Additional file 2 - Impact of protease background on bevirimat resistance. Fold increase in bevirimat $EC_{50}$ caused by single CA/p2 mutations in different protease backgrounds. All figures represent one of at least two separate experiments. (A-B) The susceptibility curves of site-directed mutants A364V and V370A in the HXB2 background are shown. (C-D) The susceptibility curves of virus with mutation V362I in wild-type vs. PR-1 and PR-2 backgrounds. (E-F) The same comparison is shown for viruses with mutation S368N.