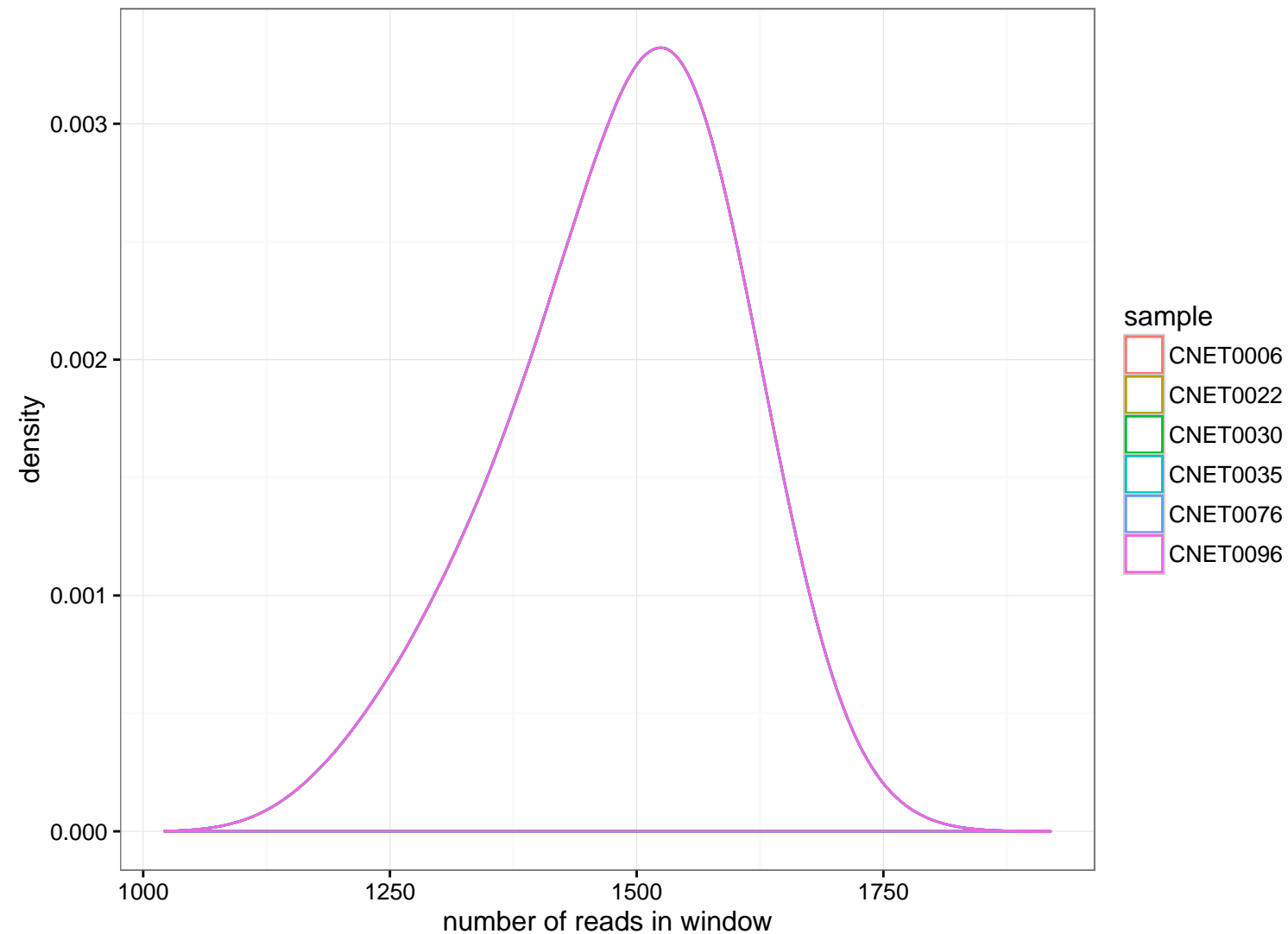
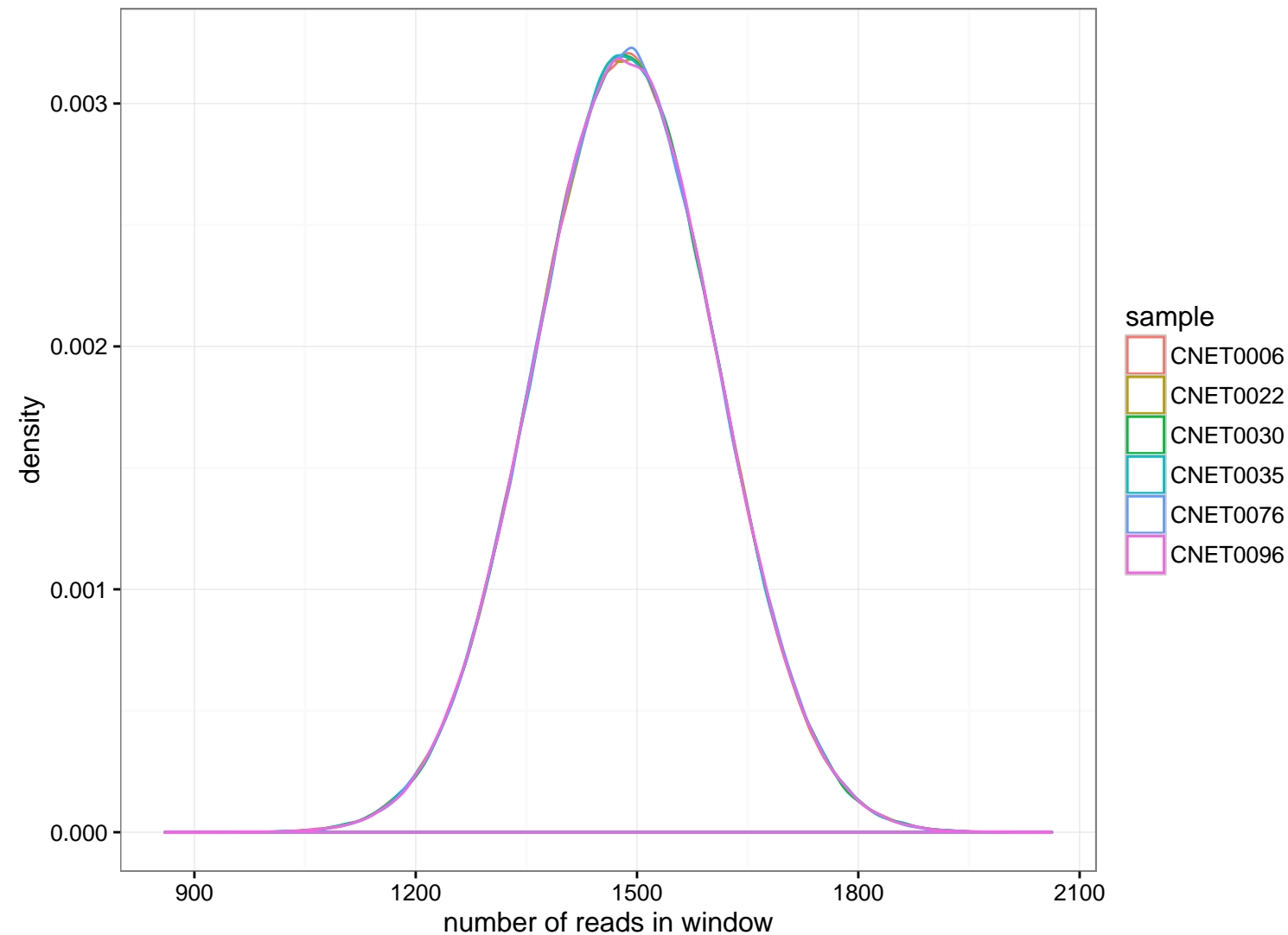


