

SUPPLEMENTARY INFORMATION¹

Mass mortality in freshwater mussels (*Actinonaias pectorosa*) in the Clinch River, USA, linked to a novel densovirus

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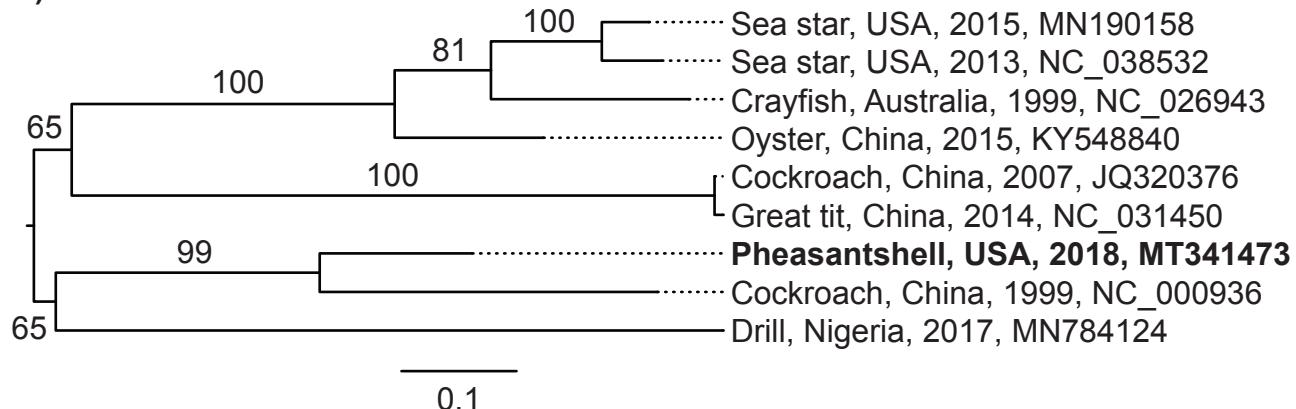
Table S1. Details of Clinch River pheasantshells (*Actinonaias pectorosa*) used in the analyses

Table S2. Loads of 17 viruses in Clinch River pheasantshells (*Actinonaias pectorosa*)

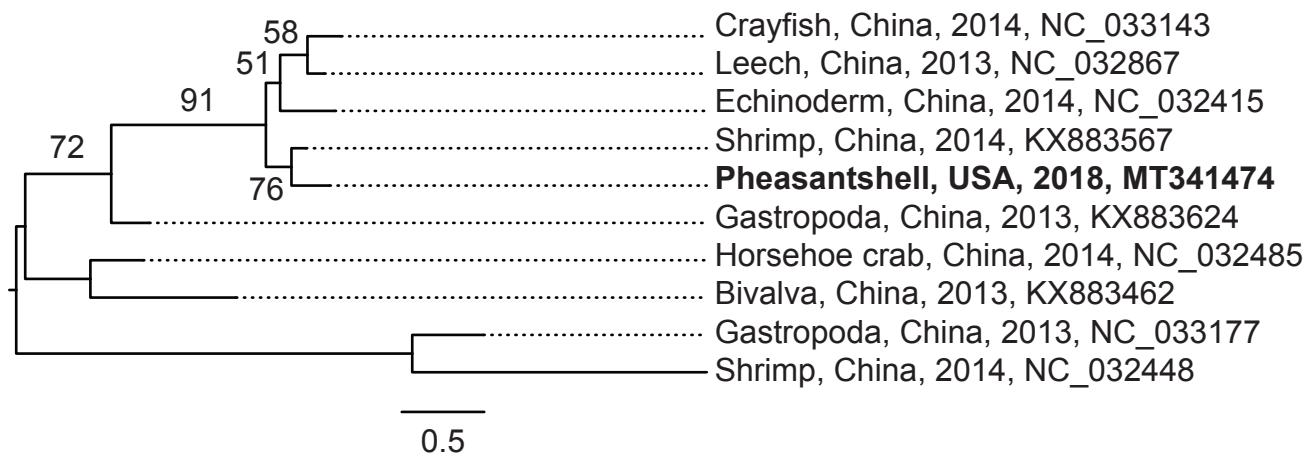
Figure S1. Maximum likelihood phylogenetic trees of 17 viruses identified in Clinch River pheasantshells (*Actinonaias pectorosa*) and their relatives in the GenBank database. Taxon names indicate host, country, year of collection, and GenBank accession number; see Table 2 for additional details. Viruses identified in the present study are shown in bold type. Numbers beside branches are bootstrap values representing statistical confidence in clades based on 1000 resamplings of the data; only bootstrap values $\geq 50\%$ are shown. Scale bars indicate nucleotide substitutions per site.

