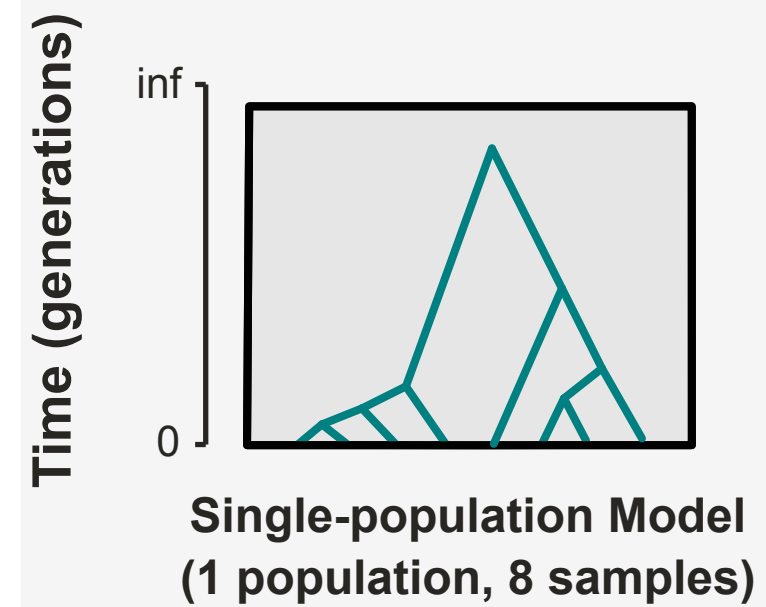
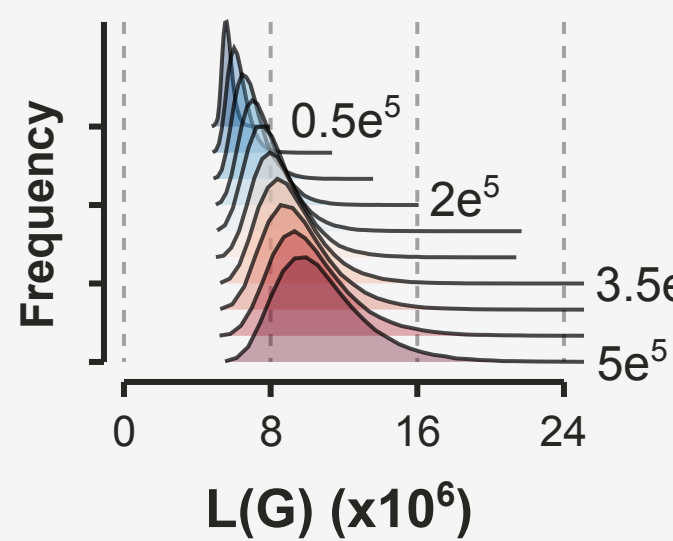
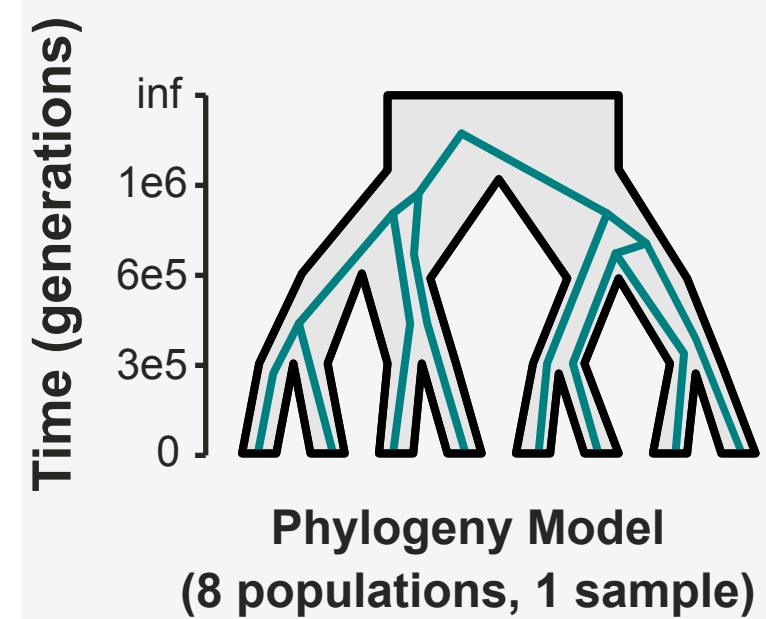
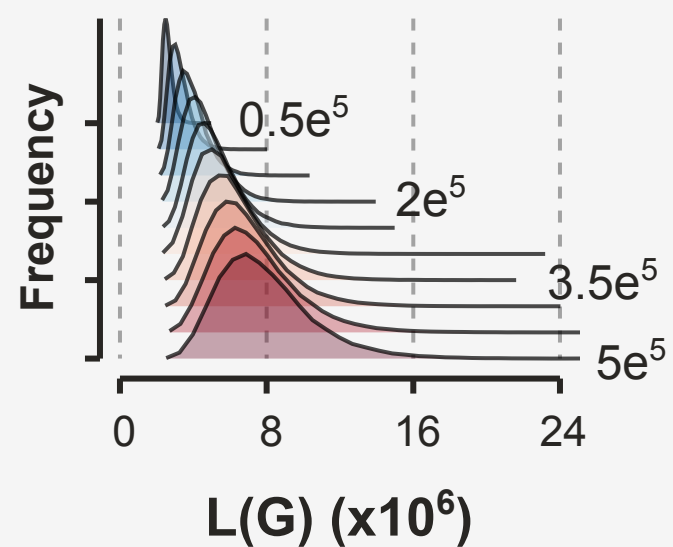
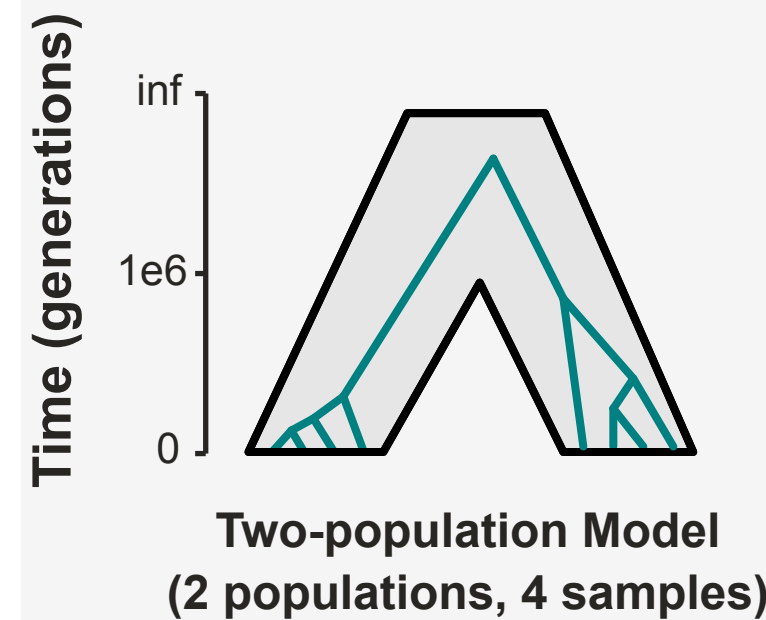
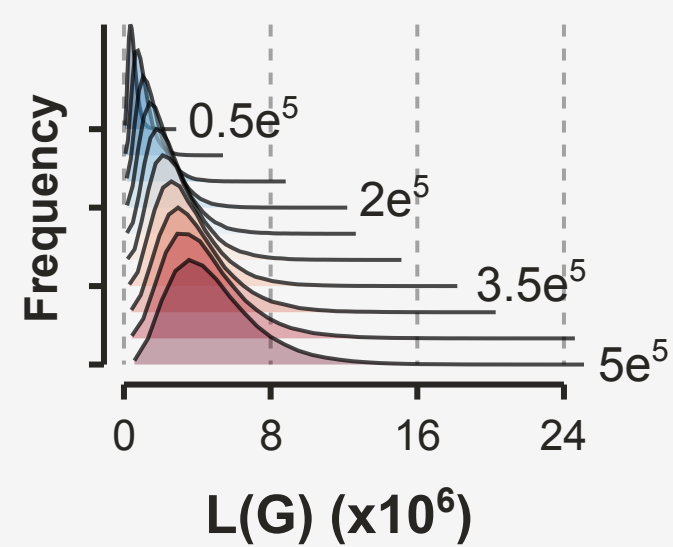


