High Inversion Mutation Rate (1e-3) Strong Selection Moderate Selection Weak Selection Amount of Additive Genetic Variance Inside Inversions  $\stackrel{\text{D}}{\circ}$ **QTN Mutation Rate** 0.000001 0.000001 0 0.001 0.01 0.1 0.25 0.4 0.5 0.001 0.01 0.1 0.25 0.4 0.5 0.001 0.01 0.1 0.25 0.4 0.5 Migration Rate