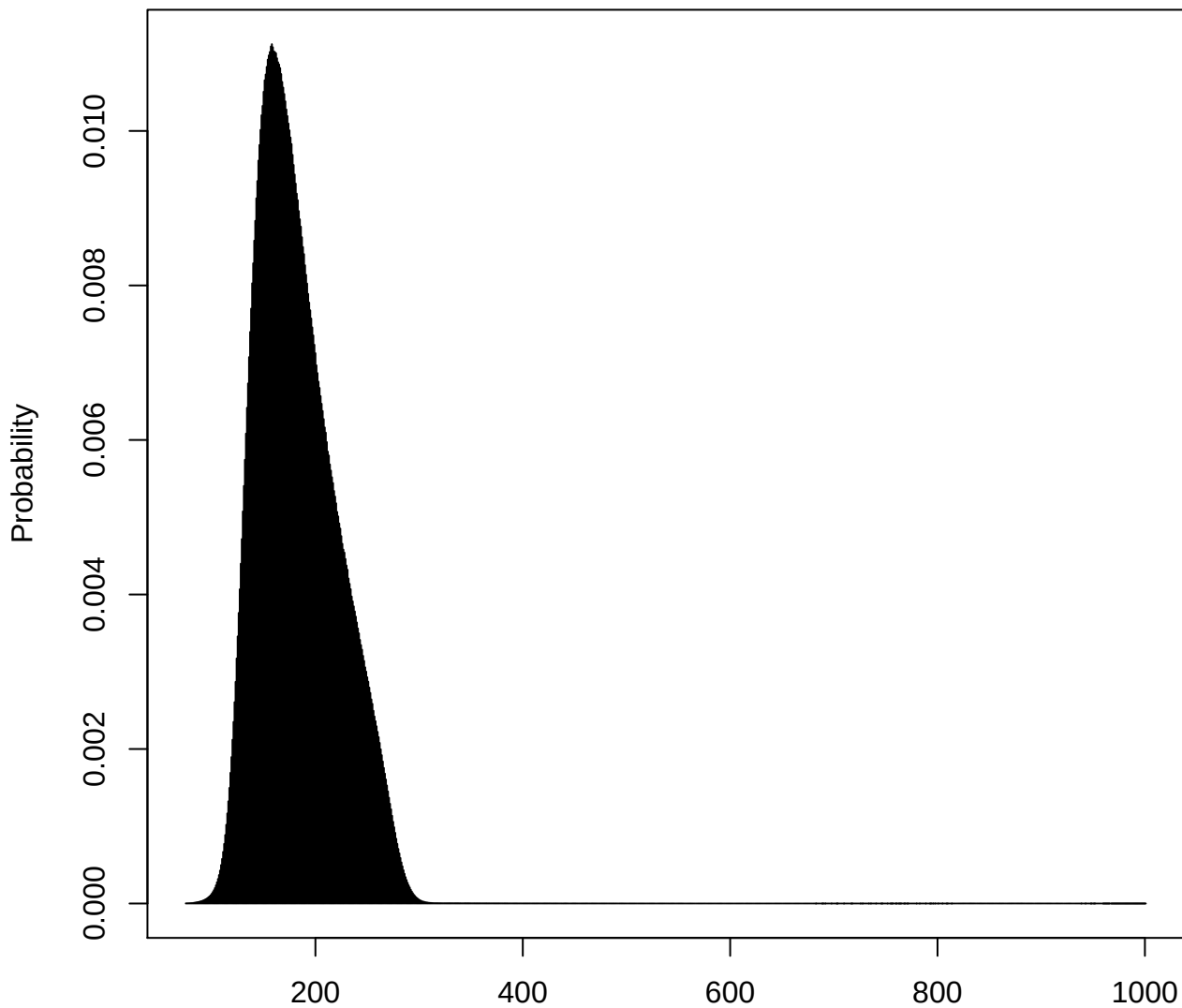


Fragment Length Distribution

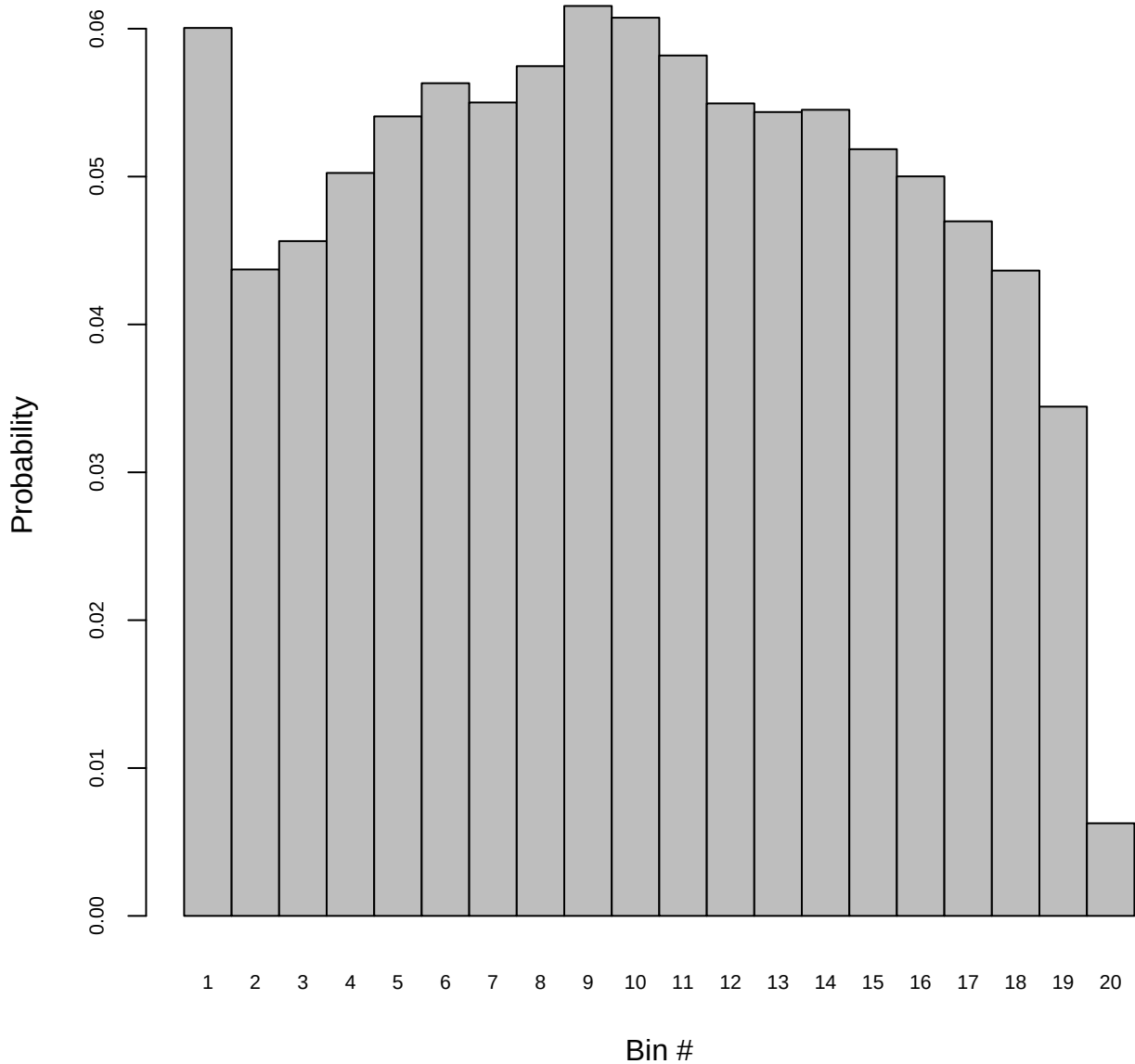


