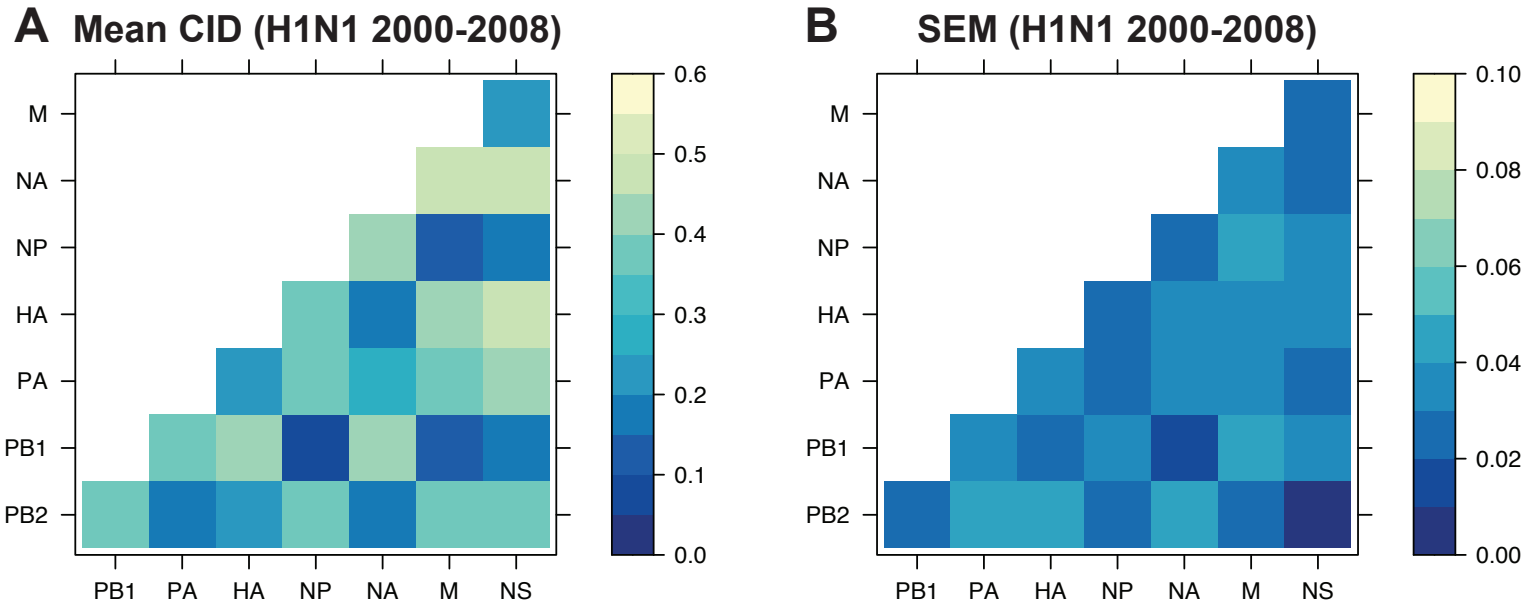

Figure 4 - figure supplement 2. The mean clustering information distance (CID) of replicate vRNA trees. The pairwise CID was calculated between each replicate vRNA tree from human H1N1 viruses from 2000-2008. The mean CID **(A)** and SEM of all pairwise CID **(B)** were visualized in a heatmap.