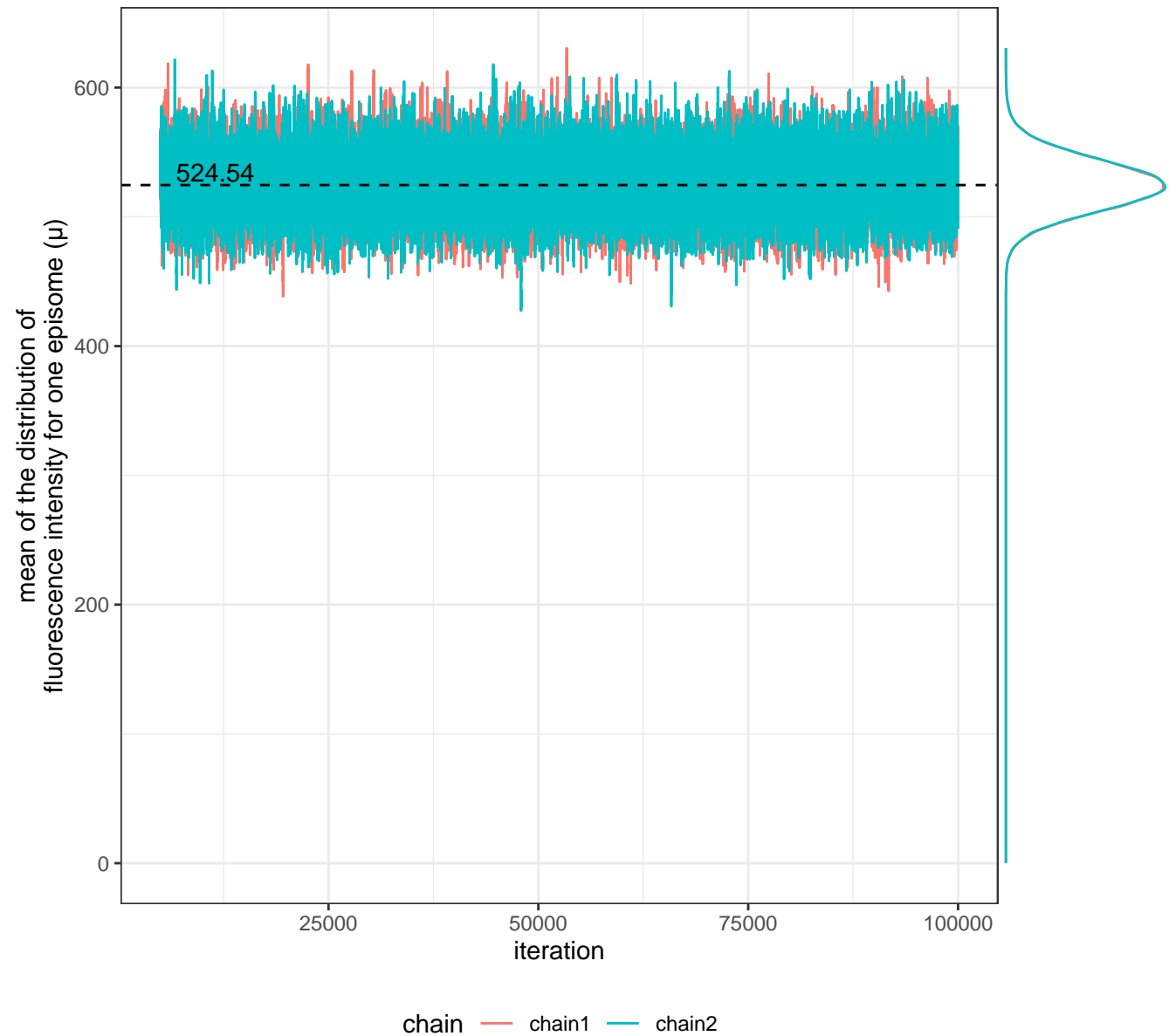
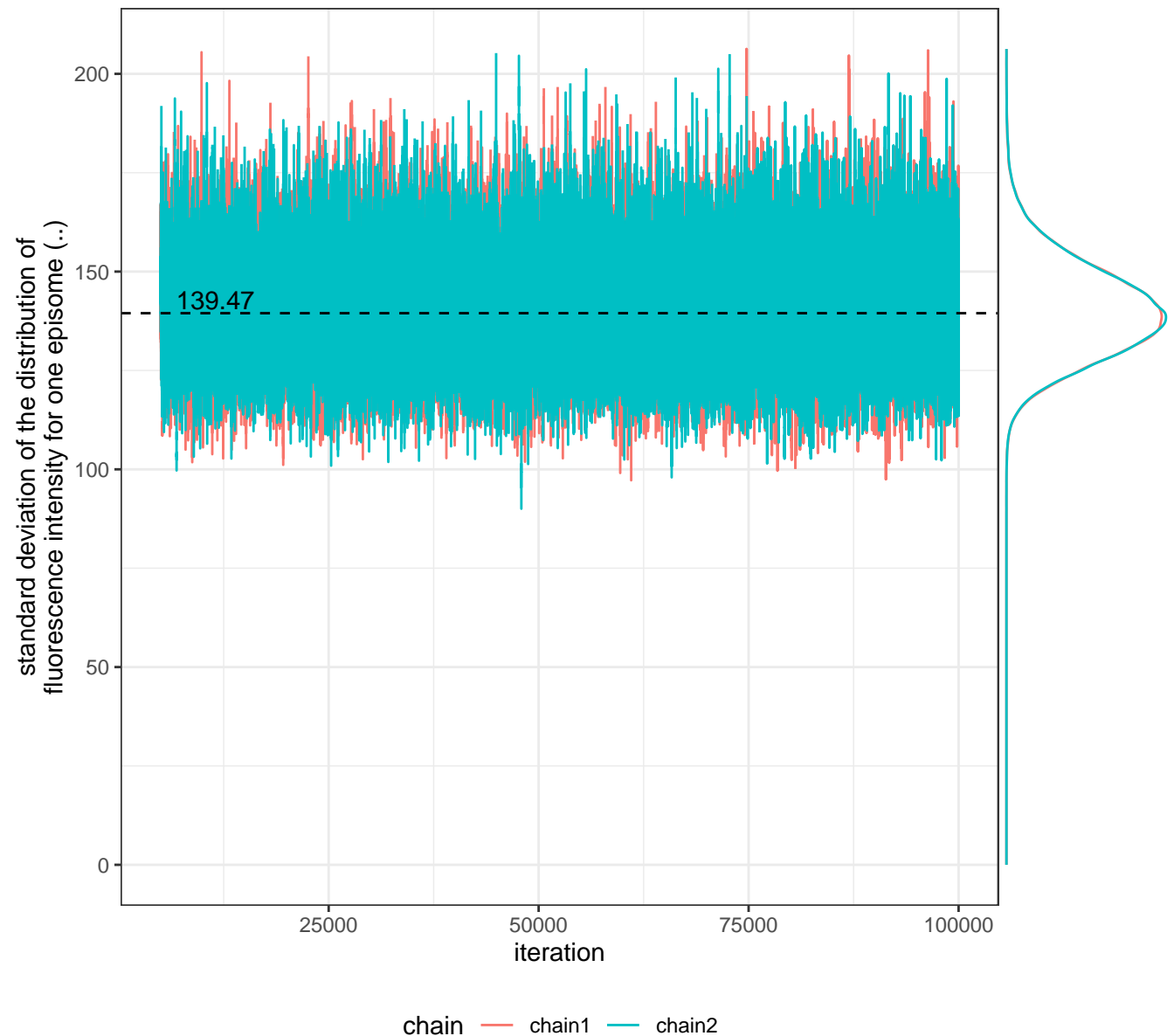