(a) Demographic Model with constant N_e

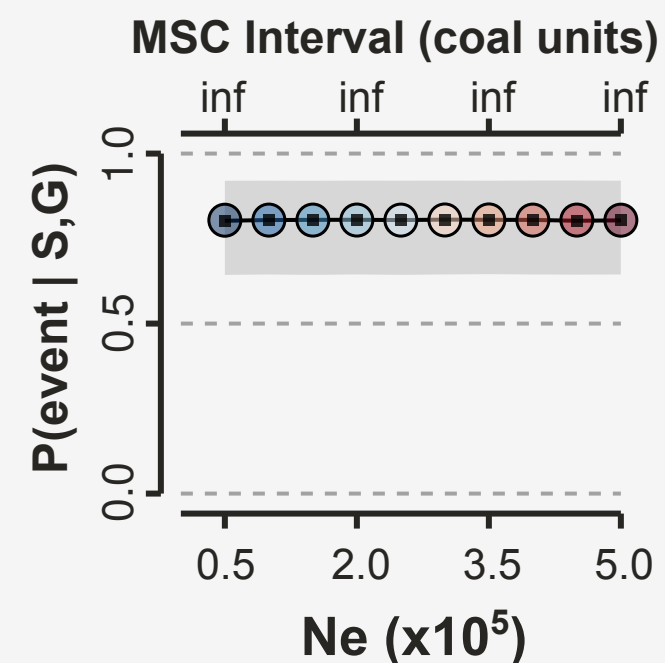


(b) Summed Genealogy Branch Lengths by N_e

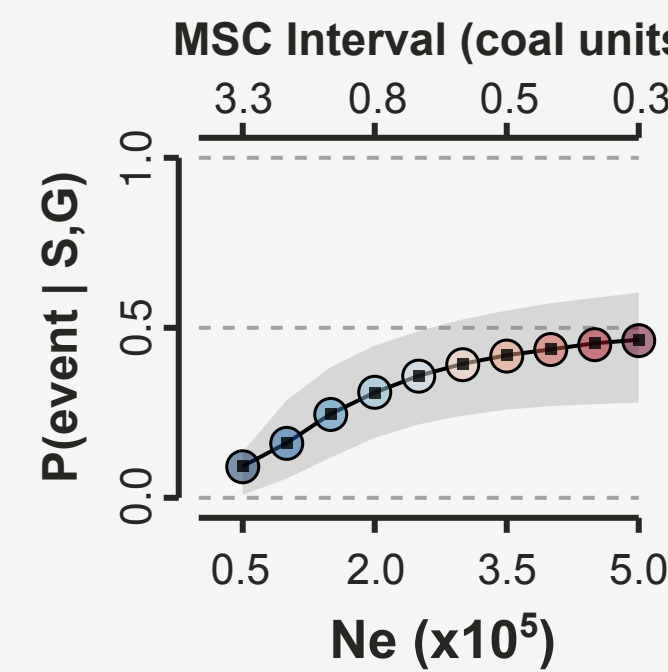
