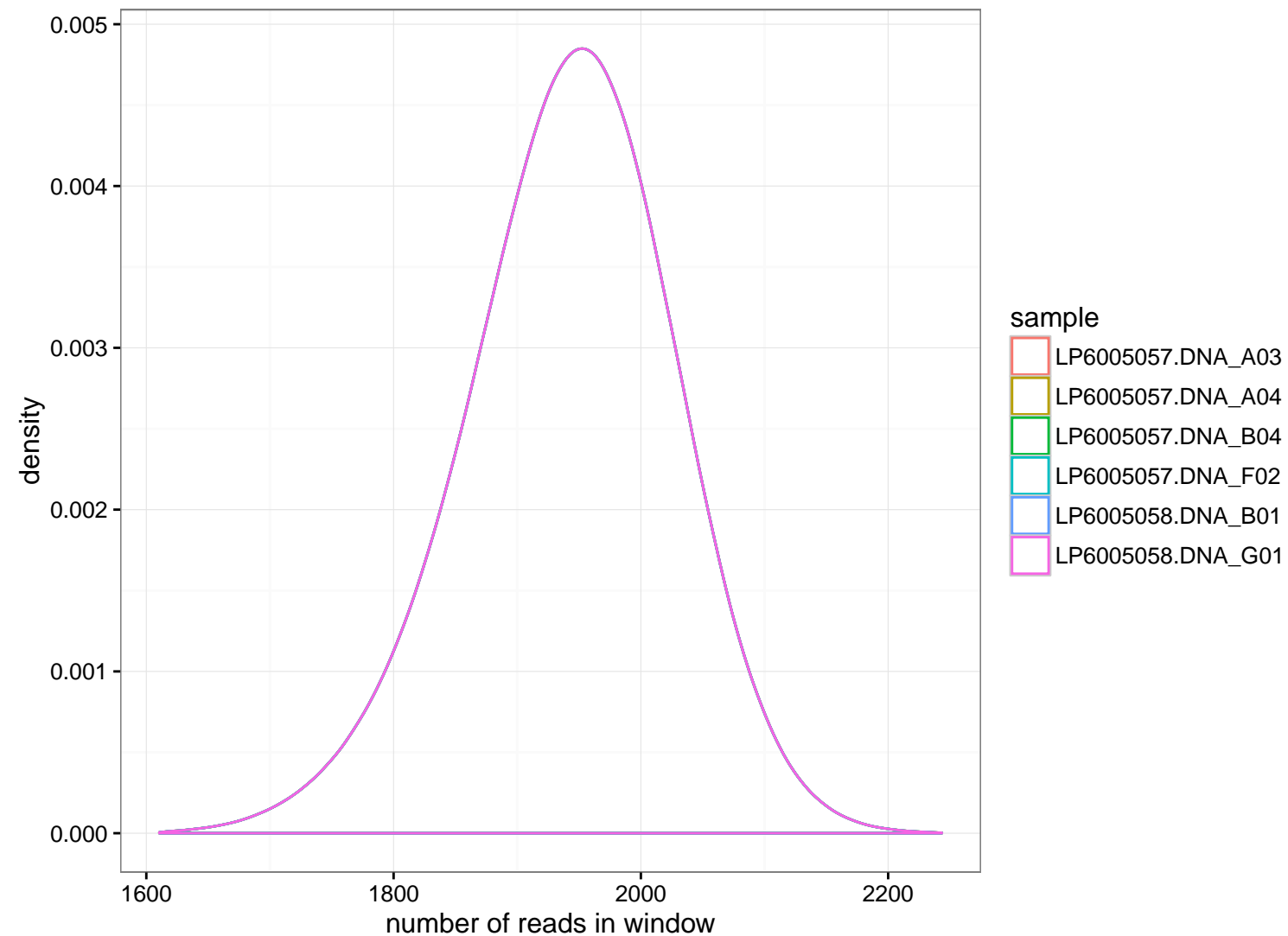
