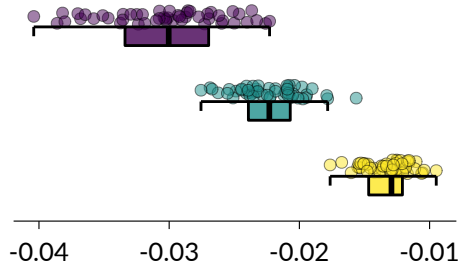
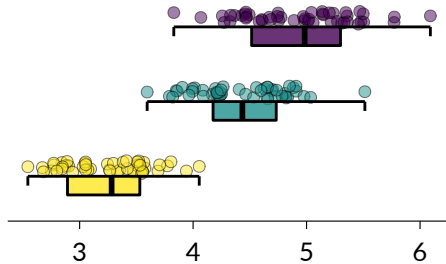


a



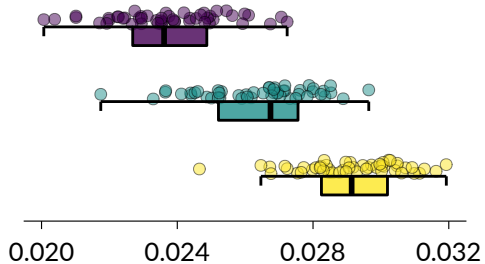
Mutation load
(Selection coefficient per cM)

b






Number of mutations per cM

c



Mean mutation frequency

ROH length class  long ($>12.5\text{cM} \mid < 4\text{g}$)  medium ($1.56\text{cM}-12.5\text{cM} \mid 4-32\text{g}$)  short ($0.39\text{cM}-1.56\text{cM} \mid 32-128\text{g}$)