**Supplementary Figure legend**

**Fig S1** Phylogenetic analysis of RdRp amplicons obtained in this study and representatives of species in the genera *Alphacoronavirus* and *Betacoronavirus* based on the Neighbor-Joining method. All sequences were classified into two groups: *Alphacoronavirus* comprised of 2 clades and *Betacoronavirus* comprised of 3 clades.