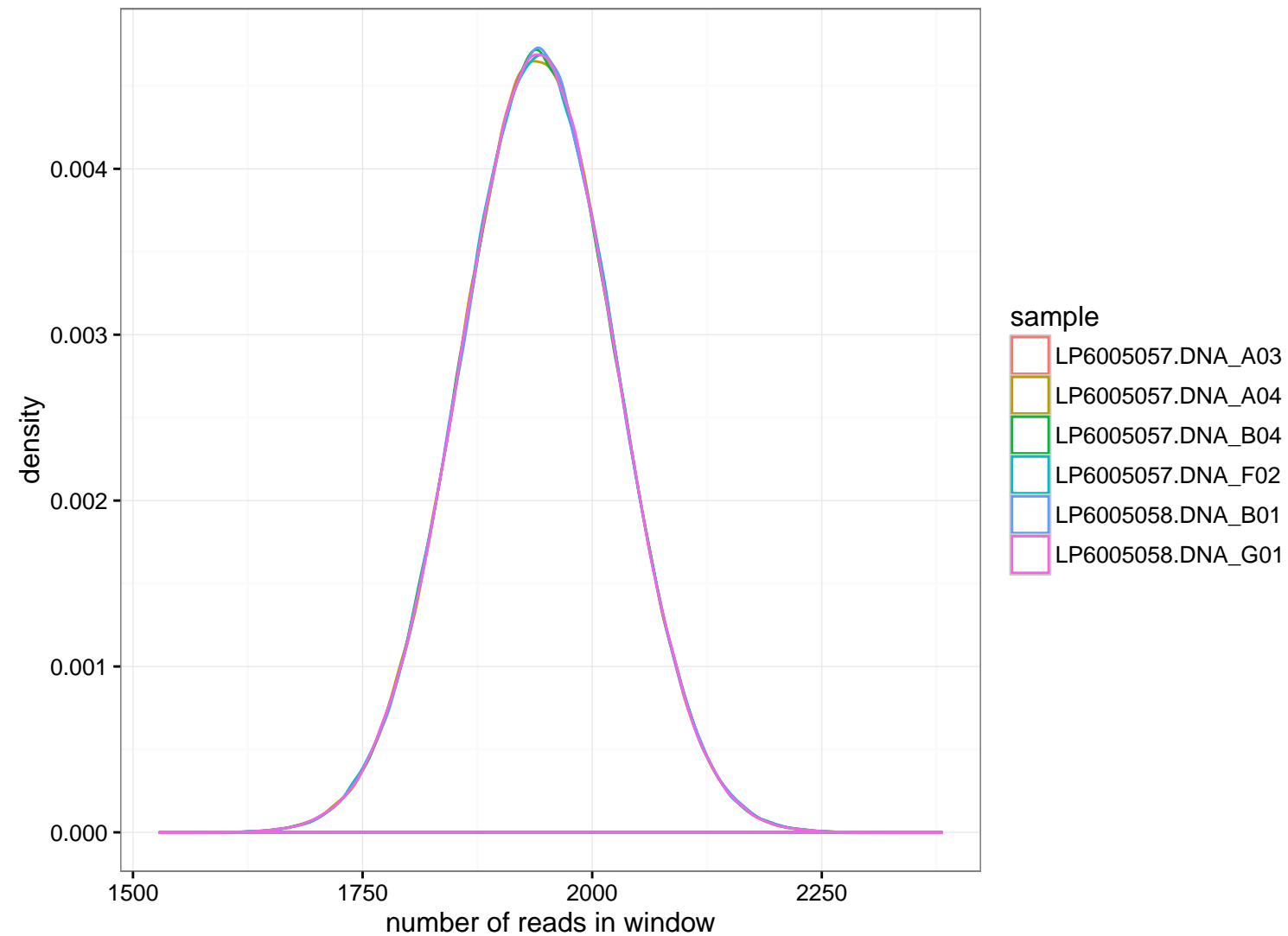


