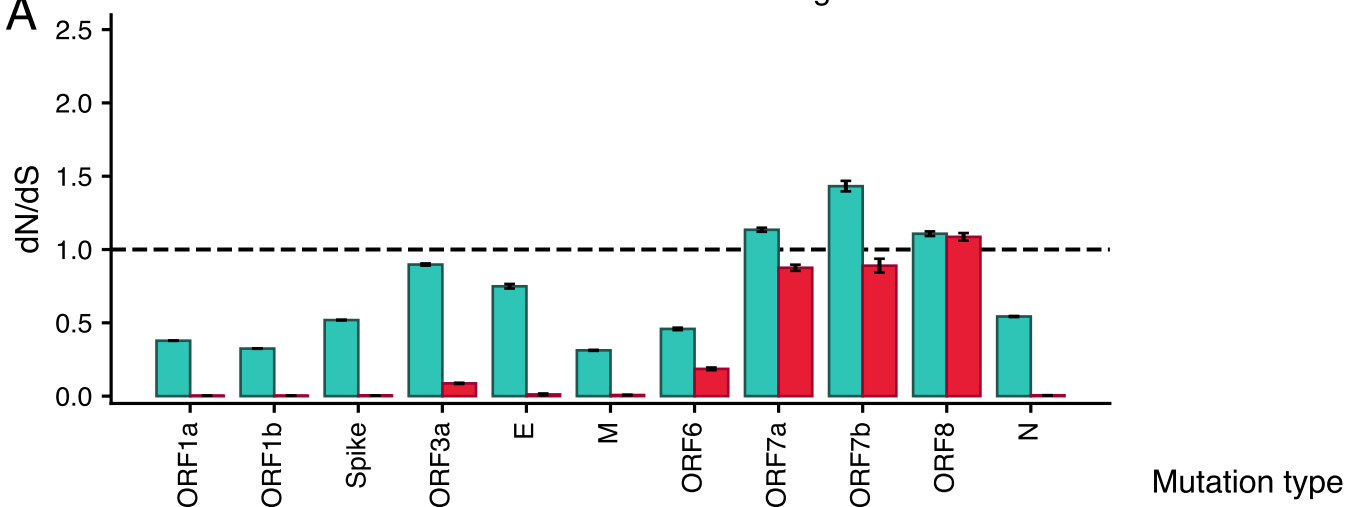


Substitution rates calculated from 4-fold degenerate sites

A



Substitution rates calculated from all sites in genome

B

