



Figure S3. RBPpb5 comparison among the four representative Tequintaviruses and Salten. **A-** MUSCLE alignment of the five aminoacid RBPpb5 sequences. **B-** RBPpb5 phylogenetic relationship reconstruction, constructed with Geneious Tree builder with Jukes-Cantor genetic distance model and neighbor-joining build method with bootstrap 1000.