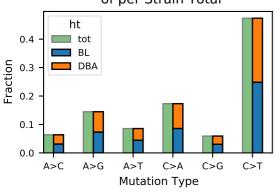
Average of Mutation Spectrum as Fraction of per Strain Total



Fraction 0.10 0.05 0.00 A>C A>G **Mutation Type** 

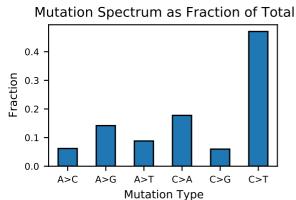
ht BL

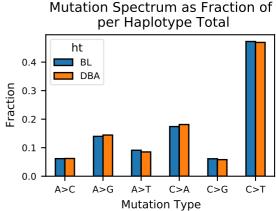
DBA

0.25

0.20

0.15





A>T

C>A

Average by Haplotype

C>G

C>T