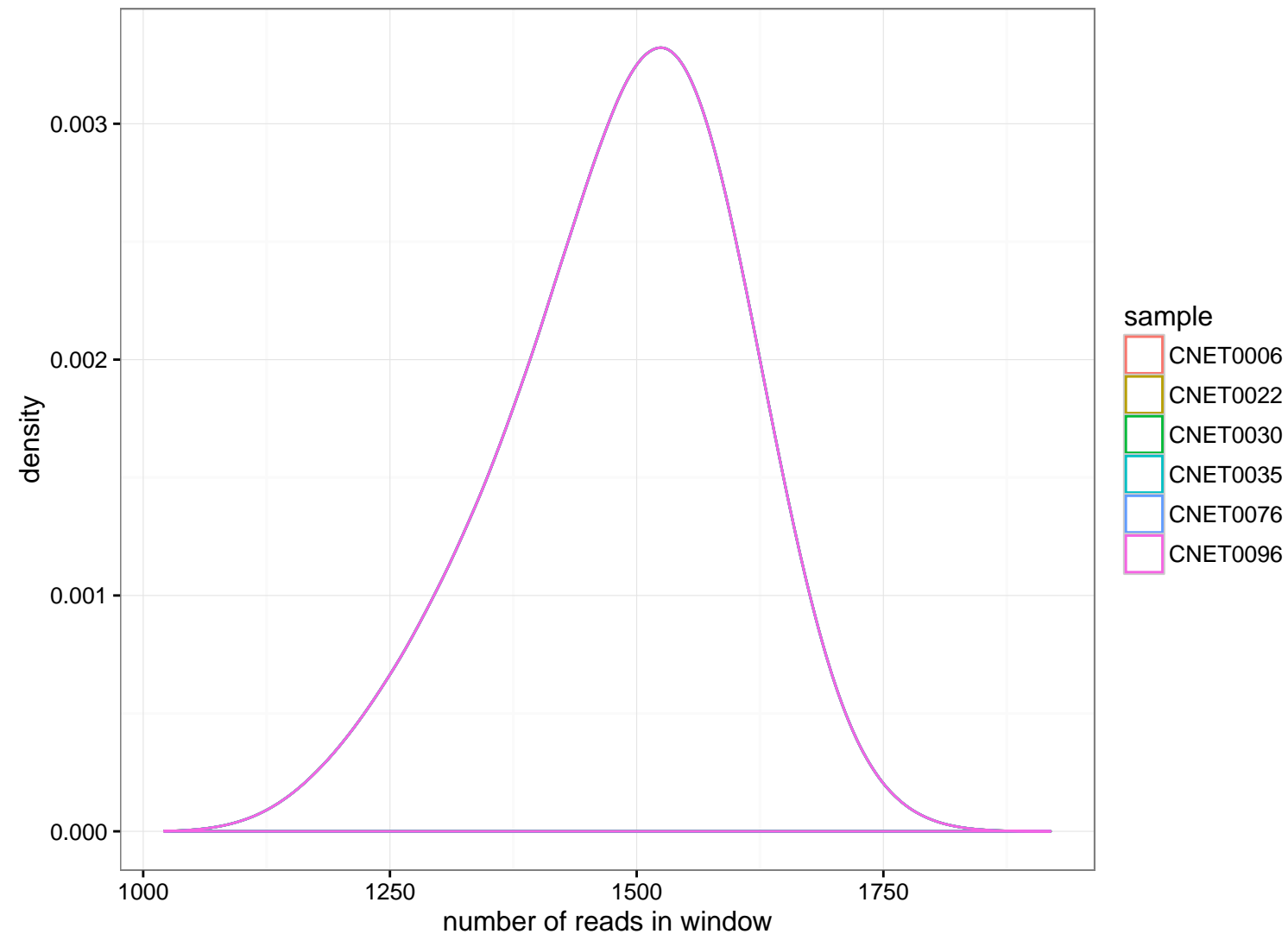
Normalized bin counts

