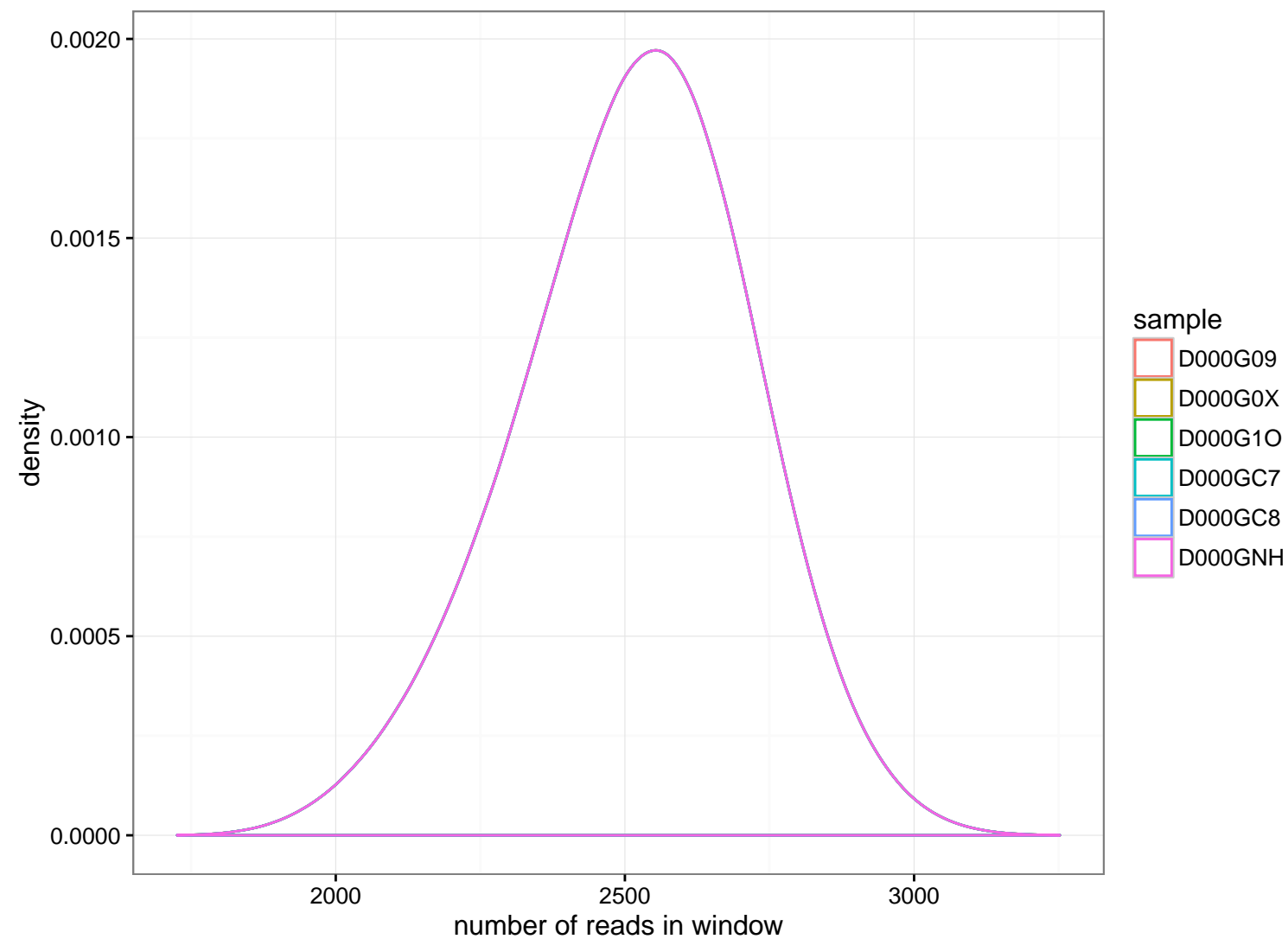