¹Any use of trade, firm, or product names is for descriptive purposes only and does not imply endorsement by the U.S. Government.

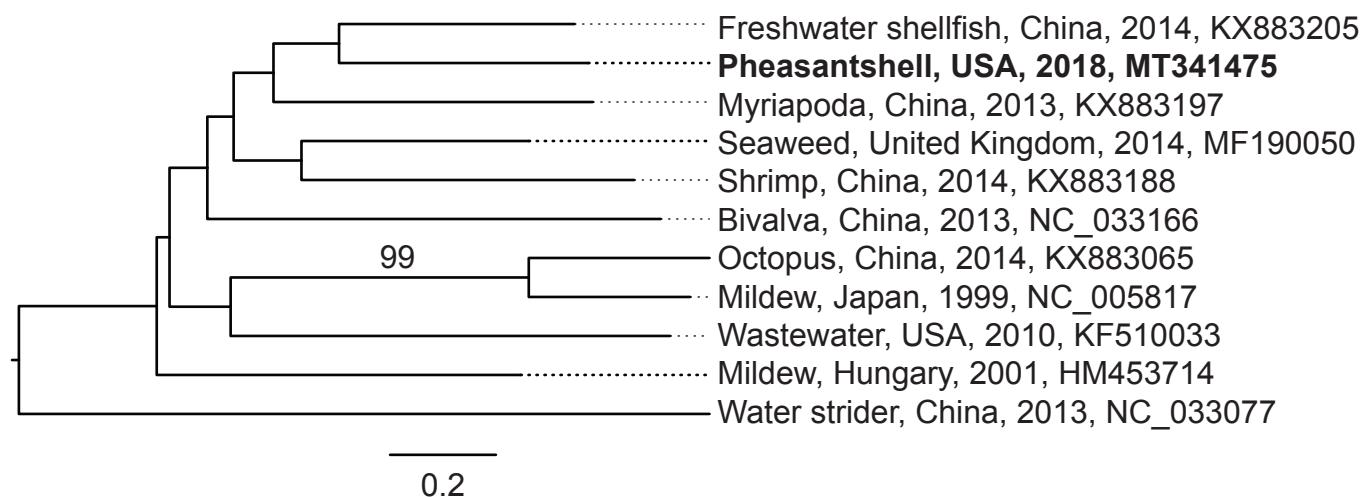
A) Densovirovuses



