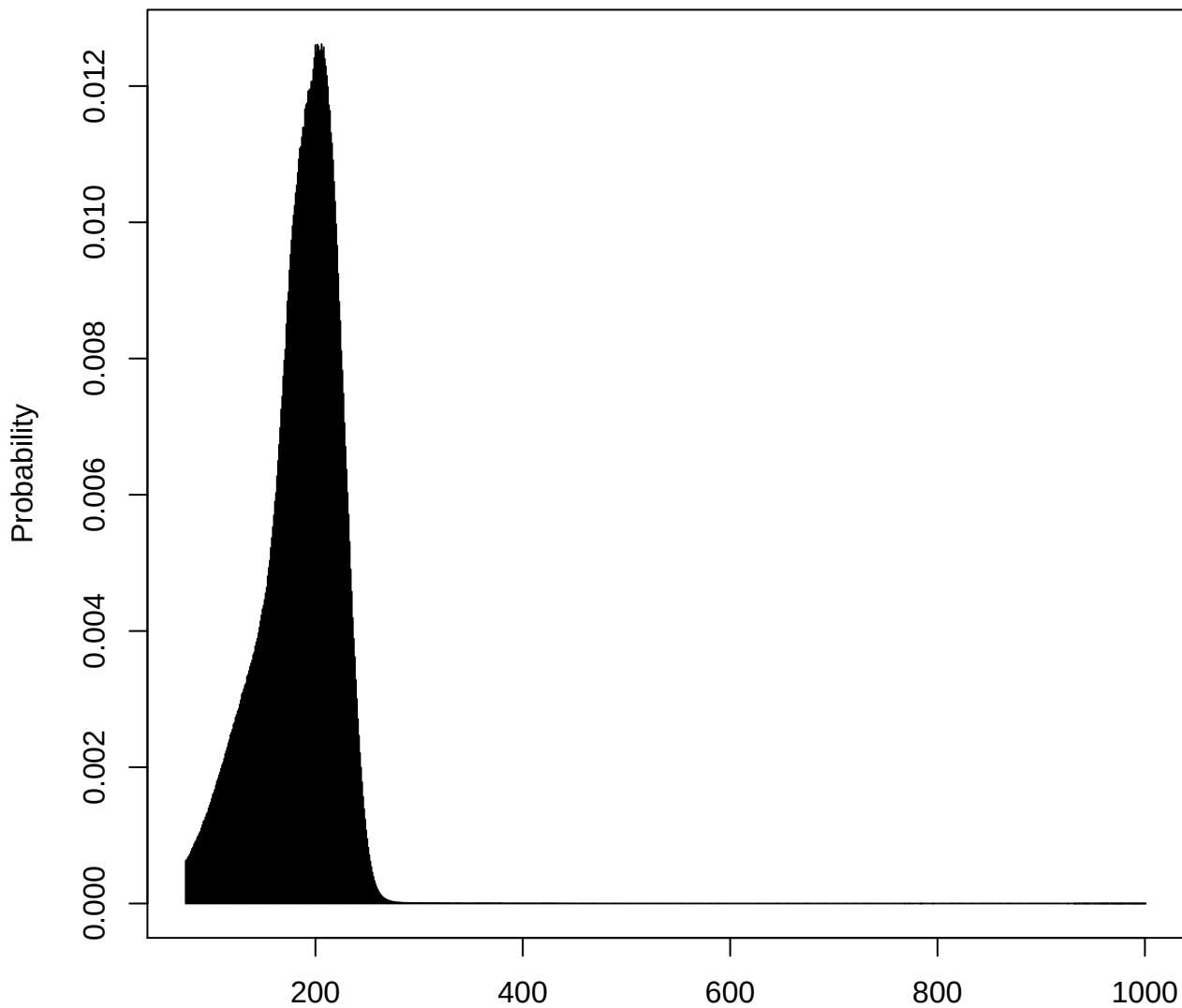


Fragment Length Distribution

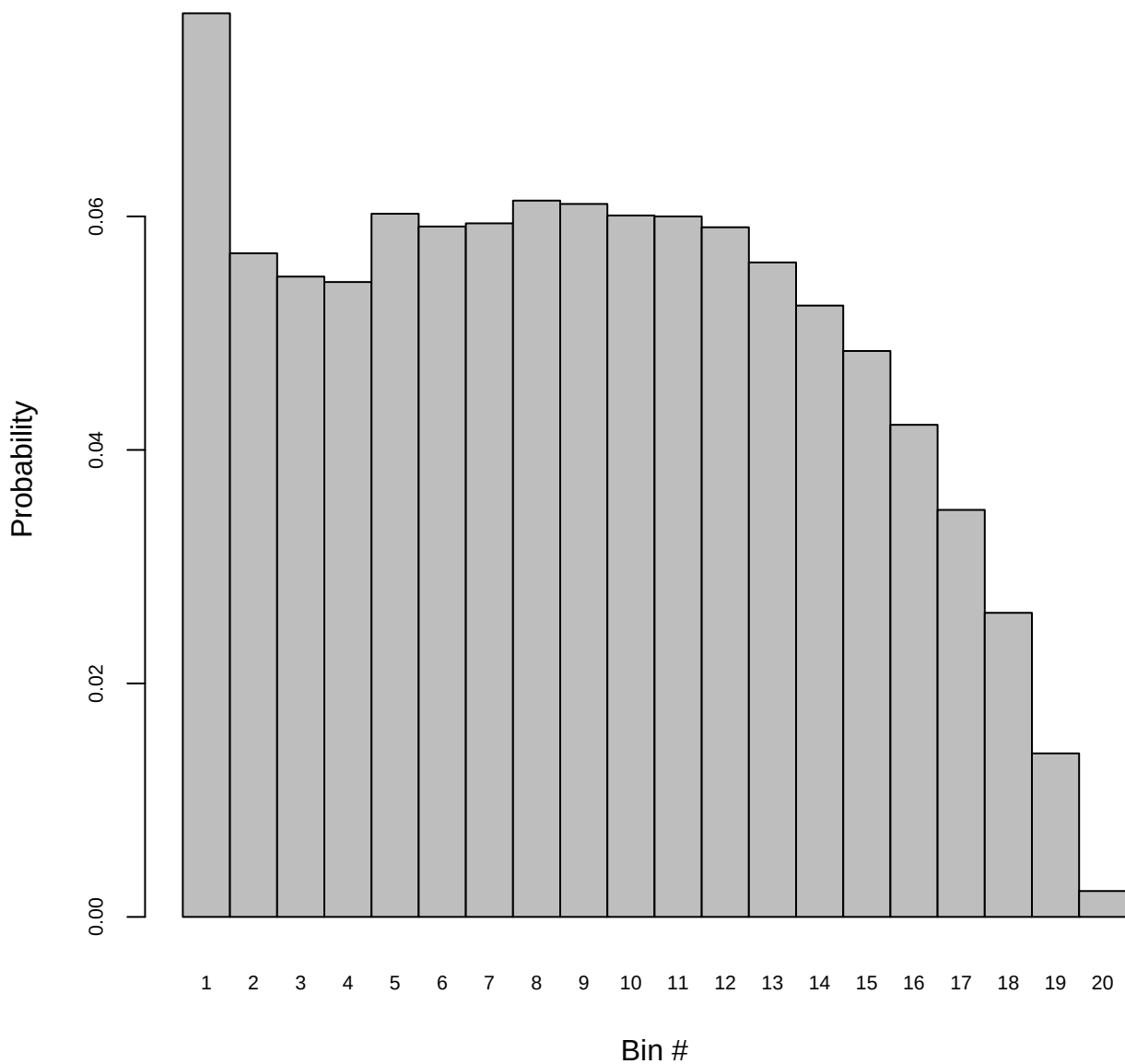


