, 20 probes, 20 in U21931mRNA 965-1373, Human fructose-1,6-biphosphatase (FBP1) gene
11.17	30.15	29.63	28.45	20.29	67.95	59.75	42.15	1.91	172.6	221.32	27.6	4.67	U67092, class C, 6 probes, 3 in all_U67092 1093-1868: 3 not in GB record, Human ataxia-telangiectasia locus protein (ATM) gene, exons 1a, 1b, 2, 3 and 4, partial cds. /gb=U67092 /ntype=DNA /annot=exon, Human ataxia-telangiectasia locus protein (ATM) gene, exons 1a, 1b, 2, 3 and 4, partial cds. /gb=U67092 /ntype=DNA /annot=exon
136.47	522.25	121.55	221.3	181.59	351.7	215.95	561.25	637.6	735.4	38.47	558.1	36.91	X05608, class A, 20 probes, 17 in X05608exon#4 172-406: 3 not in GB record, Human gene for neurofilament subunit NF-L
183.49	193.05	211.35	339.9	305.19	249.75	271.74	259.45	194.24	484.3	364.3	211.25	196.5	U66561, class A, 20 probes, 20 in U66561 2158-2614, Human kruppel-related zinc finger protein (ZNF184) mRNA, partial cds
213.69	247.25	271.03	792.9	521.7	1107.65	505.79	1027.05	103.17	495.65	199.05	753.65	113.21	M32011, class A, 20 probes, 20 in M32011mRNA 1623-2157, Human neutrophil oxidase factor (p67-phox) mRNA, complete cds
181.8	52.5	39.32	42.1	36.91	207.35	84.64	545.1	464.14	147.95	60.6	103.95	85.35	M20642, class A, 20 probes, 20 in M20642mRNA 369-898, Human alkali myosin light chain 1 mRNA, complete cds
573.96	1423.85	1047.01	2735.4	1332.9	990.8	639.65	961.7	500.91	1623.1	425.4	2656.75	130.89	Cell Division Cycle Protein 2-Related Protein Kinase (Pisslre) - Also Represents: X78342
161.08	4994.4	77.64	4228.8	3468.36	3098.05	1912.79	2994.7	763.96	2380.55	452.48	3268.9	608.82	X13546, class A, 20 probes, 20 in X13546mRNA 657-1137, put. HMG-17 protein gene extracted from Human HMG-17 gene for non-histone chromosomal protein HMG-17
50.06	180.7	38.18	453.05	382.47	1458.3	1713.6	232.3	82.02	735.35	809.14	407.9	80.61	X62744, class B, 20 probes, 12 in X62744cds 469-745: 8 in reverseSequence, 814-1018, Human RING6 mRNA for HLA class II alpha chain-like product
204.21	210.05	195.37	77.4	13.29	201.6	195.73	95.8	0.42	211.05	74.32	120.35	42.64	U71092, class B, 20 probes, 13 in U71092cds 954-1158: 7 in reverseSequence, 1611-1767, Human somatostatin receptor-like protein (SLC1) gene, complete cds

202.66	1147.65	740.84	207.7	112.43	755.3	728.32	896.85	354.61	881.35	604.22	369.5	166.17	U24685, class B, 20 probes, 10 in U24685cds 123-336: 10 in reverseSequence, 420-447, Human anti-B cell autoantibody IgM heavy chain variable V-D-J region (VH4) gene, clone E11, VH4-63 non-productive rearrangement
2225.9	8321.05	2145.43	6959.35	4692.86	11032.7	6860.35	4774.1	513.78	7966.5	1802.13	8812.1	3560.57	Heat Shock Protein, 70 Kda (Gb:Y00371)
285.32	461.25	308.23	558.8	630.74	1209.65	148.14	1069.7	816	958.1	548.86	780.45	561.8	U40571, class A, 20 probes, 20 in U40571 1695-2073, Human alpha1-syntrophin (SNT A1) mRNA, complete cds
502.05	406.15	174.73	1106.65	560.52	184.6	37.19	354.1	246.36	583.6	594.82	787.2	338.14	X85753, class A, 20 probes, 20 in X85753 1268-1646, H.sapiens mRNA for CDK8 protein kinase
53.32	336.6	353.84	1022.9	769.19	830.4	806.67	738.35	141.35	1111.3	1151.31	944.85	401.42	L33799, class A, 20 probes, 20 in L33799 1065-1455, Human procollagen C-proteinase enhancer protein (PCOLCE) mRNA, complete cds
259.86	1683.8	1110.02	1983.4	514.77	2012.7	780.36	2683.65	656.69	2396.65	716.94	2360.95	119.29	X53390, class C, 20 probes, 20 in all_X53390 2839-3093, Human mRNA for upstream binding factor (hUBF)
38.04	119.8	80.75	69.1	6.79	108.05	65.41	299.25	281.07	559.35	312.61	72.35	26.8	U17032, class A, 20 probes, 20 in U17032 4391-4961, Human p190-B (p190-B) mRNA, complete cds
93.83	714.85	15.91	861.6	197.71	359.75	299.46	438.55	404.96	1373.7	689.43	346.55	51.55	M63838, class A, 20 probes, 20 in M63838 2097-2643, Human interferon-gamma induced protein (IFI 16) gene, complete cds
149.13	210.9	36.2	166.75	122.82	411.1	402.49	188.85	104.44	452.05	231.86	536.1	517.74	U41737, class A, 20 probes, 20 in U41737 103-523, Human pancreatic beta cell growth factor (INGAP) mRNA, complete cds. /gb=U41737 /ntype=RNA
2934.21	25002.65	4439	22801.9	7926.1	17921.45	11308.69	24587.05	2896.24	22248.35	5506.45	21457.25	3753.82	X95404, class C, 20 probes, 20 in all_X95404 482-1047, H.sapiens mRNA for non-muscle type cofilin
117.87	721.7	0.85	1243.15	364.23	451.1	181.44	1035.95	1033.01	713.55	914.64	187.55	123.39	X68487, class C, 20 probes, 20 in all_X68487 1174-1667, H.sapiens mRNA for A2b adenosine receptor