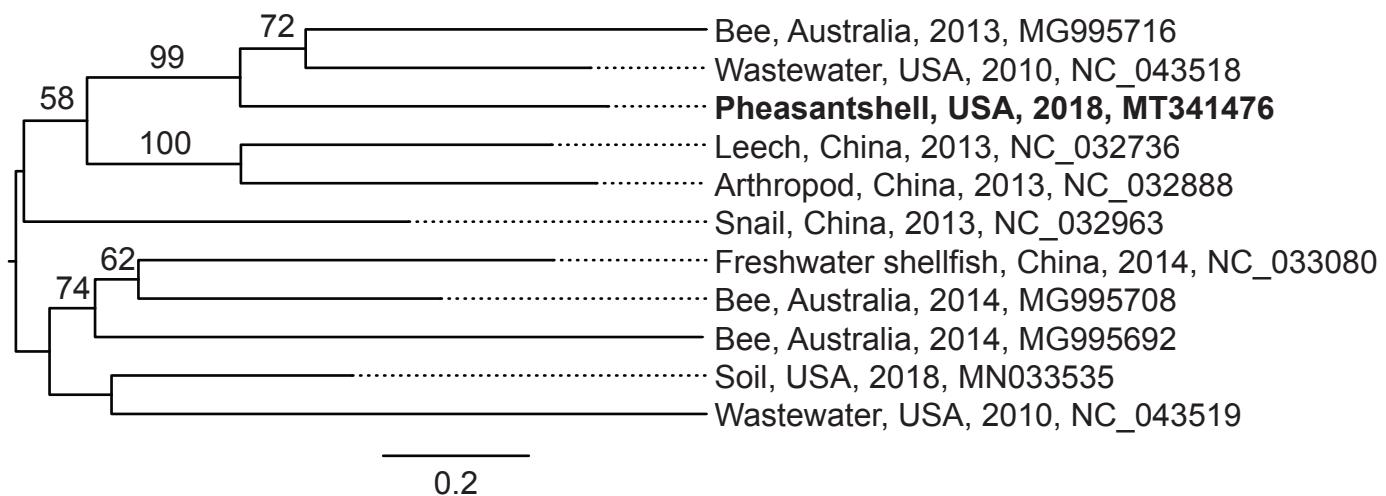
B) Narna-like viruses



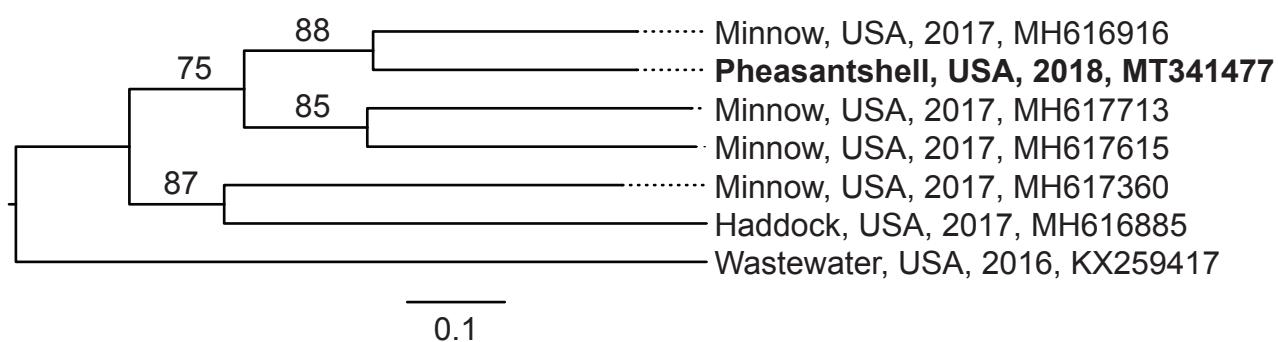
C) Noda-like viruses



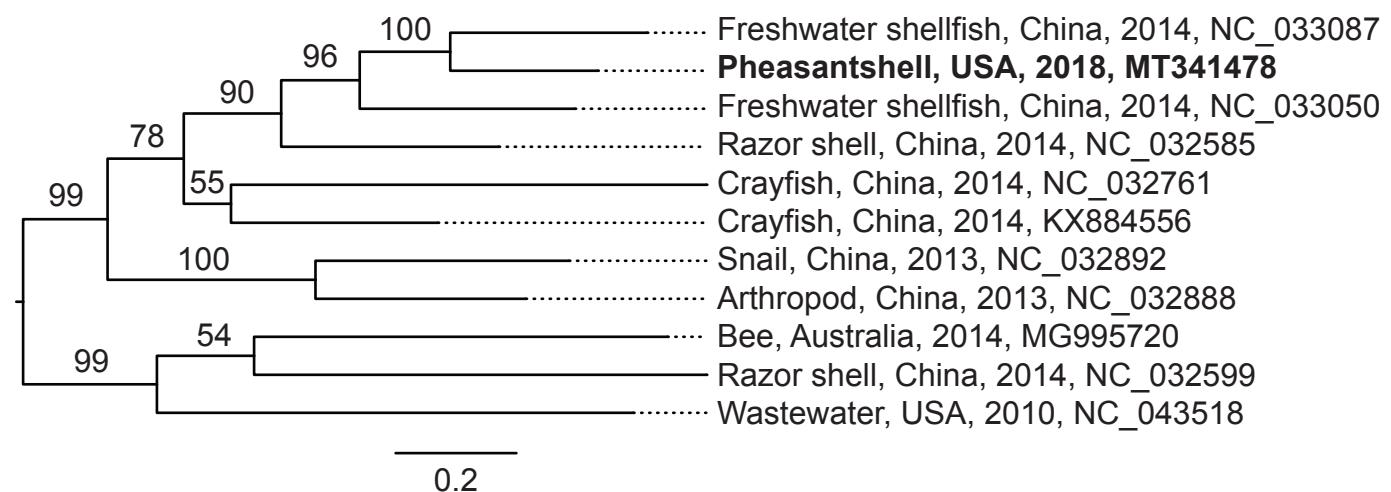
D) Picorna-like viruses (1)



