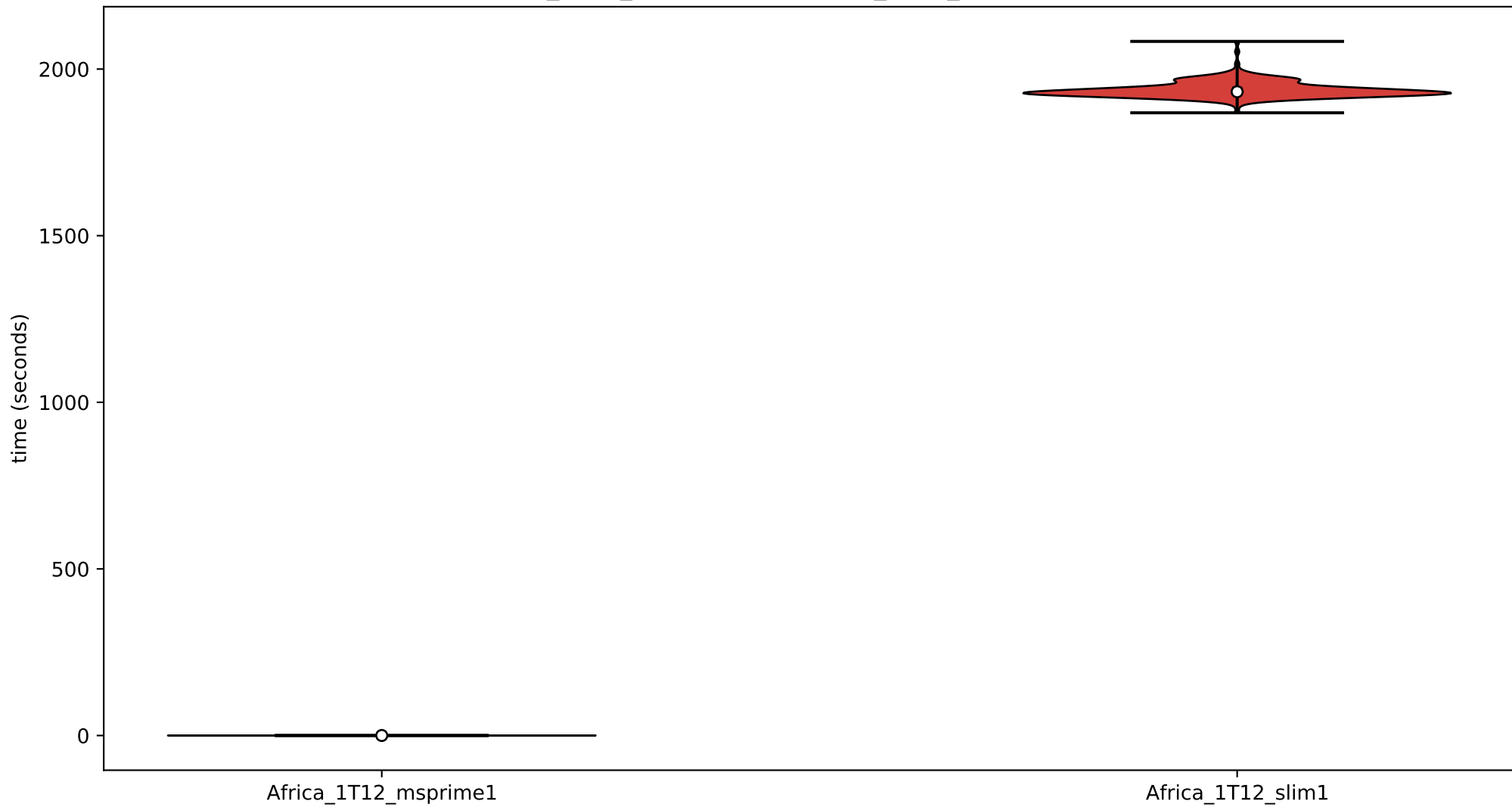
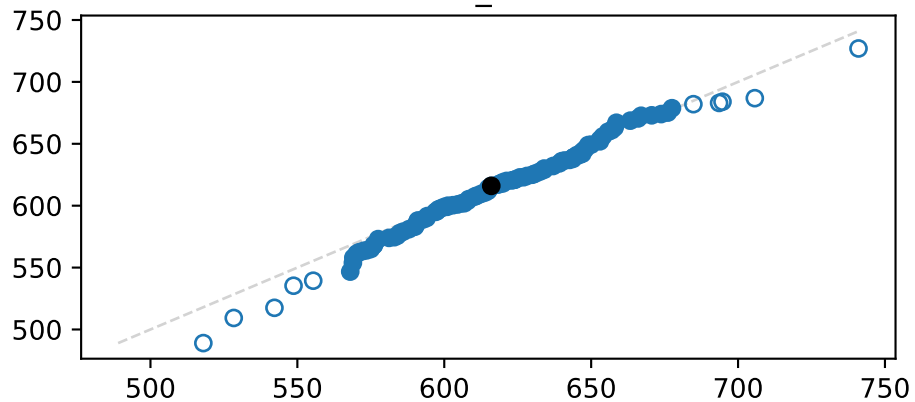