103.8	160.5	200.25	625.25	773.22	1440.95	1981.67	82.5	45.82	929.65	916.06	180.75	61.59	M29474, class A, 20 probes, 20 in M29474mRNA 5986-6490, Human recombination activating protein (RAG-1) gene, complete cds
7730.44	31449.5	3582.35	37317.9	8782.83	23630.5	16041.28	25290.15	4833.99	19574.55	1306.52	28136.95	4599.8	D78361, class A, 20 probes, 20 in D78361 504-942, Human mRNA for ornithine decarboxylase antizyme, ORF 1 and ORF 2
193.46	138.4	52.18	240.05	82.52	1321.55	1038.95	278.55	48.01	1063.15	776.05	991.15	3.04	Z29574, class A, 20 probes, 17 in Z29574exon#3 1-373: 1 in reverseSequence, 3141: 2 not in GB record, Homo sapiens gene for BCMA peptide
1260.42	1487.15	408.92	1276.3	1184.69	1178.35	733.48	1446.3	368.26	2116.1	1220.18	1399.15	85.35	M63180, class A, 20 probes, 20 in M63180 2055-2505, Human threonyl-tRNA synthetase mRNA, complete cds
757.45	14931.05	172.6	12668.9	3412.64	9848.85	5979.93	11573.3	4023.86	9945.7	6161.3	13793.65	2180.65	M84332, class A, 20 probes, 16 in M84332exon#4 764-1226: 4 in reverseSequence, 2337-2397, Human ADP-ribosylation factor 1 gene
453.96	866.3	537.12	1069.85	440.03	695.15	664.75	821.15	344.57	1462.8	0.71	463.2	433.6	U23803, class A, 20 probes, 20 in U23803 1125-1659, Human heterogeneous ribonucleoprotein A0 mRNA, complete cds
28.5	1465.25	328.17	1104.35	1370.73	868.75	845.49	2212.9	1209.58	871.55	627.7	1600.45	1557.12	U08998, class A, 20 probes, 20 in U08998 962-1280, Human TAR RNA binding protein 2 (TRBP2) mRNA, complete cds
5697.94	76693.6	12218.1	83413.45	4395.73	80388.95	70031.51	83954.05	18488.09	79581.05	16419.23	104034.6	2882.17	X03342, class A, 20 probes, 18 in X03342cds 51-375: 2 in reverseSequence, 439-445, Human mRNA for ribosomal protein L32

347.33	978.6	450.29	809.6	383.25	1011.55	174.44	1422.35	641.84	1258.9	647.14	822.35	284.75	U18543, class A, 20 probes, 20 in U18543 1853-2339, Human zinc-finger protein mRNA, complete cds
205.77	1011.45	382.76	1534.9	806.81	1472.85	974.61	1232.75	1064.55	1422.3	1328.65	1956.8	943.28	L05624, class A, 20 probes, 20 in L05624 910-1418, Homo sapiens MAP kinase kinase mRNA, complete cds
48.93	452.95	3.18	864.25	727.54	121.8	8.63	305.1	216.52	998.6	934.51	359.8	89.1	U14518, class A, 20 probes, 20 in U14518 859-1315, Human centromere protein-A (CENP-A) mRNA, complete cds
197.21	888.5	661.99	1124.1	1042.13	1926.25	2303.26	843.35	843.22	1433.1	928.01	1211.7	577.71	U67369, class A, 20 probes, 20 in U67369 2300-2720, Human growth factor independence-1 (Gfi-1) mRNA, complete cds
4.6	178.6	51.48	733.85	605.5	190.95	68.52	173.7	92.77	178.1	174.66	615.85	430.98	J04621, class A, 20 probes, 20 in J04621mRNA 2879-3347, Human heparan sulfate proteoglycan (HSPG) core protein, 3 end
76.3	84.25	66.11	259.1	315.37	139.7	91.92	142.5	126.01	180.6	106.91	201	37.05	Z23091, class C, 20 probes, 20 in all_Z23091 6853-7358, H.sapiens GPV gene encoding platelet glycoprotein V precursor
564.77	252.65	81.81	683.1	229.1	294.05	158.46	736.2	578.55	750.45	93.13	481.25	178.4	X83378, class C, 20 probes, 20 in all_X83378 4940-5523, H.sapiens mRNA for putative chloride channel
20.72	104.5	47.52	586.1	576.57	141.55	6.43	303.9	90.37	409.1	353.69	513.4	507.28	Major Histocompatibility Complex, Class Ii Beta W52
86.13	153.45	19.16	102.05	81.11	180.5	14.42	571.45	685.82	190.2	205.2	227.55	165.25	U24577, class A, 20 probes, 20 in U24577 1182-1512, Human LDL-phospholipase A2 mRNA, complete cds
54.94	360.2	97.86	480.65	142.91	356.7	174.23	247	100.97	1737.35	1737.86	797.85	909.98	X54489, class A, 20 probes, 20 in X54489mRNA 660-1034, Human gene for melanoma growth stimulatory activity (MGSA)
263.89	53.55	27.79	505.65	123.81	185.7	109.04	89.1	27.72	203.85	224.79	127.7	44.12	X14474, class A, 16 probes, 16 in X14474cds 669-710, Human mRNA for microtubule-associated tau protein
338.14	2827.4	408.99	3101.9	1220.32	1933.15	1968.8	1604.6	304.48	2396.35	8.84	2490.15	114.62	L38810, class A, 20 probes, 20 in L38810mRNA 706-1246, Homo sapiens thyroid receptor interactor (TRIP1) mRNA, complete cds
106.77	154.85	33.59	348.05	439.18	562.8	155.42	199.9	116.39	344.6	367.27	84.25	32.6	L34409, class A, 20 probes, 20 in L34409 624-990, Homo Sapiens (clone B3B3E13) chromosome 4p16.3 DNA fragment
56	97.7	32.67	174.85	170.48	207.75	181.51	151.95	52.4	613.65	697.99	85.15	15.49	X54457, class C, 17 probes, 17 in all_X54457 2328-2416, Human mRNA for bile-salt-stimulated lipase (BSSL) (EC 3.1.1.3).
