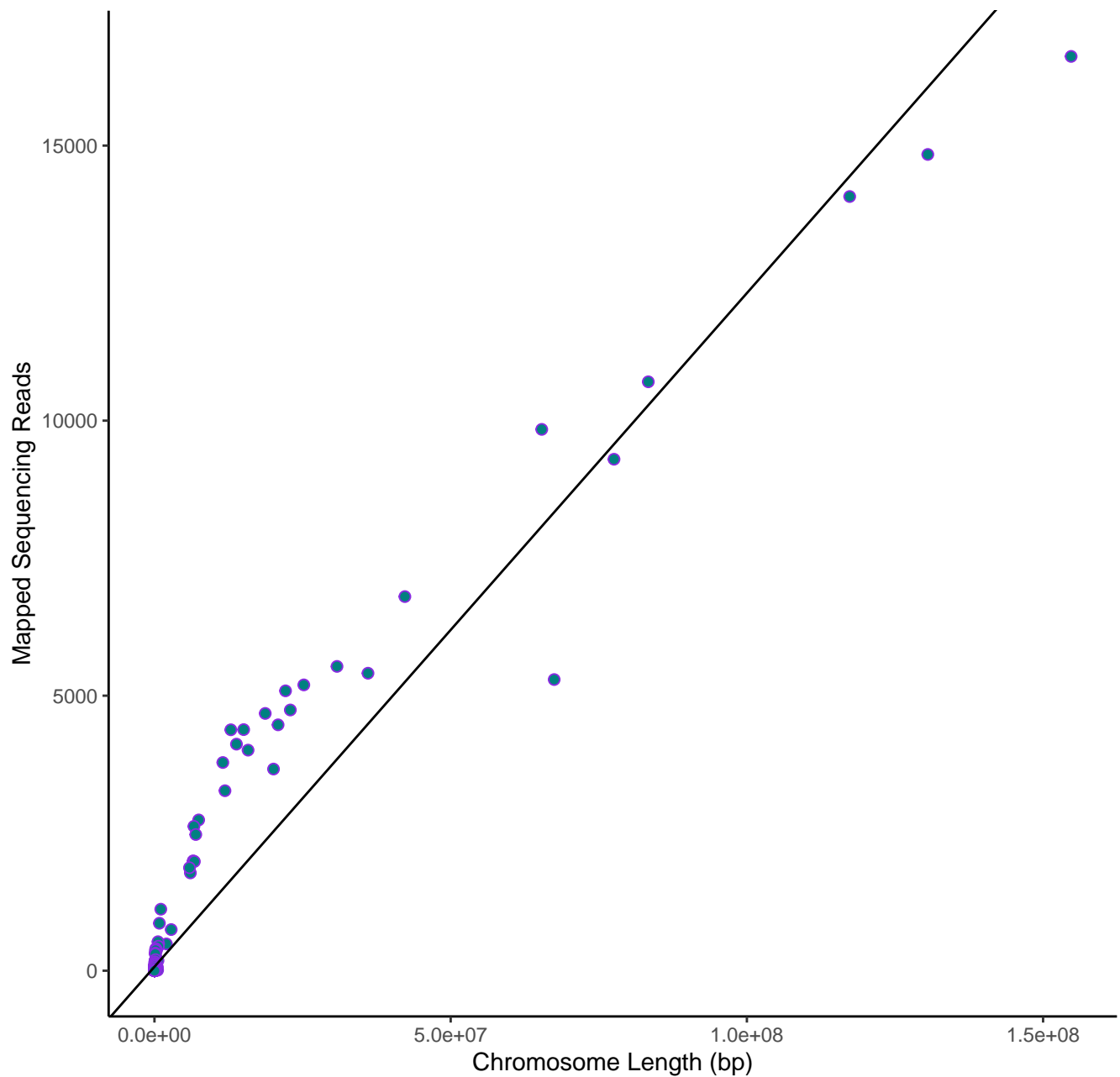


Number of sequencing reads for chromosomes of different lengths



Number of sequencing reads for chromosomes of different lengths

