

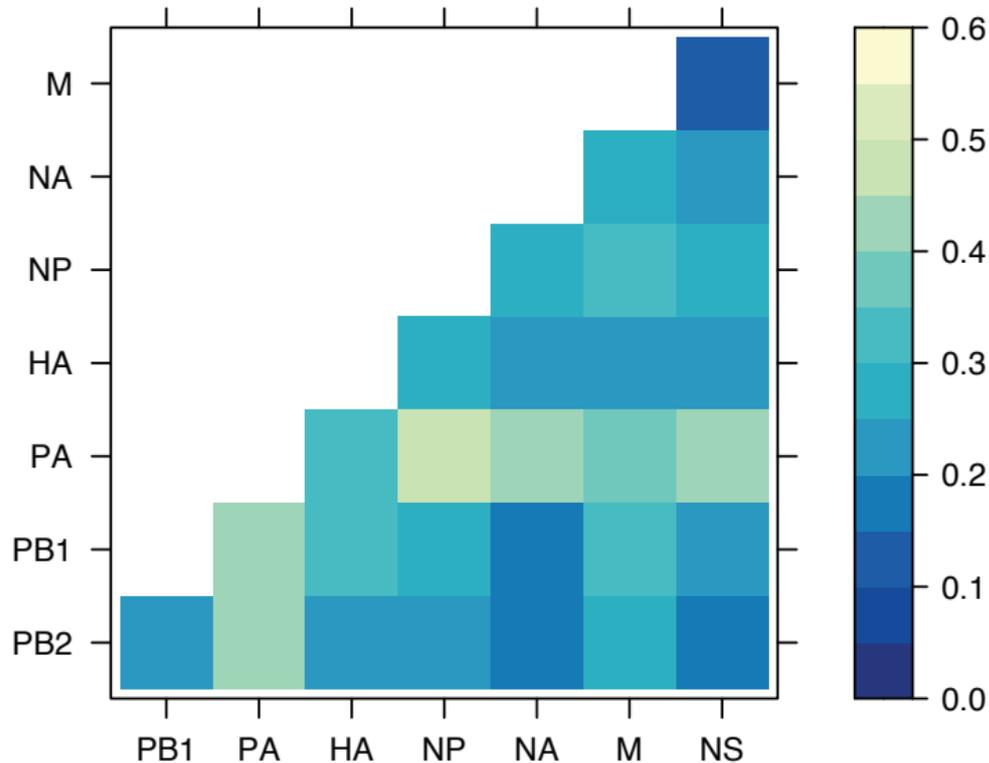
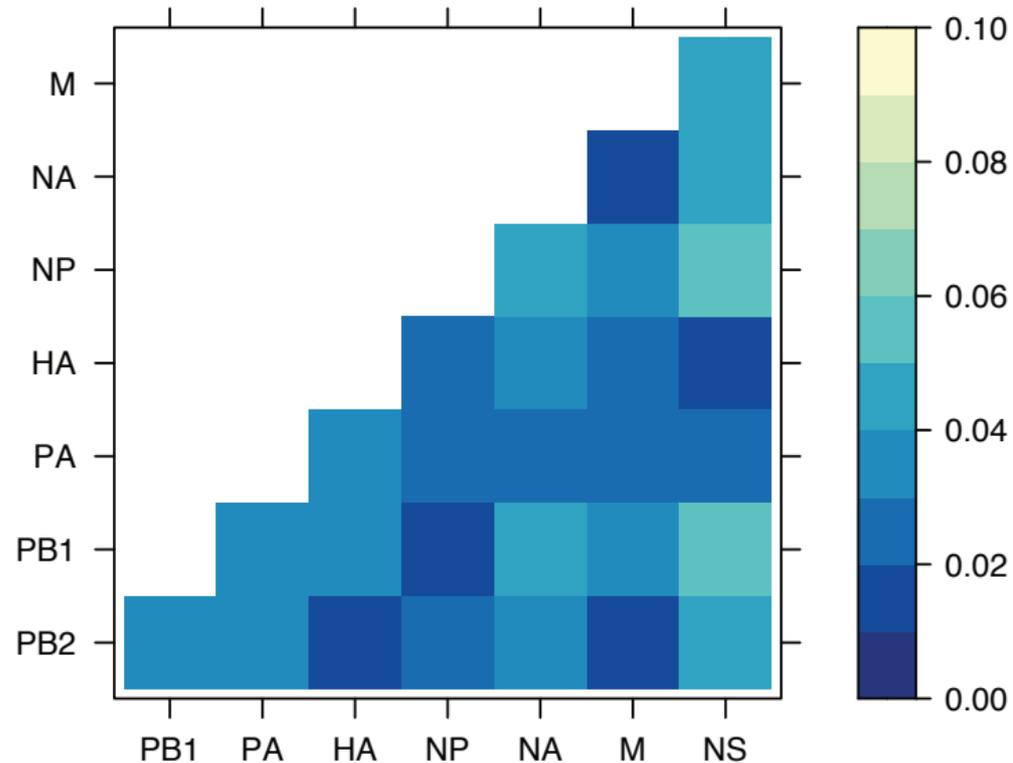
A Mean CID (H1N1 2010-2018)**B SEM (H1N1 2010-2018)**

Figure 5 - figure supplement 2. The mean clustering information distance (CID) of replicate vRNA trees. The pairwise CID was calculated between each replicate vRNA tree from human H1N1 viruses from 2010-2018. The mean CID (**A**) and SEM of all pairwise CID (**B**) were visualized in a heatmap.