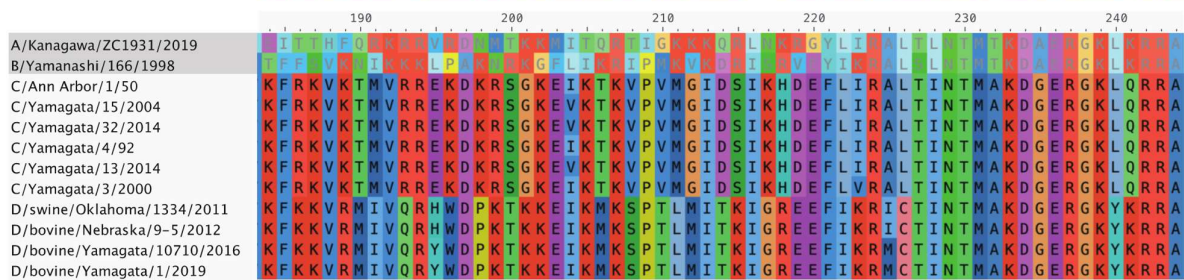
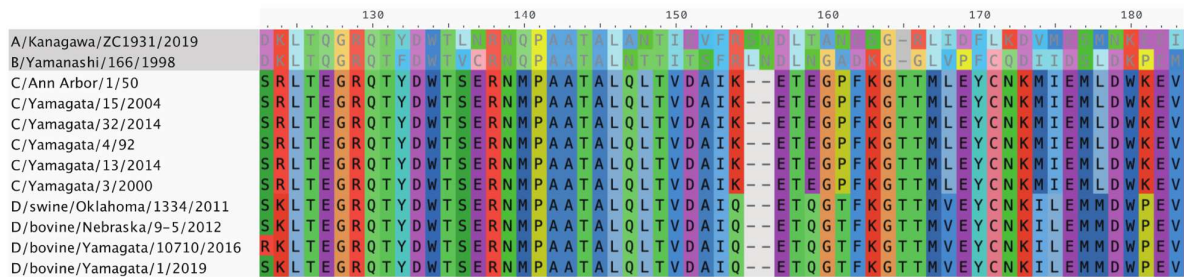
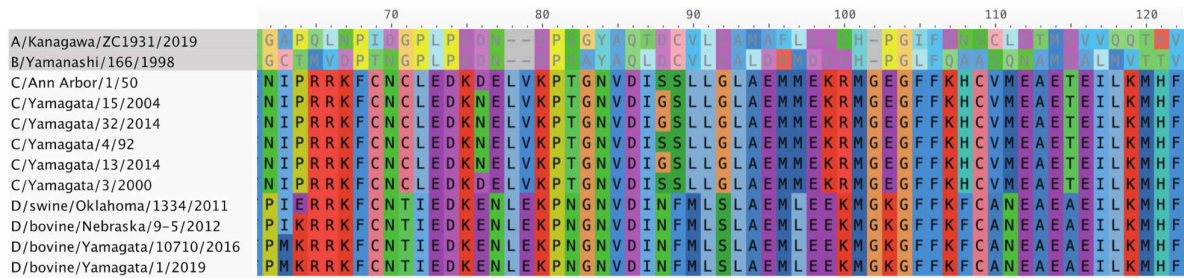
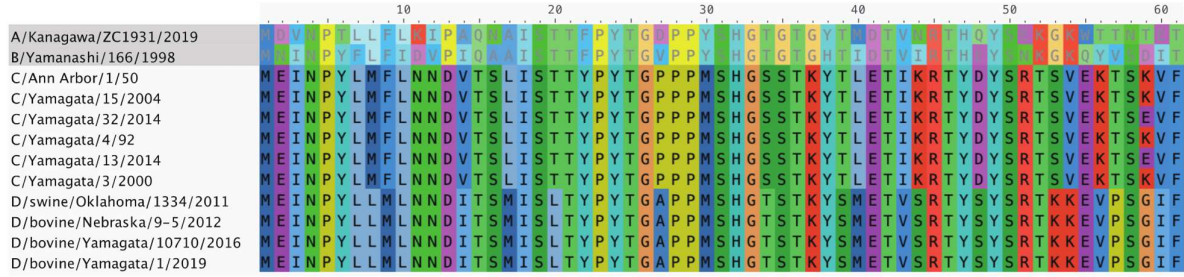


(a)



(b)

Figure S1. Amino acid alignment of selected regions of the P3 (a) and PB1 (b) proteins of the influenza C and D viruses used in this study. A/Kanagawa/ZC1931/2019(H1N1)pdm09 and B/Yamanashi/166/1998 were used as reference strains. Isolate IDs of A/Kanagawa/ZC1931/2019(H1N1)pdm09 and B/Yamanashi/166/1998 in the GISAID EpiFlu Database (<https://gisaid.org>) are EPI_ISL_403549 and EPI_ISL_20976, respectively. Arrows indicate positions 38 and 231, respectively.