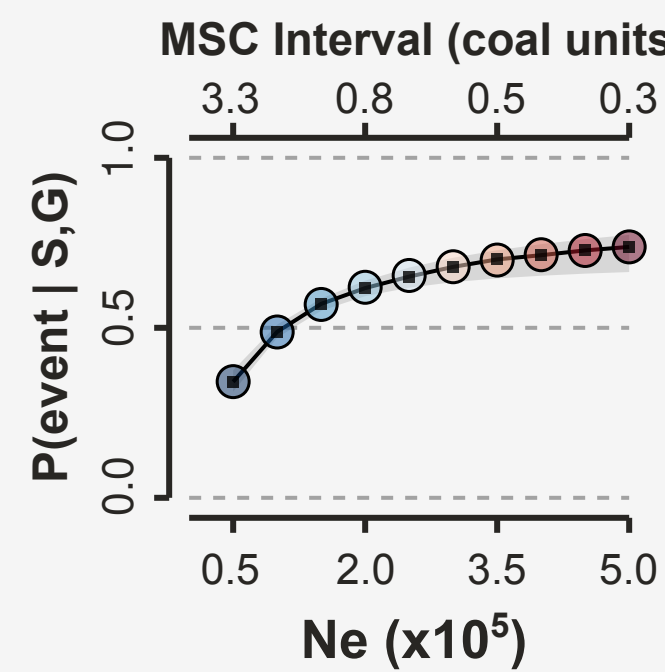
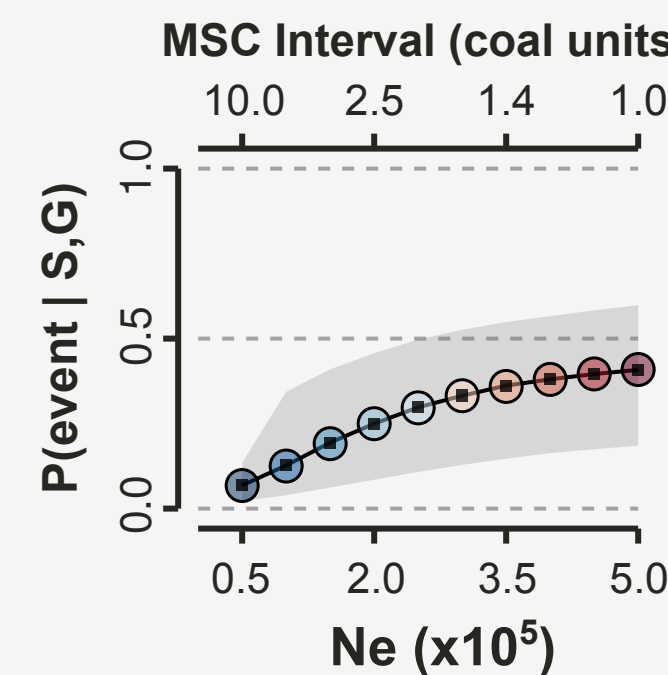
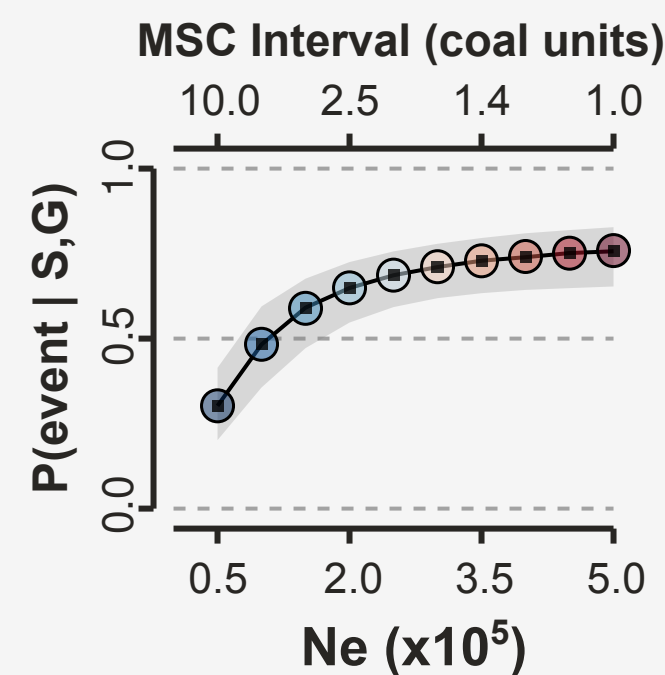
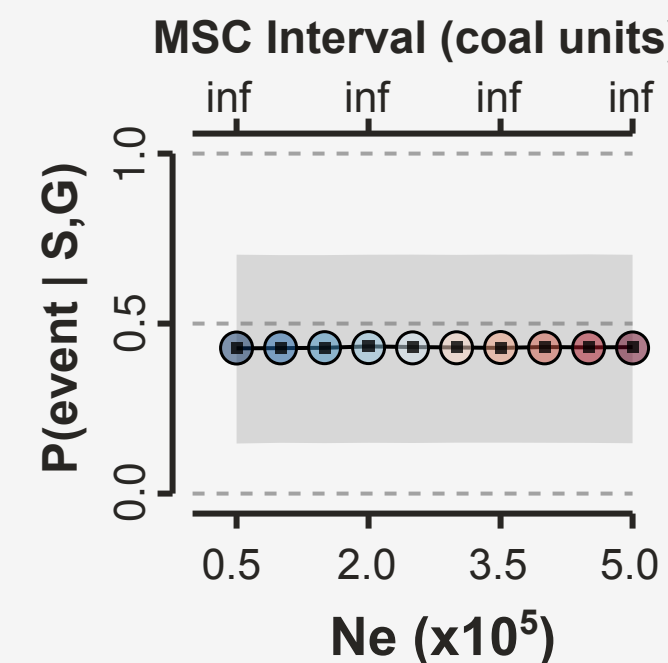


