

**A**

**B**

**Fig. S1** Conserved H(X25)D(X70–80)GXSG domain **(A)** and GXXXTXXXN(X25–40)GDD motif **(B)** determined in the P1 and P1−P2 proteins, respectively of the identified viruses. The sequence logos were generated using the webserver WebLogo 3.