Mean = 185.3 , Std = 37.7

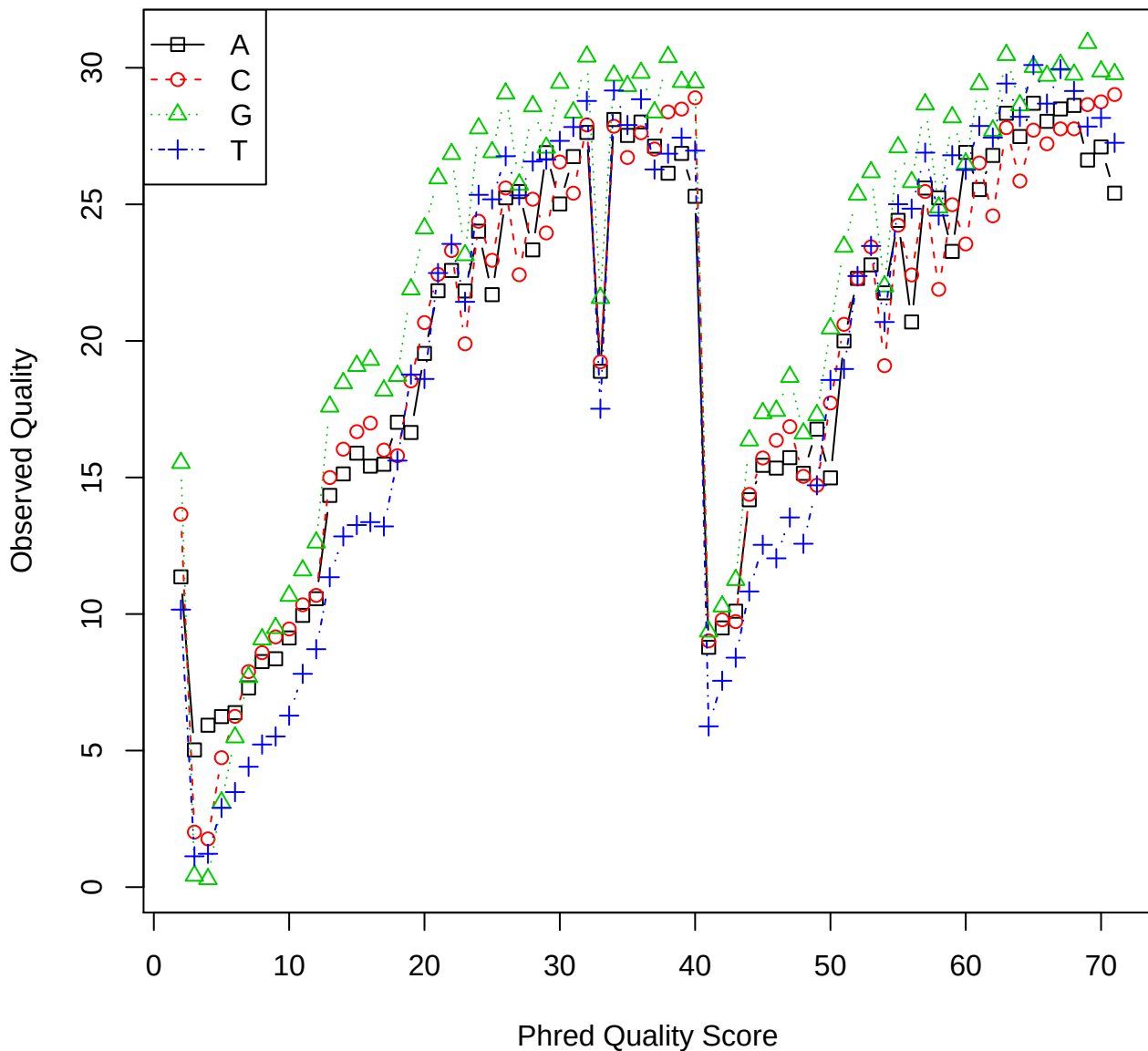
Read Length Distribution



Read Start Position Distribution



Observed Quality vs. Phred Quality Score



Histogram of Alignments per Fragment