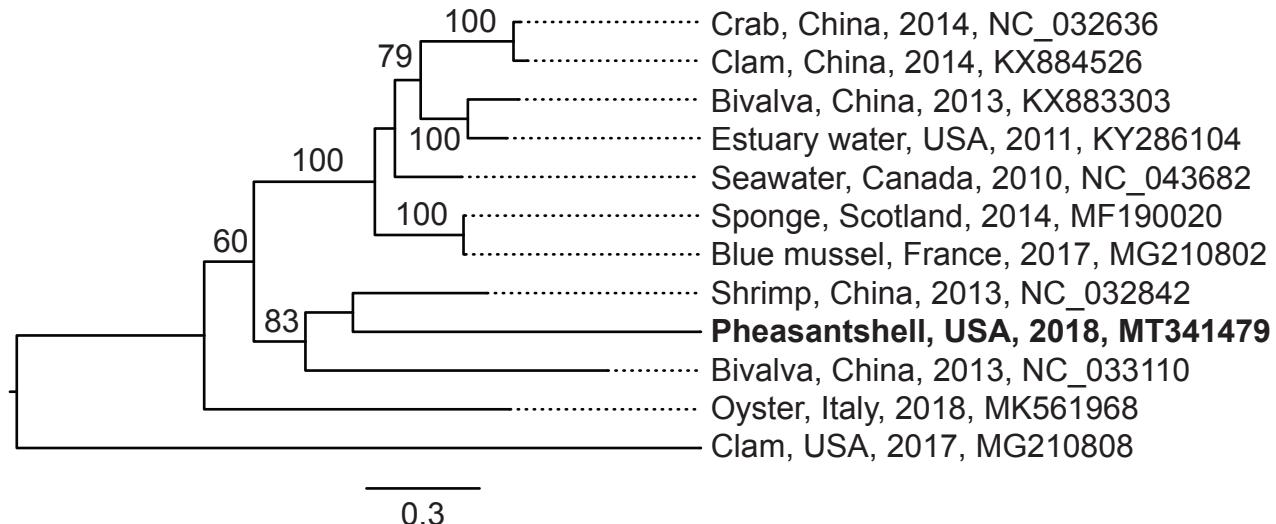
E) CRESS viruses



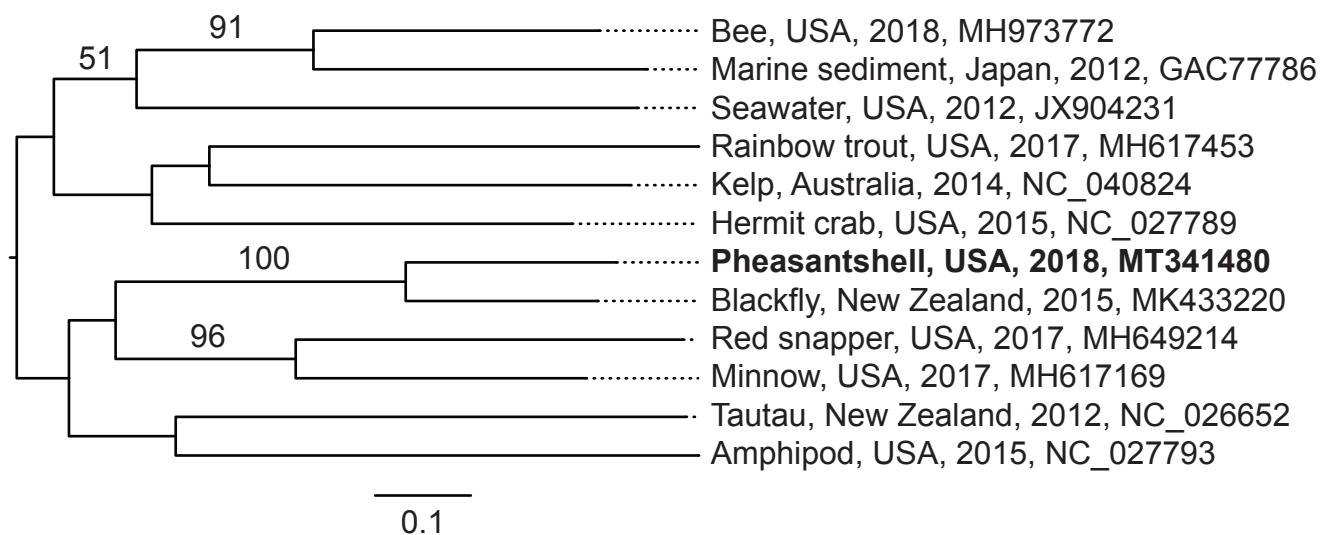
F) Picorna-like viruses (2)



