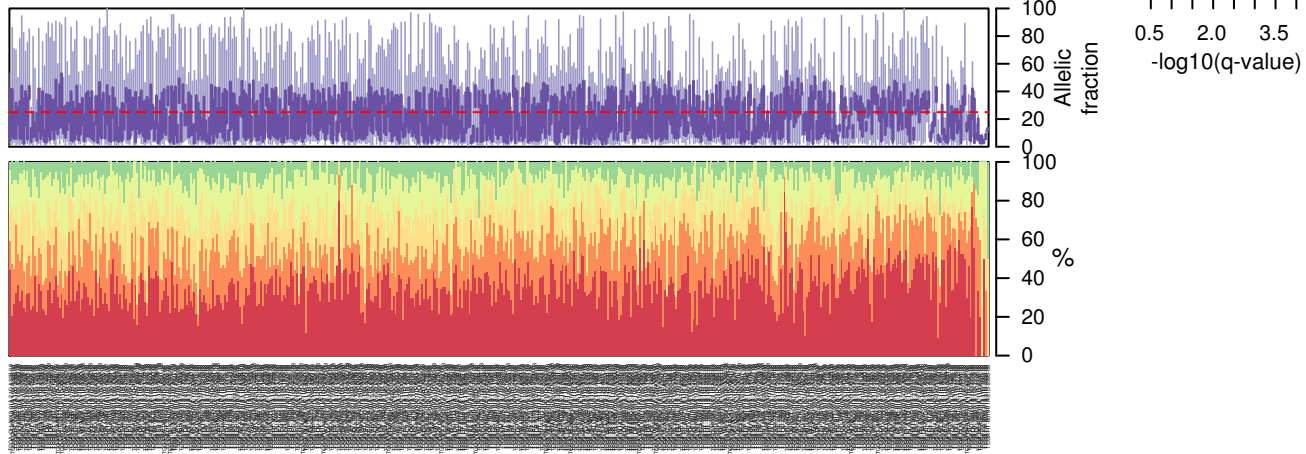


mutations

A->(T/C)
 C->(G/A)
 A->G
 *Cp(A/C/T)->T
 *CpG->T



0.5 2.0 3.5
 -log10(q-value)