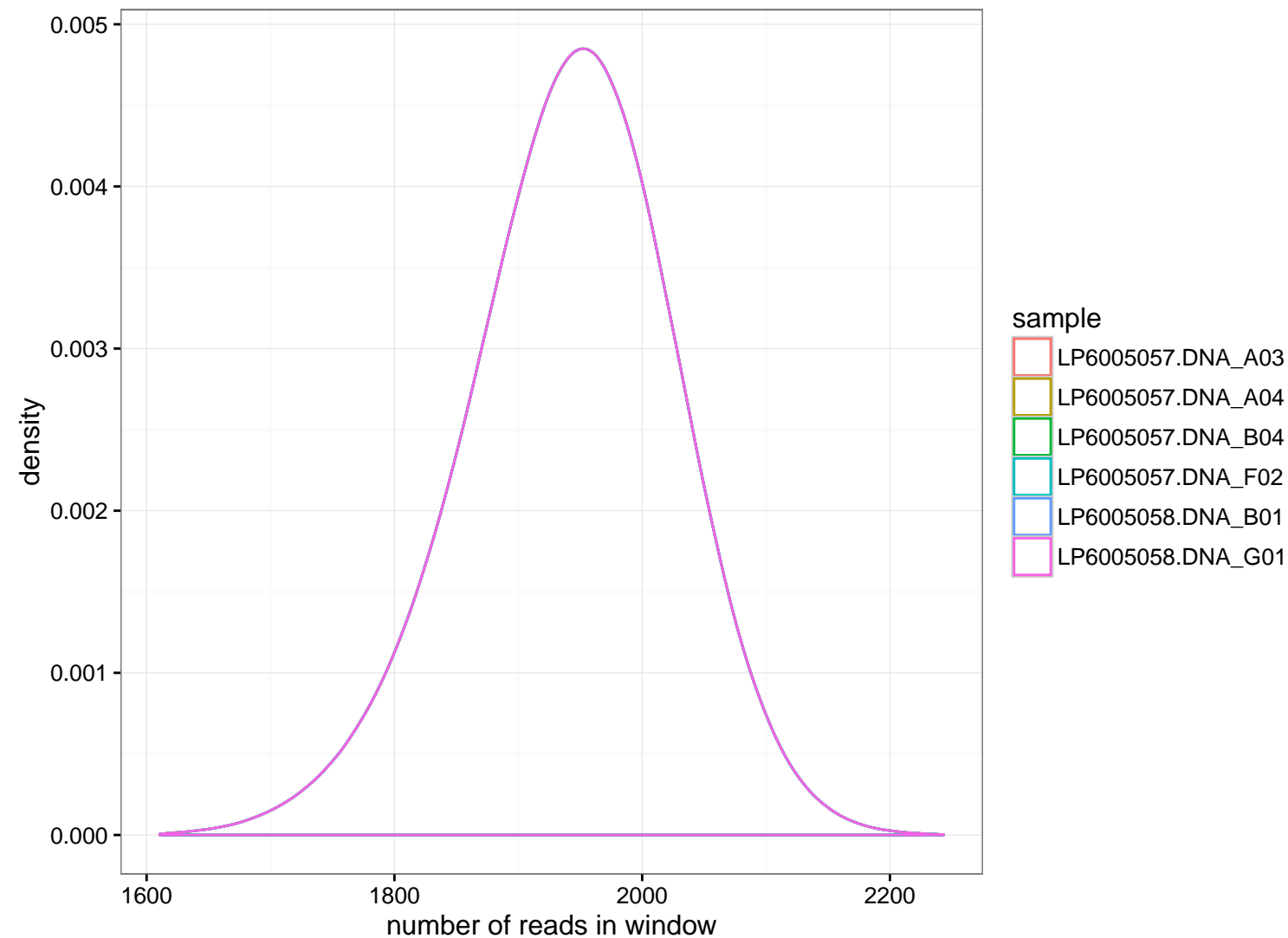
Normalized bin counts

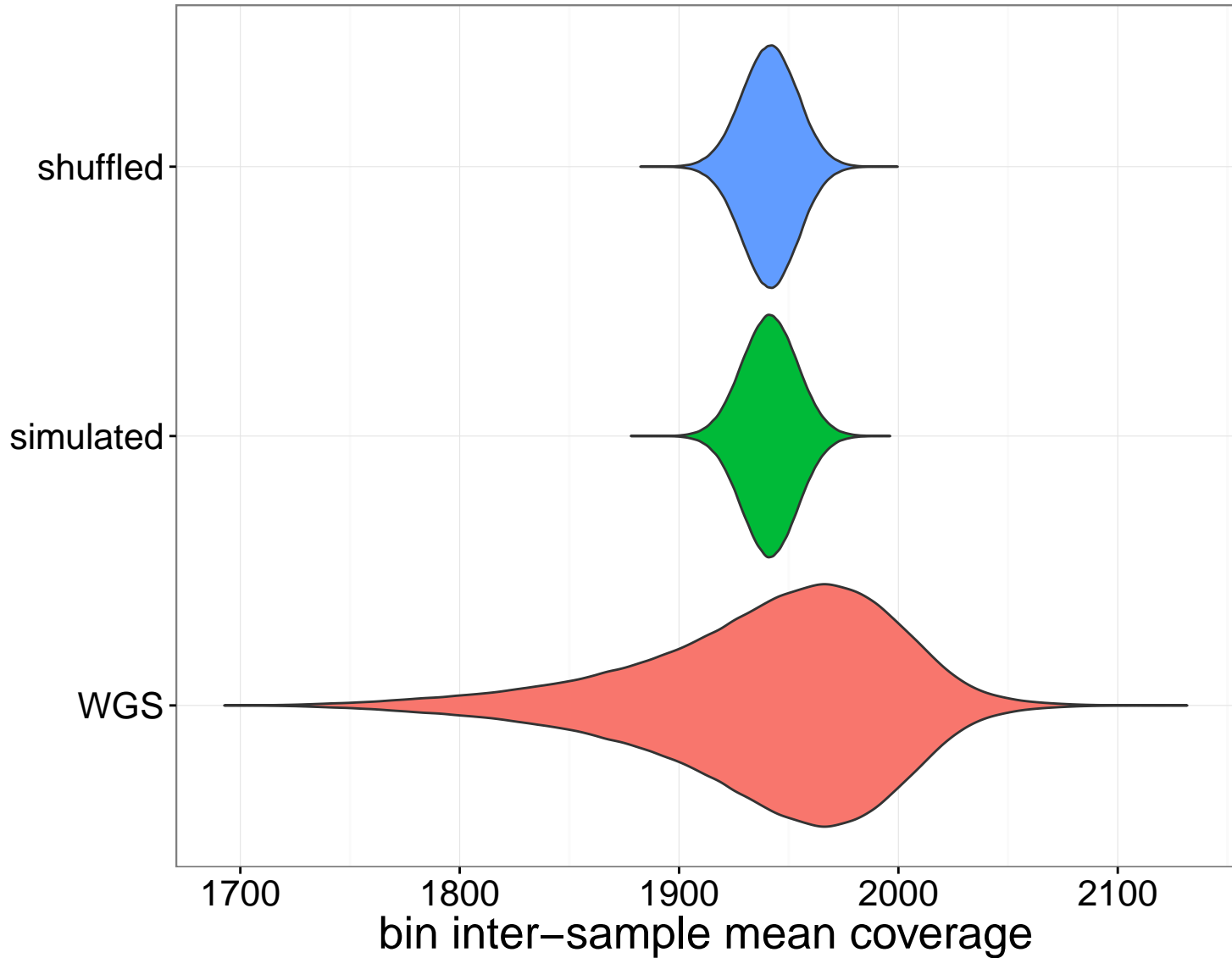