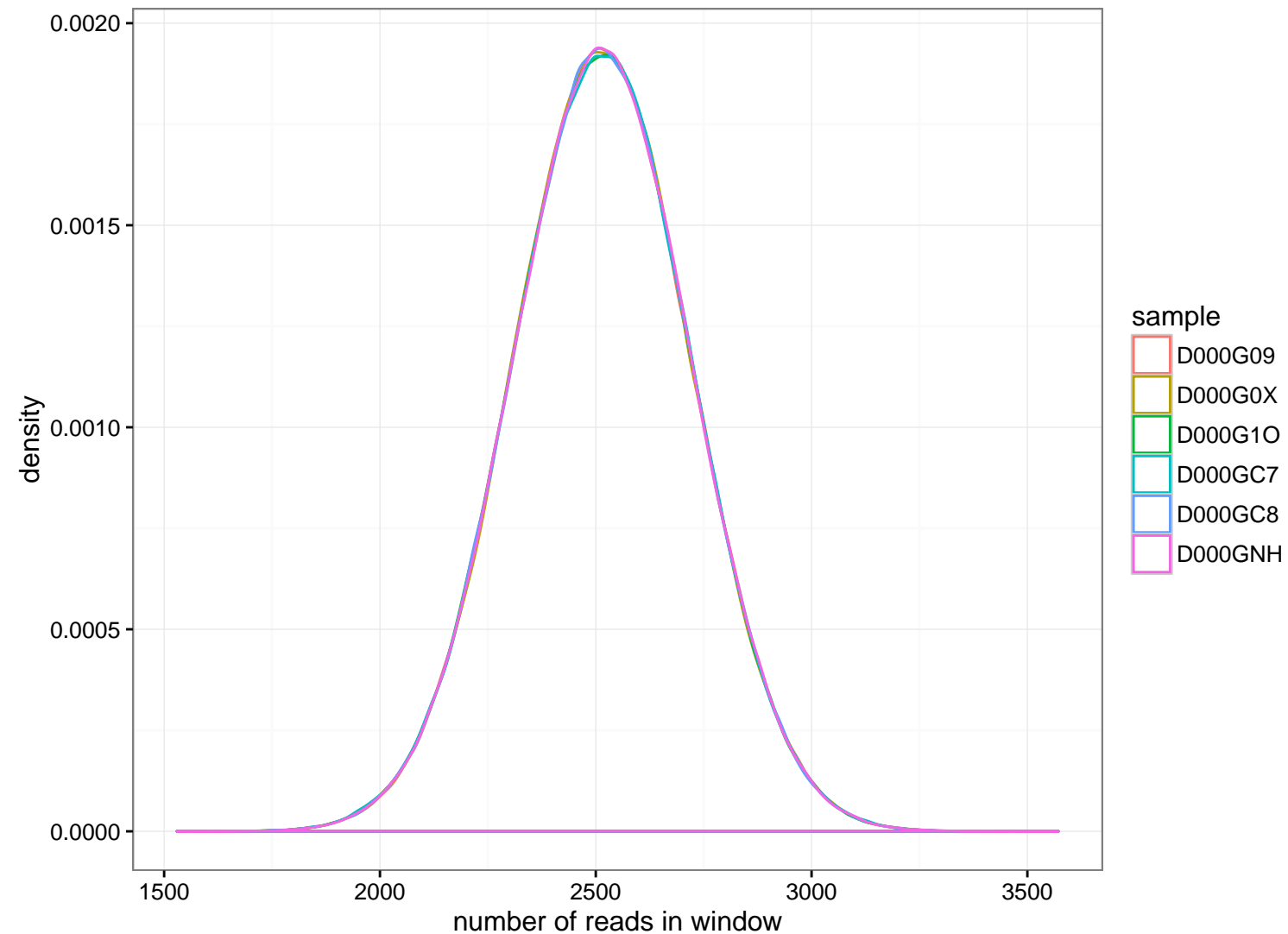