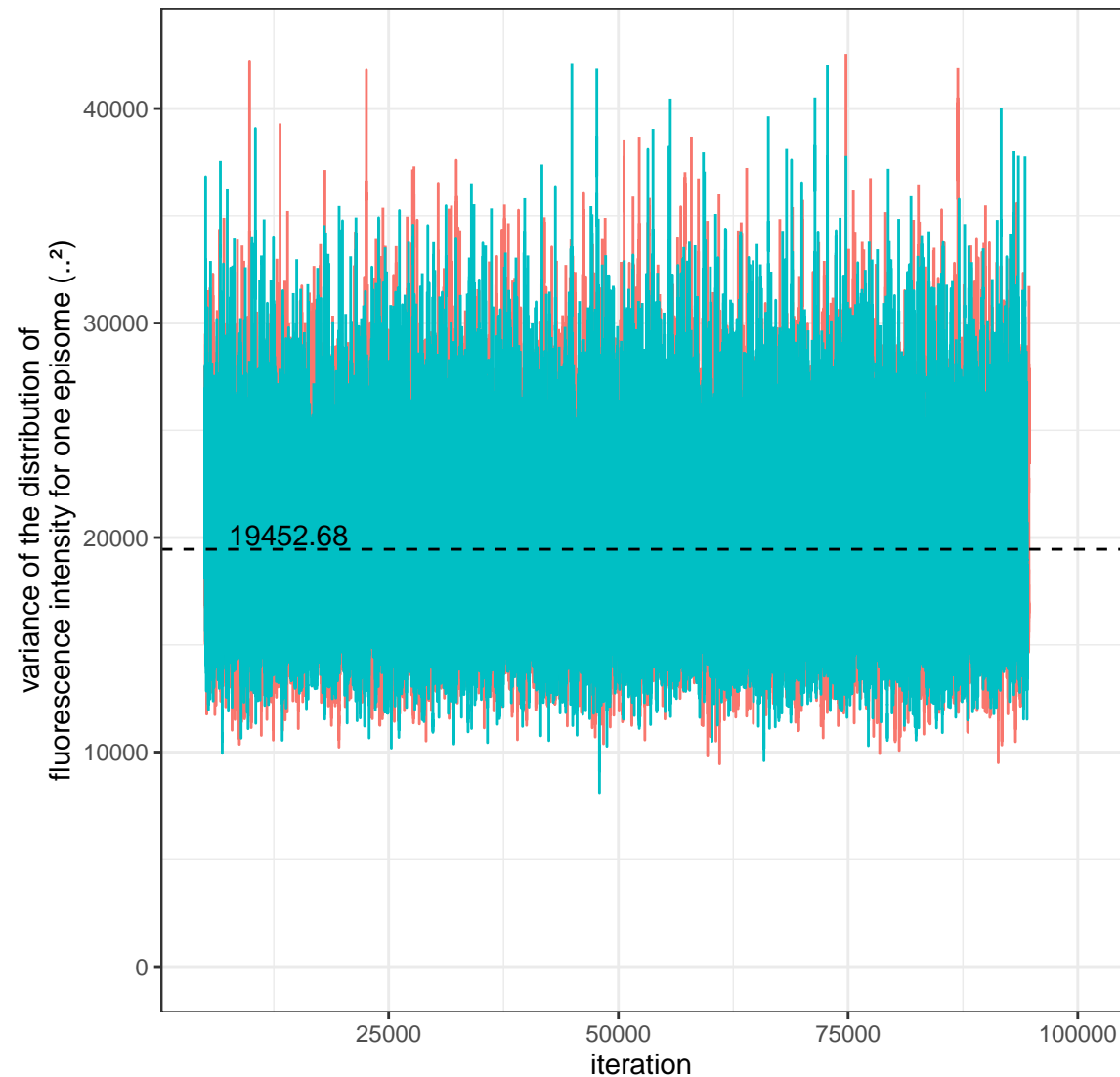