Event probabilities by N_e

(c) tree-change probability

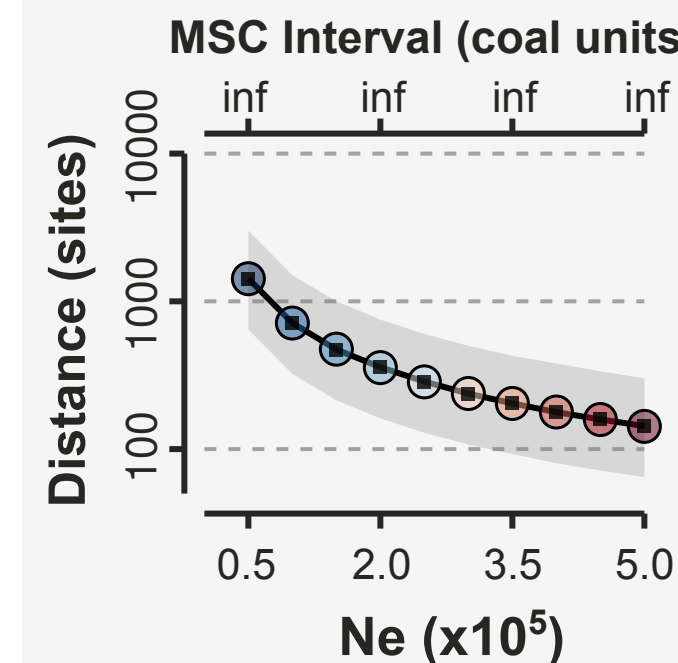


(d) topo-change probability



Waiting distances by N_e

(e) tree-change distance



(f) topo-change distance

