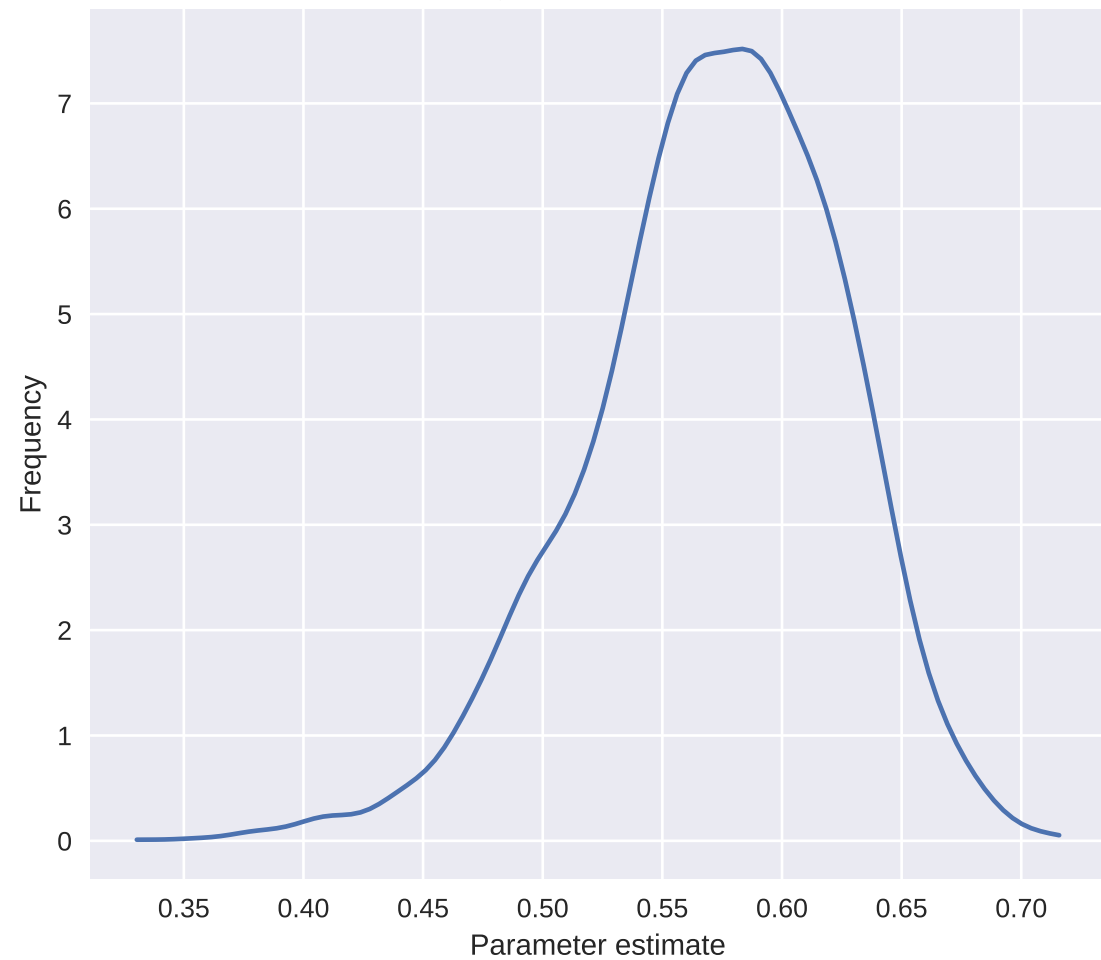