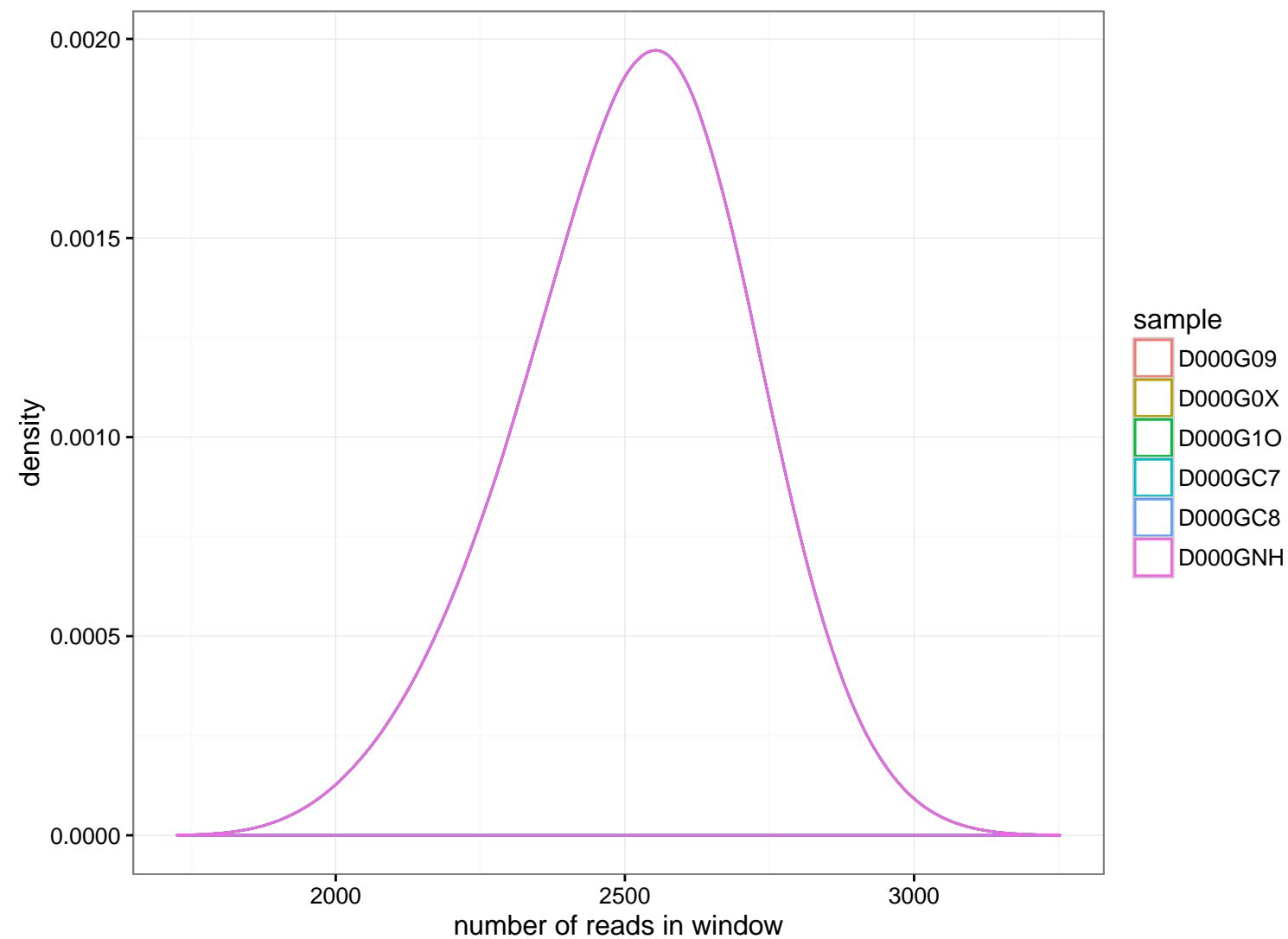
Normalized bin counts

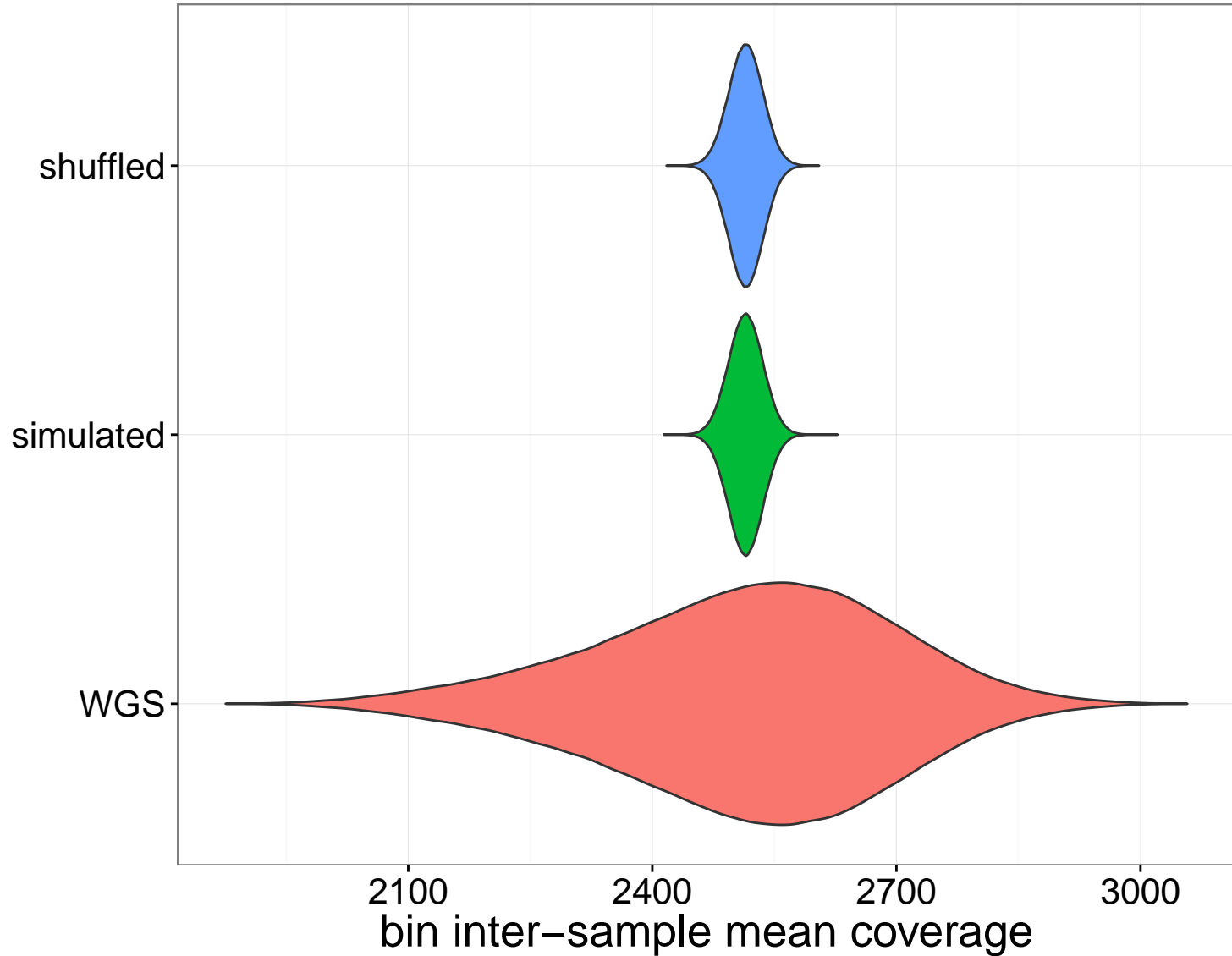