Trace of the mean (μ)

Trace of the standard deviation (..)

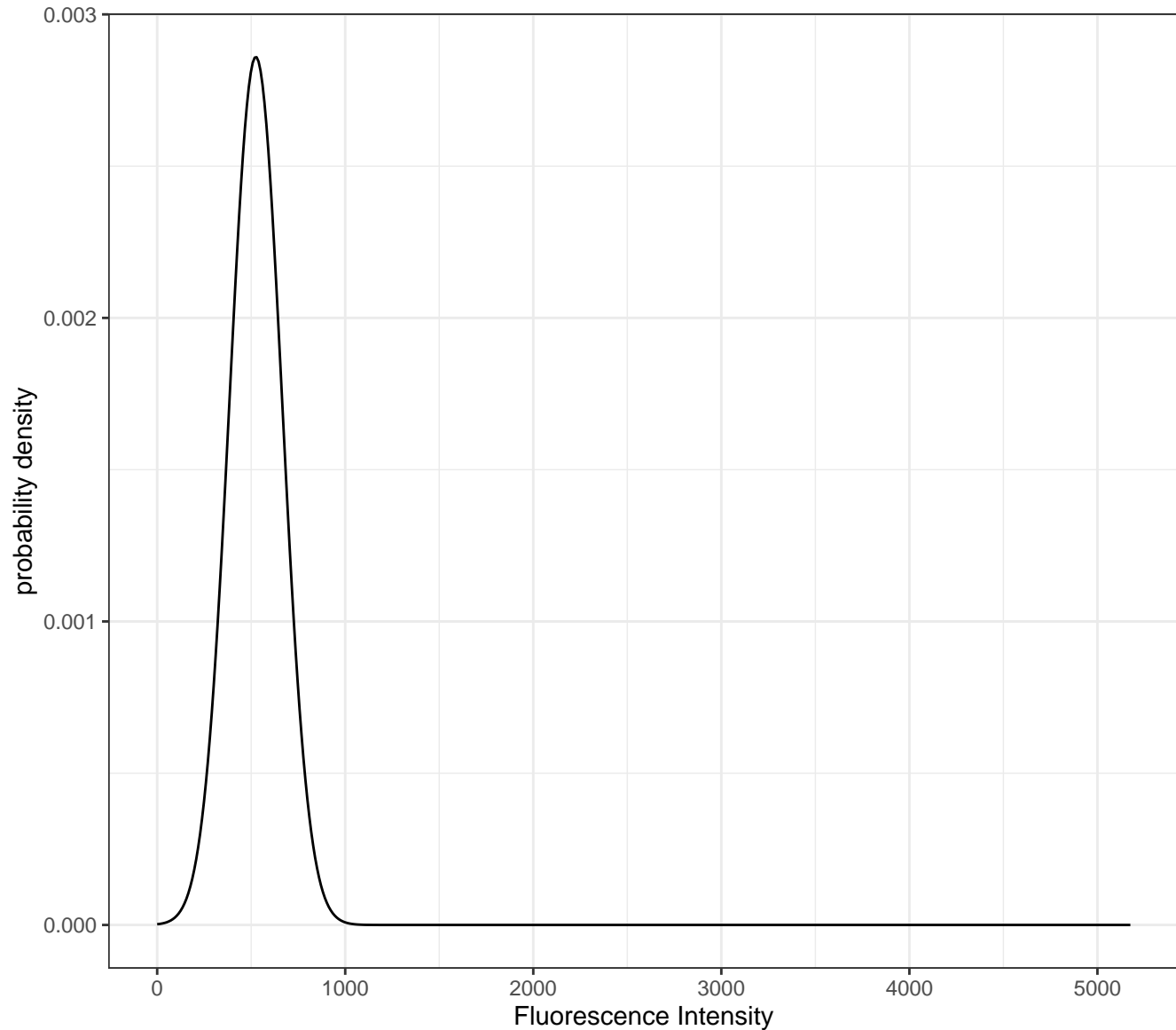


Trace of the variance (..²)