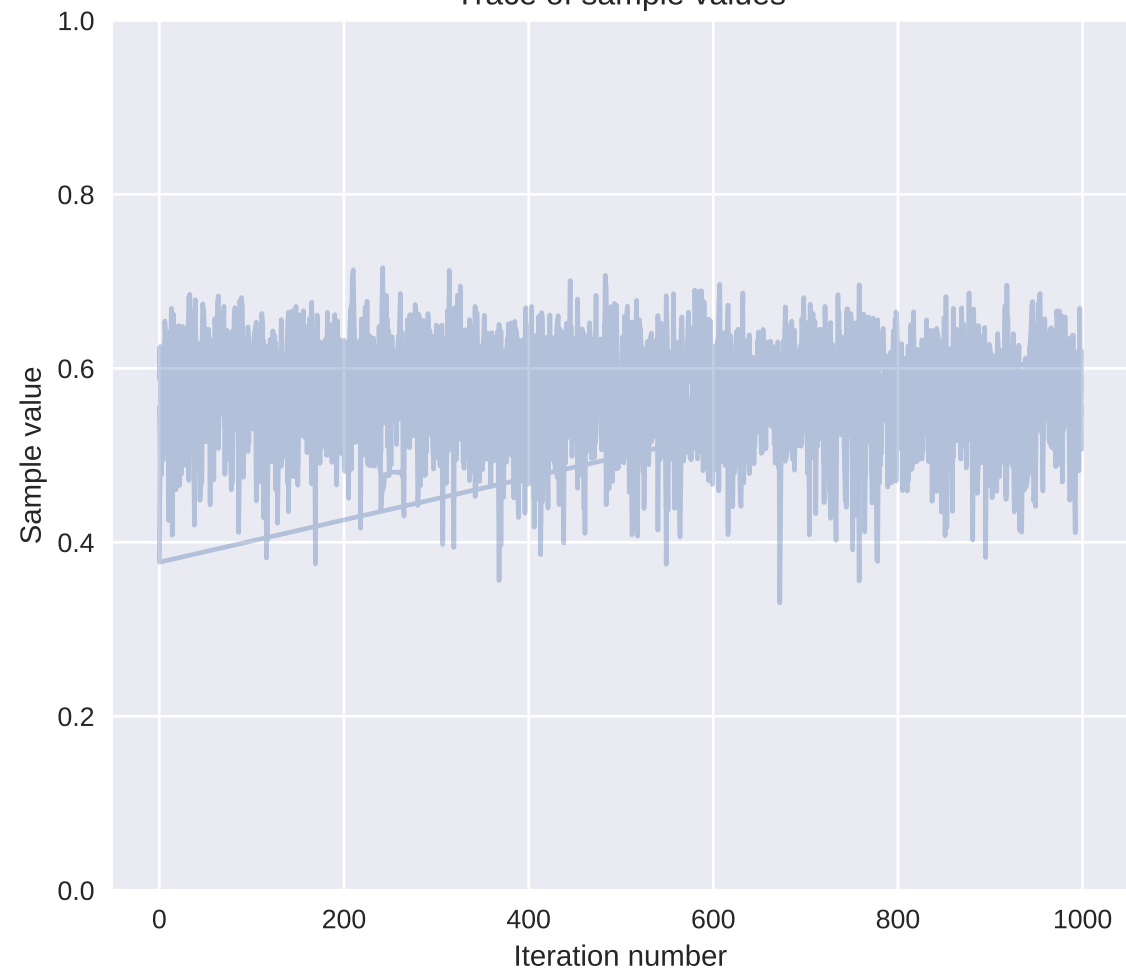