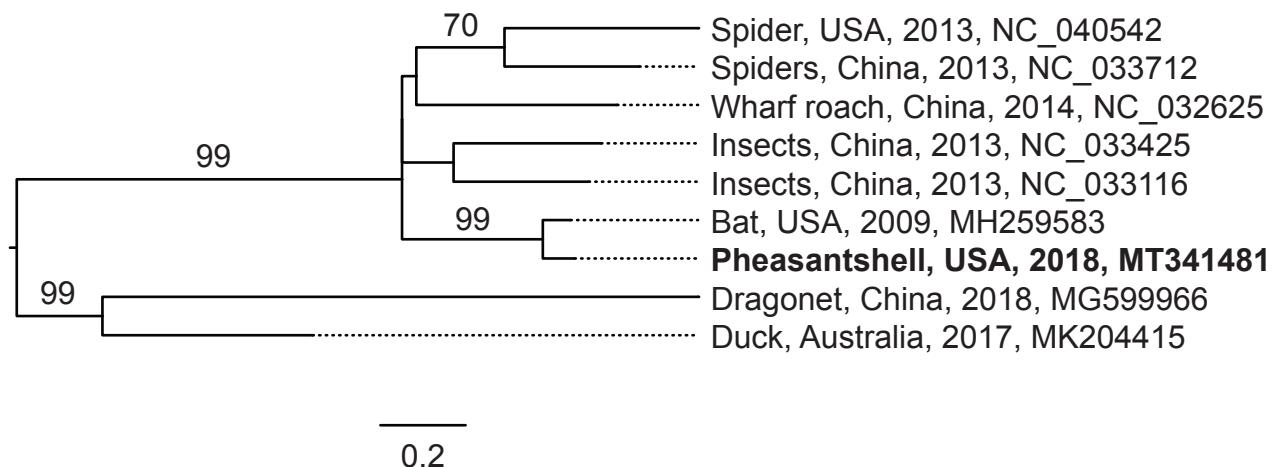
G) Picorna-like viruses (3)



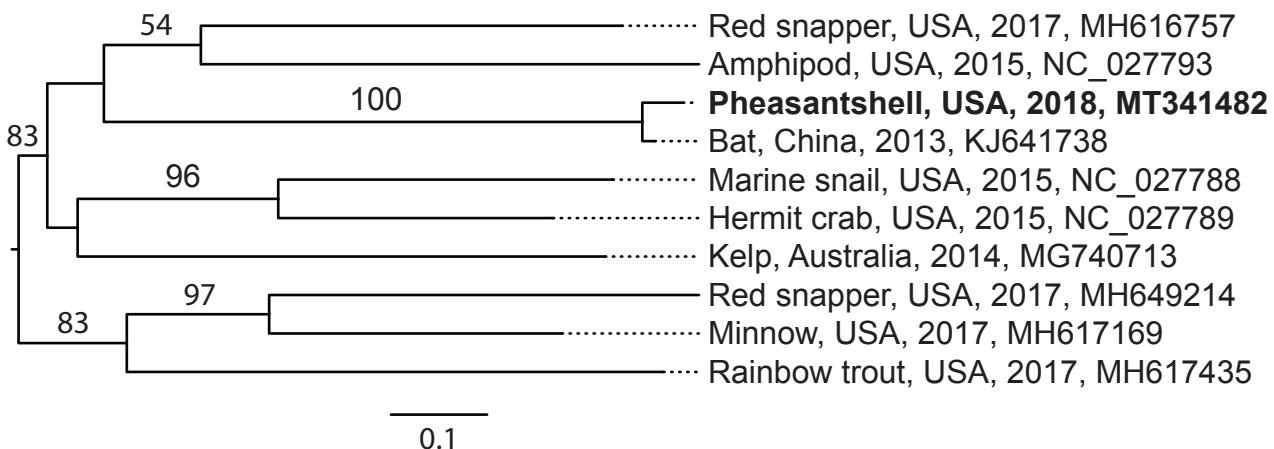
H) Circular viruses (1)