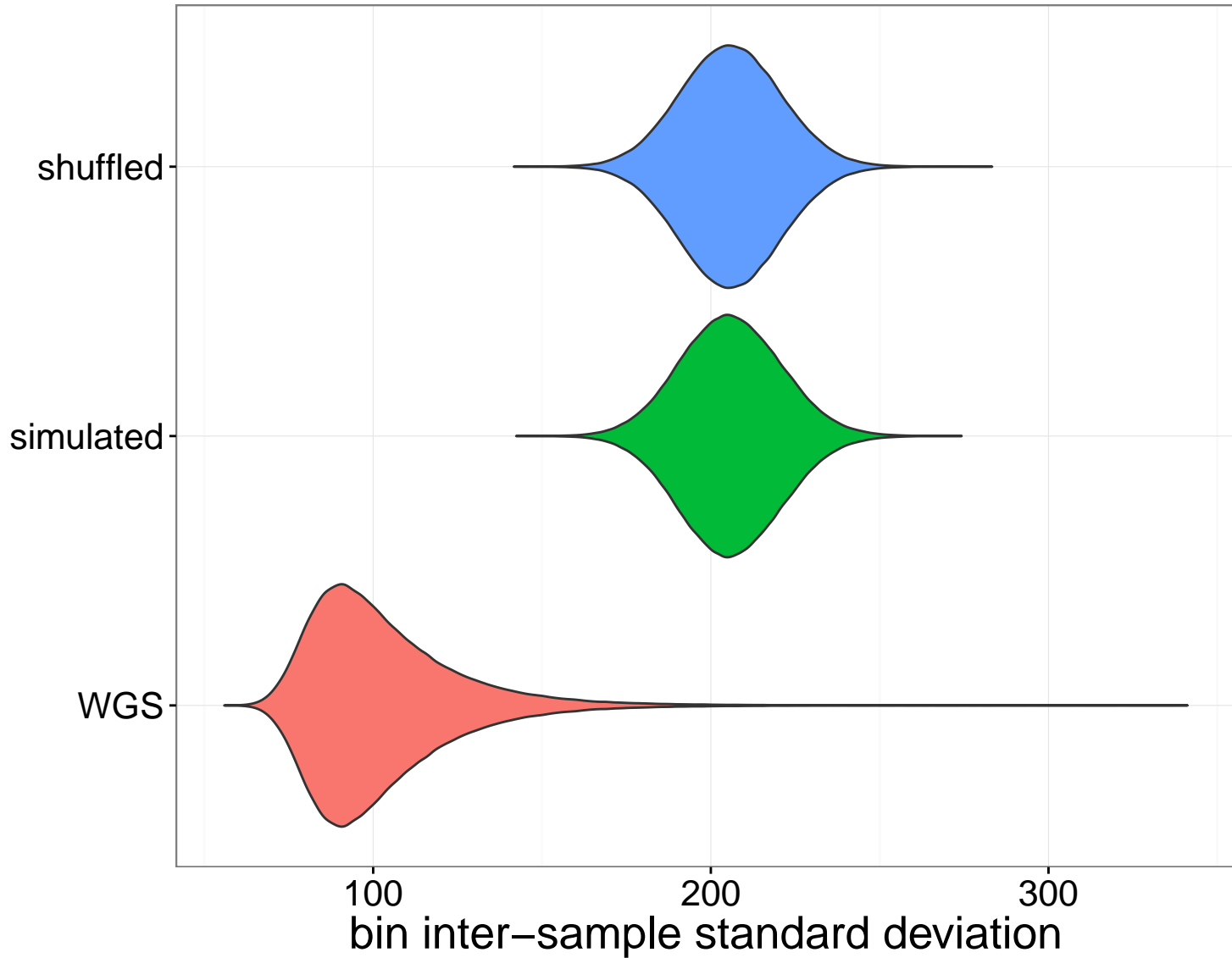
Control : simulated bin counts



Control : shuffled