

H1N1 2010-2018 (SEM)

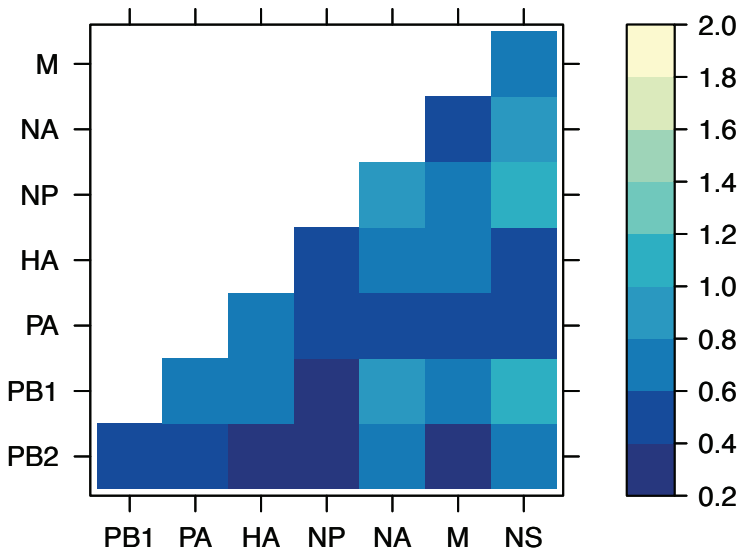


Figure 5 - figure supplement 1. The standard error of the mean (SEM) of replicate Robinson-Foulds distances. The SEM of all pairwise Robinson-Foulds distances (RF) was determined for H1N1 viruses from 2010-2018 (corresponding to the mean tree distances from Figure 5A).