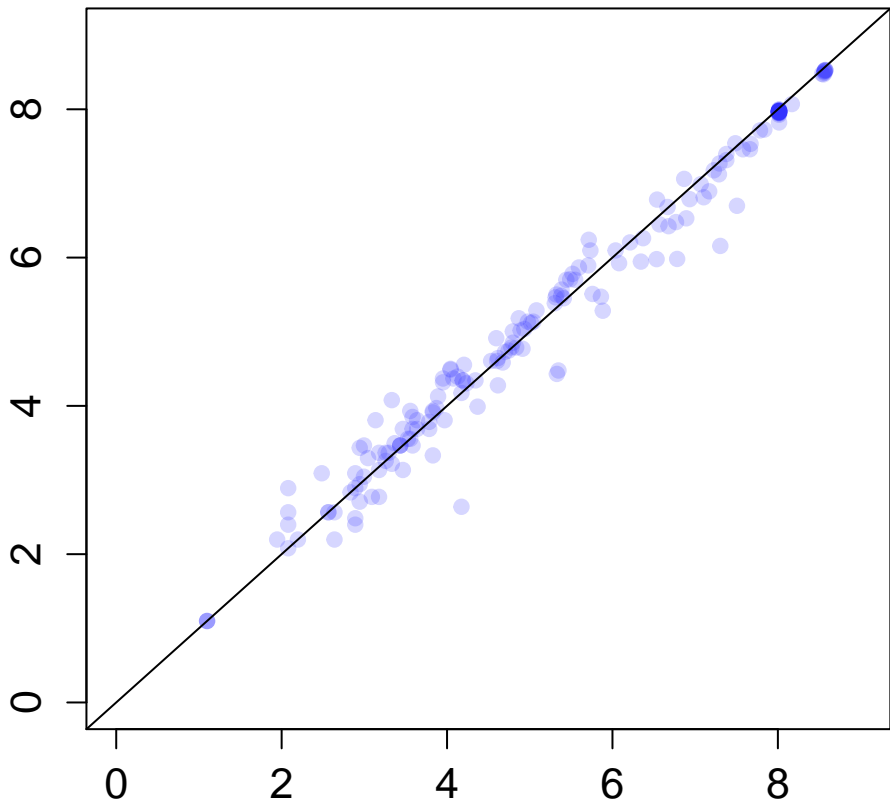


log(number of unique k-mers) in genome background



log(number of unique k-mers) in random sequences