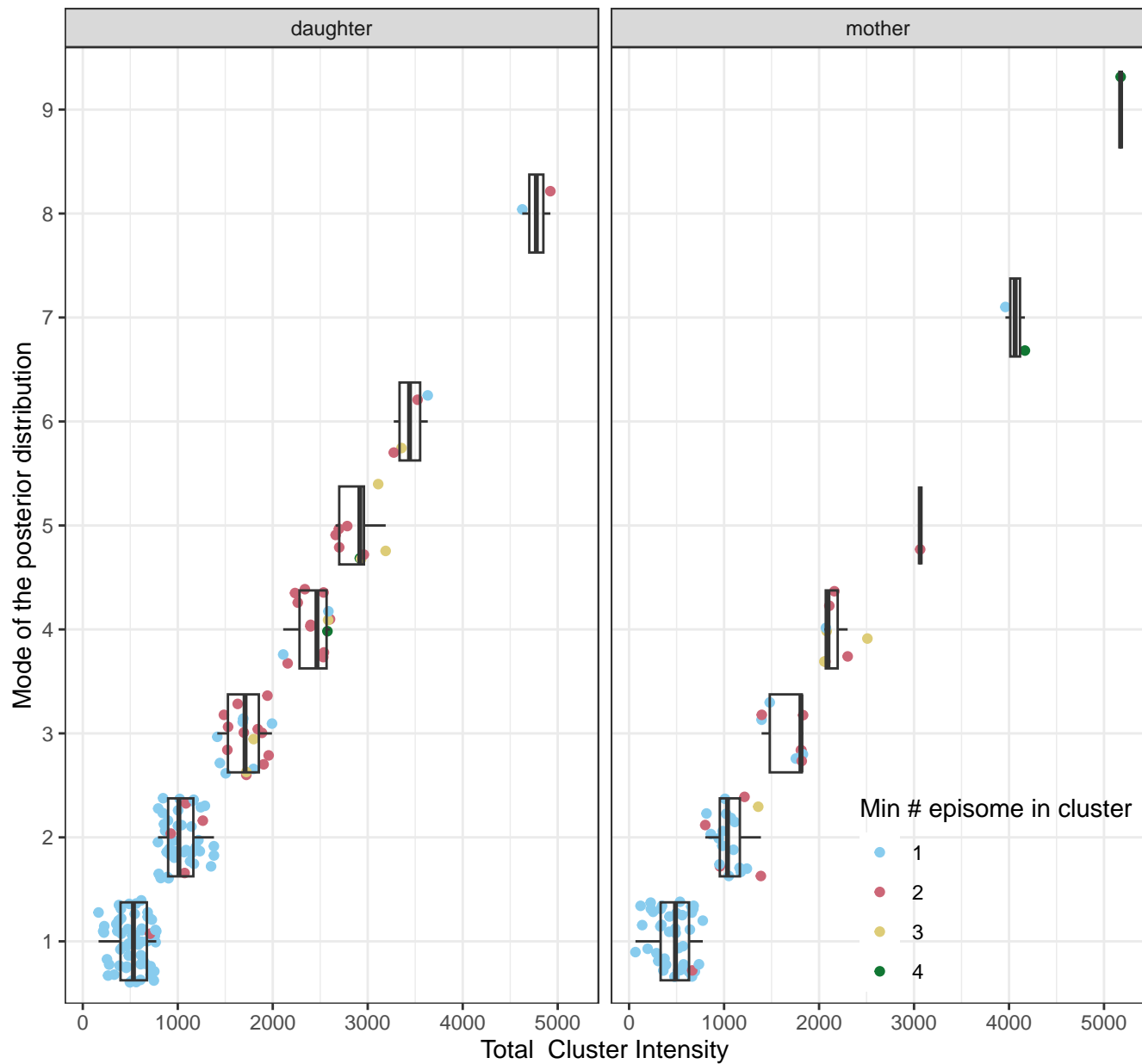
chain chain1 chain2

Inferred distribution of fluorescence intensity