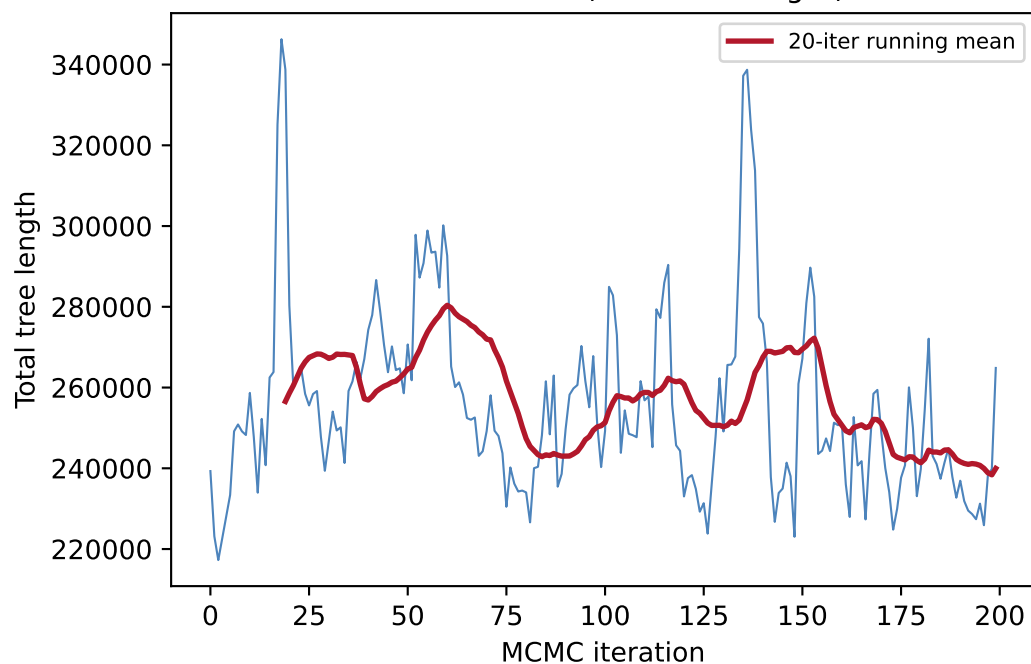
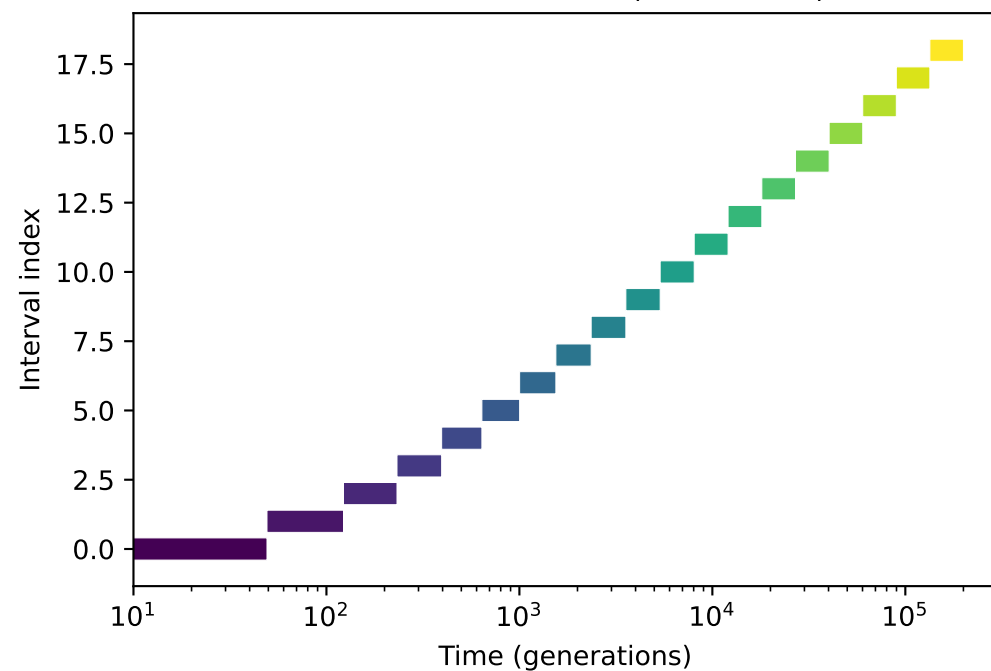


Demo: ARGweaver MCMC on Simulated Data (8 haplotypes, 50 sites)

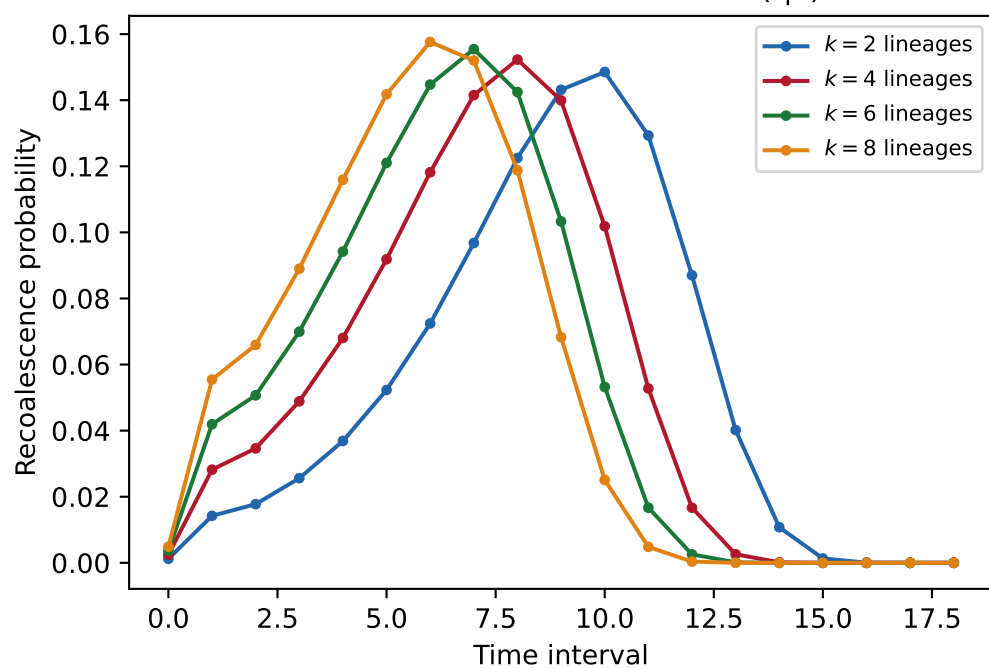
A. MCMC trace (total tree length)



B. Time discretization (19 intervals)



C. Recoalescence distribution $P(t|k)$



D. Posterior tree length distribution (post burn-in)

