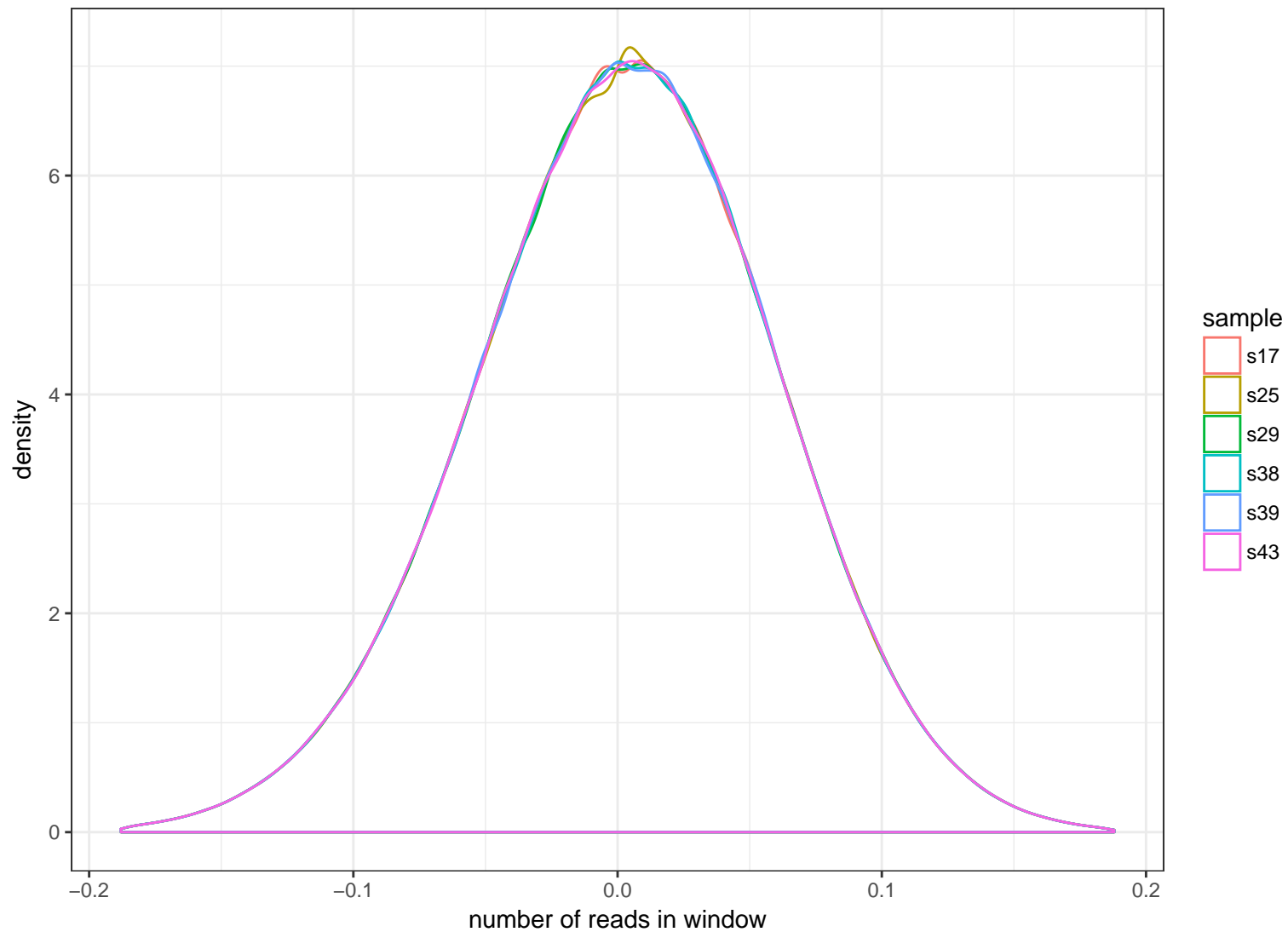