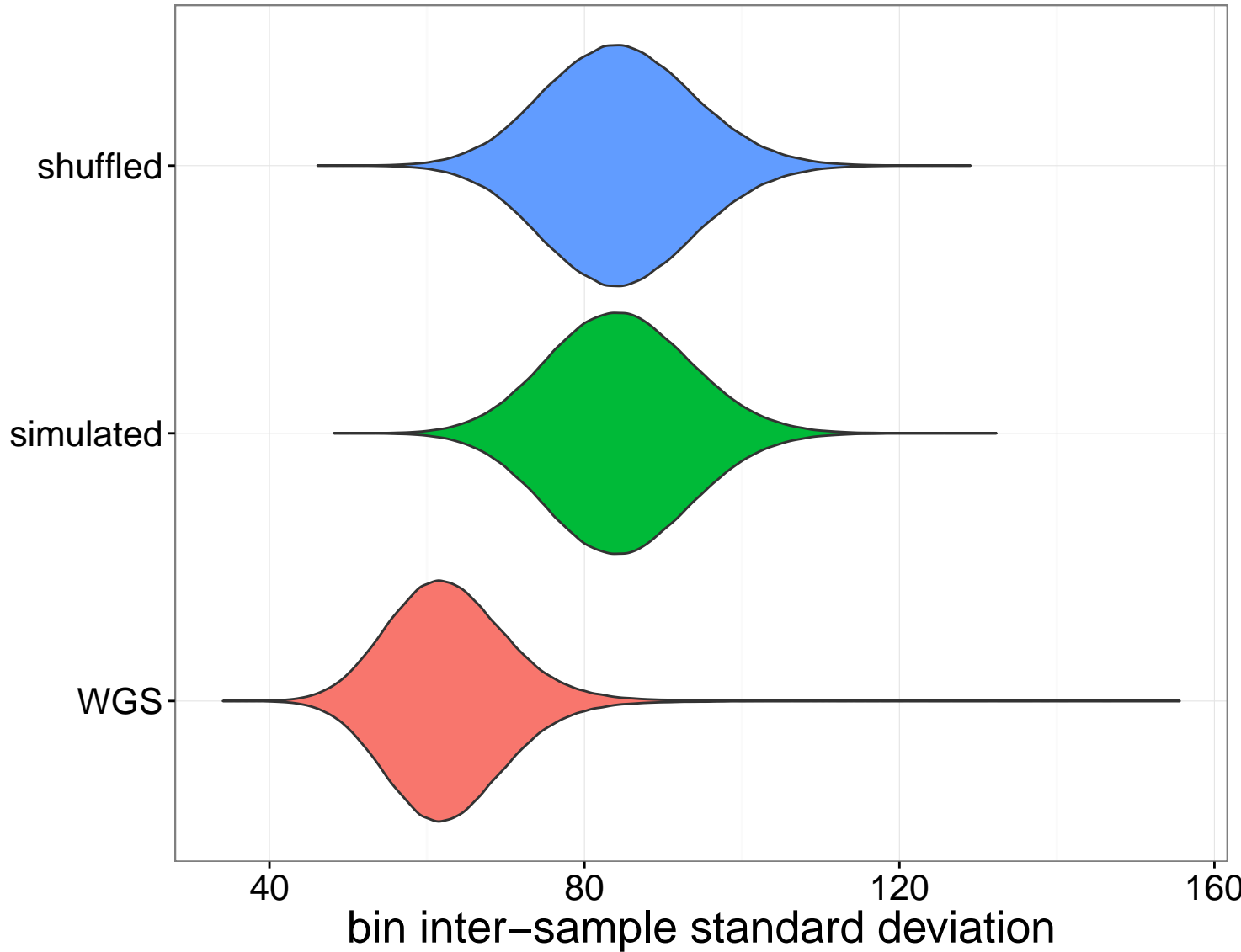
Control : simulated bin counts



Control : shuffled bin counts