Africa_1T12_msprime1 vs. Africa_1T12_slim1: run time

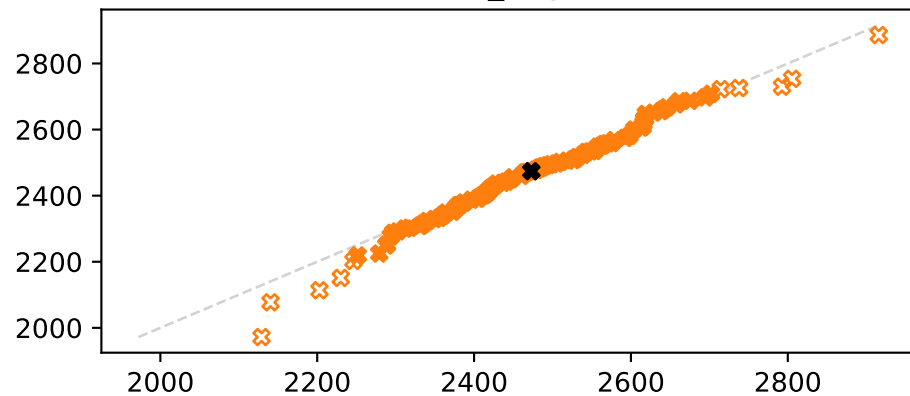


Africa_1T12_msprime1 vs. Africa_1T12_slim1: ts_properties

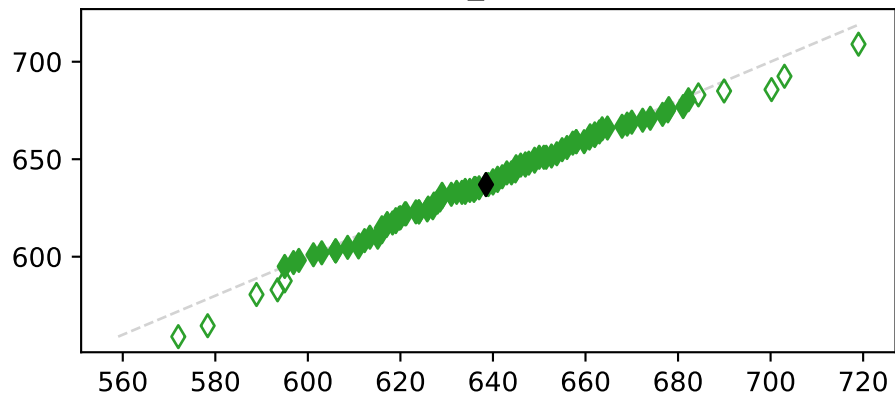
num_trees



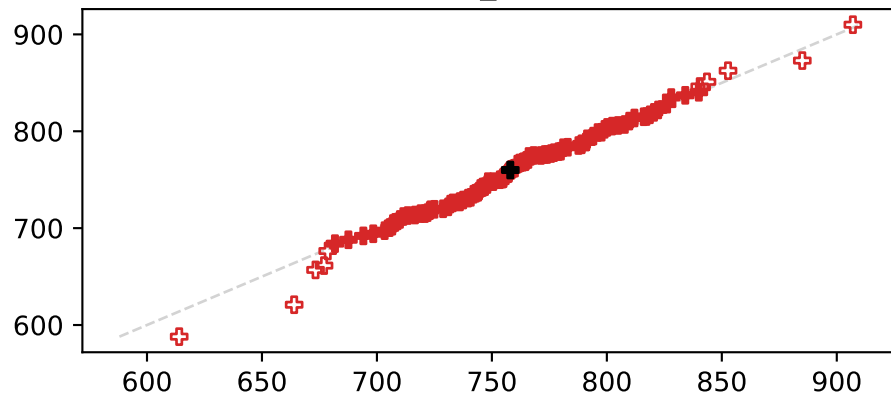
num_edges