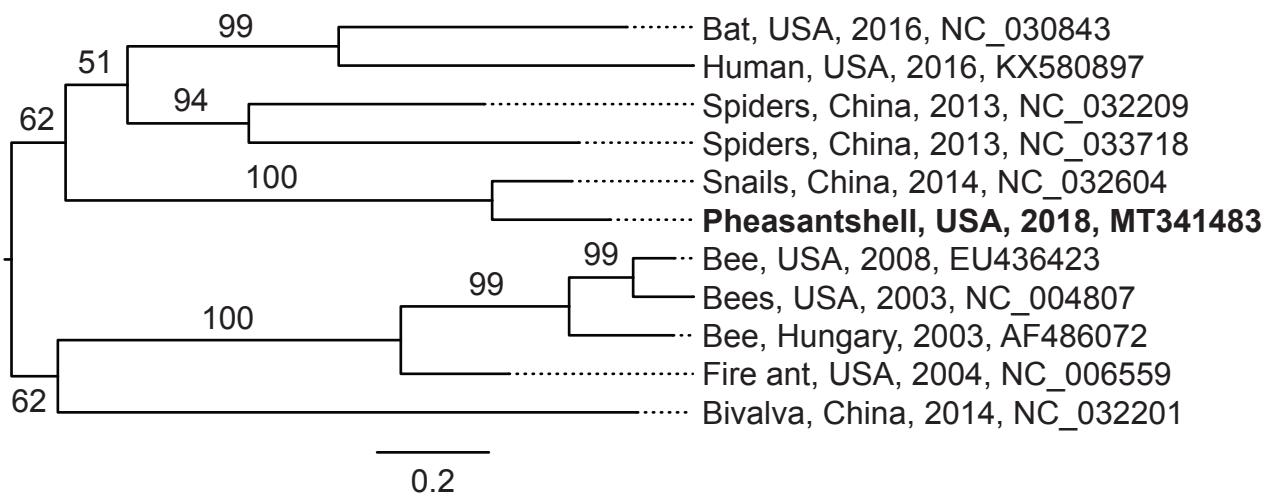
I) Caliciviruses



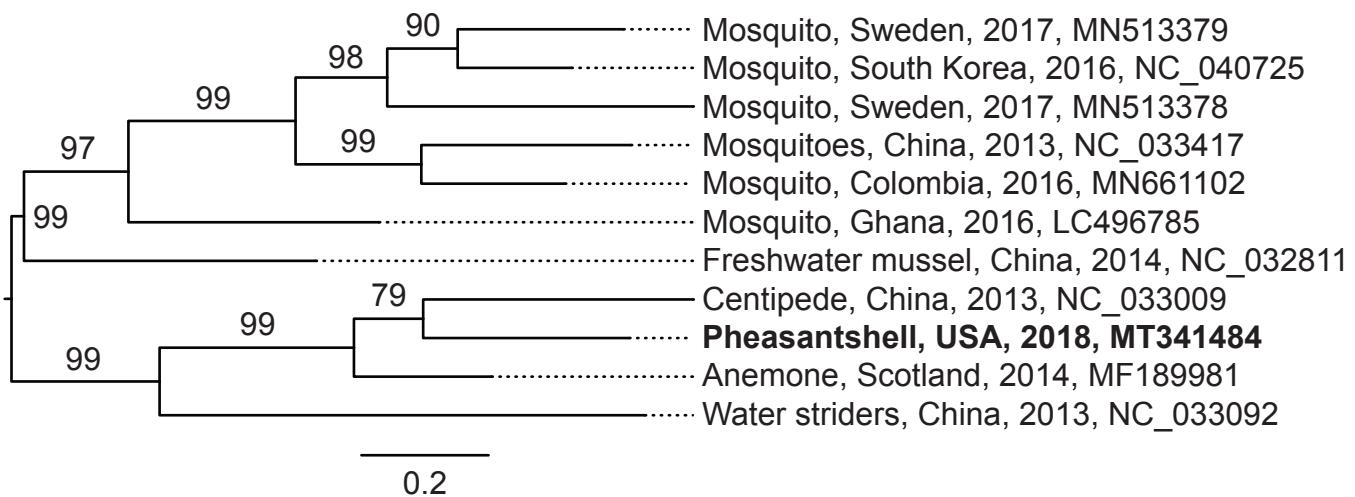
J) Circular viruses (2)



