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Sphingobium yanoikuyae Bacteremia, Japan

Appendix

16S rRNA Sample Sequences

Sample_518F Signal G:1282 A:1220 C:1612 T:1227

Lane: 45 Base Spacing: 14.643044 964 bases in 11647 scans

GGGGAAAAGGTTTTTCGGATTACTGGGCGTAAGCGCACGTAGGCGGCTATTCAAGT
CAGAGGTGAAAGCCCAGGGGCTCAACCCCGGAACTGCCTTTGAACTAGATAGCTTG
AATCCAGGAGAGGTGAGTGGAATTCGAGTGTAGAGGTGAAATTCGTAGATATTCG
GAAGAACACCAGTGGCGAAGGCGGCTCACTGGACTGGTATTGACGCTGAGGTGCGA
AAGCGTGGGGAGCAAACAGGATTAGATACCCTGGTAGTCCACGCCGTAAACGATGA
TAACTAGCTGTCAGGGCACATGGTGTTTTGGTGGCGCAGCTAACGCATTAAGTTATC
CGCCTGGGGAGTACGGTCGCAAGATTA AAACTCAAAGGAATTGACGGGGGCCTGCA
CAAGCGGTGGAGCATGTGGTTTAATTCGAAGCAACGCGCAGAACCTTACCAACGTTT
GACATCCCTATCGCGGATCGTGGAGACACTTTCCTTCAGTTCGGCTGGATAGGTGAC
AGGTGCTGCATGGCTGTCGTCAGCTCGTGTCTGAGATGTTGGGTAAAGTCCCGCAA
CGAGCGCAACCCTCGCCTTTAGTTGCCAGCATTAGTTGGGTACTCTAAAGGAACCG
CCGGTGATAAGCCGGAGGAAGGTGGGGATGACGTCAAGTCCTCATGGCCCTTACGC
GTTGGGCTACACACGTGCTACAATGGCGACTACAGTGGGCAGCCACCTCGCGAGAG
GGAGCTAATCTCCAAAAGTCGTCTCAGTTCGGATCGTTCTCTGCAACTCGAGAGCGT
GAAGGCGGAATCGCTAGTAATCGCGGATCAGCATGCCGCGGTGGAATACCGTTCCC
AGGCCTTGTACACACCGCCCGTCACACCATGGGAGTTGGATTCACTCGAAGGCGTTG
AGCTAACCGTAAGGAGGCAGGCGACCACAGTGGGTTTAGCGACTGGGGTGAGCTAG
AGGGA

Sample_800R Signal G:1287 A:1340 C:2903 T:1931

Lane: 43 Base spacing: 14.576325 706 bases in 8462 scans

2GGCCCTTTTTCGCCCTCAGCGTCATAACCAGTCCAGTGAGCCGCCTTCGCCACTGGTG
TTCTTCCGAATATCTACGAATTCACCTCTACACTCGGAATTCACCTCACCTCTCCTG
GATTCAAGCTATCTAGTTTCAAAGGCAGTTCGGGGTTGAGCCCCGGGCTTTCACCT
CTGACTTGAATAGCCGCCTACGTGCGCTTTACGCCAGTAATTCGAACAACGCTAG
CTCCCTCCGTATTACCGCGGCTGCTGGCACGGAGTTAGCCGGAGCTTATTCTCCCGG
TACTGTCATTATCATCCCGGGTAAAAGAGCTTTACAACCCTAAGGCCTTCATCACTC
ACGCGGCATTGCTGGATCAGGCTTTCGCCATTGTCCAATATTCCCTACTGCTGCCTC
CCGTAGGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCTGATCATCCTCTCAGACCA
GCTAAGGATCGTCGCCTTGGTGAGCCTTACCTACCAACTAGCTAATCCTACGCGG

GCTCATCCTTGGGCGATAAATCTTTGGACTTACGTCATCATCCGGTATTAGCTTCCGT
TTCCAGAAGTTATTCCGAACCCAAGGGCAGATTCCCACGCGTTACGCACCCGTGCGC
CACTATCTCCGAAGAGATCGTTCGACTTGCATGTATTAGGCATGCCGCCAGCGTTCG
TTCTGAGCAGTTCAAAAAA