

The impacts of antivirals on the coronavirus genome structure and subsequent pathogenicity, virus fitness and antiviral design

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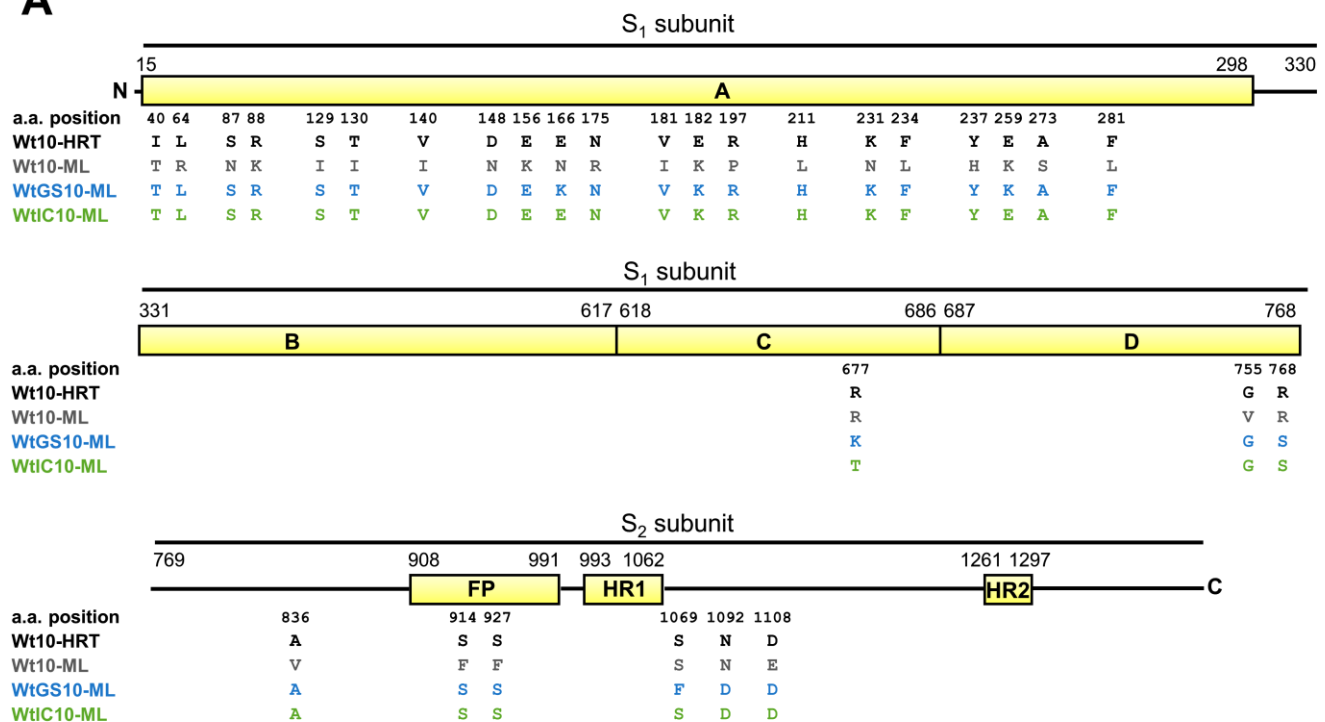
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Supplementary Figures

A



B

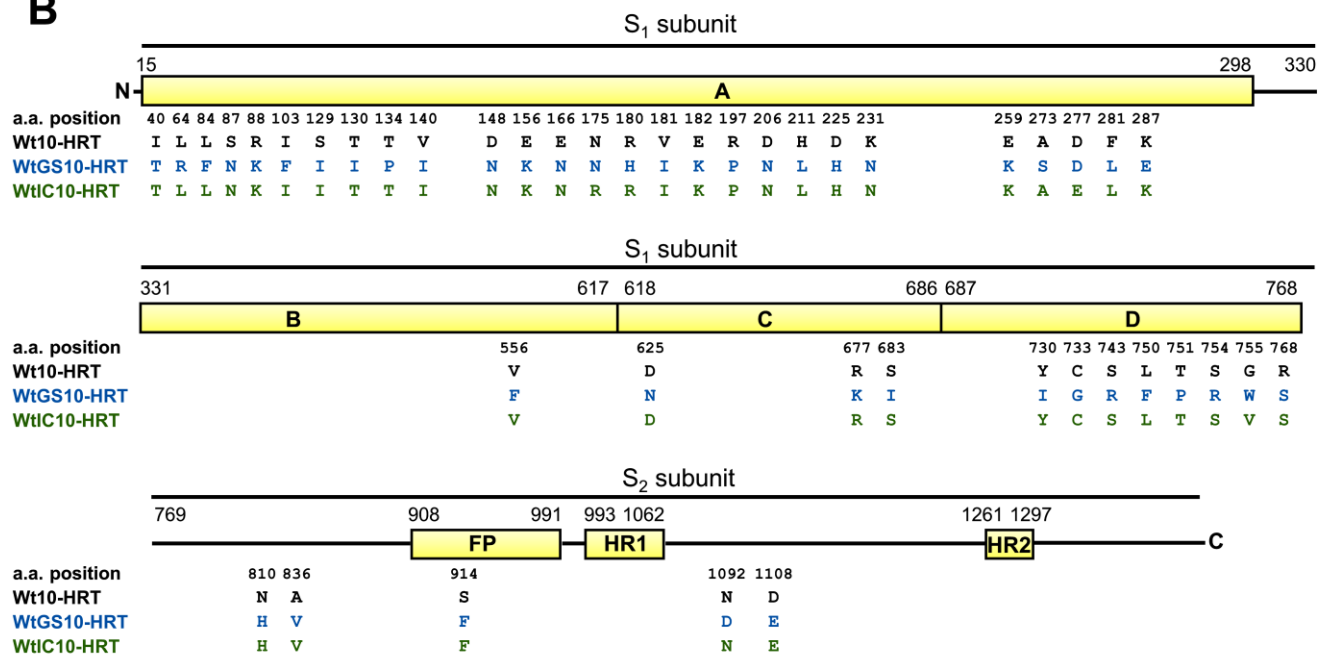


Figure S1. Comparison of the S protein aa sequences identified from HRT-18 cells and ML cells infected with wt BCoV and its variants. (A) Linear schematic of BCoV S protein showing the comparison of the aa sequences for Wt10-ML, WtGS10-ML and WtIC10-ML identified from ML cells infected with Wt10, WtGS10 and WtIC10, respectively, at VP10. **(B)** Linear schematic of BCoV S protein showing the comparison of the aa sequences identified from HRT-18 cells under the treatments of antiviral GS-5734 (WtGS10-HRT, i.e., WtGS10 in Figure 1) and poly IC-induced innate immunity (WtIC10-HRT, i.e., WtIC10 in Figure 4) at VP10. Wt10-HRT (i.e., Wt10 in Figures 1 and 4): Viral RNA collected from HRT-18 cells infected with wt BCoV at VP10.