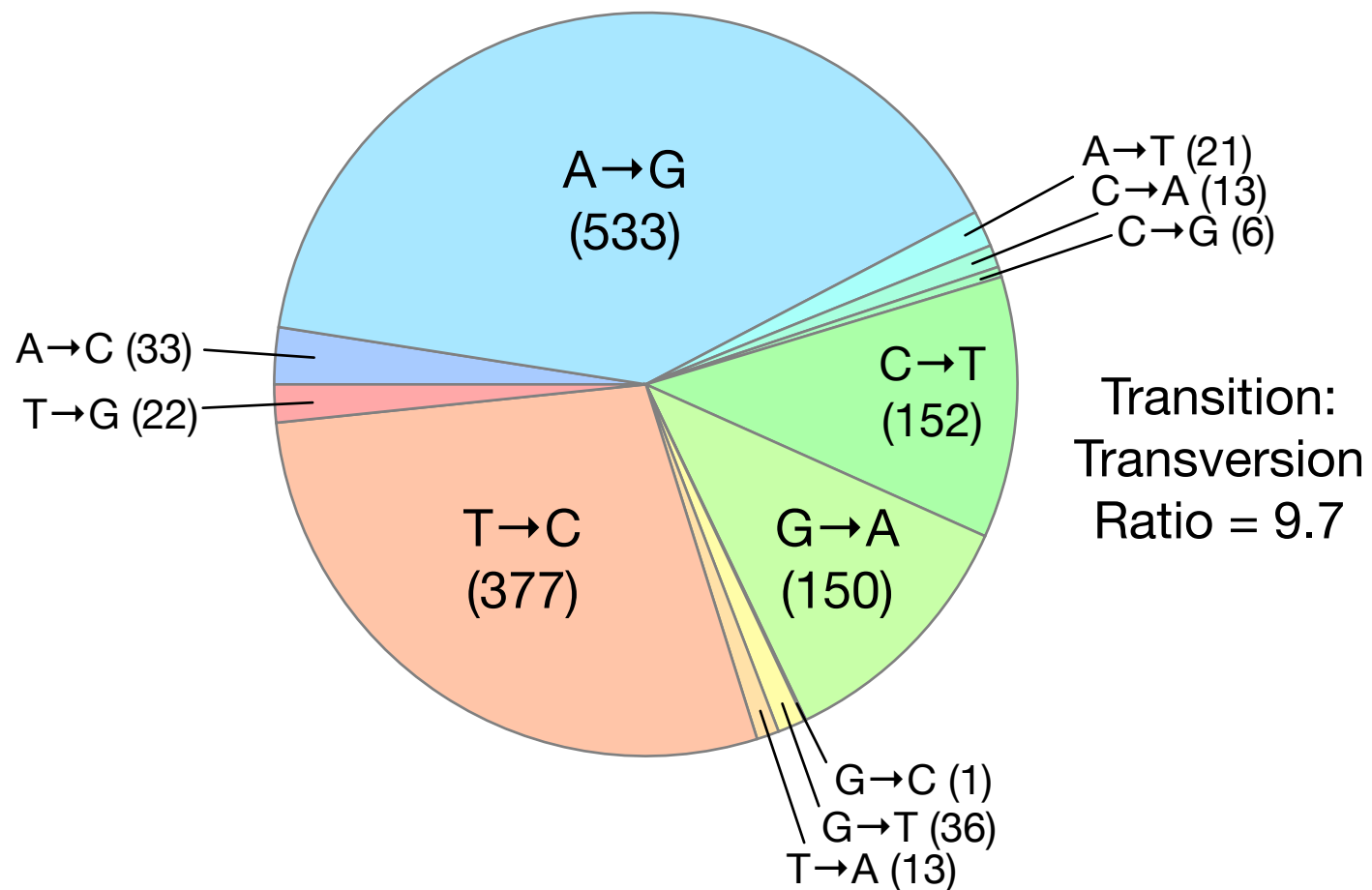


(A)

Number of observed mutations



(B)

Fraction of observed mutations
(normalised by sequence content)