bin counts







proportion of the studied genome

coverage

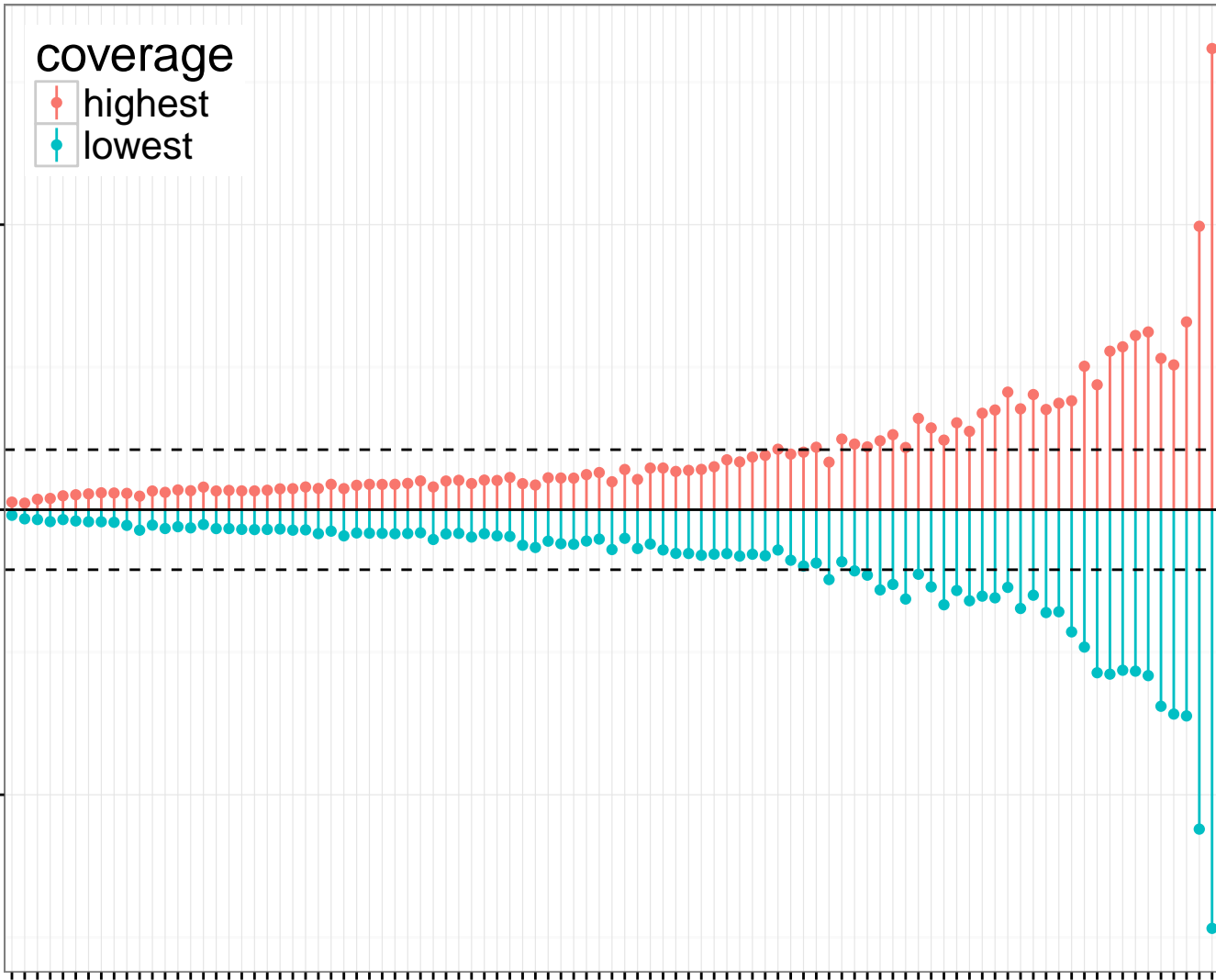