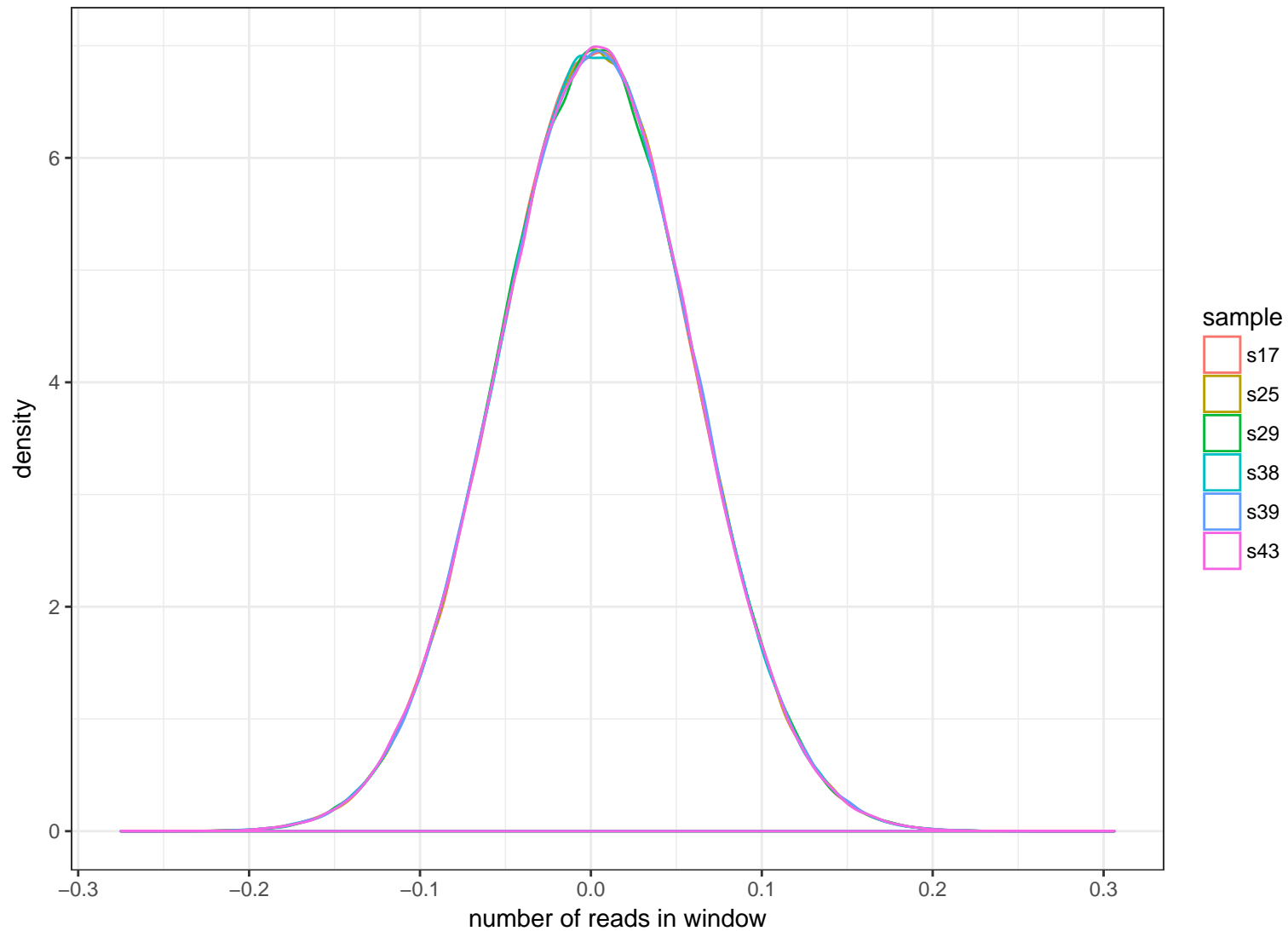