num_nodes



num_sites

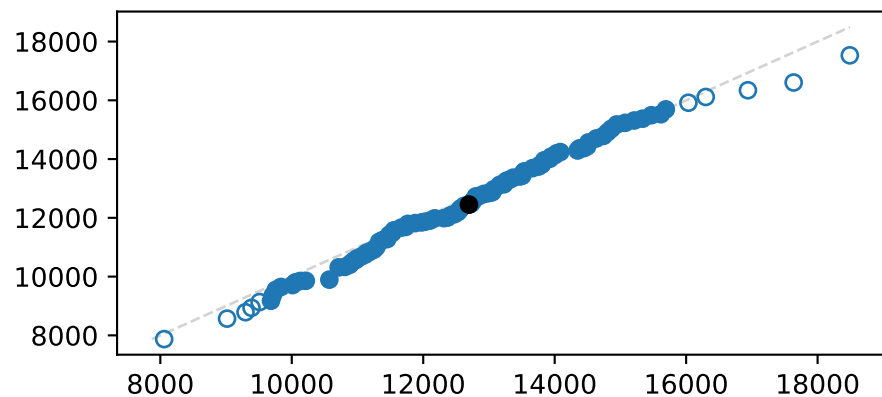


Africa_1T12_msprime1

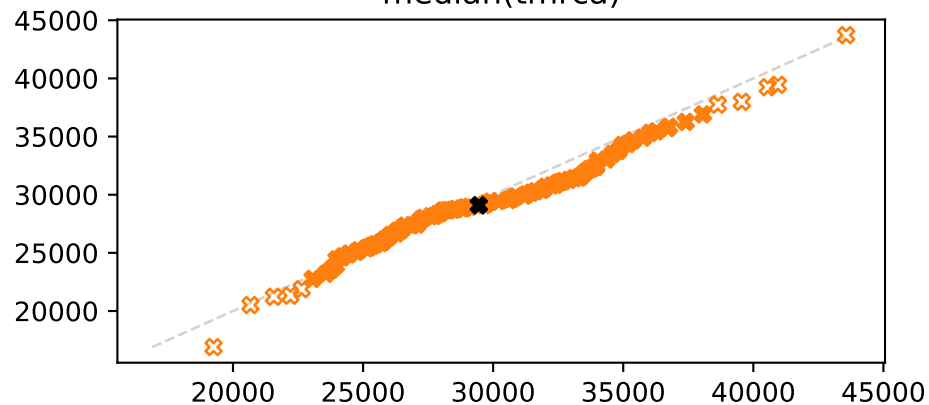
Africa_1T12_slim1

Africa_1T12_msprime1 vs. Africa_1T12_slim1: tmrca

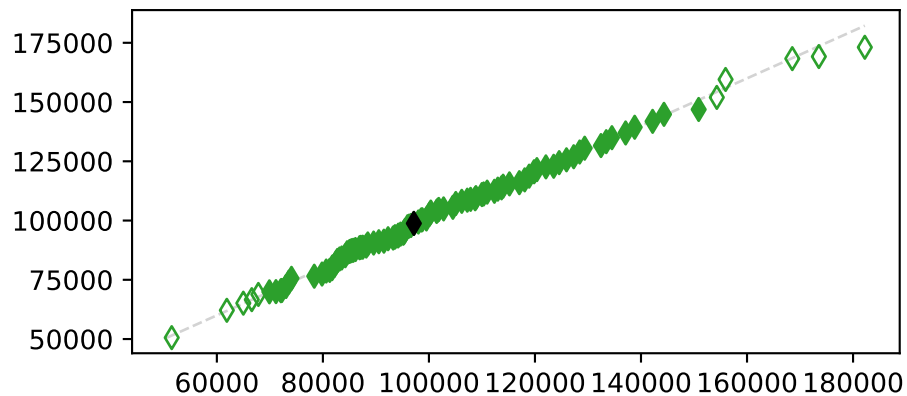
min(tmrca)



median(tmrca)