of a single cluster

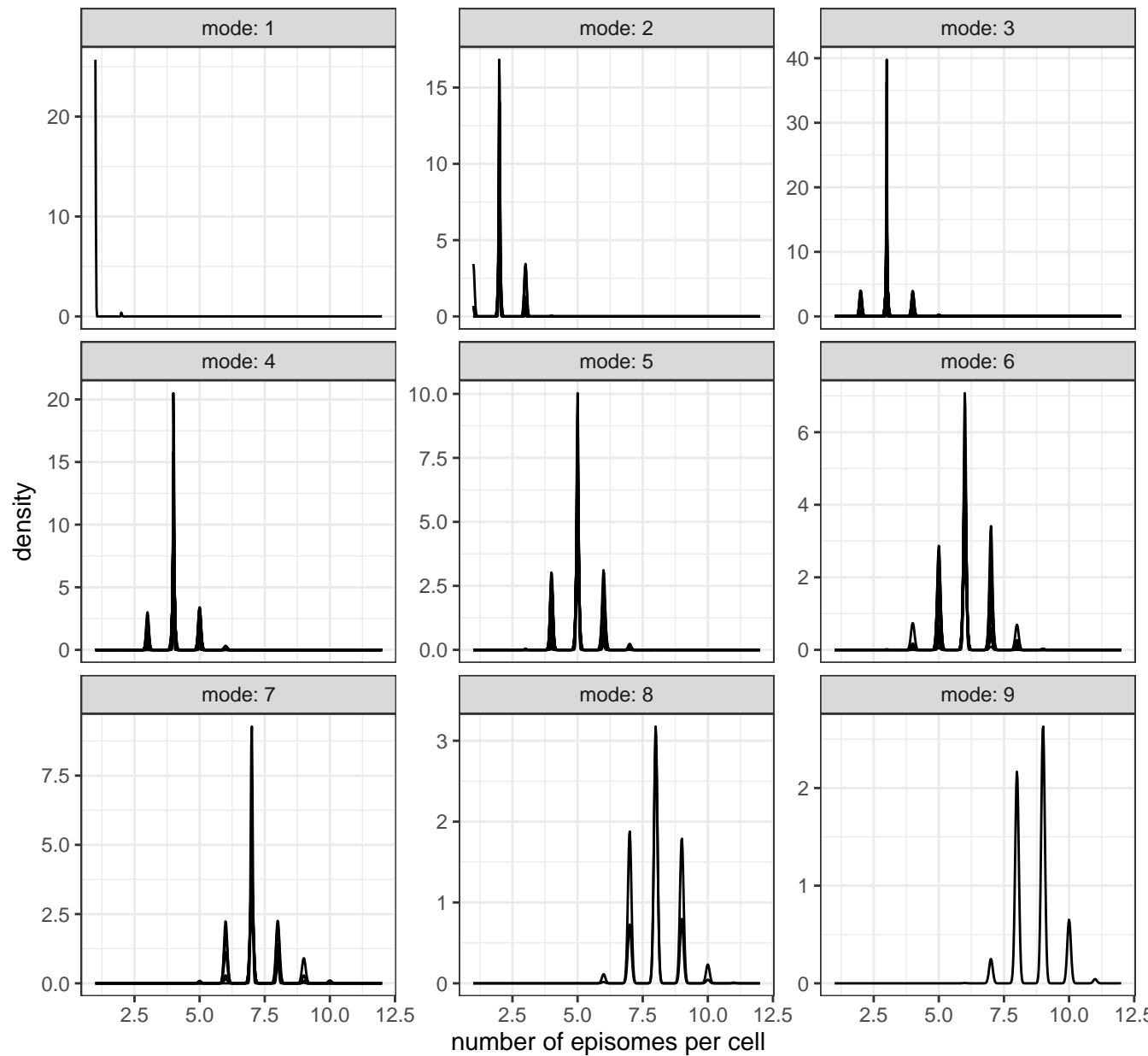
mean = 524.54, standard deviation = 139.47