proportion of the studied genome

coverage

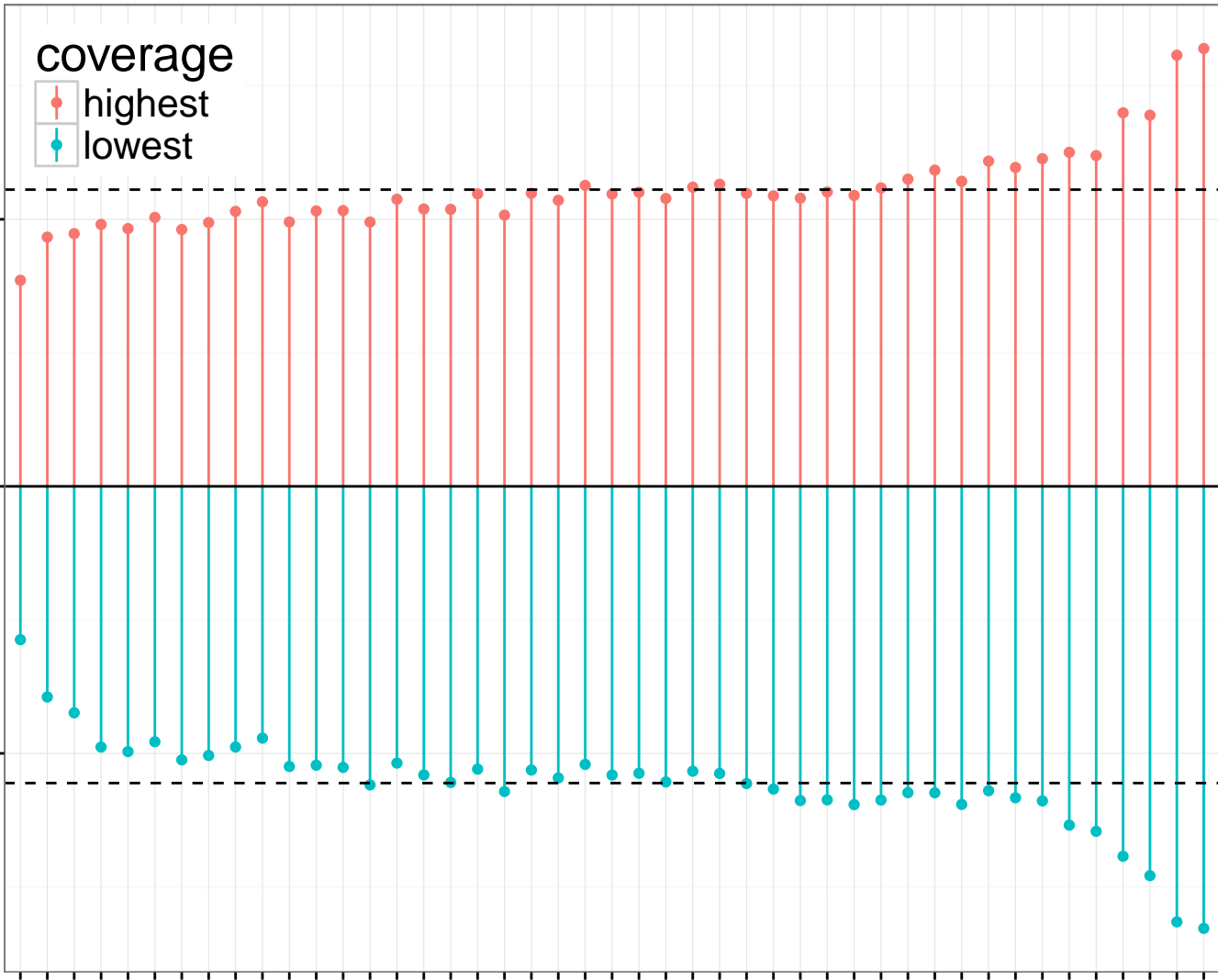
highest  
