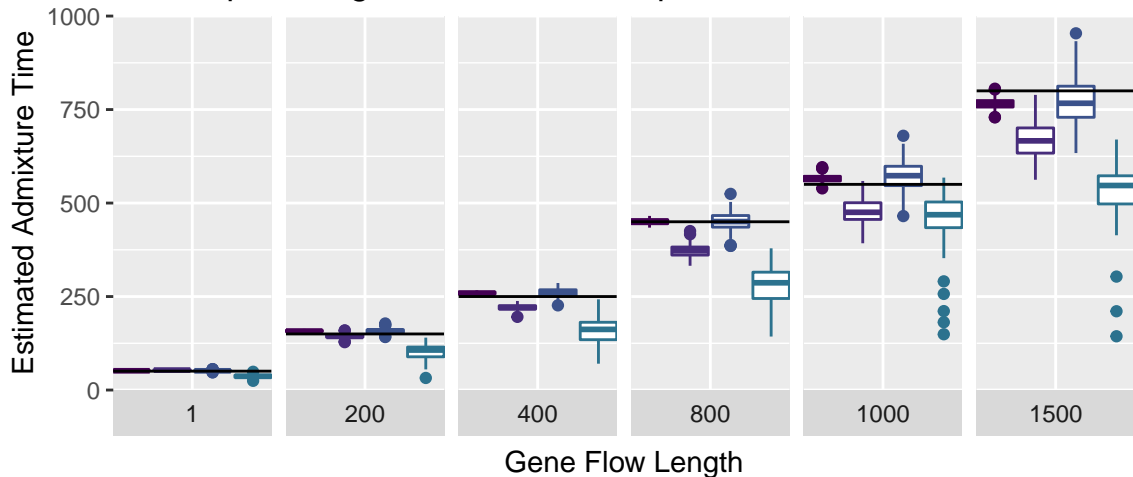
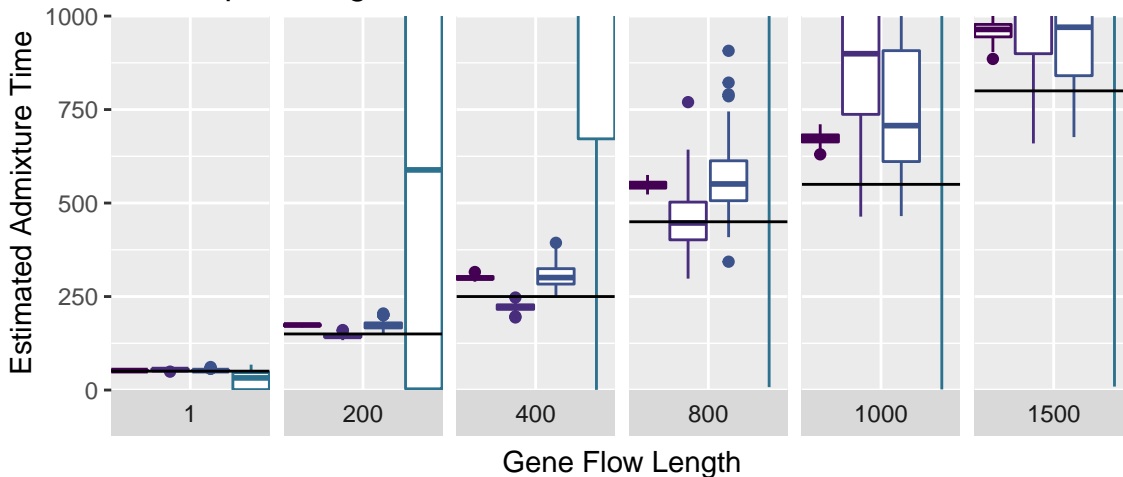
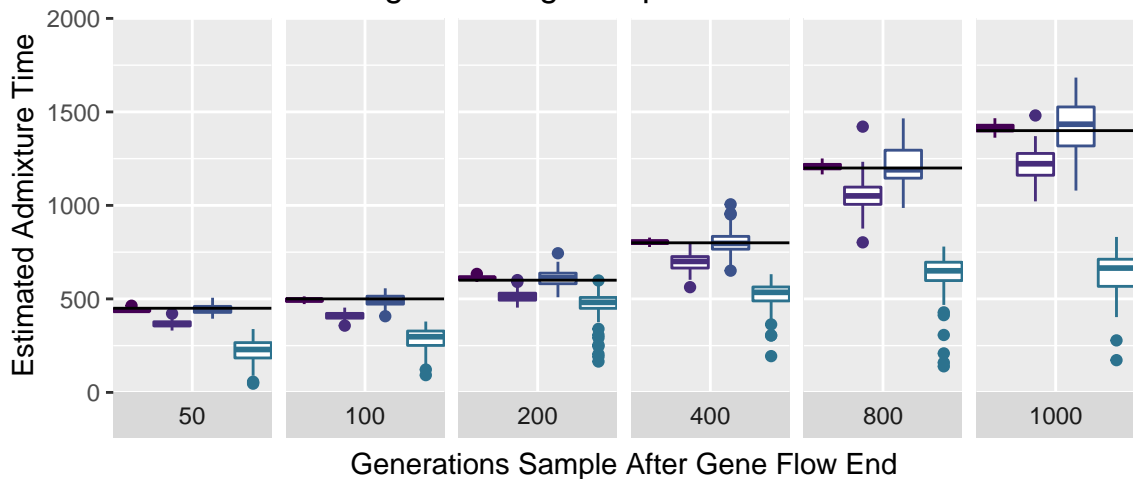
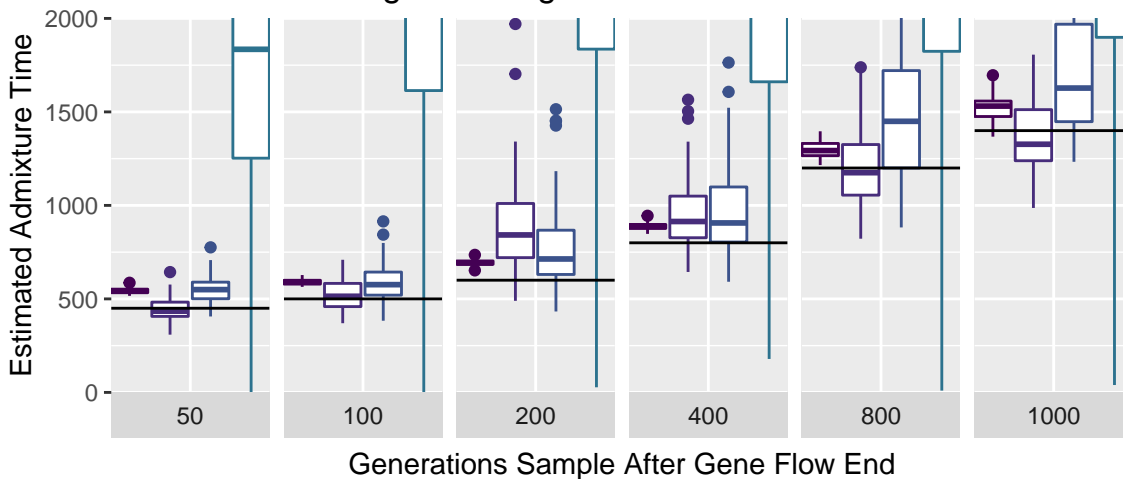


**A** All sampled 50 gen from GF end exponential fit**B** All sampled 50 gen from GF end lomax fit**C** All Gene Flow length = 800 gen exponential fit**D** All Gene Flow length = 800 gen lomax fit

Simulations: Constant Recombination, HapMap AMap corrected, HapMap HapMap corrected, HapMap not corrected