



S2 Fig. Reassortment inference between unpassaged sequences. Histograms of reported HA-NA reassortment events between unpassaged sequences for different core distances δ (red bars) are compared to expected number of false positives (blue bars), which decays exponentially with increasing δ . This result is qualitatively comparable with the distance dependence of real events and false positives that we find including in the analyses also strains subjected to passaging.