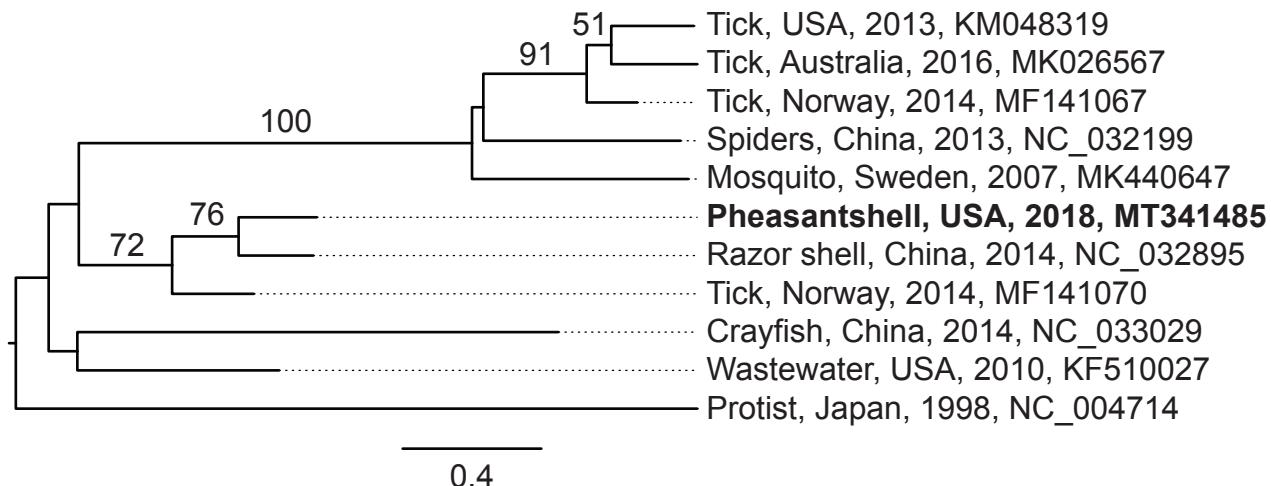
K) Dicistro-like viruses (1)