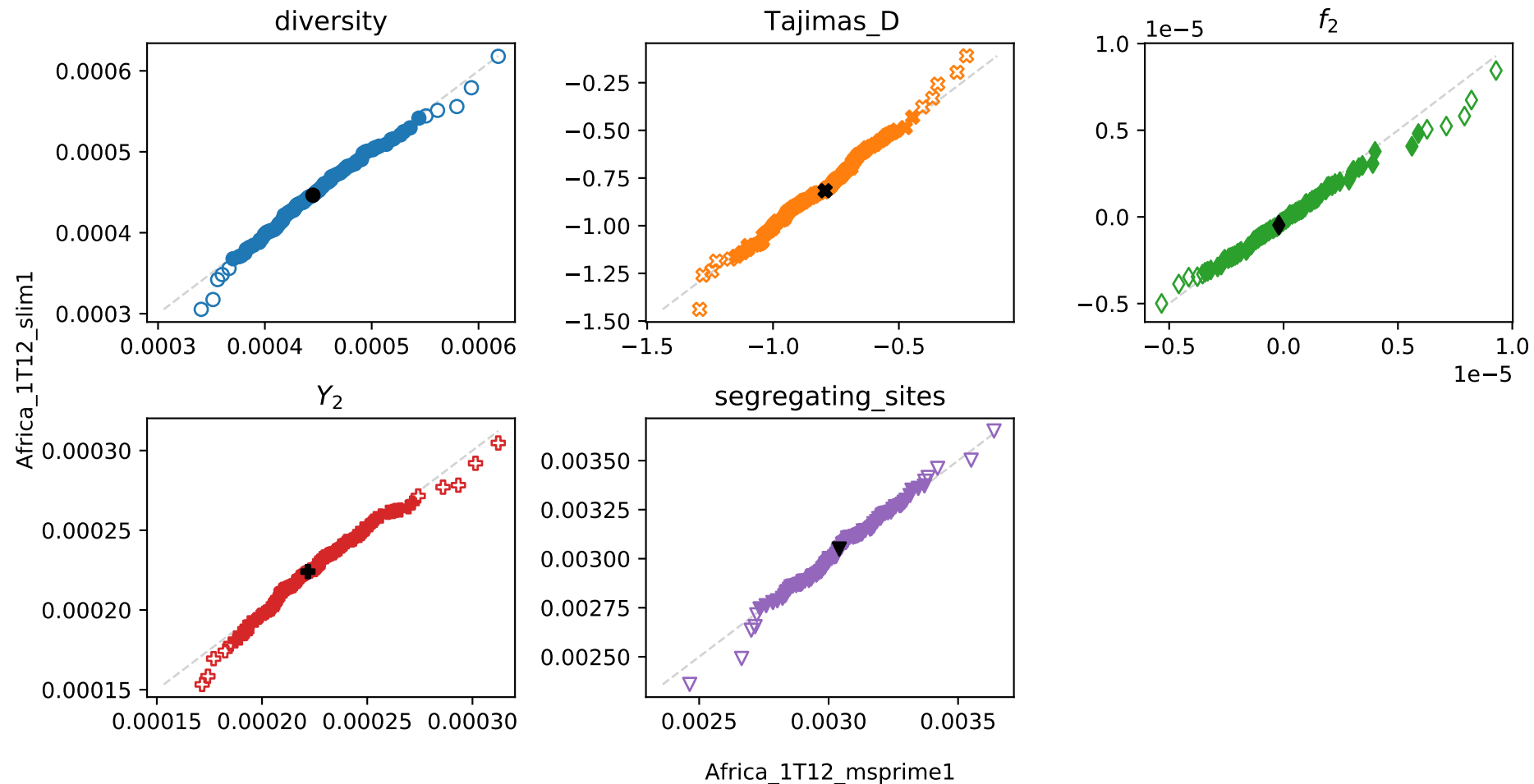


max(tmrca)