highest  
lowest

0.05

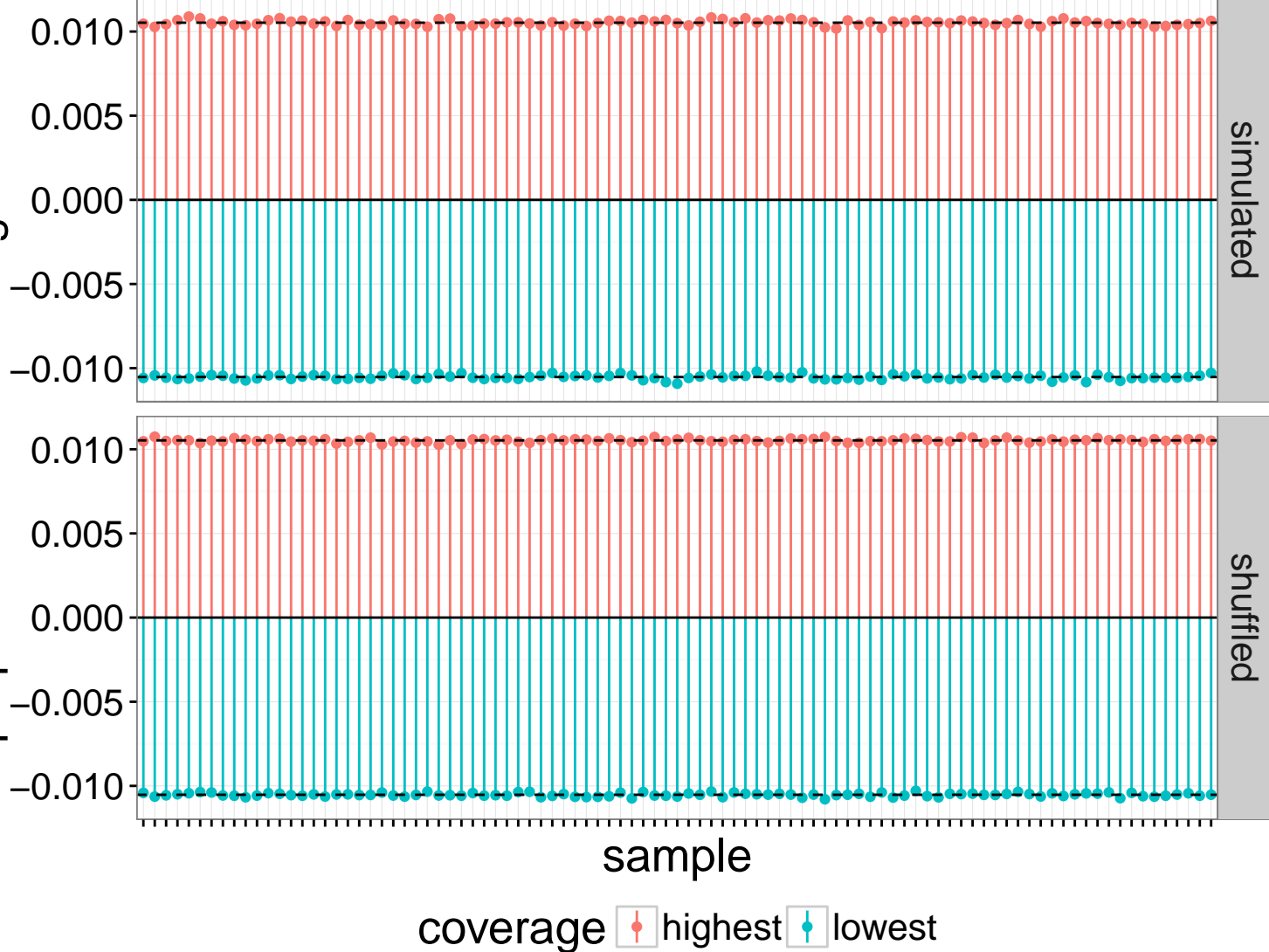
0.00

-0.05

sample