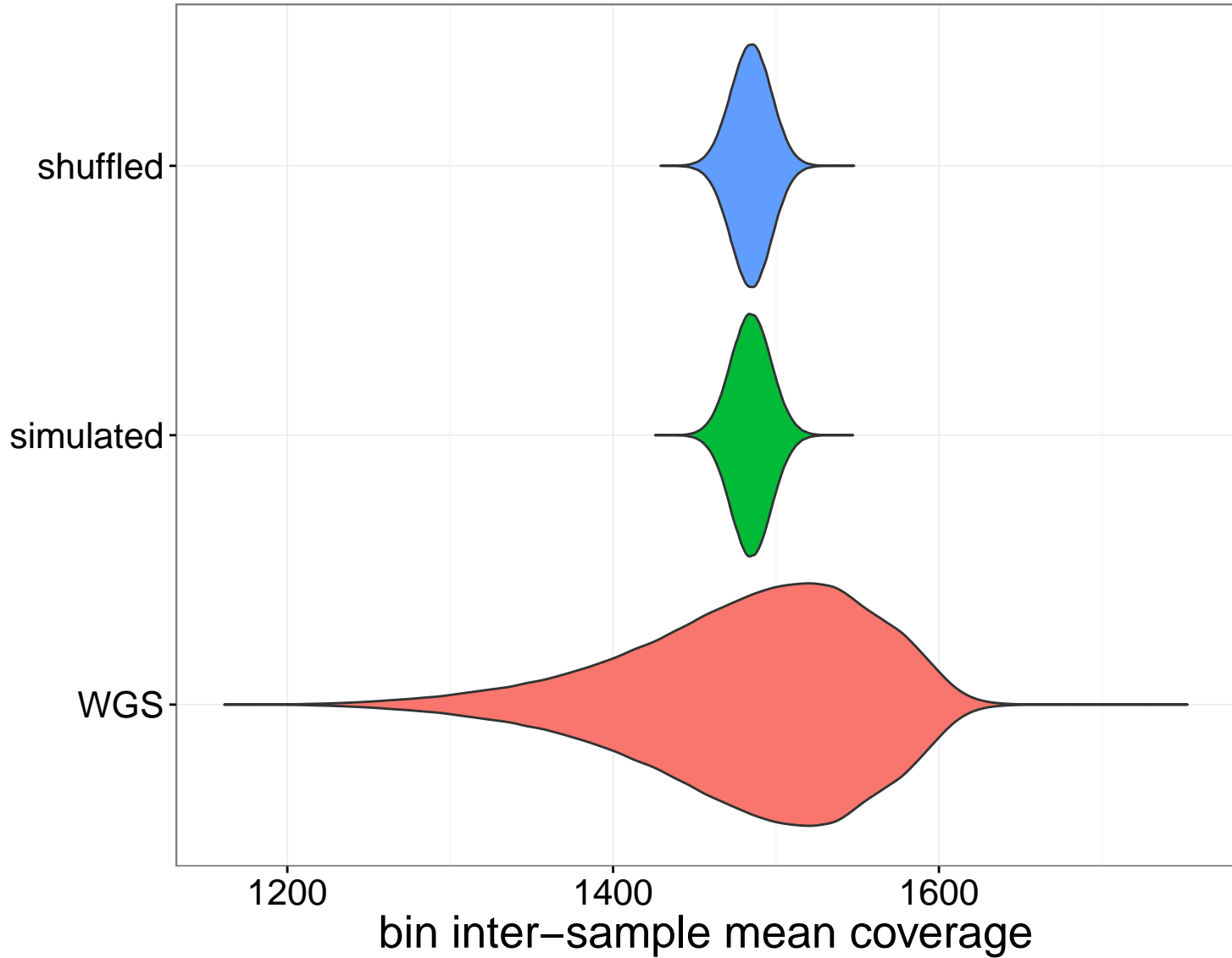


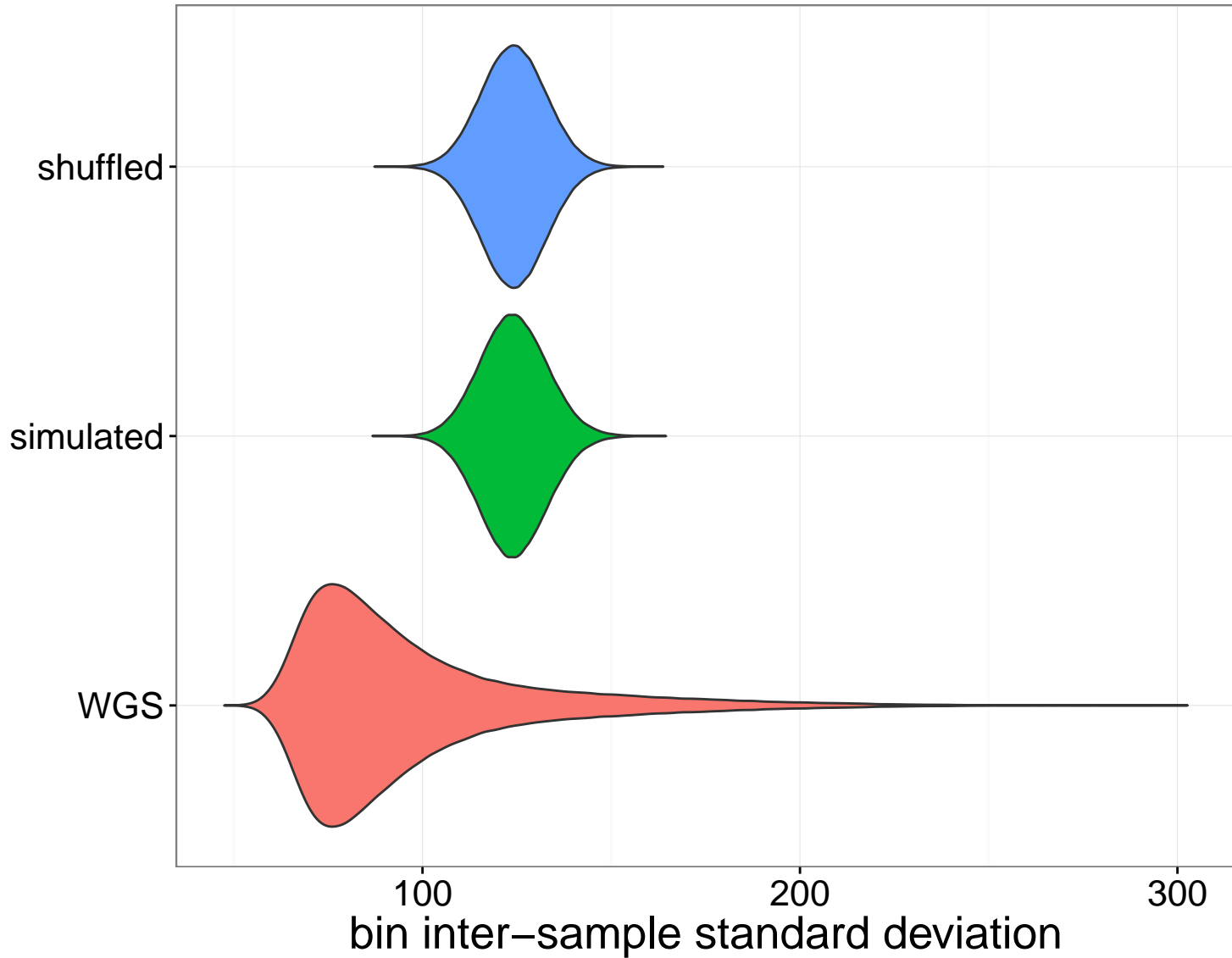
