

T Mutation rates at A/G/N reference nucleotides

prop

0.4
0.2
0.0

0% I at A site

10% I at A site

100% I at A site

50% I at A site

A at A site

I at A site

I at A site, TGIRT

Multiple Inosines: A at A site

Multiple Inosines: I at A site

Multiple Inosines: I at A,B sites

Multiple Inosines: I at A,B,C sites

Multiple Inosines: I at all sites

Multiple Inosines: I at B site

cond

color

A-site

B-site

C-site

D-site

E-site

NA

treat

CE2

untreated