proportion of the studied genome



proportion of the studied genome

coverage

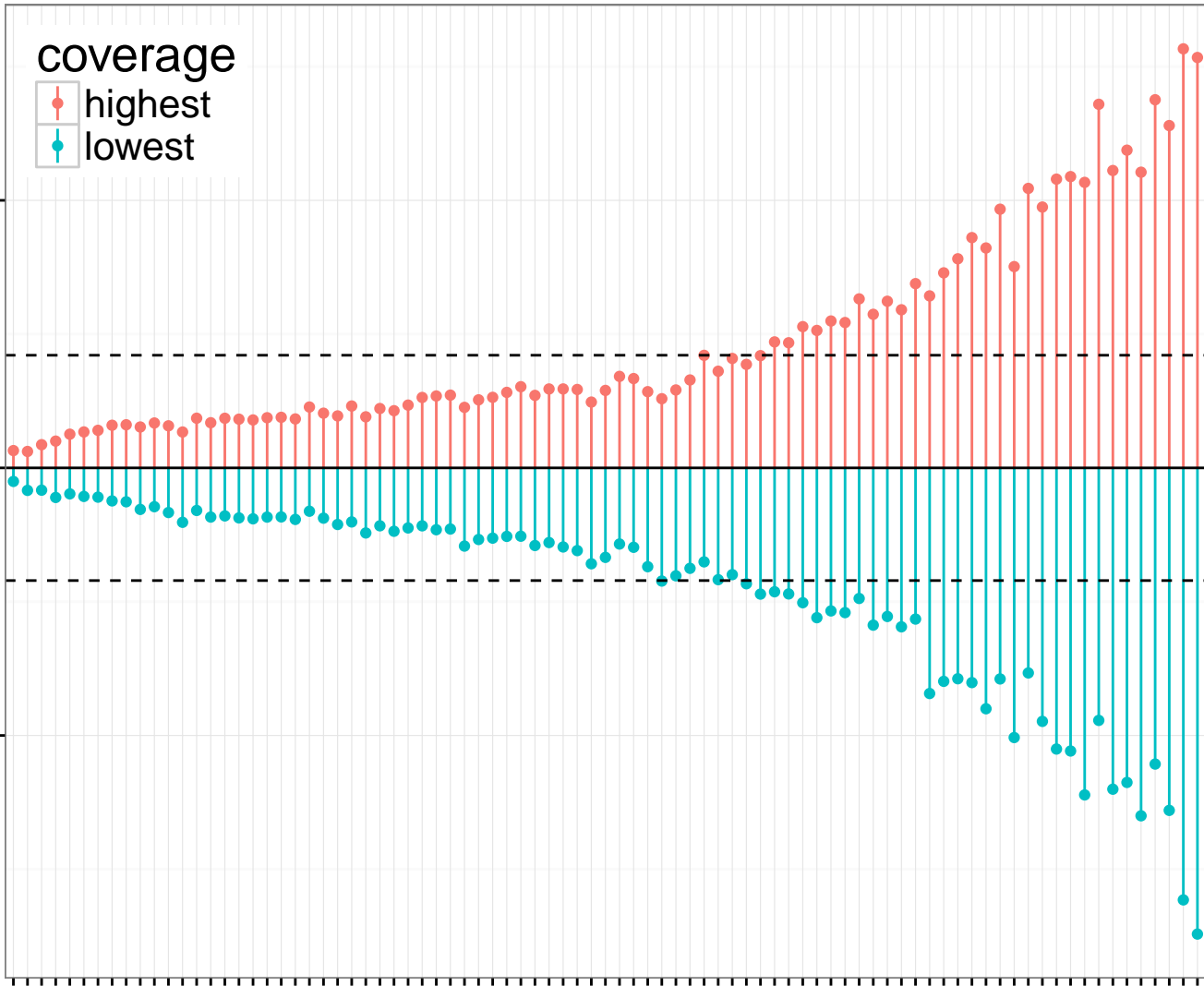
highest  
lowest

0.025

0.000

-0.025

sample





proportion of the studied genome