Control : simulated bin counts



Control : shuffled bin counts







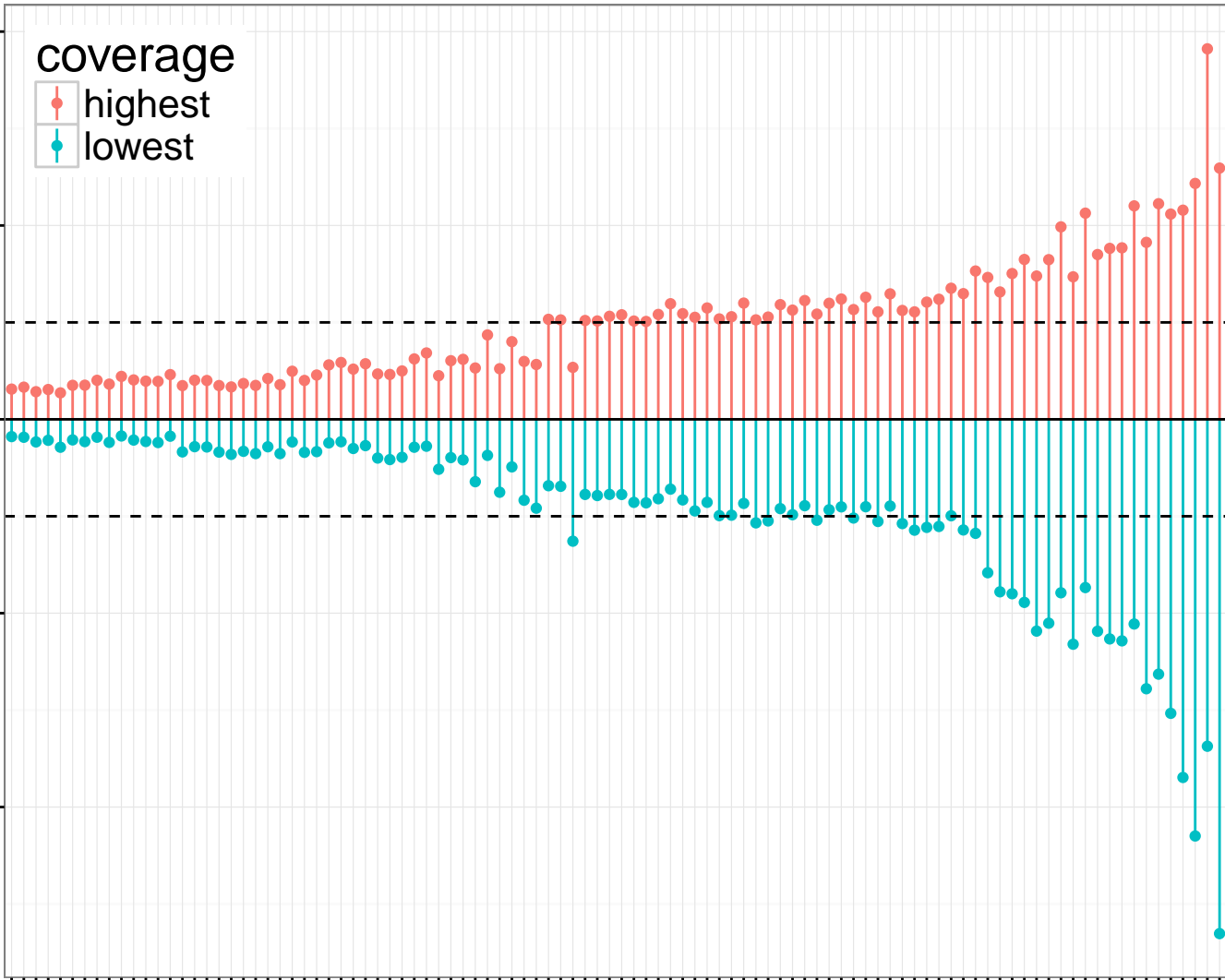
proportion of the studied genome