88.25	508.5	223.59	249.6	29.7	1399.15	1529.83	294.2	133.08	673.5	59.4	837.45	702.37	U52154, class A, 20 probes, 20 in U52154 2352-2610, Human G protein-coupled inwardly rectifying potassium channel Kir3.4 mRNA, complete cds
684.41	765	635.26	863.5	960.25	389.3	101.26	459.65	439.04	346.45	96.66	133.35	46.88	D42055, class A, 20 probes, 20 in D42055 5178-5718, Human mRNA for KIAA0093 gene, partial cds
11.03	545.85	95.11	935.5	458.35	982.85	71.06	967.85	372.29	1001.6	619.28	942.6	96.31	U93867, class A, 20 probes, 20 in U93867 1248-1764, Human RNA polymerase III subunit (RPC62) mRNA, complete cds. /gb=U93867 /ntype=RNA
9309.49	13145.4	5640.87	15906.75	9521.97	10238.5	8481.46	11052.7	3517.01	8346.5	2238.98	10641.95	637.88	U46025, class A, 20 probes, 17 in U46025cds 2254-2710: 3 in reverseSequence, 2777-2843, Human translation initiation factor eIF-3 p110 subunit gene, complete cds
768.34	2352.7	390.75	2096.9	1347.75	989.7	207.47	1146.55	900.5	2573.8	1196.85	1419.85	543.55	U51333, class A, 20 probes, 20 in U51333 2437-3005, Human hexokinase III (HK3) mRNA, complete cds
266.3	310	386.36	494.4	514.92	305.2	232.36	270.35	289.7	98.05	48.44	430.2	146.65	X57025, class A, 20 probes, 20 in X57025mRNA 6635-7151, Human IGF-I mRNA for insulin-like growth factor I

37.97	59.9	1.56	386.9	207.47	175.75	65.27	720.45	877.73	534.75	603.09	593.75	205.7	Y00067, class A, 20 probes, 20 in Y00067mRNA 2655-3207, Human gene for neurofilament subunit M (NF-M)
69.79	81.85	91.99	469.5	636.96	413.8	150.76	99.7	61.38	792.8	1012.15	197.8	49.64	U28368, class A, 20 probes, 20 in U28368 841-1249, Human Id-related helix-loop-helix protein Id4 mRNA, complete cds
482.18	1498.55	594.61	1345	120.77	1835.95	400.86	2096.45	233.13	1769.5	658.46	1916.9	1006.92	Insulin-Like Growth Factor 2
4.95	178	15.41	251.35	194.1	693.95	299.88	258.05	137.67	312.6	68.31	1006.2	317.92	U49742, class B, 20 probes, 12 in U49742cds 744-984: 8 in reverseSequence, 5287-5473, Human rhodopsin gene, complete cds
119.22	732.2	41.3	1088.1	647.85	116.9	39.17	516	626.64	1224.1	779.66	705.9	768.06	Zinc Finger Protein, Kruppel-Like
73.4	326.7	83.58	1215.75	1348.38	507.9	195.3	420.2	120.21	292.45	204.85	394.1	163.06	J03890, class A, 20 probes, 19 in J03890mRNA#1 482-1022: 1 not in GB record, SP-C1 gene (pulmonary surfactant protein SP-C) extracted from Human pulmonary surfactant protein C (SP-C) and pulmonary surfactant protein C1 (SP-C1) genes, complete cds
80.89	205.85	26.8	292.4	213.83	267.05	62.01	425.1	153.87	184.05	60.6	575.85	172.46	D84239, class A, 20 probes, 20 in D84239 15949-16339, Human mRNA for IgG Fc binding protein, complete cds
96.52	873.6	73.4	914.95	632.22	464.95	87.47	165.7	157.54	331.7	323.15	765.85	14.92	S72008, class A, 20 probes, 20 in S72008 1731-2229, hCDC10=CDC10 homolog [human, fetal lung, mRNA, 2314 nt]
32.67	51.3	40.45	421.75	541.15	553.95	315.3	642.7	852.63	667.45	745.64	357.3	287.65	U10492, class A, 20 probes, 20 in U10492 1894-2266, Human Mox1 protein (MOX1) mRNA, complete cds
160.44	335.55	120.84	210.65	111.37	954.45	1274.28	330.95	336.79	255.8	103.1	252.55	155.78	U20350, class A, 20 probes, 20 in U20350 2697-3045, Human G protein-coupled receptor V28 mRNA, complete cds
112.15	7091	2694.5	6956.2	3756.72	18985.6	16708.79	6669.95	1151.66	12301.65	1759.07	10166.95	5813.34	AB001325, class A, 20 probes, 20 in AB001325 967-1387, Human AQP3 gene for aquaporine 3 (water channel), partail cds.
219.06	734.05	127.07	1097.7	122.05	444.5	487.76	1021.4	174.51	986.95	80.54	1243.65	773.65	M64174, class A, 20 probes, 20 in M64174 3012-3468, Human protein-tyrosine kinase (JAK1) mRNA, complete cds
1927.64	10157.35	3978.68	10799.6	2425.09	7509.55	5923.5	7286.15	593.62	8209.15	2119.69	7720.3	2666.36	V00572, class C, 20 probes, 20 in all_V00572 1364-1731, Human mRNA encoding phosphoglycerate kinase
11.6	186.6	17.25	113.1	19.09	197.15	54.52	346.65	99.49	240	80.75	164.7	33.66	U90304, class A, 20 probes, 20 in U90304 1272-1782, Human iroquois-class homeodomain protein IRX-2a mRNA, complete cds.