Mean = 182.8 , Std = 39.2

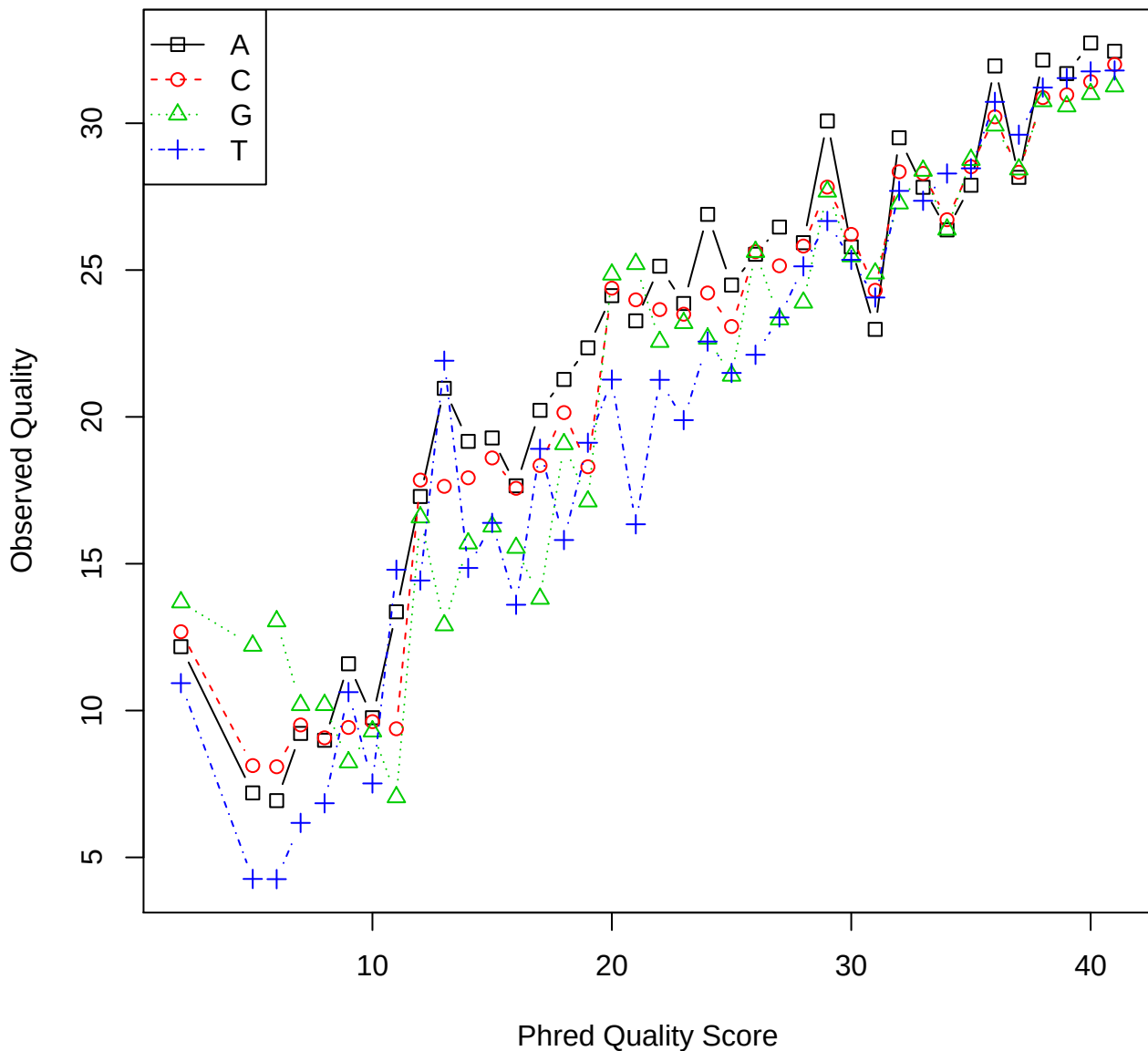
Read Length Distribution



Read Start Position Distribution



Observed Quality vs. Phred Quality Score



Histogram of Alignments per Fragment