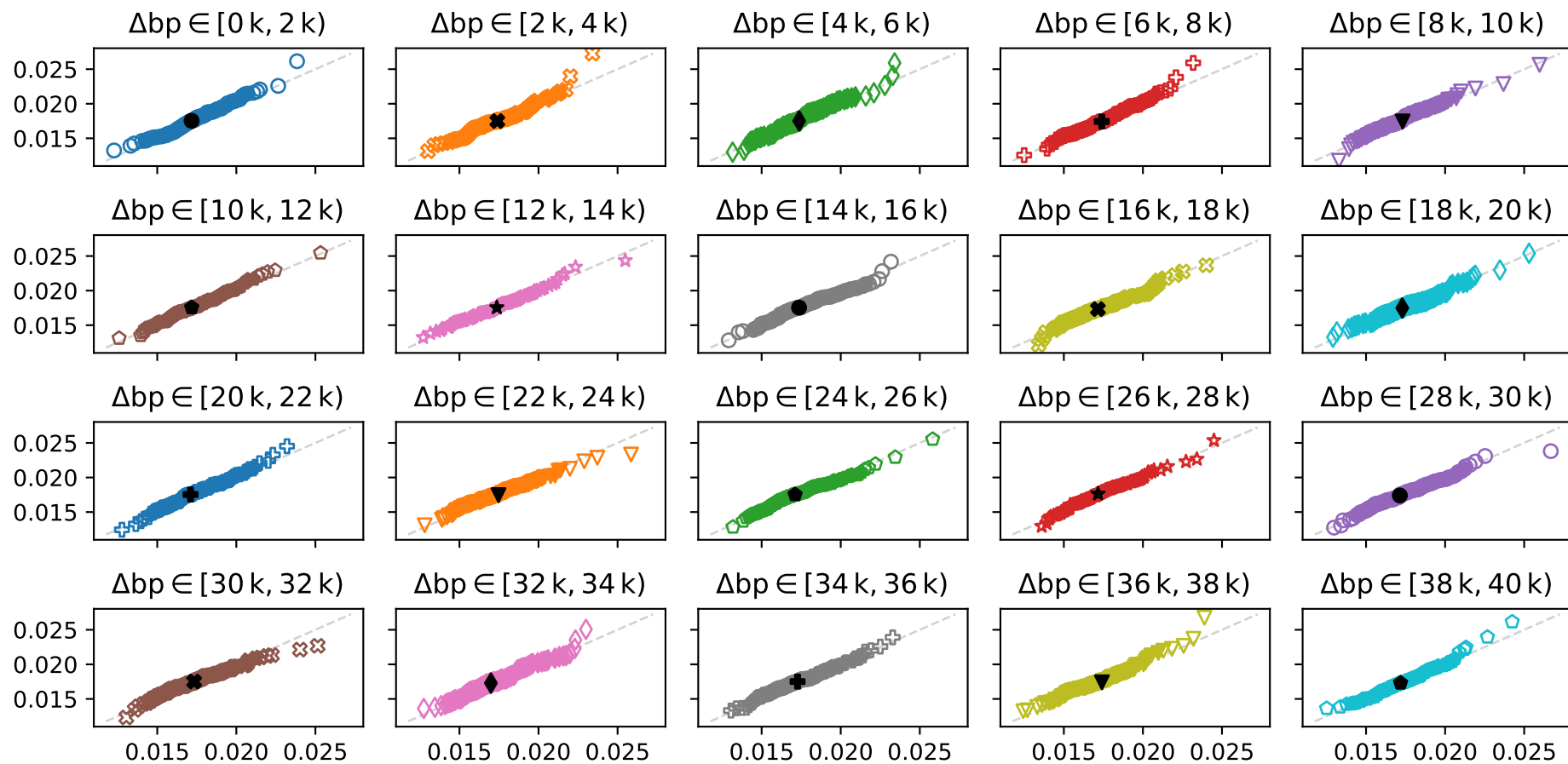
$\text{Africa_1T12_msprime1}$

Africa_1T12_msprime1 vs. Africa_1T12_slim1: pooled_pop_stats



Africa_1T12_msprime1 vs. Africa_1T12_slim1: linkage_disequilibrium

Africa_1T12_slim1



Africa_1T12_msprime1

Africa_1T12_msprime1 vs. Africa_1T12_slim1: allele_frequency_spectrum

Africa_1T12_slim1

