**Supplementary figure 1. Alignment of HeV-G2 M gene sequences amplified by PCR**

Consensus    GCCCCCTTCACCATCTCTTACCATGGAAAAAAGTTTTGACTGGCGGGTCAATATTCAATG  60

2013-01     ............................................................

2019-01     ............................................................

2019-02      ......................G.....................................

2020-01      ............................................................

2020-02      ............................................................

2020-03      ............................................................

Consensus    CTGTCAAGGTTTGCCGTAACGTAGATCAGATTCAACTGGAAAAGCAACAATCATTGAGGA  120

2013-01      ............................................................

2019-01      ............................................................

2019-02     ............................................................

2020-01     ............................................................

2020-02      ............................................................

2020-03      ............................................................

Consensus    TTTTTTTCCTAAGTATTACCAAATTGAATGATTCCGGTATCTACATGATCCCGAGAACAA  180

2013-01     ....................................................A.......

2019-01     ....................................................A.......

2019-02      ............................................................

2020-01      ............................................................

2020-02      ............................................................

2020-03      ............................................................

Consensus    TGTTGGAATTCAGAAGGAACAATGCCATTGCTTTTAATCTTCTGATATACCTCAAAATTG  240

2020-02      ............................................................

2019-02      ............................................................

2013-01     ............................................................

2019-01     ............................................................

2020-01      ............................................................

2020-03      ............................................................

Consensus    ATGCAGATCTTGCAAAAGCCGGAATCCAAGGGAGTCTCGACAAGGACGGAGCTAAAGTGG 300

2013-01      ............................................................

2019-01      ............................................................

2019-02      ............................................................

2020-01     ............................................................

2020-02      ..................................C.........................

2020-03     ..................................C.........................