

A Virological and Phylogenetic Analysis of the Emergence of New Clades of Respiratory Syncytial Virus

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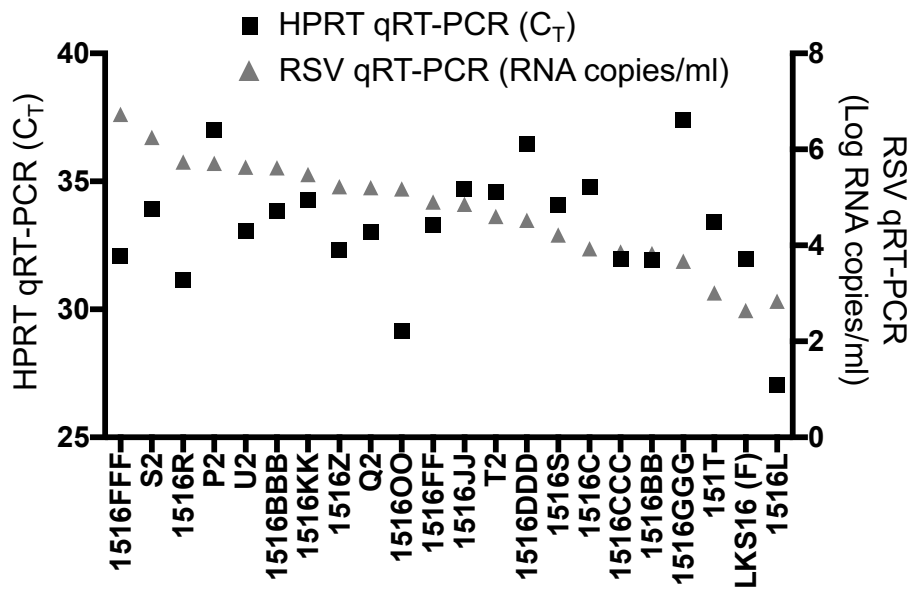
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Supplementary Figure 1. Patient HPRT gene mRNA levels in nasopharyngeal swabs. Patient nasopharyngeal swabs were used for viral load determination (denoted by a ▲) corresponding to log FFU/mL on the right axis. HPRT housekeeping gene levels were determined to monitor host patient derived cellular RNA levels. HPRT levels in terms of C_T (denoted by a ■) on the left axis. Patient samples are ordered from highest to lowest viral load.