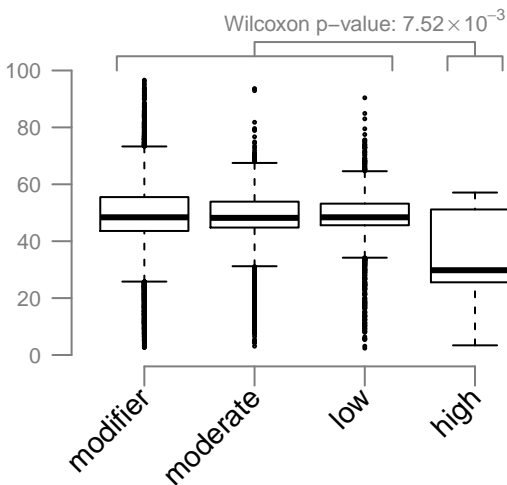


a**Expected functional impact**

Estimated percentage alternative allele

**b****SNP Effect**

Estimated percentage alternative allele