8.34	2131.75	287.72	2246.1	18.1	913.95	1042.2	2133.45	234.55	2029.7	247.06	1857.15	296.49	X83425, class C, 20 probes, 20 in all_X83425 1831-2390, H.sapiens LU gene for Lutheran blood group glycoprotein
73.26	151.55	53.1	140.7	55.72	482.4	438.26	142.3	45.11	210.2	81.74	343.5	21.5	S62696, class A, 19 probes, 19 in S62696 39-119, EBV/C3d receptor {alternatively spliced, exons 8a,9,10} [human, Jurkat T cells, mRNA Partial, 151 nt]
17.75	75.95	7.57	299.6	365.57	311	274.5	173.55	30.19	226.05	150.12	227.95	133.57	U10473, class A, 20 probes, 20 in U10473 333-895, Human clone p4betaGT/3 beta-1,4-galactosyltransferase mRNA, partial cds. /gb=U10473 /ntype=RNA
37.41	317.85	54.66	359.7	173.52	405.6	259.51	329.15	199.19	256.05	42.36	360.45	17.18	U43959, class A, 20 probes, 20 in U43959 758-815, Human beta 4 adducin mRNA, alternatively spliced partial cds

476.02	2426.6	650.4	2880.25	1287.99	3324.5	2101.38	1937.85	516.26	2969.15	834.6	2848.75	1167.22	D13146, class A, 20 probes, 20 in D13146mRNA#1 2083-2551, 2 ,3 -cyclic-nucleotide 3 -phosphodiesterase gene extracted from Human 2 ,3 -cyclic-nucleotide 3 -phosphodiesterase gene
81.88	325.7	79.05	288	197.99	783.6	163.62	455.8	159.52	79.2	59.4	327	127.28	D50927, class A, 20 probes, 20 in D50927 3955-4411, Human mRNA for KIAA0137 gene, complete cds
525.52	425	532.17	316.65	13.51	308.1	26.02	163.9	69.72	121.85	54.52	192.8	68.31	M31651, class B, 20 probes, 11 in M31651cds 900-1146: 9 in reverseSequence, 5912-6098, Human human sex hormone-binding globulin (SHBG) gene, complete cds
351.64	1075.7	539.52	1528.4	437.13	1301.35	329.72	2305.8	1571.76	2339.45	1374.54	921.2	560.74	M16424, class C, 20 probes, 20 in all_M16424 135-664, Human beta-hexosaminidase alpha chain (HEXA) gene
71.21	181.45	48.86	132.25	17.18	515.4	112.01	266.15	4.74	476.4	458.77	953.4	1122.74	U12259, class B, 19 probes, 14 in U12259cds 492-808: 5 in reverseSequence, 16275-16368, Human paired box homeotic protein (PAX3) gene
55.08	75.75	59.47	110.1	8.77	120.6	65.34	117.9	52.75	125.05	40.09	121.4	7.21	X93512, class B, 20 probes, 11 in X93512cds 61-157: 9 in reverseSequence, 175-211, H.sapiens mRNA for telomeric DNA binding protein (orf2)
8456.43	82597.3	16393.7	86418.4	2998.7	86321.95	30033.44	103089.45	12926.55	80835.7	29213.83	106112.6	4048.75	X67247, class A, 20 probes, 20 in X67247mRNA 116-662, H.sapiens rpS8 gene for ribosomal protein S8
858.85	16273.8	1250.17	13505.85	5539.4	9196.25	5104.82	13302.55	6456.8	9851.4	6756.83	11555.6	832.12	U73377, class A, 20 probes, 20 in U73377 3059-3363, Human p66shc (SHC) mRNA, complete cds.
83.65	197.1	115.26	96.2	52.18	980.75	1228.03	436.45	368.47	326.6	46.95	275.85	32.46	U77413, class A, 20 probes, 20 in U77413 2543-2975, Human O-linked GlcNAc transferase mRNA, complete cds. /gb=U77413 /ntype=RNA
185.19	857.4	433.46	1137.95	1070.77	1869.3	580.25	840.35	599.56	770.5	499.64	1035.05	321.38	Y08836, class A, 20 probes, 20 in Y08836mRNA 37-181, H.sapiens mRNA for HRX-like protein. /gb=Y08836 /ntype=RNA
1395.97	3754.4	186.53	3695.05	2187.86	4151.85	95.81	6057.3	1919.51	8431.85	6398.68	4656.55	1166.94	Y08265, class C, 20 probes, 20 in all_Y08265 1306-1834, H.sapiens mRNA for DAN26 protein, partial
203.86	81.65	30.48	107.45	47.45	376.35	430.7	106.05	96.52	158.95	74.32	211.95	97.93	U07804, class A, 20 probes, 20 in U07804 1857-2384, Human DNA topoisomerase I mRNA, partial cds
1033.72	1538.3	549.14	1275.6	815.72	1652.6	556.21	1980.3	861.54	1536.8	484.51	1150	183.56	X95097, class A, 20 probes, 20 in X95097mRNA 924-1503, H.sapiens mRNA for VIP2 receptor
78.28	274.5	230.09	457.45	398.74	1098.15	1265.51	310.8	177.91	586.3	30.26	499.35	51.55	X78678, class C, 20 probes, 20 in all_X78678 1384-1871, H.sapiens KHK mRNA for ketohexokinase, clone pHKHK3a
1329.93	2033.95	260.14	3526.45	881.55	1013.25	949.29	564.55	473.69	1245.45	164.83	1439.7	494.83	M93651, class A, 20 probes, 20 in M93651 1973-2519, Human set gene, complete cds
127.21	167.65	90.58	508.35	553.31	173.3	70.43	100.05	22.42	667.15	524.6	298.95	57.49	U21936, class A, 20 probes, 20 in U21936 2630-3038, Human peptide transporter (HPEPT1) mRNA, complete cds
122.19	350.4	239.99	420.5	100.41	466	245.93	284	67.46	218.45	56.36	716.45	747.34	M76665, class A, 20 probes, 20 in M76665mRNA 792-1332, Human 11-beta-hydroxysteroid dehydrogenase (HSD11) gene
95.81	108.75	84.36	67.75	45.89	315.8	365.29	130.6	94.33	103.15	68.24	86.8	4.95	X57129, class B, 20 probes, 12 in X57129cds 373-583: 8 in reverseSequence, 1216-1366, H.sapiens H1.2 gene for histone H1
1577.14	81893.8	16342.65	73127.45	7771.18	82083.3	62005.35	81295.15	21188.1	70975.45	20281.17	93363.45	28441.03	Ribosomal Protein S20
69.79	696.7	55.01	718.9	177.34	1460.8	1433.16	942.85	654.85	965.6	311.83	1000.75	417.41	U73936, class A, 20 probes, 20 in U73936 5049-5523, Human Jagged 1 (HJ1) mRNA, complete cds

99.49	366.85	119.15	732.15	1003.31	1343.8	1452.11	172.45	23.55	1071.7	1019.93	185	47.66	M88579, class A, 20 probes, 20 in M88579 1225-1615, Human zinc finger protein (SRE-ZBP) mRNA, 3 end
619.85	1298.15	774.21	750	507.7	996.75	275.56	831.65	462.8	1162.85	621.76	924.9	345.35	J05158, class A, 20 probes, 20 in J05158mRNA 2315-2825, Human carboxypeptidase N mRNA, 3 end
38.96	77.1	84.43	788.6	1101.11	146.8	134.35	435.35	569.43	205.45	253.21	80.25	8.27	D17570, class A, 20 probes, 20 in D17570 565-1083, Human mRNA for zona-pellucida-binding protein (sp38), complete cds.