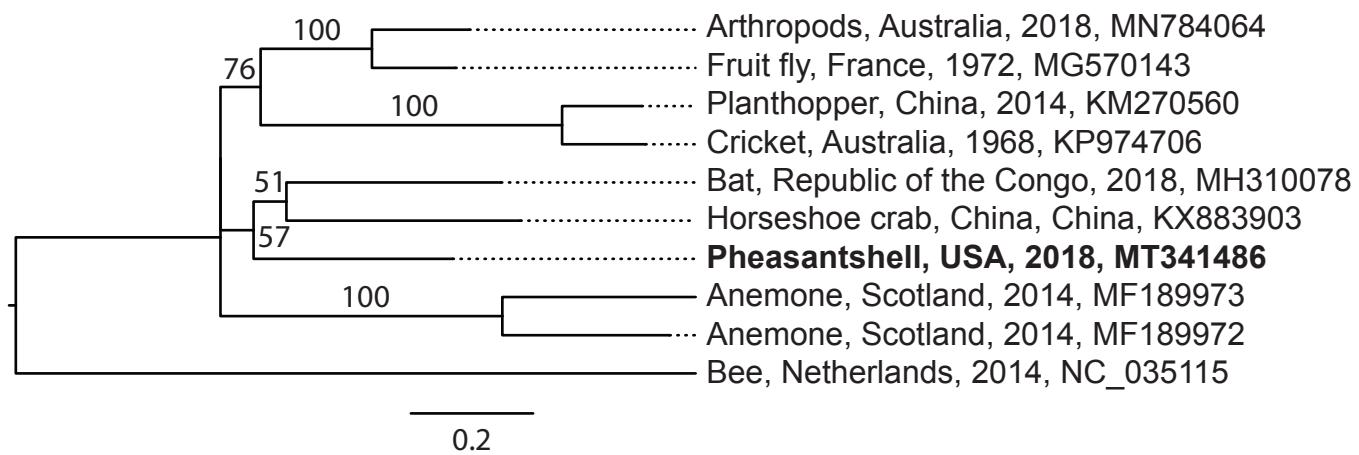
L) Tombus-like viruses



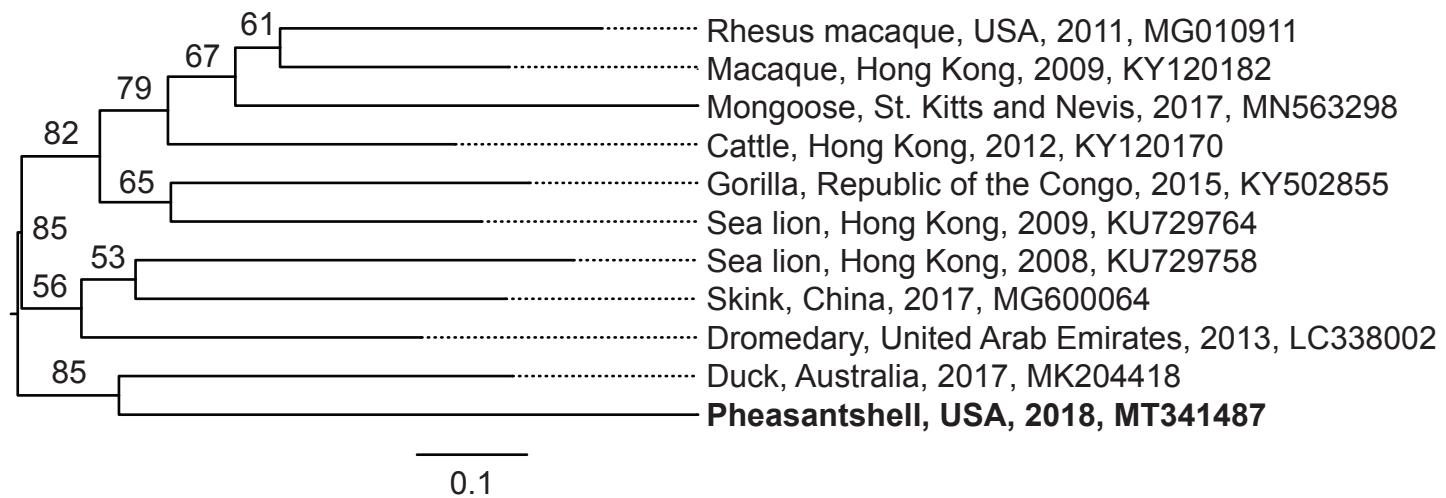
M) Sobemo-like viruses



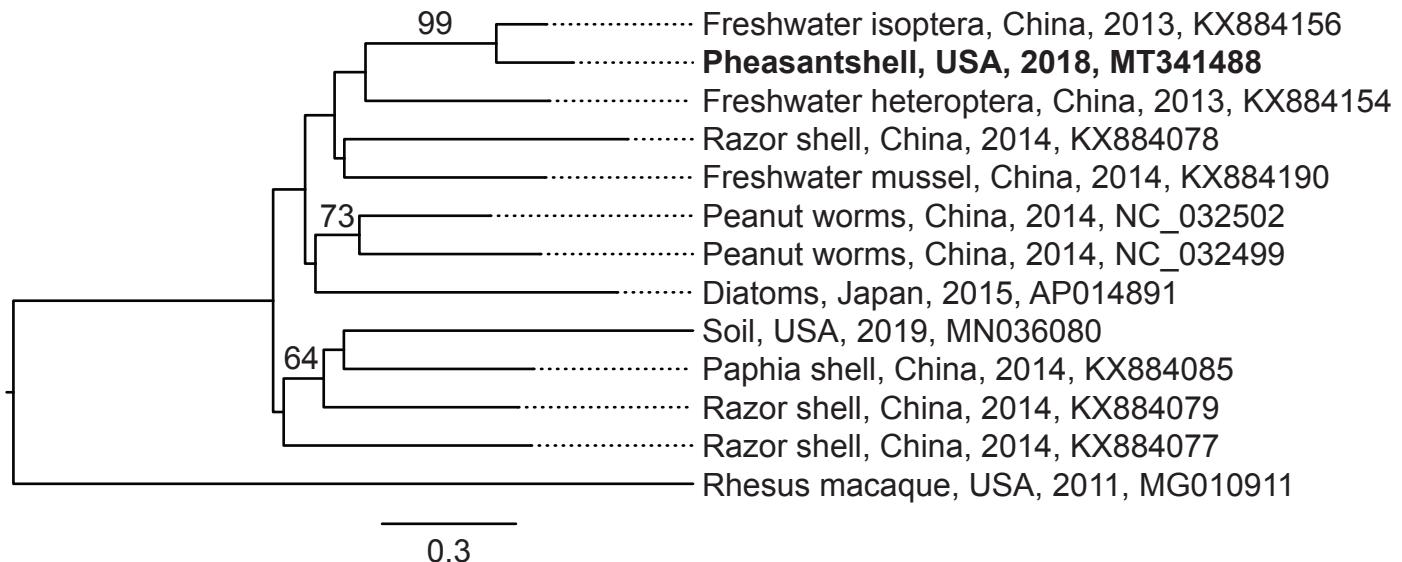
N) Dicistro-like viruses (2)



O) Picobirnaviruses



P) Picobirna-like viruses



Q) Totiviruses

