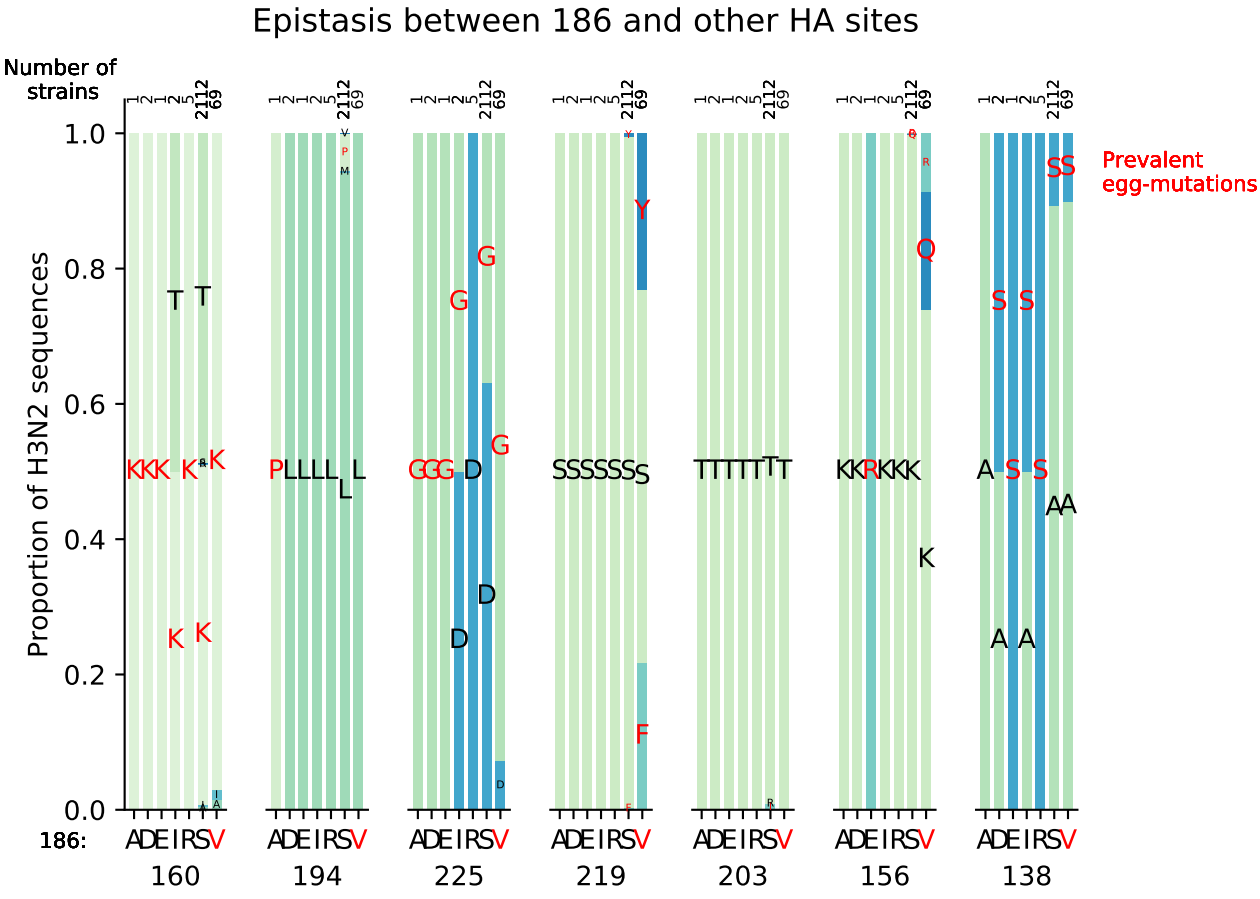
Epistasis between 186 and other HA sites
Number of strains
Proportion of H3N2 sequences
Prevalent egg-mutations
186:
ADEIRSV
160
194
225
219
203
156
138