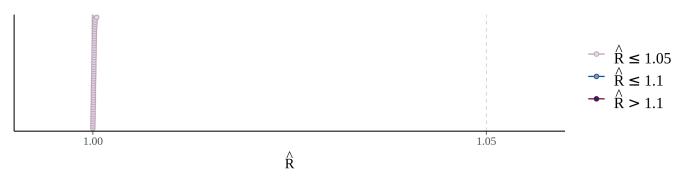
## Convergence report for Lung SCC signatures 20000ROO RData

Date of inference: 2020-06-13

Explanation of parameters Rhat indicates the ratio of variance within a change over variance pooling all chains. We recommend running at least four chains by default and only using the sample if R-hat is less than 1.05. ESS bulk/tail indicate the bulk/tail effective sample size estimate, i.e. show the sampling efficiency of mean and median estimates. We recommend running at least four chains by default and only using the sample if R-hat is less than 1.05. Both bulk-ESS and tail-ESS should be at least 100 (approximately) per Markov Chain in order to be reliable and indicate that estimates of respective posterior quantiles are reliable.

Max Rhat: 1.00048307270755ess bulk: 943.271014615656

 $\bullet$  ess tail: 1022.31084285855



In the plots below ESS should have high values