305.82	411.45	510.32	239.4	150.76	144	6.93	369.6	40.87	367.4	264.74	99	30.83	U09646, class A, 20 probes, 20 in U09646exon 358-874, Human carnitine palmitoyltransferase II precursor (CPT1) gene
962.51	758.85	930.76	110.8	81.32	269.95	298.75	748.15	473.83	83.45	29.77	212.25	105.85	Transcription Factor Btf3 Homolog (Gb:M90355)
9.83	64.45	49.29	188.5	122.33	194.25	154.08	66.6	60.39	703.45	869.95	120	83.3	J04970, class A, 20 probes, 20 in J04970 1397-1715, Human carboxypeptidase M, 3 end
1752.21	4304.3	801.15	1498.2	1258.23	3903.7	2764.79	3682.1	367.84	2626.25	1520.49	3223.75	1959.18	U27333, class B, 11 probes, 11 in U27333 2523-2728, Human alpha (1,3) fucosyltransferase (FUT6) mRNA, major transcript I, complete cds, Human alpha (1,3) fucosyltransferase (FUT6) mRNA, major transcript I, complete cds
223.09	460.95	139.79	509.6	316.64	1327.4	1251.15	2660.65	595.17	1888	2118.77	531.9	102.67	J00268, class C, 20 probes, 20 in J00268gene 270-1415, Human insulin gene
31.11	48.1	34.93	48.35	19.16	255.55	10.39	45.35	24.54	194.1	213.97	151	178.76	S81944, class A, 20 probes, 20 in S81944 1173-1689, gamma-aminobutyric acid type A receptor alpha 6 subunit [human, cerebellum, mRNA Partial, 1732 nt]
1115.81	1218.7	1107.47	608.05	36.7	516.3	37.48	460.7	413.66	1824.9	345.35	1024.3	50.63	M81933, class A, 20 probes, 20 in M81933 1920-2394, Human cdc25A mRNA, complete cds
729.66	2172.05	76.58	968.9	390.18	1989.25	1763.31	290.5	162.21	511.25	244.73	279.85	24.4	D84557, class A, 20 probes, 20 in D84557 2412-2874, Human mRNA for HsMcm6, complete cds
24048.49	53387.25	25941.84	44637.1	20446.28	44536.45	36104.09	52108.5	18064.18	47591.2	27584.95	56591.8	15748.68	Z19554, class C, 18 probes, 18 in all_Z19554 1243-1734, H.sapiens vimentin gene
275.28	340.4	250.17	165.6	23.19	466.85	238.22	395.25	218.99	274.25	131.17	256.8	99.56	M15958, class A, 20 probes, 17 in M15958cds 12-282: 3 in reverseSequence, 6718-6778, Human gastrin gene, complete cds
1355.24	1783.85	1417.25	2020	705.83	1778.4	318.48	1786.05	440.32	1779.3	1221.6	2479.95	781.42	U14391, class A, 20 probes, 20 in U14391 4095-4623, Human myosin-IC mRNA, complete cds
1110.72	2141.8	152.31	643.95	57.2	558.3	24.32	1322.45	931.9	1014.6	827.03	1328.15	1165.52	M55153, class A, 20 probes, 20 in M55153 2794-3232, Human transglutaminase (TGase) mRNA, complete cds
66.33	161.15	32.03	126.35	15.91	210.35	70.92	1656.4	1566.95	148.7	129.26	168.9	101.68	X83412, class B, 20 probes, 10 in X83412cds 225-412: 10 in reverseSequence, 507-539, H.sapiens B1 mRNA for mucin.
397.39	949.8	365.57	1078.55	712.13	719.8	598.07	172.5	56	1143	85.56	1263.75	265.94	M86528, class C, 20 probes, 20 in all_M86528 954-1357, Human neurotrophin-4 (NT-4) gene, complete cds
197.42	278.55	274	285.65	345.56	172.65	155.21	539.55	628.55	427.7	474.61	320.3	325.41	J05125, class A, 20 probes, 20 in J05125 1038-1422, Human triglyceride lipase mRNA, complete cds
320.67	1013.85	270.47	732.75	184.2	1850.3	783.19	1162.2	295.29	1170.9	605.42	772.55	668.85	X77909, class B, 20 probes, 13 in X77909cds 888-1122: 7 in reverseSequence, 1202-1406, H.sapiens IKBL mRNA
166.45	1283.85	659.24	1343.4	343.65	853.6	377.17	2326.8	86.69	1031.7	629.89	1770.9	14.28	Beta-Hexosaminidase, Alpha Polypeptide, Abnormal Splice Mutation
437.63	1938.25	734.19	1769.4	127.14	1428.35	677.2	1038.35	124.1	1319.8	235.47	1226.95	535.92	J04031, class A, 20 probes, 20 in J04031 2529-3057, Human methylenetetrahydrofolate dehydrogenase-methenyltetrahydrofolate cyclohydrolase-formyltetrahydrofolate synthetase mRNA, complete cds

6422.86	6849.85	3862.01	8760.2	5070.1	4427.15	3827.92	8361.05	3648.88	6339.25	5253.59	6901.7	1676.41	U09510, class A, 20 probes, 20 in U09510 1878-2425, Human glycyl-tRNA synthetase mRNA, complete cds
2101.1	4175	35.64	6078.6	2055.14	5515.4	3510.08	5331.15	1327.03	4493.4	1675.7	4619.4	610.52	L37368, class A, 20 probes, 20 in L37368 1849-2359, Human (clone E5.1) RNA-binding protein mRNA, complete cds
99.77	135.35	28.5	575.15	617.8	142.4	8.63	309.35	120	567.55	154.93	662.1	325.55	X55889, class B, 20 probes, 15 in X55889cds 264-546: 5 not in GB record, Human gene for ciliary neurotrophic factor, exon 1
446.54	2982.3	611.79	1775.9	842.87	1563.8	892.79	3041.85	233.27	2476.75	389.26	1933.25	63.71	X65784, class A, 18 probes, 18 in X65784cds 58-399, H.sapiens CAR gene
34.29	111.65	27.51	168.75	141.49	433.75	400.01	160.05	59.89	222.85	169.92	274	106.63	M64710, class B, 20 probes, 11 in M64710cds 64-353: 9 in reverseSequence, 1226-1353, Human C-type natriuretic peptide gene, complete cds
92.42	229.05	69.37	760.75	470.58	607.6	30.83	1070.1	459.9	162.35	69.65	1029.4	427.52	U66061, class C, 20 probes, 20 in all_U66061 134531-176445, Human germline T-cell receptor beta chain TCRBV17S1A1T, TCRBV2S1, TCRBV10S1P, TCRBV29S1P, TCRBV19S1P, TCRBV15S1, TCRBV11S1A1T, HVB relic, TCRBV28S1P, TCRBV34S1, TCRBV14S1, TCRBV3S1, TCRBV4S1A1T, TRY4, TRY5, TRY6, TRY7, TRY8, TCRBD1, TCRBJ1S1, TCRBJ1S2, TCRBJ1S3, TCRBJ1S4, TCRBJ1S5, TCRBJ1S6, TCRBC1, TCRBD2, TCRBJ2S1, TCRBJ2S2, TCRBJ2S3, TCRBJ2S4, TCRBJ2S5, TCRBJ2S6, TCRBJ2S7, TCRBC2, TCRBV20S1A1N2 genes from bases 452324 to 684973 (section 3 of 3).
