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Supplemental information

Distinct codon usage bias evolutionary patterns between weakly and strongly virulent respiratory viruses Feng Chen and Jian-Rong Yang

Supplementary figures

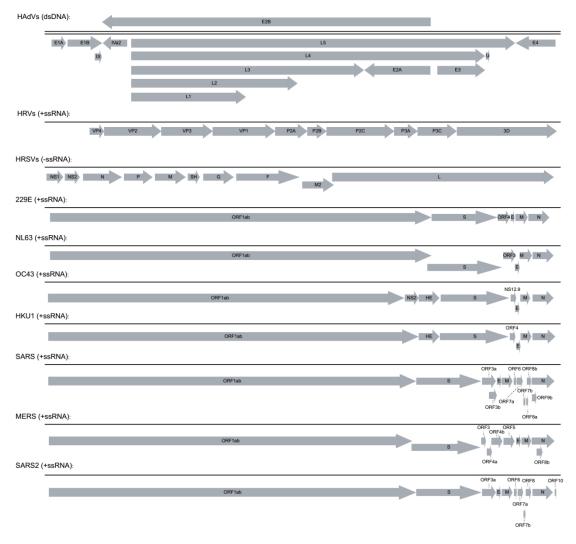


Figure S1. The genome structure of viruses analyzed in this paper, related to Figure 1-4. The gray boxes represent loci of genes in each virus.