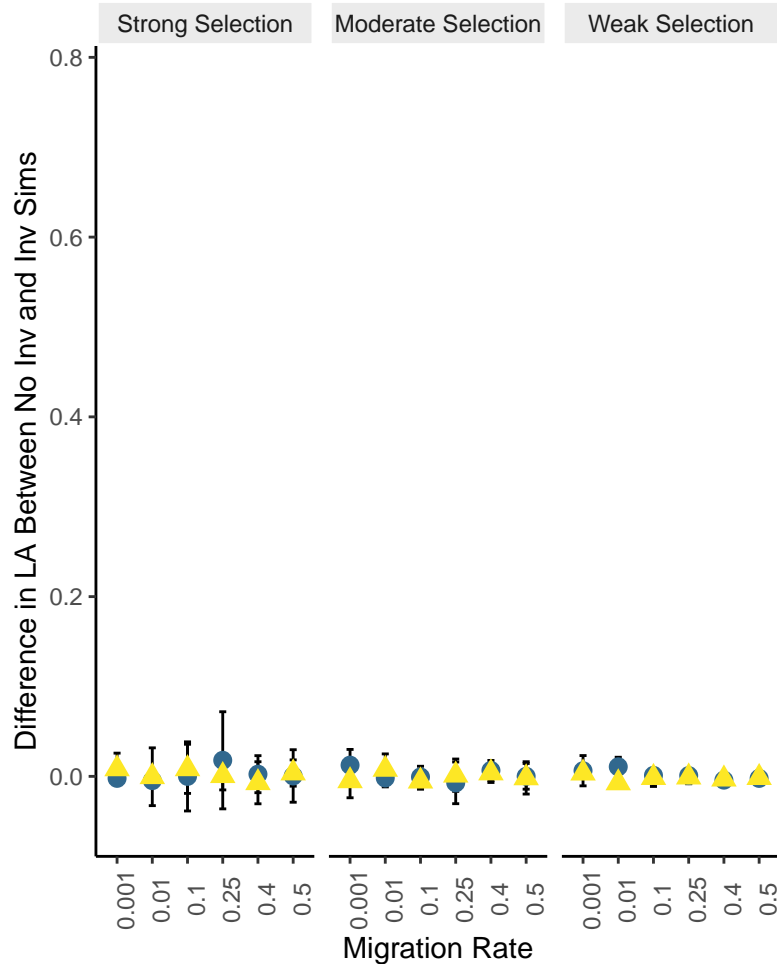
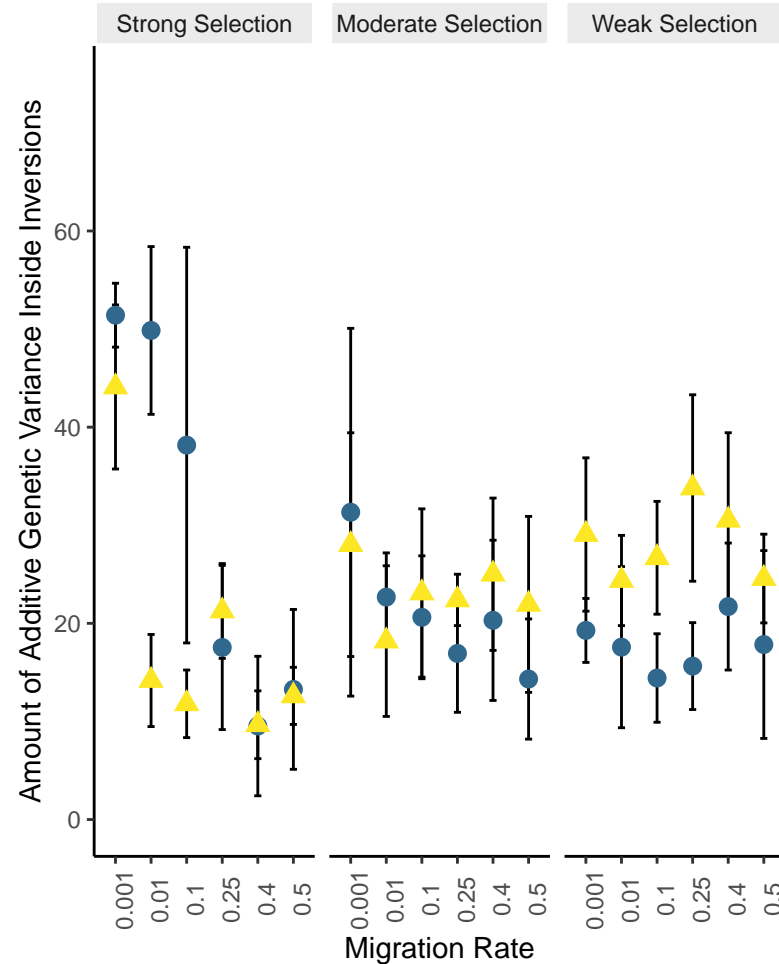


