## Asian Musk Shrew as a Reservoir of Rat Hepatitis E Virus, China

## Technical Appendix

Technical Appendix Table 1. Rat hepatitis E virus IgG and IgM positivity rates among Asian musk shrews (Suncus murinus) trapped in Zhanjiang City, Guangdong Province, China, December 2011-September 2012

|  | $\%$ (no. positive/no. tested) |  |
| :--- | :---: | :---: |
| Geographic region | $\operatorname{lgG}$ | $\operatorname{lgM}$ |
| Total | $10.4(27 / 260)$ | $4.6(12 / 260)$ |
| Chikan district | $2.7(3 / 113)$ | $1.0(1 / 113$ |
| Mazhang district | $16.3(24 / 147)$ | $7.5(11 / 147)$ |
| Huangwai village | $13.3(2 / 15)$ | $0(0 / 15)$ |
| Houyang village | $71.4(5 / 7)$ | $28.6(2 / 7)$ |
| Nanpan village | $23.8(5 / 21)$ | $9.5(2 / 21)$ |
| Chiling village | $8.3(1 / 12)$ | $0(0 / 12)$ |
| Beigou village | $16.7(1 / 6)$ | $16.7(1 / 6)$ |
| Chaofa village | $9.5(6 / 63)$ | $7.9(5 / 63)$ |
| Pig farm | $17.4(4 / 23)$ | $4.3(1 / 23)$ |

Table 2. Rat hepatitis E virus-positive strains isolated from Asian musk shrews (Suncus murinus) trapped in Zhanjiang City, Guangdong Province, China, December 2011-September 2012*

| Isolate name | Date of sampling | Sex | Village where trapped | Optical density value |  | Cluster or subcluster | GenBank accession no. |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  |  | $\lg G$ | IgM |  |  |
| CHZ-sRat-E-739 | 2012 May14 | M | Nanpan | 2.254 | 0.492 | C | KC473527 |
| CHZ-sRat-E-1086 | 2012 Sep 19 | F | Chaofa | 2.767 | 0.289 | C | KC473528 |
| CHZ-sRat-E-1107 | 2012 Sep 19 | F | Chaofa | 0.123 | 0.253 | A1 | KC473529 |
| CHZ-sRat-E-1129 | 2012 Sep. 20 | F | Chaofa | 0.104 | 0.561 | C | KC473530 |
| CHZ-sRat-E-1133 | 2012.Sep 20 | M | Chaofa | 0.101 | 0.505 | A2 | KC473531 |

*Positive results were obtained by reverse transcription PCR.

Table 3. Percent shared nucleotide sequence identities between rat hepatitis E virus strains from rats and Asian musk shrews (Suncus murinus) trapped in Zhanjiang City, Guangdong Province, China, December 2011-September 2012

| Rat HEV strain by <br> GenBank accession no. | \% Identity with rat hepatitis E virus strains from Asian musk shrews |  |  |  |  |
| :--- | :---: | :---: | :---: | :---: | :---: |
|  | KC473529 | KC473531 | KC473528 | KC473530 | KC473527 |
| KC465999 | 91.4 | 97.2 | 79.3 | 78.9 | 80.0 |
| KC465997 | 89.3 | 96.8 | 79.3 | 78.9 | 80.0 |
| KC465998 | 89.6 | 97.2 | 80.0 | 79.6 | 80.7 |
| KC465990 | 97.5 | 89.3 | 79.1 | 78.7 | 79.4 |
| KC465991 | 97.5 | 87.9 | 78.2 | 77.9 | 78.6 |
| KC465993 | 99.6 | 90.0 | 78.6 | 78.2 | 78.9 |
| KC466000 | 83.3 | 84.6 | 77.5 | 78.3 | 77.6 |
| KC465996 | 85.1 | 85.0 | 79.3 | 80.0 | 78.9 |
| KC466001 | 86.1 | 87.5 | 79.3 | 80.0 | 78.9 |
| KC465994 | 79.6 | 81.1 | 97.5 | 96.4 | 95.4 |
| KC465995 | 80.0 | 81.4 | 97.2 | 96.1 | 96.4 |
| KC465992 | 79.3 | 81.1 | 94.0 | 94.3 | 97.2 |
| JX120573 | 90.3 | 93.2 | 81.0 | 80.6 | 81.0 |
| GU345042 | 77.5 | 80.7 | 80.3 | 79.9 | 79.2 |
| GU345043 | 74.6 | 77.9 | 78.9 | 78.9 | 78.9 |
| KC473529 | 100.0 | 90.4 | 78.9 | 78.6 | 79.3 |
| KC473531 | 90.4 | 100.0 | 80.0 | 79.6 | 80.7 |
| KC473528 | 78.9 | 80.0 | 100.0 | 98.9 | 95.7 |
| KC473530 | 78.6 | 79.6 | 98.9 | 100.0 | 96.1 |
| KC473527 | 79.3 | 80.7 | 95.7 | 96.1 | 100.0 |