145.88	3012.4	791.11	3124.6	1034.36	3477.85	195.09	3298.6	434.3	2908.5	1037.04	2593.7	481.26	U64105, class A, 20 probes, 20 in U64105 2540-3104, Human guanine nucleotide exchange factor p115-RhoGEF mRNA, partial cds
241.19	2046.65	167.09	2503.6	490.31	730.1	432.61	1453.3	87.26	2383	928.29	1582.3	968.45	Z22555, class C, 20 probes, 20 in all_Z22555 2041-2516, H.sapiens encoding CLA-1 mRNA
232.07	213.05	117.31	802.85	228.61	404.7	249.18	1232	436.57	1358.15	617.94	805.5	180.88	X78711, class B, 20 probes, 12 in X78711cds 1553-1638: 8 in reverseSequence, 1665-1735, H.sapiens mRNA for glycerol kinase testis specific 1
31.68	206.15	162.14	118	54.59	392.2	185.97	449.5	215.67	1023.25	606.06	856.9	388.2	S77893, class A, 20 probes, 20 in S77893 121-203, GPSAT=glycophorin SAT [human, peripheral bloods, mRNA Partial, 407 nt]
144.32	432.9	198.84	252.05	23.26	673.35	737.72	311.35	211.64	388.35	71.35	687.7	892.23	U30930, class A, 20 probes, 20 in U30930 1877-2423, Human UDP-Galactose ceramide galactosyl transferase (CGT) mRNA, complete cds
31.61	74.5	0.99	98.65	71.91	104.65	66.4	96.6	45.54	1142.95	1478.07	186.35	110.1	X99268, class C, 20 probes, 20 in all_X99268 928-1367, H.sapiens mRNA for B-HLH DNA binding protein
566.46	1079.4	488.19	408.55	381.34	374.35	105.43	1660.9	690.98	431.25	241.19	580.65	170.06	D88155, class B, 20 probes, 13 in D88155cds 1025-1357: 7 in reverseSequence, 439-576, Human DNA for Ad4BP (SF-1) gene
232.14	323.6	147.22	348.3	225.14	868.35	1057.34	265.25	176.42	413.85	212.2	617	355.96	U90543, class B, 12 probes, 12 in U90543 2445-2739, Human butyrophilin (BTF1) mRNA, complete cds, Human butyrophilin (BTF1) mRNA, complete cds
511.59	489.05	614.26	703.9	553.66	109.05	91	62.6	53.88	132	7.21	406.7	244.09	J04093, class A, 20 probes, 20 in J04093 1842-2342, Homo sapiens phenol UDP-glucuronosyltransferase (UDPGT) mRNA, complete cds
31.4	354.1	28.43	380.75	165.68	398.25	108.68	674.35	328.59	507.65	57.63	561.6	20.79	D79998, class A, 20 probes, 20 in D79998 3100-3562, Human mRNA for KIAA0176 gene, partial cds

1063.98	1580.95	815.79	2388.75	26.66	2913.35	1633.63	1170.1	977.93	2538.75	64.98	2573.15	641.56	D79996, class A, 20 probes, 20 in D79996 1759-2305, Human mRNA for KIAA0174 gene, complete cds
134.99	380.45	16.9	1417.3	1849.08	610.3	457.92	360.15	132.02	1124.5	1064.34	862.8	174.94	X13766, class C, 20 probes, 20 in all_X13766 14-551, Human beta-casein mRNA 3 -terminal fragment
108.89	114.9	121.48	61.1	55.86	241.85	187.03	498.55	629.4	98.8	43.98	152.6	153.73	Z46788, class C, 20 probes, 20 in all_Z46788 1637-2082, H.sapiens mRNA for cylicin II
79.34	2555.4	348.04	2641.8	842.87	1105.3	974.53	1156.35	1152.23	1452.65	1623.16	1632.25	859.49	J05249, class A, 20 probes, 20 in J05249 941-1409, Human replication protein A 32-kDa subunit mRNA, complete cds
25.81	40.4	22.06	50.7	52.61	223.45	202.02	206.2	225	111.4	84	331.8	393.86	X01057, class A, 20 probes, 20 in X01057mRNA 1492-1738, Human mRNA for interleukin-2 receptor
101.54	225.25	8.56	500.6	490.87	753.85	892.86	673.1	803.27	87.5	12.73	396	349.17	U47677, class A, 20 probes, 20 in U47677mRNA 2495-2636, Human transcription factor E2F1 (E2F1) gene, promoter and
755.4	1022.75	699.54	953.55	84.64	1299.85	323.22	547.1	248.9	578.05	177.55	1030.95	774.35	M93036, class A, 20 probes, 18 in M93036mRNA 987-1353: 2 in reverseSequence, 527-545, Human (clone 21726) carcinoma-associated antigen GA733-2 (GA733-2) mRNA
39.24	119.5	61.38	583.7	659.59	192.85	65.55	244.8	90.23	271.55	142.62	233.5	89.94	U41898, class A, 20 probes, 20 in U41898 73-505, Human sodium cotransporter RKST1 mRNA, partial cds. /gb=U41898 /ntype=RNA
192.47	646.2	549.85	492.5	31.54	1176.45	510.46	719.35	561.23	699.05	95.95	695.6	238.58	Y12670, class A, 20 probes, 20 in Y12670mRNA 531-1011, H.sapiens OB-RGRP gene. /gb=Y12670 /ntype=RNA
58637.18	128598.8	46403.04	113996.3	39744.06	132117.55	80910.91	169375.56	9917.67	149954.55	113252.98	163077.66	3411.87	Z19574, class A, 20 probes, 20 in Z19574mRNA 1039-1479, H.sapiens gene for cytokeratin 17
8.63	999.45	177.98	860.85	268.06	442.15	92.14	914.6	378.58	454.3	499.22	971.25	471.43	M15990, class A, 20 probes, 20 in M15990 3916-4390, Human c-yes-1 mRNA
165.25	559.2	126.29	609.25	165.68	2765.8	2563.26	585.65	109.96	543.25	9.55	491.05	20.29	U28131, class A, 20 probes, 20 in U28131 88-283, Human HMGI-C chimeric transcript mRNA, partial cds.
