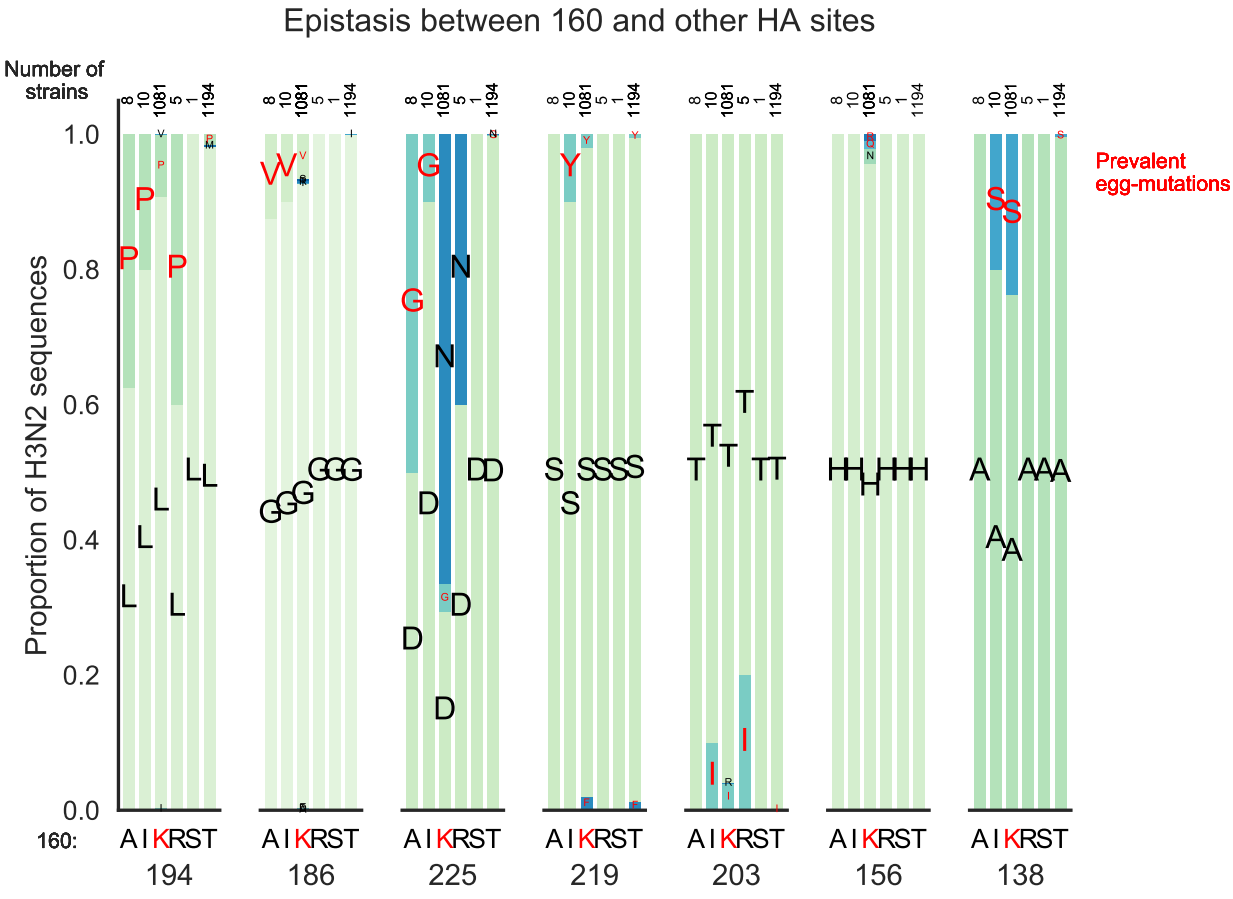
Epistasis between 160 and other HA sites
Number of strains
8 10 1081 5 1 1194
Proportion of H3N2 sequences
1.0
0.8
0.6
0.4
0.2
0.0
160: A I K R S T
194
186
225
219
203
156
138
Prevalent egg-mutations