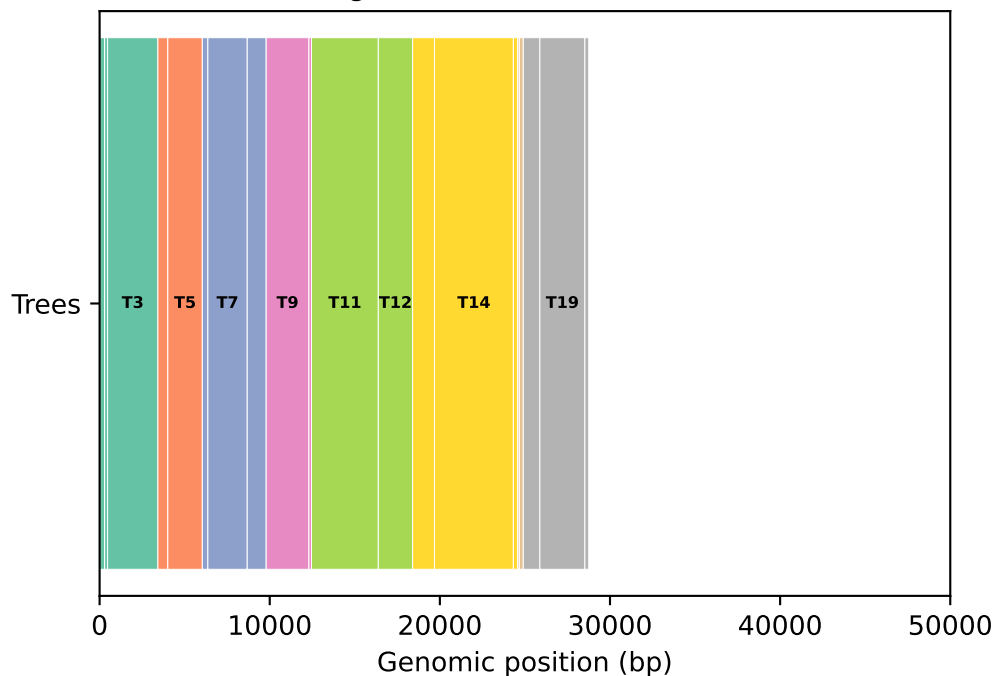
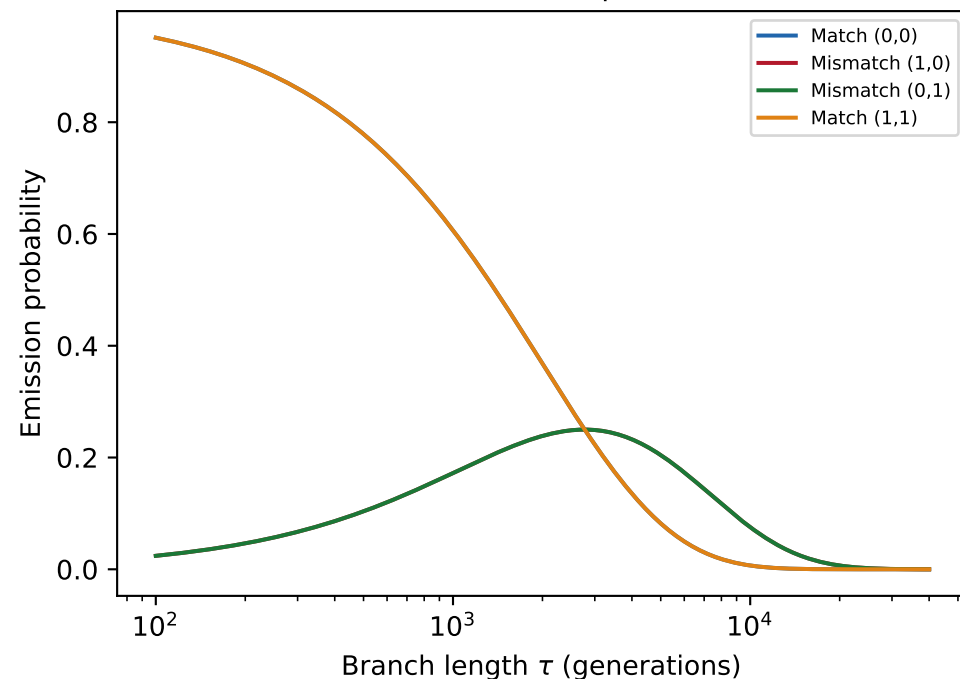


Demo: SINGER on msprime-simulated Data (8 haps, 54 sites)

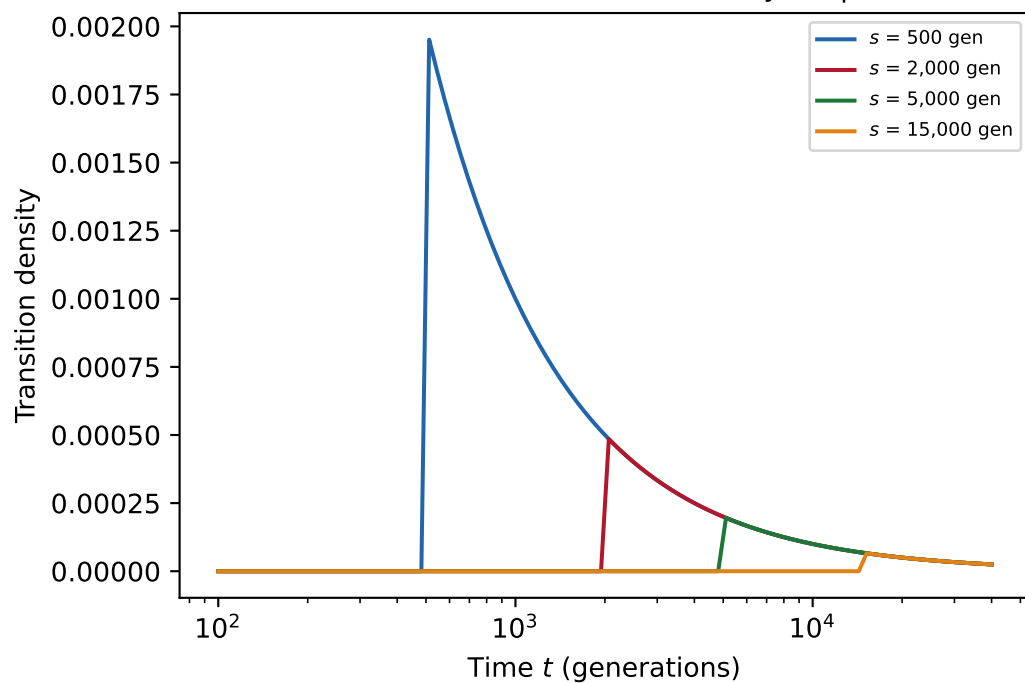
A. True marginal trees (44 trees, 54 mutations)



B. SINGER emission probabilities



C. PSMC-like transition density $P(t'|s)$



D. Tree height vs branch length (true trees)