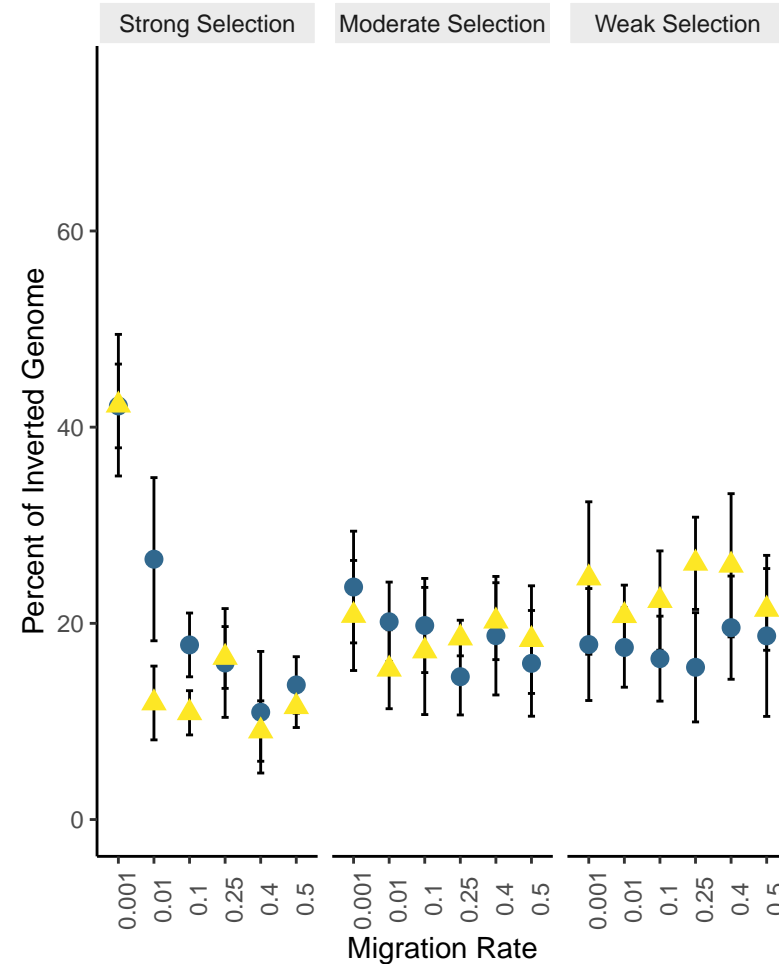
Difference in Amount of Local Adaptation



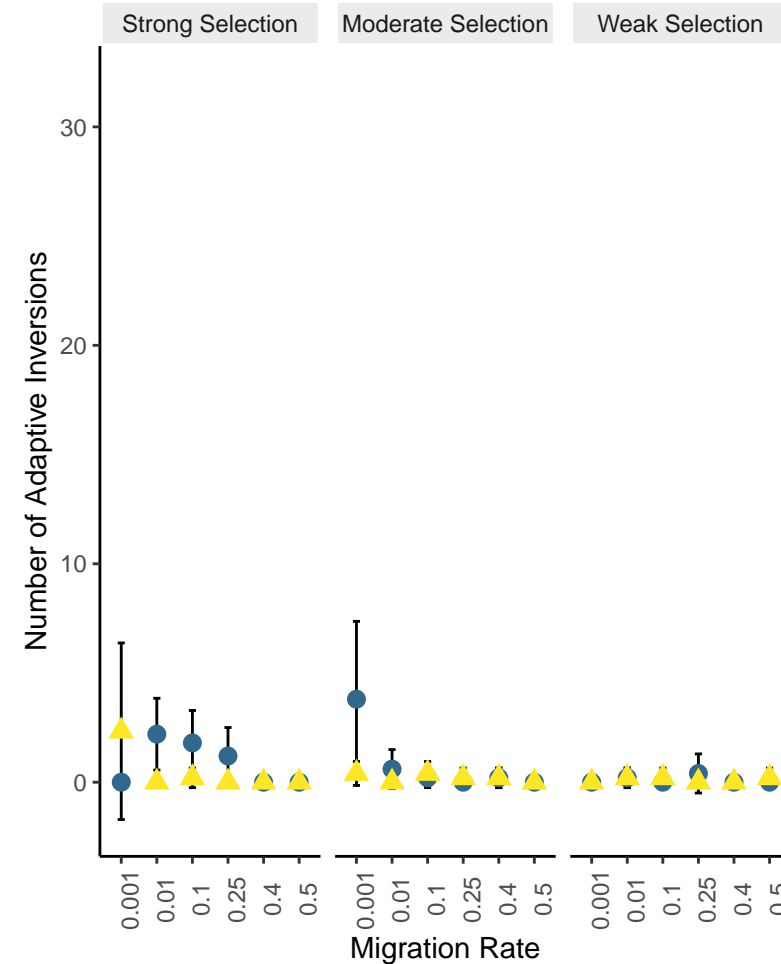
Percent of Additive Genetic Variance in Inversions



Percent of Inverted Genome



Number of Adaptive Inversions



QTN Mutation Rate (Ne*mu)

- 0.0002
- 0.002