coverage

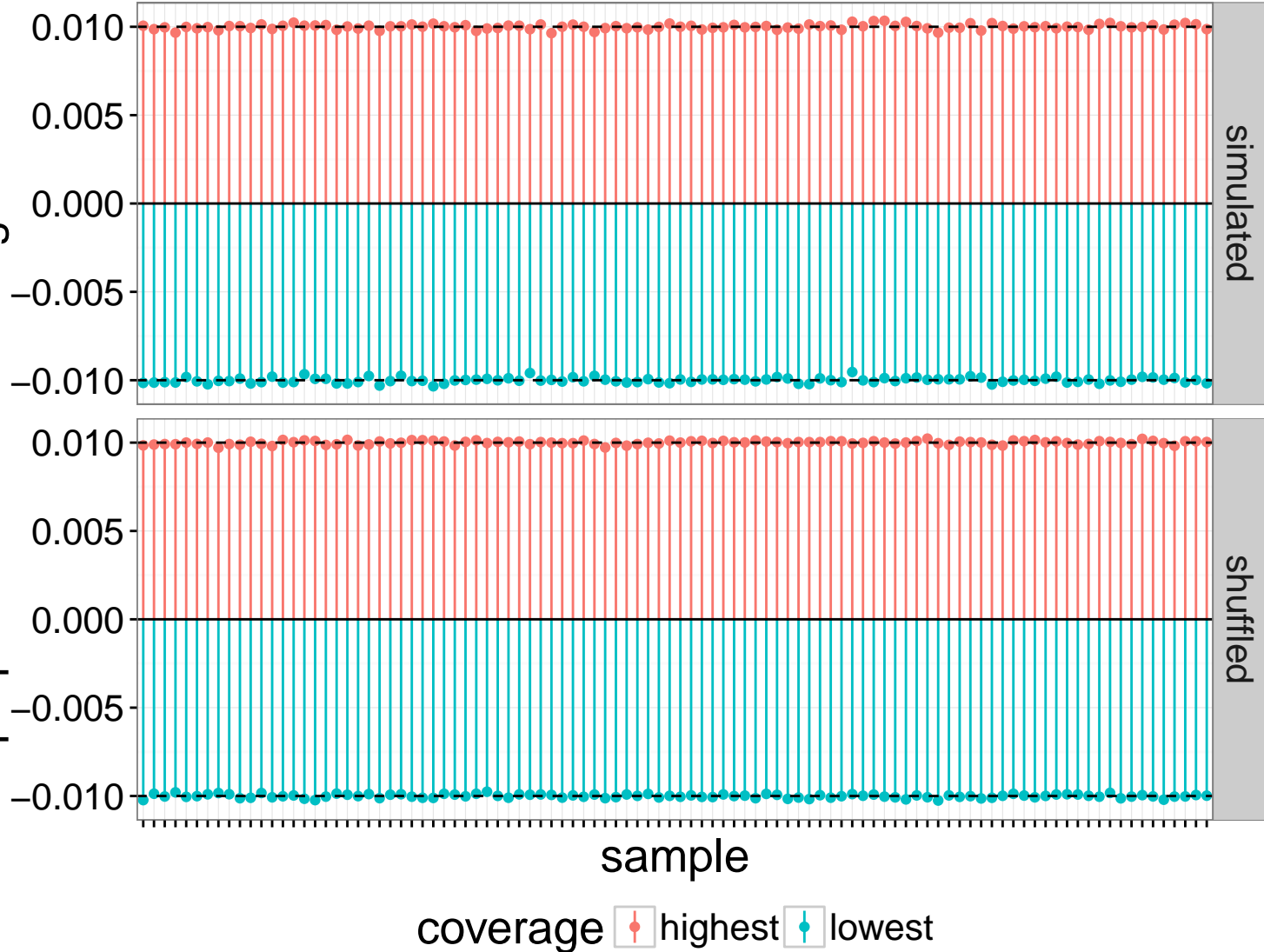
highest

lowest