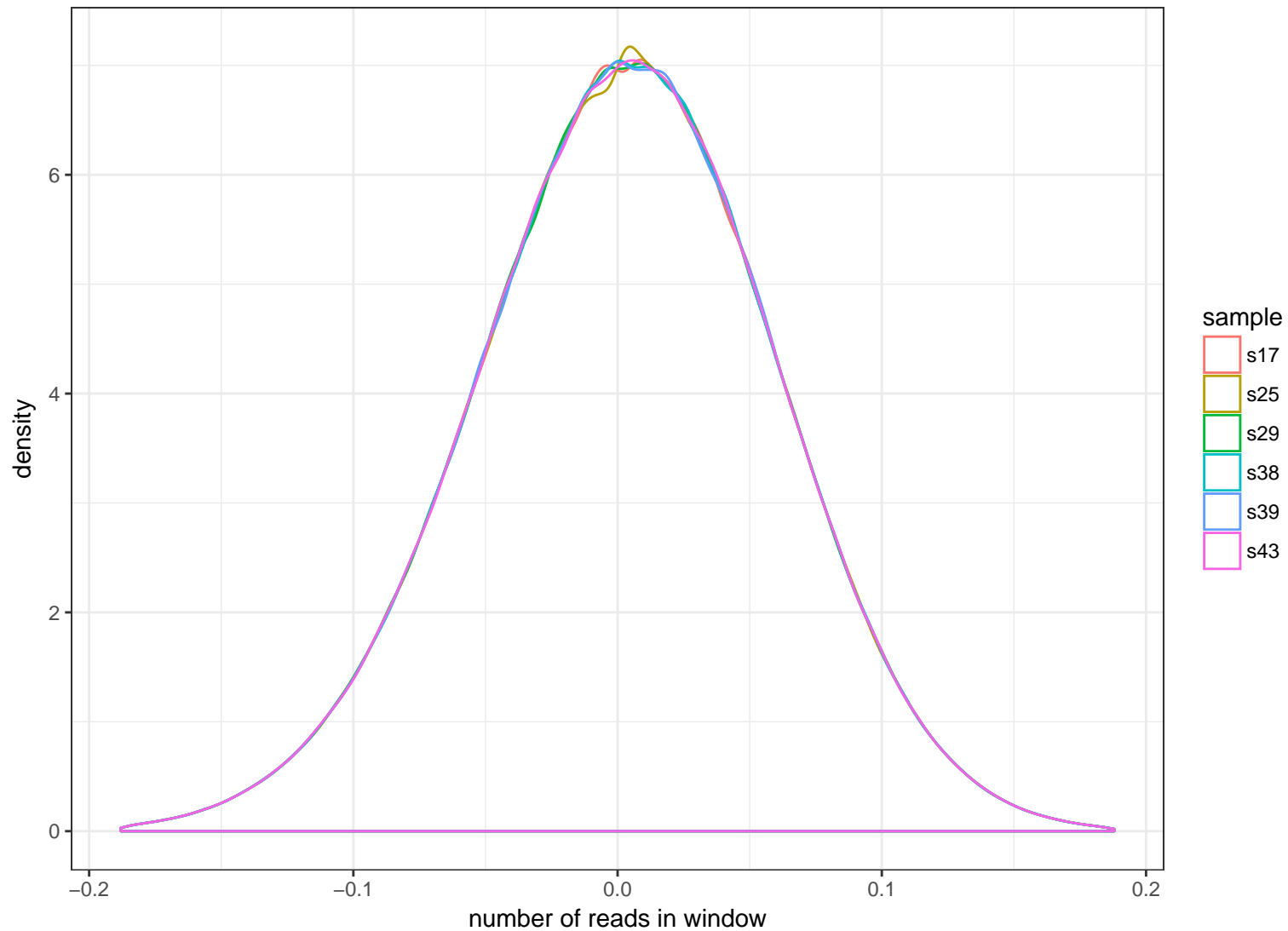
Normalized bin counts

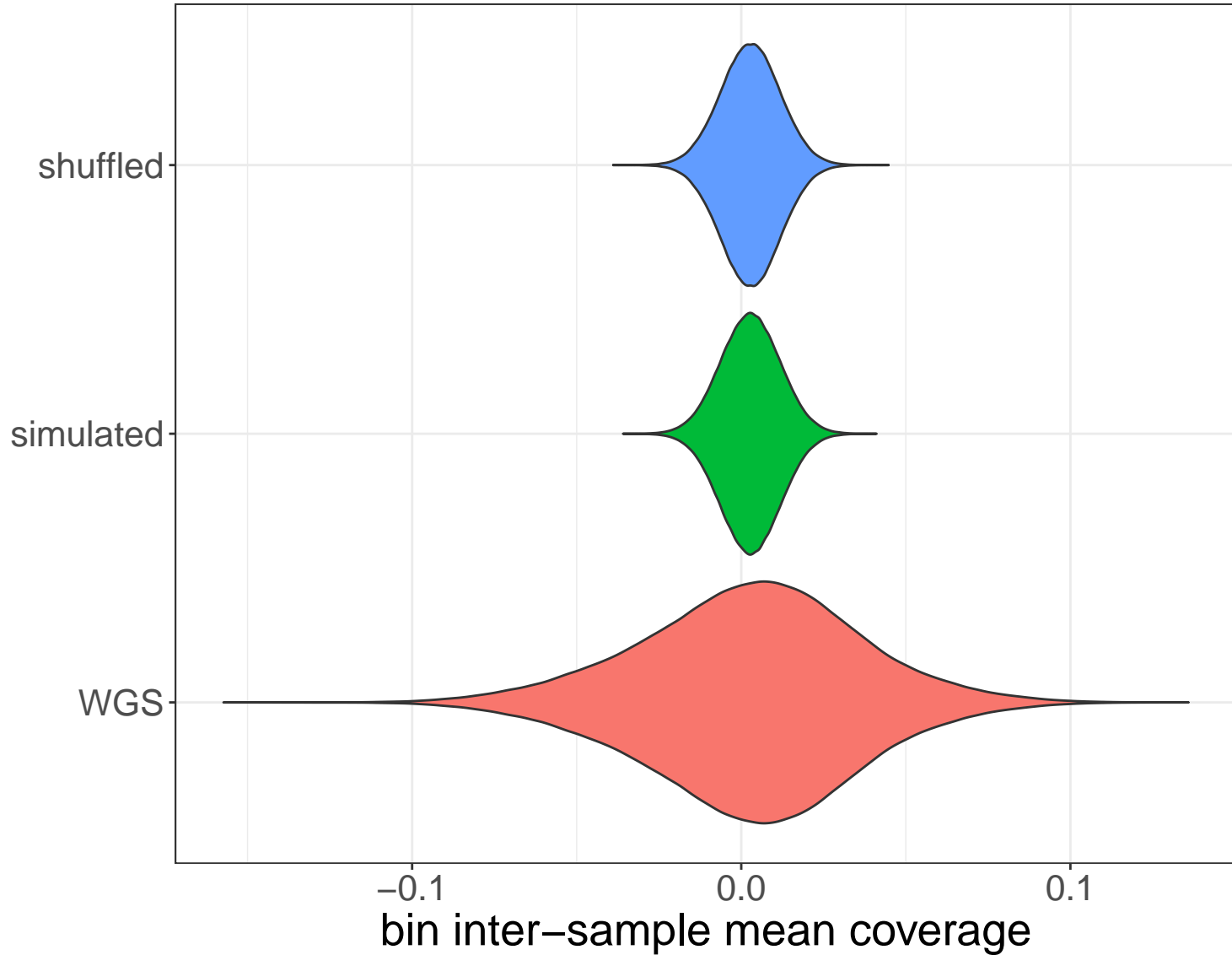


