

Supplementary Figure 4: Maximum likelihood phylogenetic trees of consensus sequences for (A) HA, 2004-2005 (B) NA, 2004-2005 (C) HA 2005-2006 (D) NA 2005-2006 HA with tips coded by vaccine status and pre-season HAI (blue >40, magenta <40). Black, no data (note no NAI data). Outgroups are A/Wyoming/03/2003 (A,B) and A/California/7/2004 (C,D). Bootstrap values (n=1000 bootstraps) are shown and nodes with bootstrap values <50 are collapsed for easier visualization.

