

FIGURE S1

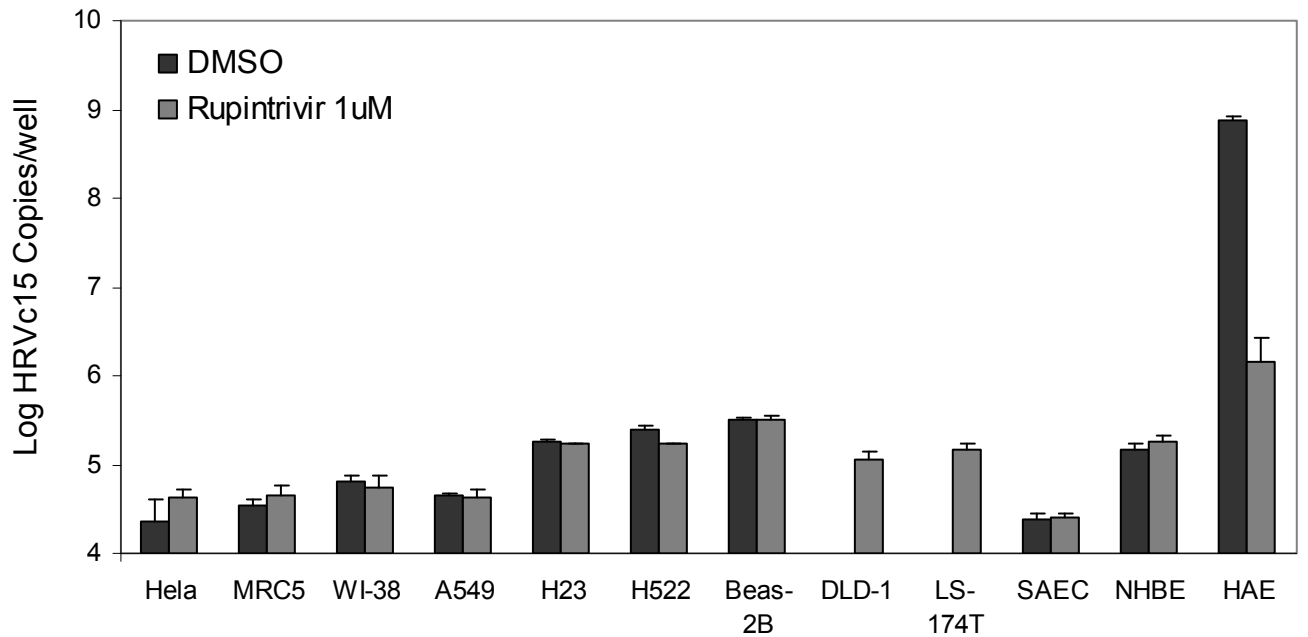


FIG S1. Permissivity of tested cell lines to HRVc15 virus infection. Selected transformed and primary respiratory cell lines were infected with HRVc15 virus (3.8×10^7 copies) in the presence of 0.5% DMSO or 1uM rupintrivir and incubated for 5-days at 33°C. Total viral RNA was isolated and analyzed by RT-qPCR. Error bars represent the standard deviation of the mean from triplicate tests.

FIGURE S2

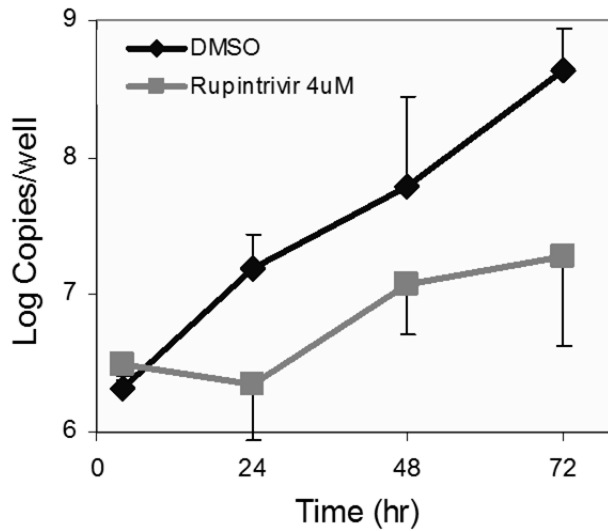


FIG S2. Serial re-infection and growth of HRVc15 virus in HAE cultures. Apical wash from HRVc15-infected HAE culture (5.4×10^6 copies RNA/well) was collected and used to infect fresh HAE cultures in the presence of 0.5% DMSO or 4uM rupintrivir and incubated at 33°C. At selected time points, virus shed into the HAE apical mucosa was collected and quantified by RT-qPCR. Error bars represent the standard deviation of the mean from triplicate tests.