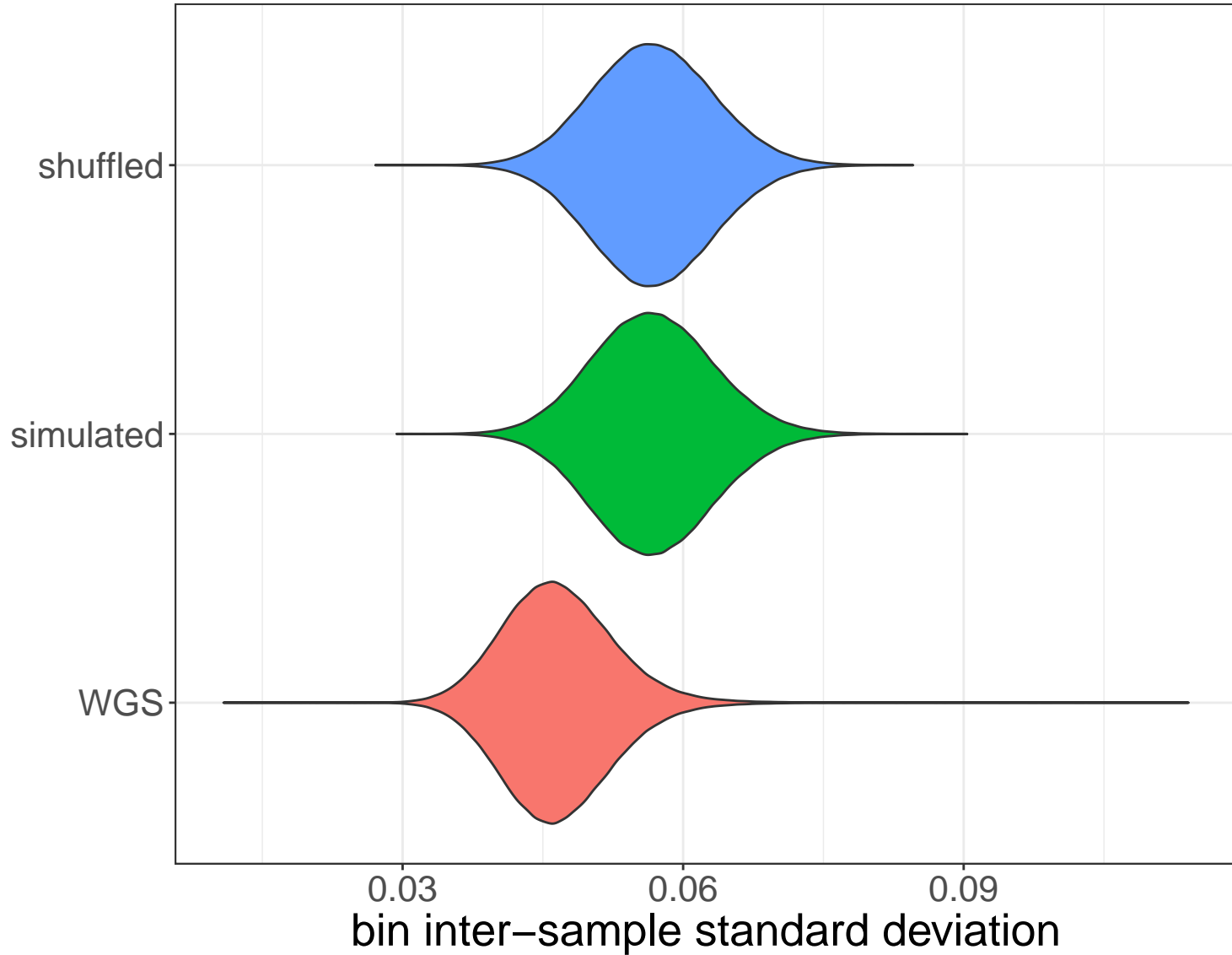
Control : simulated bin counts



Control : shuffled bin counts