lowest

0.02

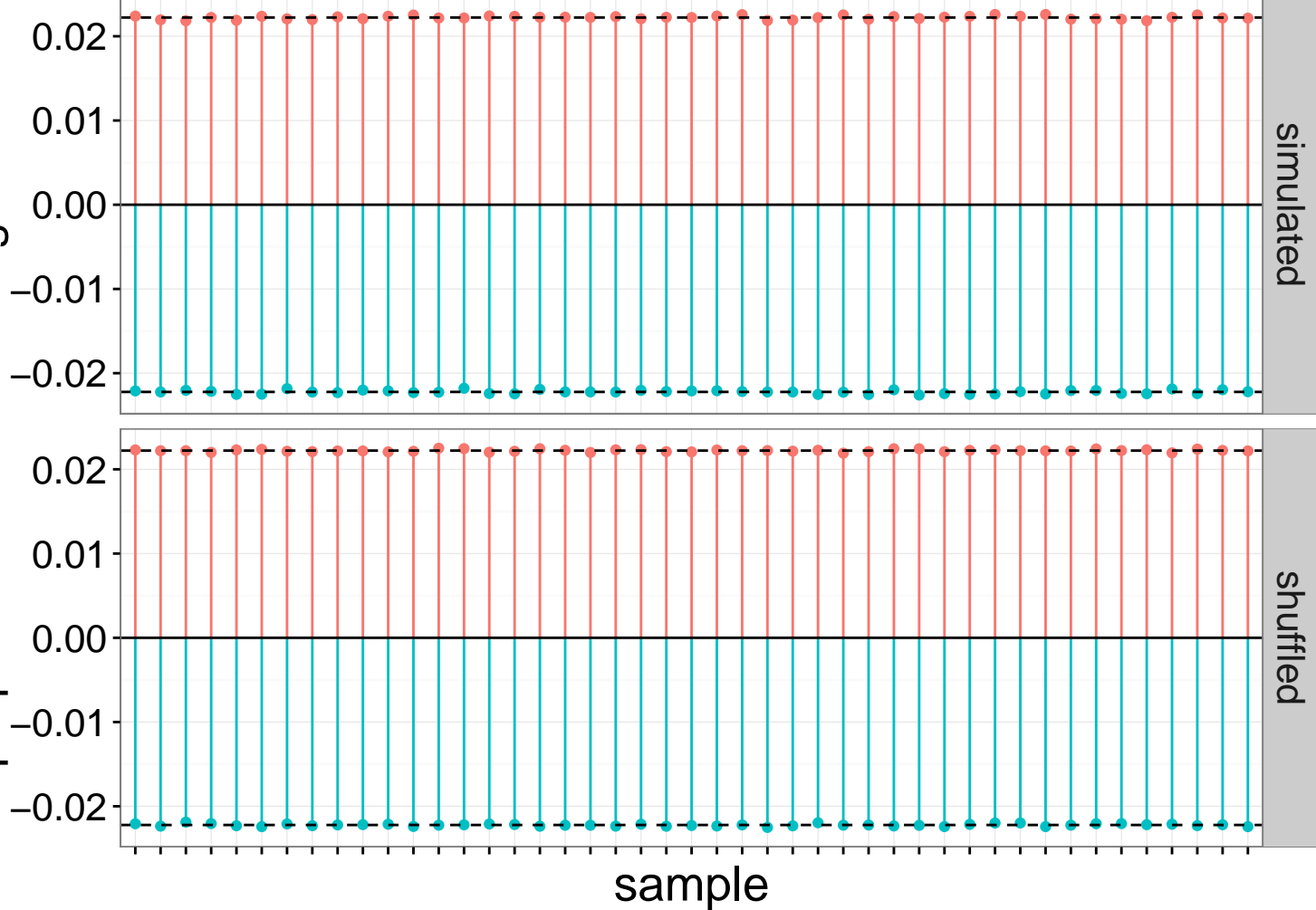
0.00

-0.02

sample



proportion of the studied genome



proportion of the studied genome

coverage

highest  
lowest

0.02

0.00

-0.02

sample