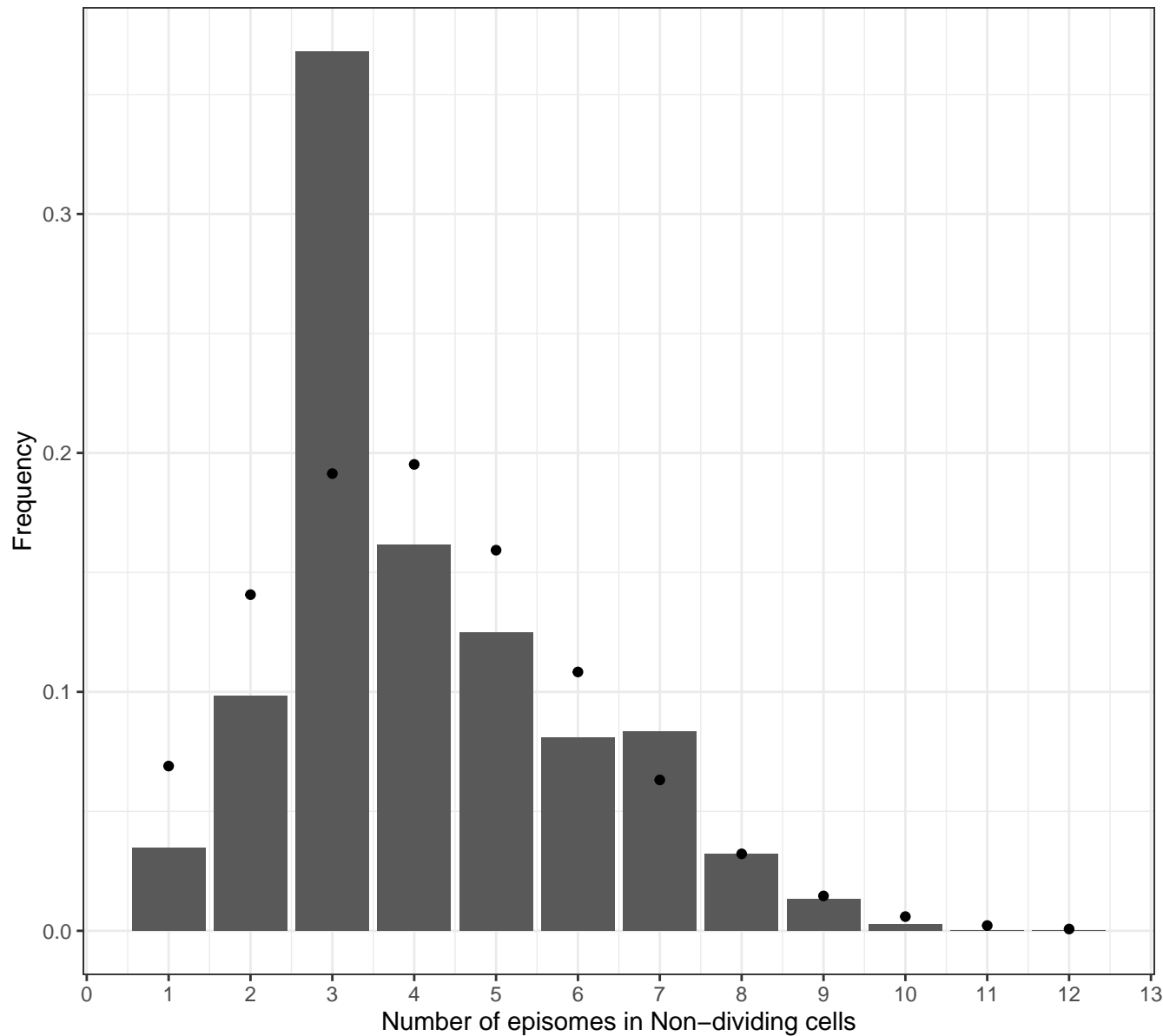
Distribution of cluster intensities

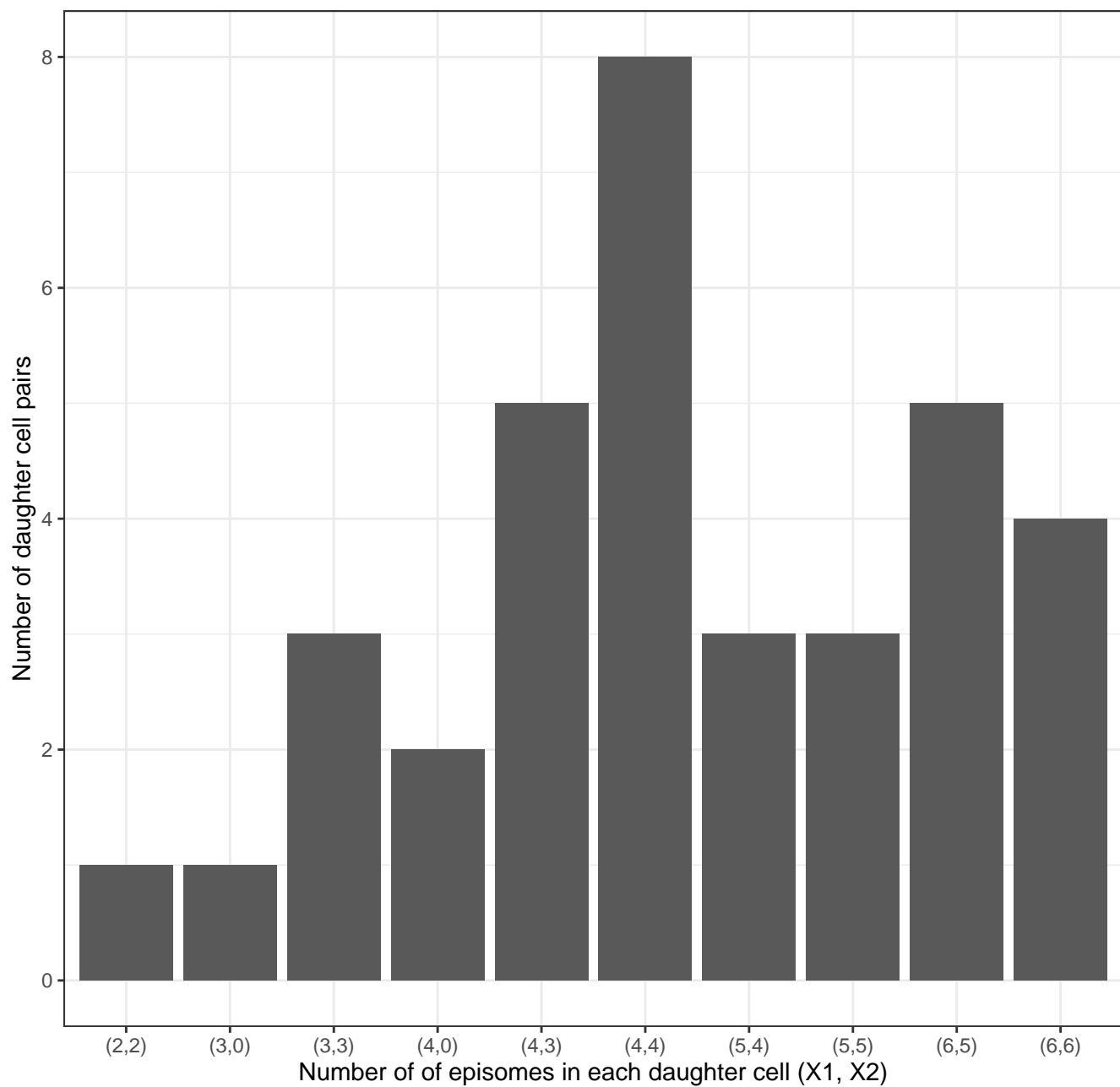


Posterior probabilities for