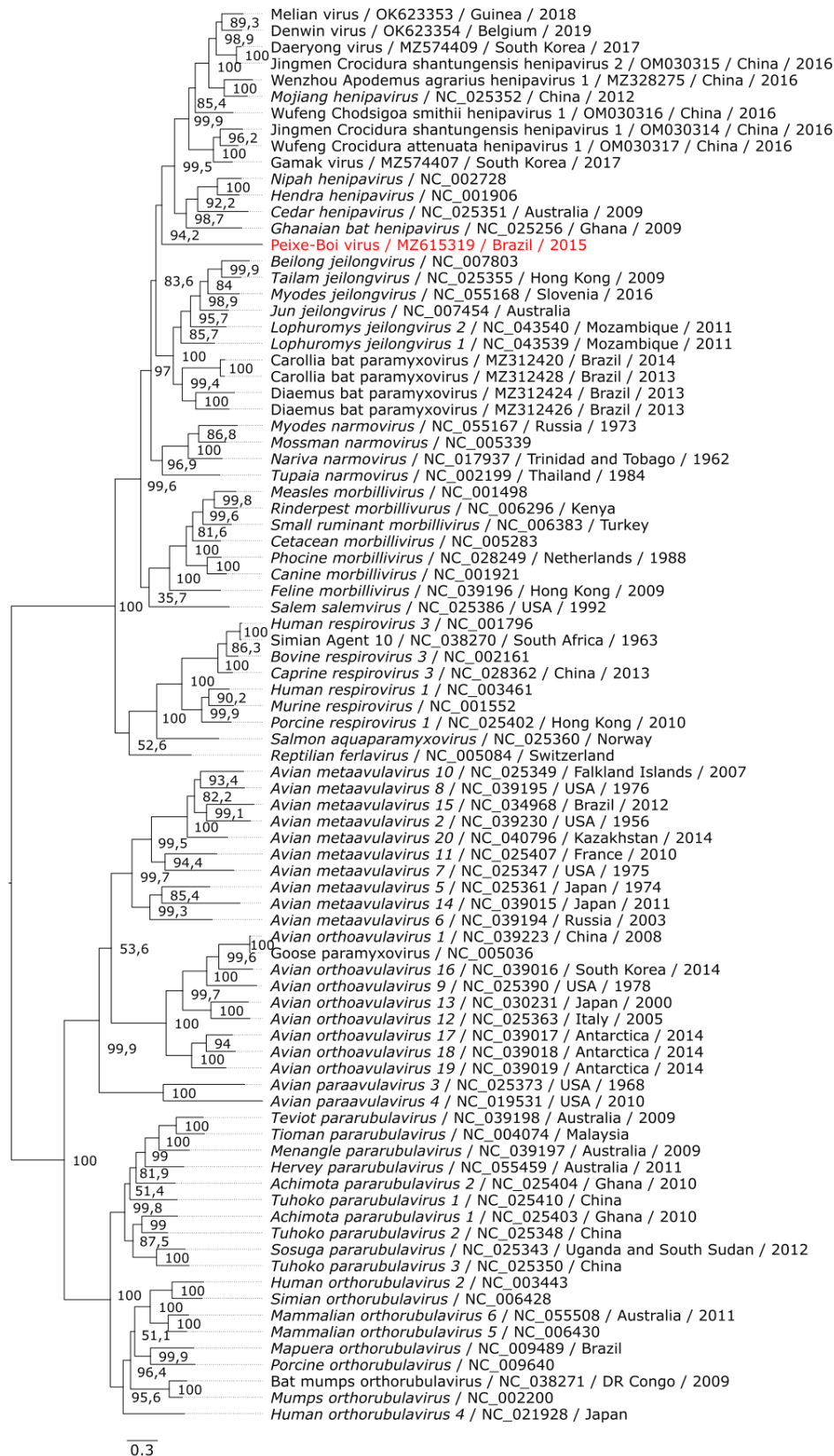


## Supplementary Figure



**Figure S1.** Phylogenetic tree based on the partial nucleotide *Paramyxoviridae* L gene sequences, novel rodent and shrew-borne henipaviruses, Brazilian jeilongviruses, and PBV (in red) sequences. Each record consists of the virus species/name, accession number, country and year of detection/isolation.