S8 Fig. The divergence of adhesin proteins encoded by viruses from the family Hafnaviridae.

(A) Pairwise sequence alignments of adhesins encoded by viruses from the genera Haloferacalesvirus and Mincapvirus. The hypervariable segments (HVSs) which are predicted to interact with host receptor are indicated.

(B) Phylogeny of the adhesins. Viruses that were used for efficiency of plating (EOP) assay in this study are indicated in black, otherwise in grey. Nodes with bootstrap support values greater than 90% are indicated with dots.