proportion of the studied genome

coverage

highest

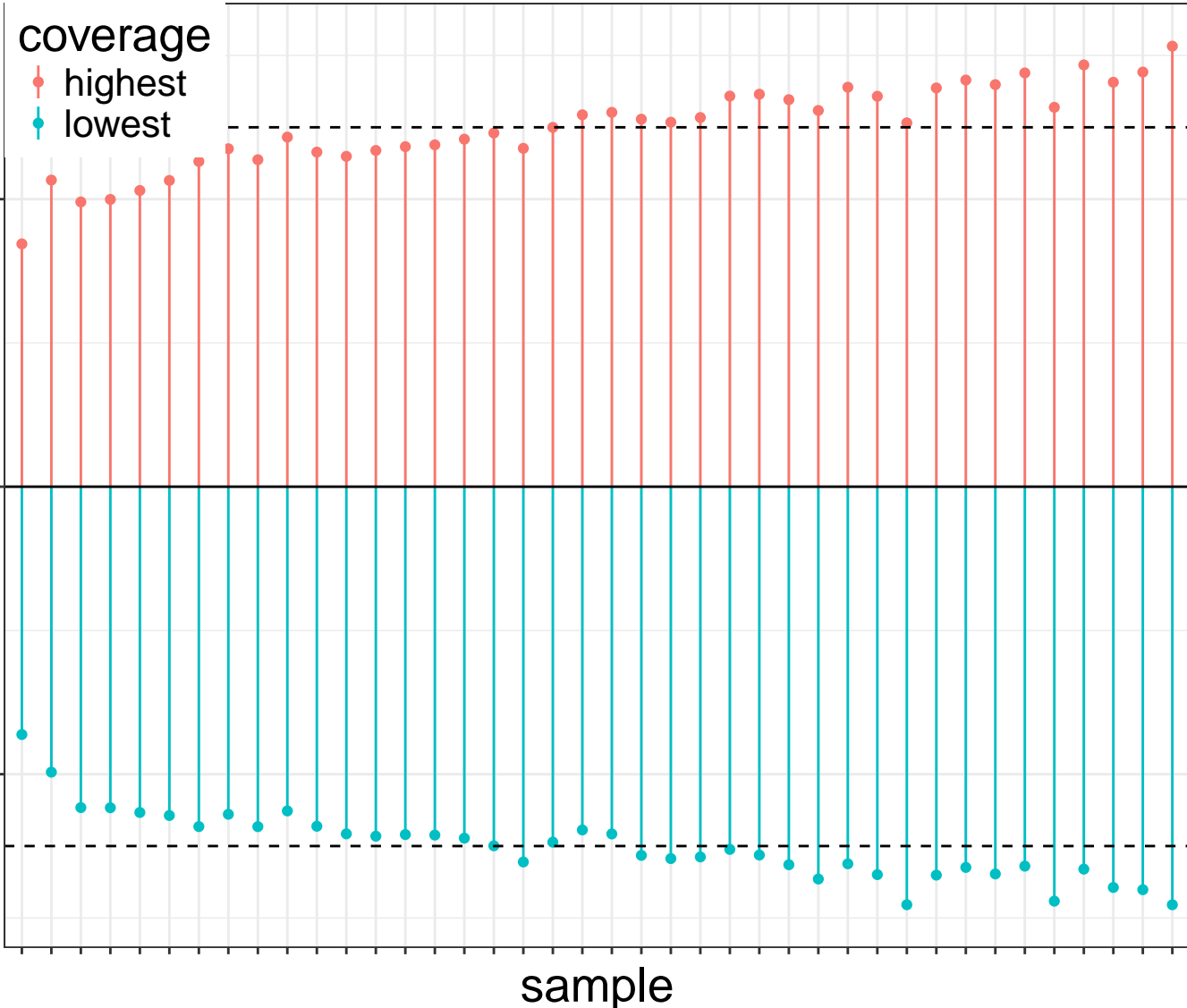
