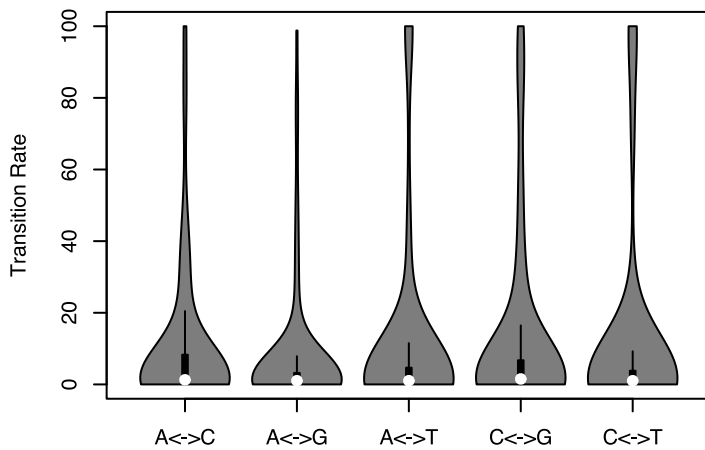
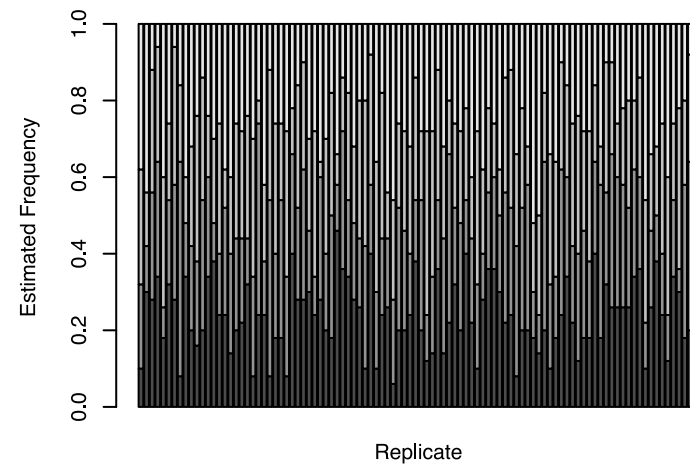


**A**

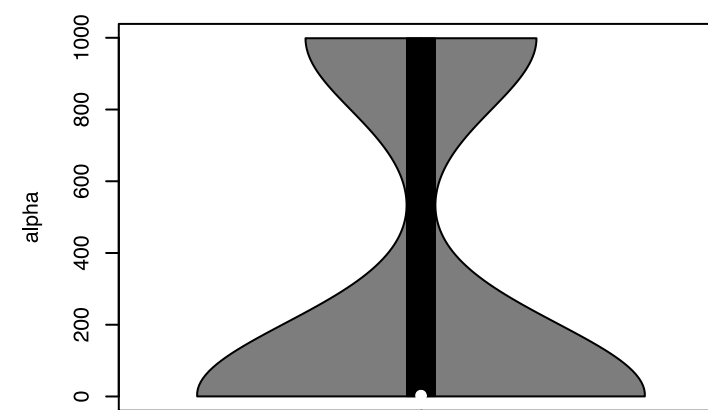
Transition rate estimates

**B**

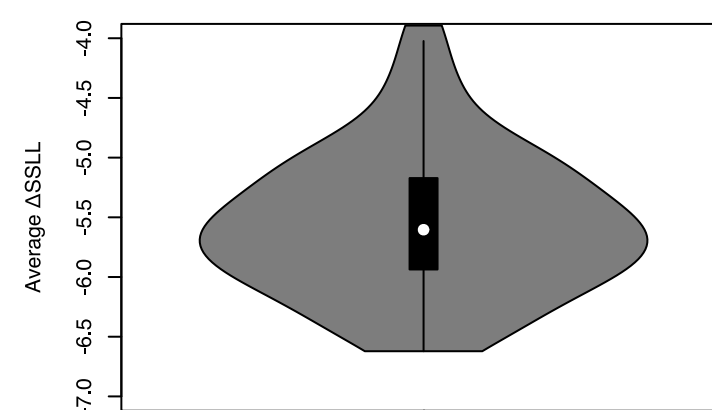
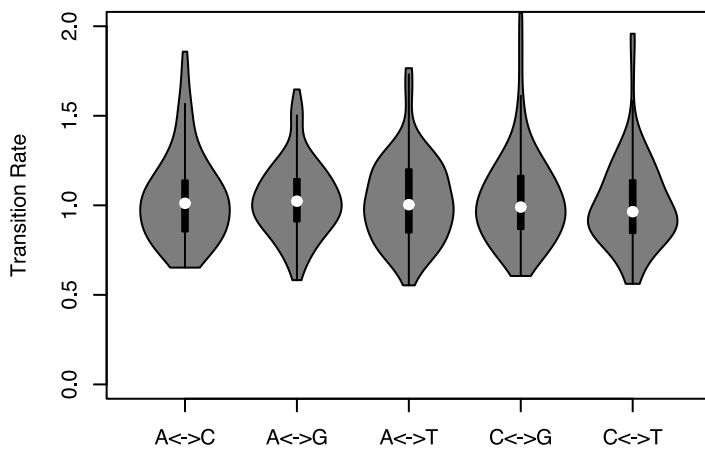
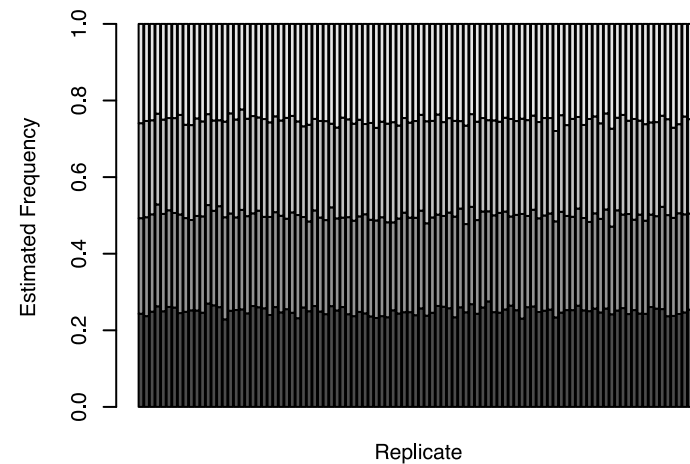
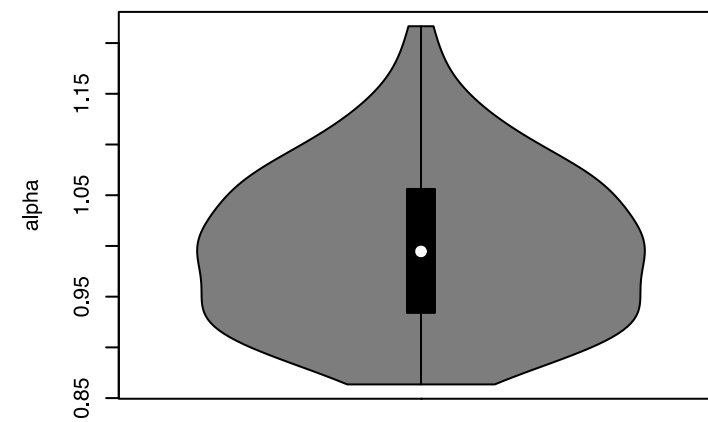
Nucleotide frequency estimates

**C**

Gamma estimate

**D**

Likelihood Estimates

**E****F****G****H**