number of episomes per cell

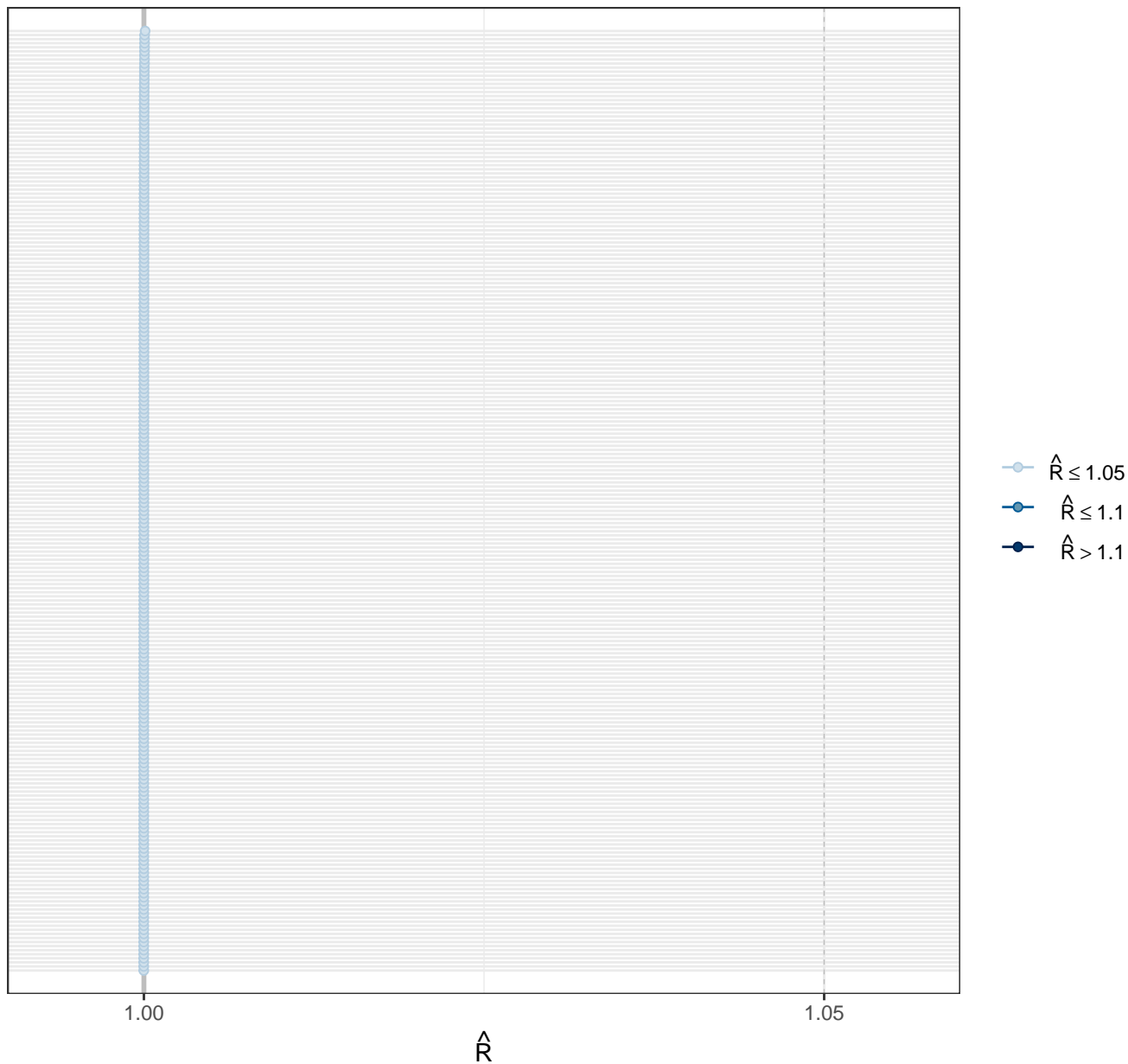


Sampled distribution of initial number of episomes

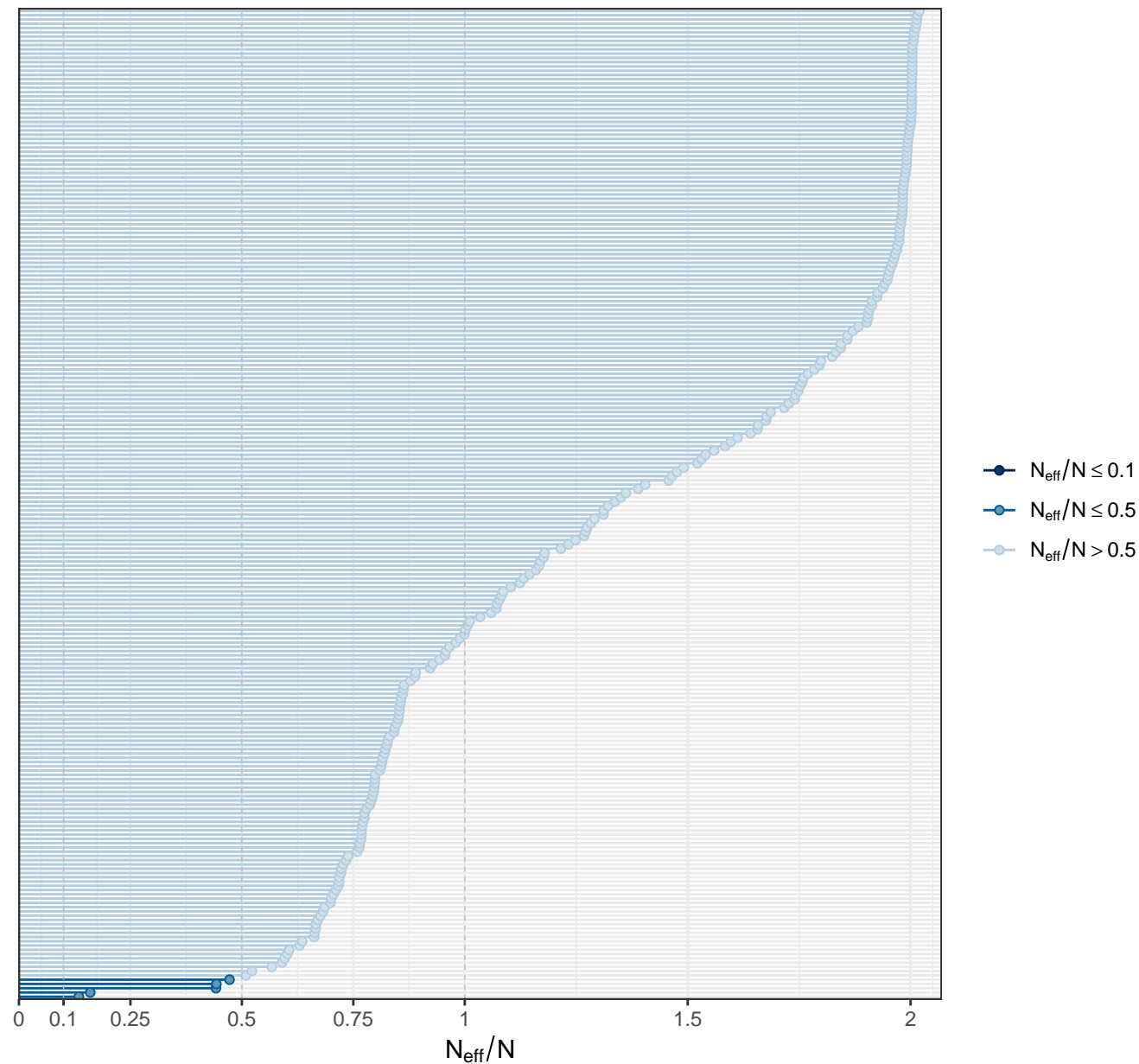
 $X_0 \sim \text{poisson}(\text{lambda} = 4.08)$, median = 3



Rhat



Bulk Effective Sample Size Ratio



Tail Effective Sample Size Ratio