lowest

0.02

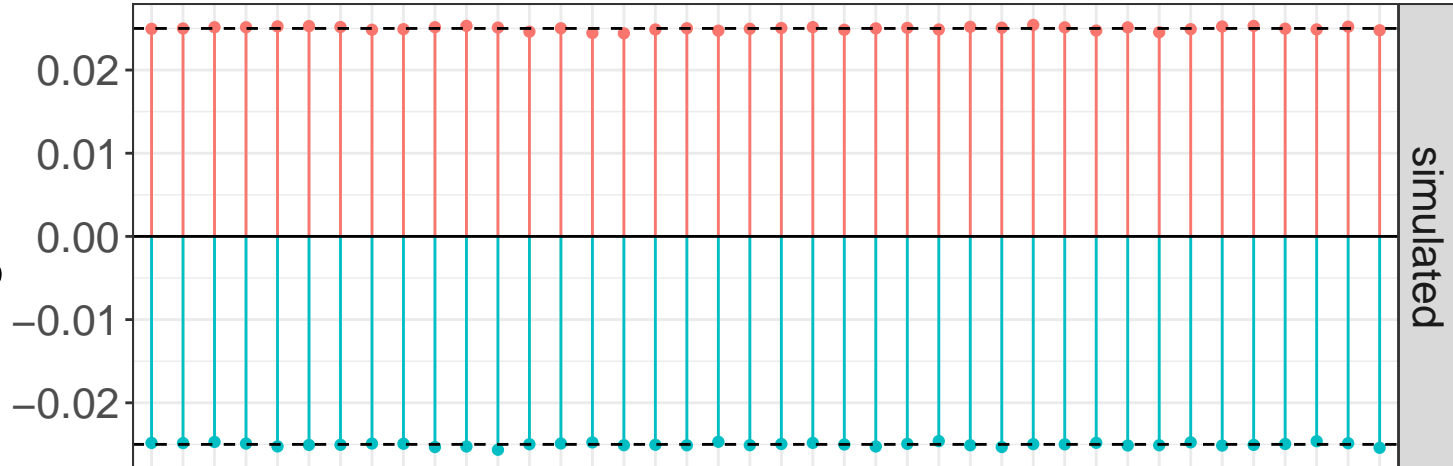
0.00

-0.02

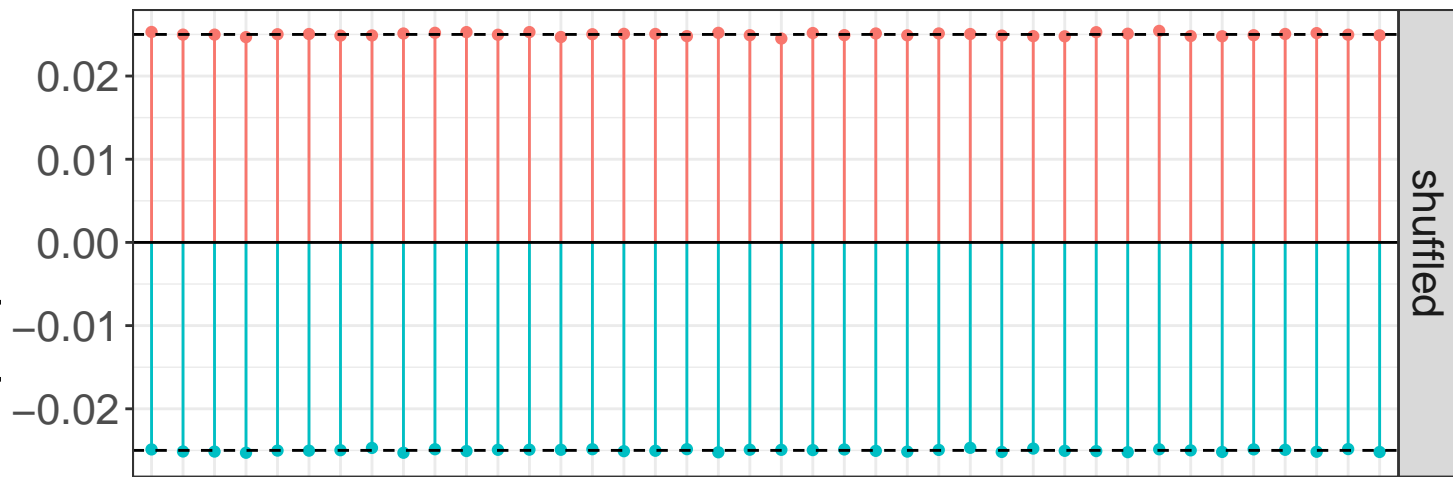
sample



proportion of the studied genome



simulated



shuffled

sample

coverage highest lowest