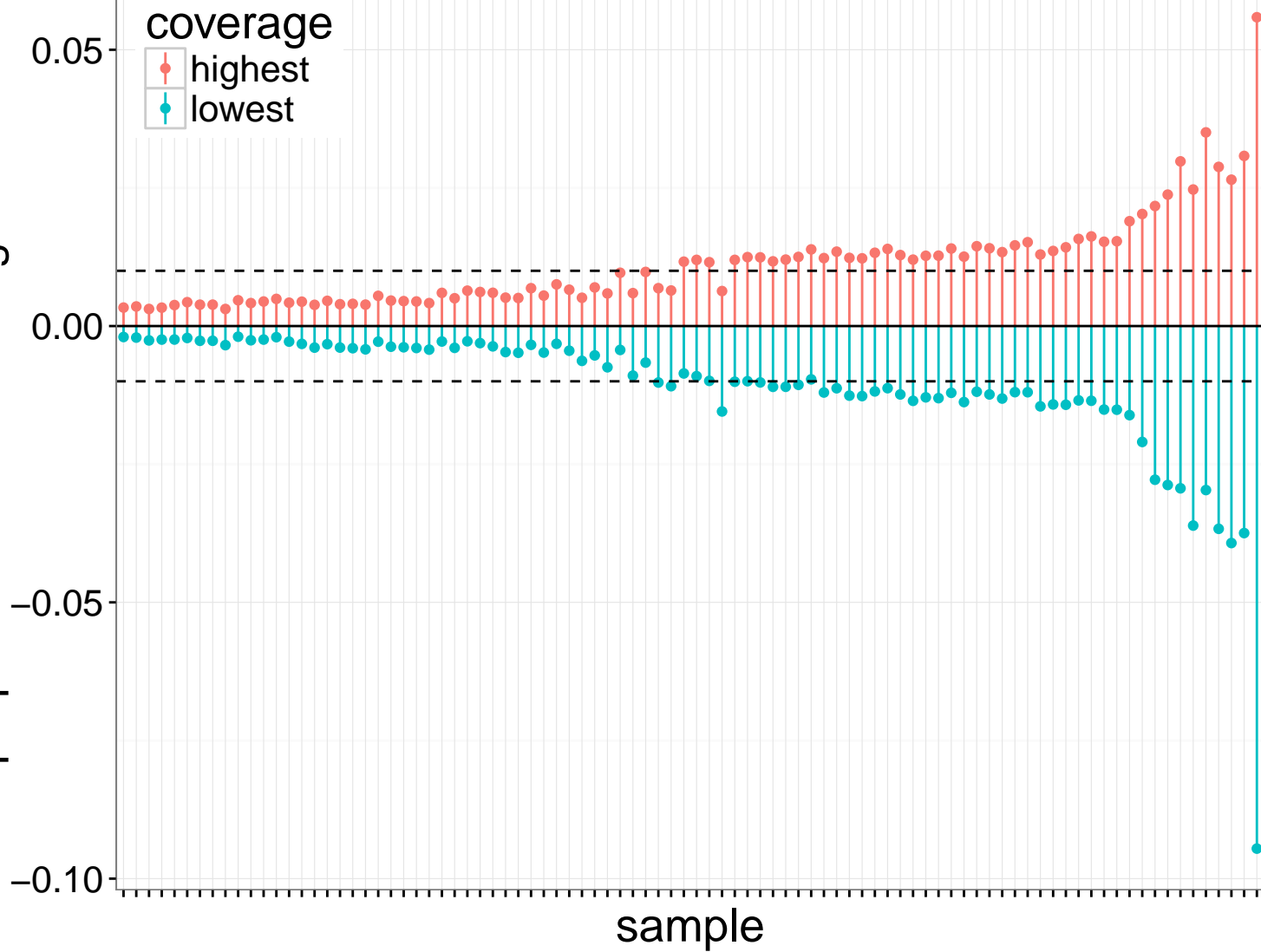
sample



proportion of the studied genome



proportion of the studied genome





proportion of the studied genome

