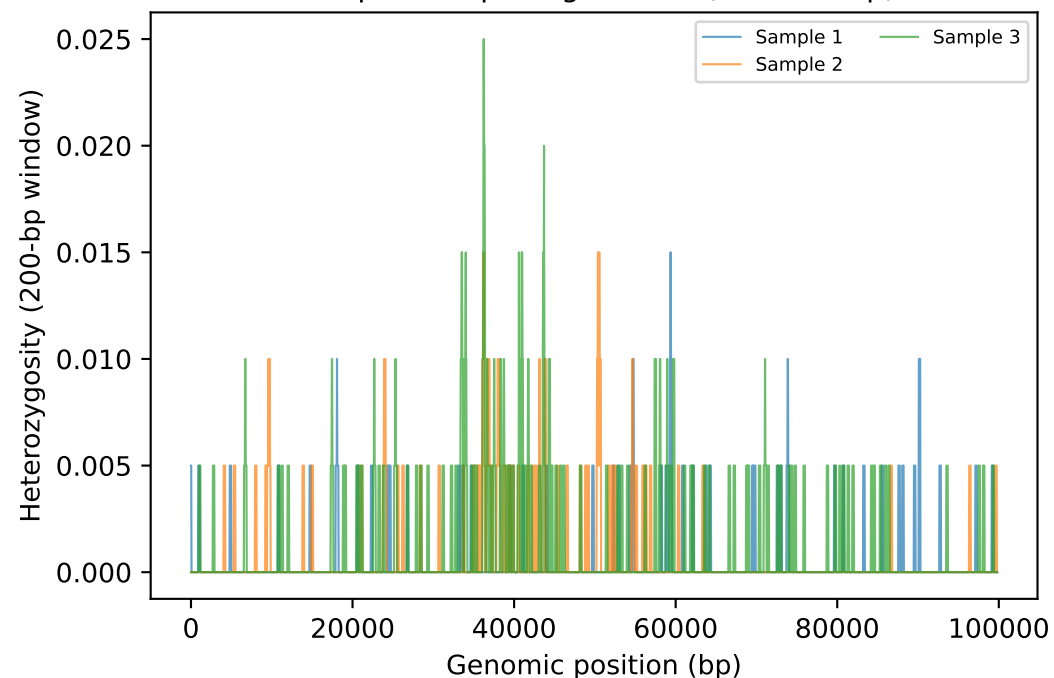
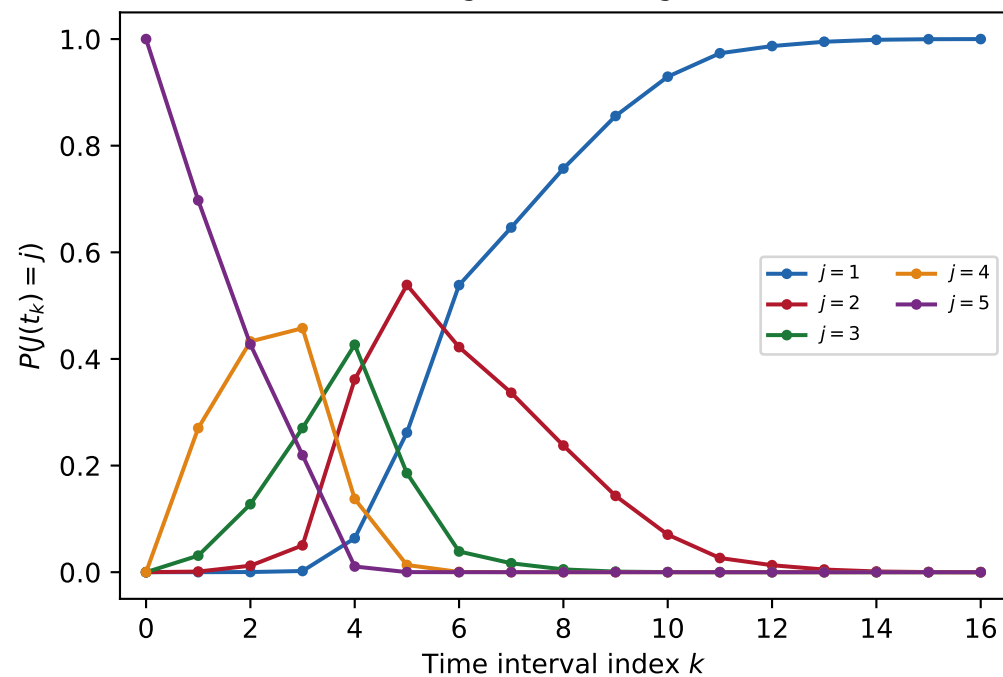


Demo: SMC++ on Multi-sample Diploid Data (Bottleneck History)

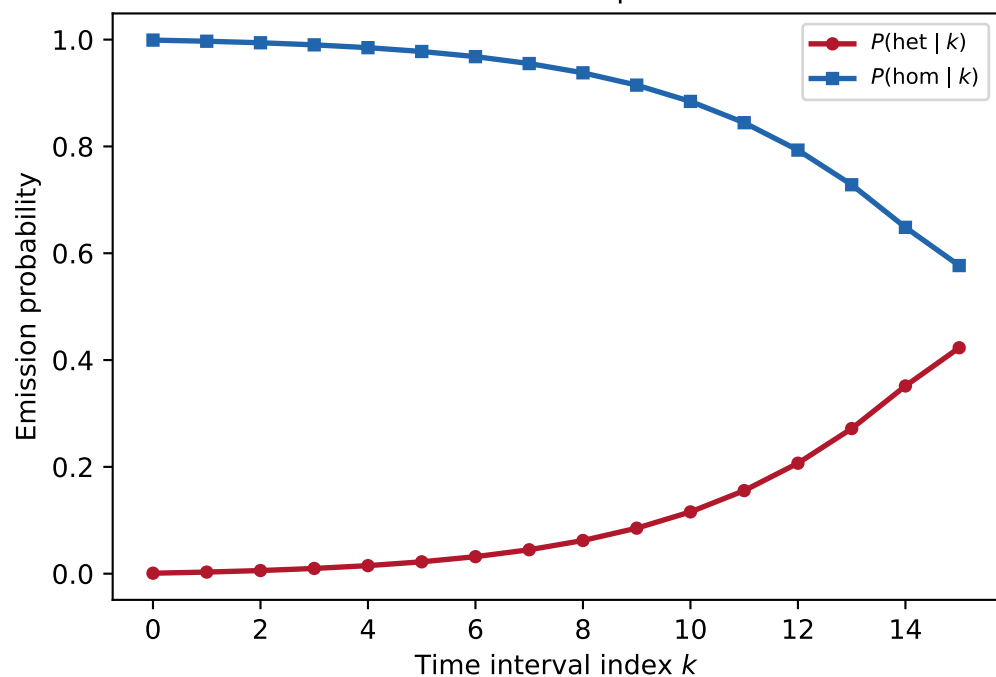
A. Input: 3 diploid genomes (100,000 bp)



C. ODE: undistinguished lineage count ($n_U = 5$)



B. SMC++ emission probabilities



D. Coalescence rate $h(t)$ under different demographies