250.03	1017.35	645.38	296.05	111.23	450.65	155.35	281.35	93.41	393	217.08	302.75	46.88	X51698, class B, 19 probes, 15 in X51698cds 39-343: 4 in reverseSequence, 387-465, H.sapiens spasmolytic polypeptide (SP) mRNA
680.02	993.8	837.64	826.15	696.15	962.85	105.15	741.8	223.16	1348.1	353.98	1154.05	407.65	U69141, class A, 20 probes, 20 in U69141 1311-1719, Human glutaryl-CoA dehydrogenase mRNA, complete cds
472.06	845.65	305.26	1695.05	1797.68	4688.85	5318.65	2223.8	160.8	647.1	766.79	2017.6	2207.02	Mucin 6, Gastric (Gb:L07518)
8004.59	28305.8	4243.21	35675.85	20785.47	18815.9	16362.45	28294.6	11678.86	17112.6	3439.37	21990.95	3114.17	M80244, class A, 20 probes, 20 in M80244 3401-3869, Human E16 mRNA, complete cds
379.72	1948.35	302.01	1599.55	194.38	1599.9	443.07	1033.6	785.45	1787.35	146.58	2025.85	646.65	S82470, class A, 20 probes, 20 in S82470 1398-1836, BB1=malignant cell expression-enhanced gene/tumor progression-enhanced gene [human, UM-UC-9 bladder carcinoma cell line, mRNA, 1897 nt]
67.53	218.2	161.79	531.3	646.86	469.4	339.41	519.1	172.25	254.2	203.36	521.1	507.7	X67098, class A, 20 probes, 20 in X67098exon#8 40-454, H.sapiens rTS alpha mRNA containing four open reading frames
1131.37	4485.8	998.43	1775.85	1454.31	2738.05	1832.47	4135.1	680.1	5642.8	507	3914.8	1499.63	U78521, class A, 20 probes, 20 in U78521 655-1111, Homo sapiens immunophilin homolog ARA9 mRNA, complete cds
226.27	472.45	111.65	414.95	391.24	768.6	642.48	225.25	30.76	624.65	79.27	301.3	279.45	S62539, class A, 20 probes, 20 in S62539 5366-5756, insulin receptor substrate-1 [human, skeletal muscle, mRNA, 5828 nt]

460.89	982.5	217.51	413.15	36.7	452.5	132.65	512.95	75.31	610.1	93.76	658	56.29	M83308, class A, 20 probes, 20 in M83308 31-343, Human mitochondrial cytochrome-c oxidase subunit VIa (COX6A) mRNA, complete cds
846.62	1203.1	525.1	686	396.12	643.1	393.29	540.85	463.51	348.8	305.33	151.9	101.4	Helix-Loop-Helix Protein Sef2-1d - Also Represents: M74719
280.65	589.95	231.29	1187.45	1147	2113.3	2158.23	1560.35	807.87	565.25	49.14	901.25	453.33	U40371, class A, 20 probes, 20 in U40371 2129-2591, Human 3 ,5 cyclic nucleotide phosphodiesterase (HSPDE1C1A) mRNA, complete cds
120.28	344.4	47.23	79.95	43.63	111.7	71.84	405.1	432.61	255.65	8.13	187.1	111.02	U37359, class A, 20 probes, 20 in U37359 1964-2414, Human MRE11 homolog hMRE11 mRNA, complete cds
163.41	159.55	147.86	148.2	68.45	179.45	89.31	225.45	144.6	205.55	101.61	123	58.55	Cellular Retinol Binding Protein Ii
134.21	193.2	91.08	270.65	225.64	248.8	140.71	359.55	226.06	124.6	80.19	288.5	147.36	X63522, class C, 20 probes, 20 in all_X63522 1800-1997, H.sapiens mRNA DAUDI6 for retinoic acid X receptor b
130.11	1010.35	147.01	1435.65	412.17	1378.35	324.07	954.8	769.19	833.1	923.62	449.3	402.49	L38616, class A, 20 probes, 20 in L38616mRNA 1184-1634, Homo sapiens brain and reproductive organ-expressed protein (BRE) gene, complete cds
141.07	434.65	69.93	993.75	1091.42	103.65	36.27	836.3	1028.27	414.85	231.44	942.15	569.86	Y10313, class A, 20 probes, 16 in Y10313 1352-1730: 4 not in GB record, H.sapiens mRNA for nerve growth factor-inducible PC4 homologue
728.54	77454.41	8130.03	57305	14195.59	56647.65	47453.72	67853.05	12458.73	67938.1	12685.5	64051.5	30706.82	J04456, class A, 20 probes, 20 in J04456 31-469, Human 14 kd lectin mRNA, complete cds
141.85	607.3	57.7	627.35	75.31	1783.3	1703.56	669.7	69.44	878.5	200.68	1218.45	778.6	U48213, class A, 20 probes, 20 in U48213mRNA 1031-1601, Human D-site binding protein gene, promoter region and
220.69	1421.1	115.26	715.2	650.96	499.4	40.59	1623.45	196.79	963.95	1191.55	1750.85	4.88	X98248, class A, 20 probes, 20 in X98248mRNA 3140-3698, H.sapiens mRNA for sortilin
68.45	189.2	39.74	700.95	655.84	602.15	764.88	340.1	39.17	360.4	32.24	609.5	407.29	X17648, class B, 20 probes, 11 in X17648cds#2 1069-1177: 9 in reverseSequence, 1341-1605, Human mRNA for granulocyte-macrophage colony-stimulating factor receptor (hGM-CSF-R)
462.94	2094.15	714.11	1772.85	379.36	1171.6	1225.27	1138.4	406.87	2470.55	128.2	1935.95	397.46	U24152, class A, 20 probes, 20 in U24152 1879-2215, Human p21-activated protein kinase (Pak1) gene, complete cds
687.8	570.05	657.54	631.25	775.2	772.9	132.37	357.95	282.91	145.3	66.47	553.85	341.46	Y13896, class A, 20 probes, 20 in Y13896 4-179, Homo sapiens skeletal muscle alternate 5 end of gene Kir4.2 5 UTR. /gb=Y13896 /ntype=RNA
220.26	1743.3	617.02	1983.9	151.32	1320.65	90.58	1313.9	165.89	940.2	521.7	2314.9	975.95	X97160, class A, 20 probes, 20 in X97160mRNA 2016-2532, TFE3 transcription factor gene extracted from H.sapiens TFE3 gene, exons 1,2,3 (and joined CDS)