Low serial correlation in simulation trace

CD4+ Treg cell fraction distribution

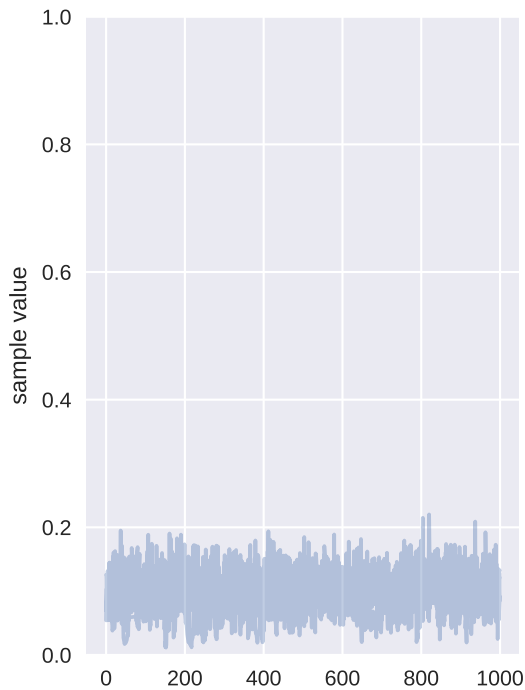
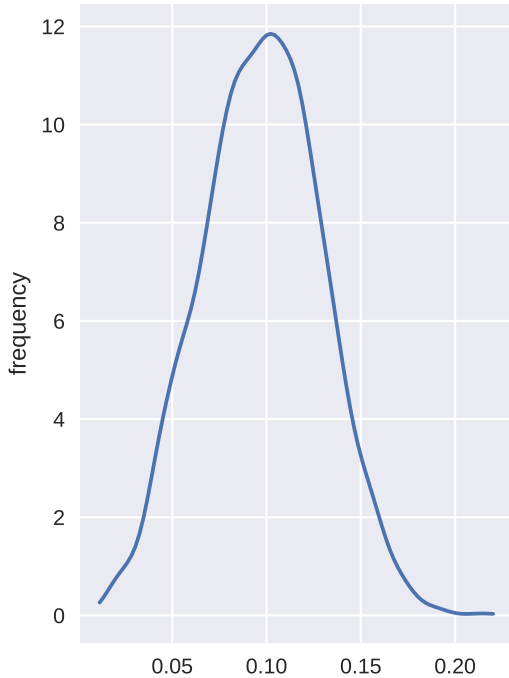


Trace of sample values

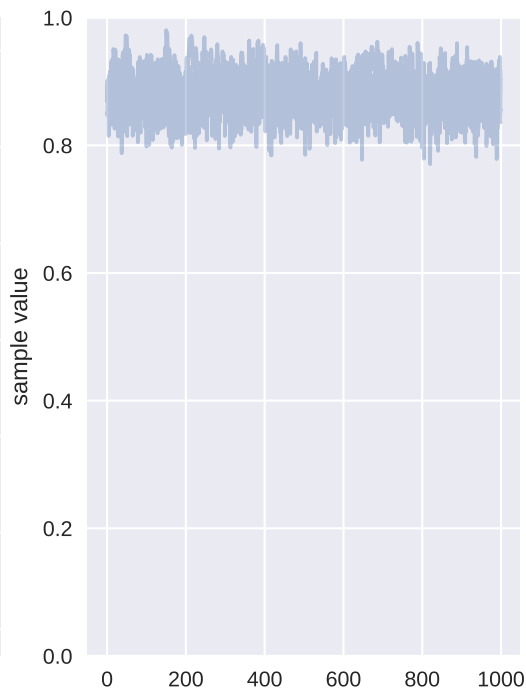
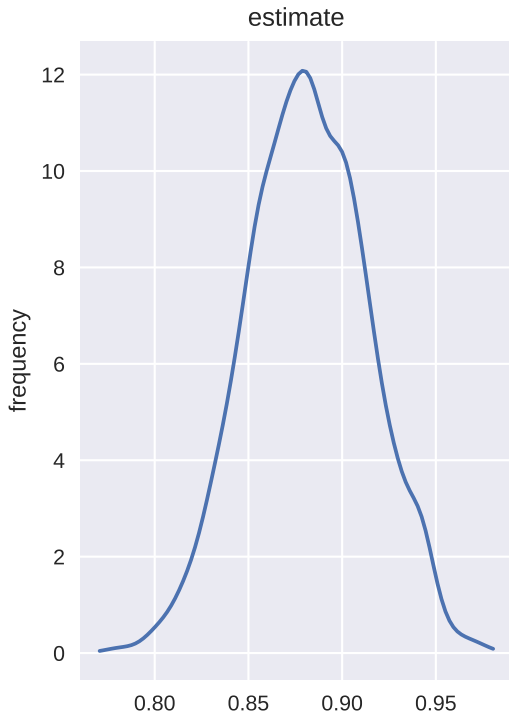


Mixture 1, Memory B cell fraction estimate

estimate



Mixture 1, Naive B cell fraction estimate



Model's estimated correlation matrix recovers known biological similarities between cell types