68.59	479.7	188.66	1114.35	958.48	186.15	47.02	185	12.45	145.45	137.82	555	352.56	U80628, class A, 20 probes, 20 in U80628 2216-2588, Human thymidine kinase 2 isoform B (TK2) mRNA, alternatively spliced, partial cds
310.91	708.6	22.06	1686.1	1120.48	1579.3	561.02	968.75	237.8	876.65	141.21	891.9	989.95	U23435, class A, 20 probes, 20 in U23435 832-1319, Human Abl interactor 2 (Abi-2) mRNA, complete cds
166.81	98	53.17	270.3	94.75	880.85	912.1	313.35	253.5	103.25	47.16	187.1	19.09	X05360, class A, 20 probes, 20 in X05360cds 353-785, Human CDC2 gene involved in cell cycle control
64.35	183.65	142.76	206.15	119.01	866.9	617.59	365.95	444.28	166.95	41.79	231.65	213.76	D79983, class A, 20 probes, 20 in D79983 5024-5498, Human mRNA for KIAA0161 gene, complete cds
415.99	473.05	312.05	260.75	265.8	450.25	278.81	540.95	580.75	261.9	314.1	404.6	463.58	U52828, class A, 20 probes, 20 in U52828 1418-1882, Human Cri-du-chat region mRNA, clone NIBA2

19.16	97.85	40.94	228.35	23.97	418.75	415.85	153.4	91.08	282.9	254.13	118.65	37.83	U79271, class A, 20 probes, 20 in U79271 596-1130, Human clones 23920 and 23921 mRNA sequence
746	3106.05	907.43	2900.15	2003.45	1529.5	1674.85	2326.05	984.36	1707.2	1810.05	1684.3	1332.47	X99325, class C, 20 probes, 20 in all_X99325 1482-1927, H.sapiens mRNA for Ste20-like kinase
98.01	1393.15	504.8	1440.8	1078.76	2057.4	62.65	2233	887.7	1458	144.25	1328.75	412.88	X74614, class A, 20 probes, 16 in X74614exon#2 17-377: 3 in reverseSequence, 1279-1387: 1 not in GB record, H.sapiens ODF2 (allele 2) gene for outer dense fiber protein
296.35	4764.15	201.03	2623.85	664.04	5106.3	2846.81	6083.9	1126.56	3246.4	790.69	4081.2	1300.65	U51127, class A, 20 probes, 20 in U51127 1706-2084, Human interferon regulatory factor 5 (Humirf5) mRNA, complete cds
3454.78	3819.45	3035.96	2189.05	1202.72	3243.6	772.87	1997.35	935.57	2346.8	183	2990.1	264.32	U77846, class B, 10 probes, 10 in U77846mRNA 979-1356, Human elastin gene, partial cds and partial 3 UTR, Human elastin gene, partial cds and partial 3 UTR
14.35	338.1	106.63	345.55	267.78	879.05	133.57	398.55	326.47	961.4	340.12	119.8	106.91	D63940, class A, 20 probes, 20 in D63940 153-656, Human mRNA for Mxi1 protein, complete cds
32.03	285.3	0.57	470.1	412.67	330.75	56.07	349.7	194.17	531.25	278.11	322.65	11.81	Homeotic Protein Emx1
575.73	938.55	361.54	1026	386.5	2388.65	2173.43	845.35	60.6	155.15	1.48	615.45	481.61	Transcription Factor Oct-1a/1b, Alt. Splice 2, Oct-1b - Also Represents: X13403
197.49	647.7	248.76	1084.8	1163.61	2847.1	2617.43	593.9	380	1657.35	970.08	1154.15	800.09	U52830, class A, 20 probes, 20 in U52830 19-271, Homo sapiens Cri-du-chat region mRNA, clone CSC8.
106.7	131.3	108.61	85.55	81.53	1014.1	1382.82	473.5	533.44	53.95	13.22	106.65	49.99	Z00010, class A, 20 probes, 20 in Z00010exon#2 75-158, H.sapiens germ line pseudogene for immunoglobulin kappa light chain leader peptide and variable region (subgroup V kappa I).
99.77	1880.25	42.21	2146.5	510.67	1915.4	405.46	1829.8	87.12	3243.9	608.68	2008.1	335.59	M83221, class A, 20 probes, 20 in M83221 1788-2262, Homo sapiens I-Rel mRNA, complete cds
238.51	298.95	378.8	83.1	12.02	409.35	134.28	314.9	314.95	515.9	631.3	702.7	826.61	D63412, class A, 20 probes, 20 in D63412 1299-1713, Human mRNA for aquaporin, complete cds
485.15	982.9	506.01	620.65	60.32	914.2	968.31	863.1	50.63	940.05	81.11	1559.15	274.29	AC000064, class A, 20 probes, 17 in AC000064cds#1 1287-1581: 3 in reverseSequence, 16950-17160, WUGSC:H_RG083M05.2 gene extracted from Human BAC clone RG083M05 from 7q21-7q22, WUGSC:H_RG083M05.2 gene extracted from Human BAC clone RG083M05 from 7q21-7q22
55.3	61.25	15.49	93.7	49.64	125.5	29.56	753.8	437.27	472.25	530.4	59.7	5.94	X85750, class C, 20 probes, 20 in all_X85750 1935-2500, H.sapiens mRNA for transcript associated with monocyte to macrophage differentiation
153.8	177.25	171.9	258.4	94.61	96	29.13	149	71.84	452	533.72	124.1	84.29	L23808, class A, 20 probes, 20 in L23808 1297-1717, Human metalloproteinase (HME) mRNA, complete cds
21.92	3630.25	20.86	2433.45	1470.43	1850.8	771.17	2576.3	105.92	2004.75	2232.12	1622.1	1302.21	AF006087, class A, 20 probes, 20 in AF006087 237-777, Homo sapiens Arp2/3 protein complex subunit p20-Arc (ARC20) mRNA, complete cds. /gb=AF006087 /ntype=RNA
4.81	397.6	153.02	354.1	78.35	970.8	658.6	1668.85	1723.29	608.2	170.98	532.7	183.85	M91368, class A, 20 probes, 20 in M91368 2656-3189, Human Na+/Ca+ exchanger (CNC) mRNA, complete cds
315.23	468.85	319.97	215	217.51	498.4	290.34	50.7	32.81	165.4	35.07	595.4	362.6	U19765, class A, 20 probes, 20 in U19765exon#5 630-1194, Human nucleic acid binding protein gene, complete cds