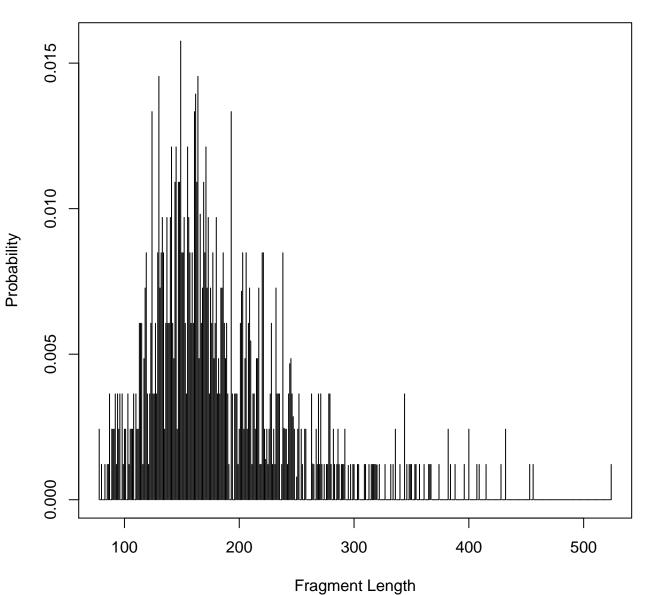
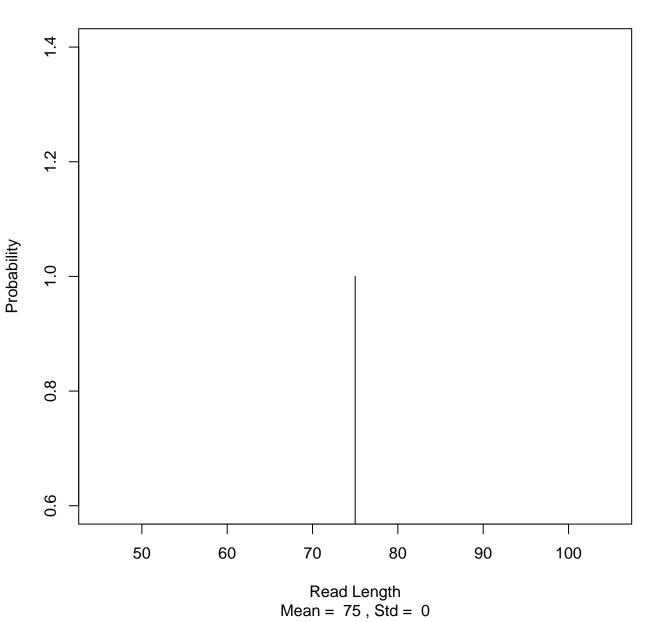
Fragment Length Distribution

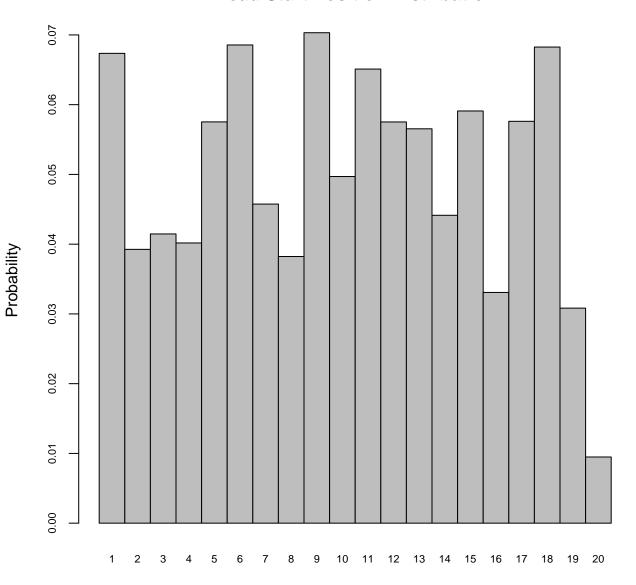


Mean = 183.6 , Std = 64.9

Read Length Distribution

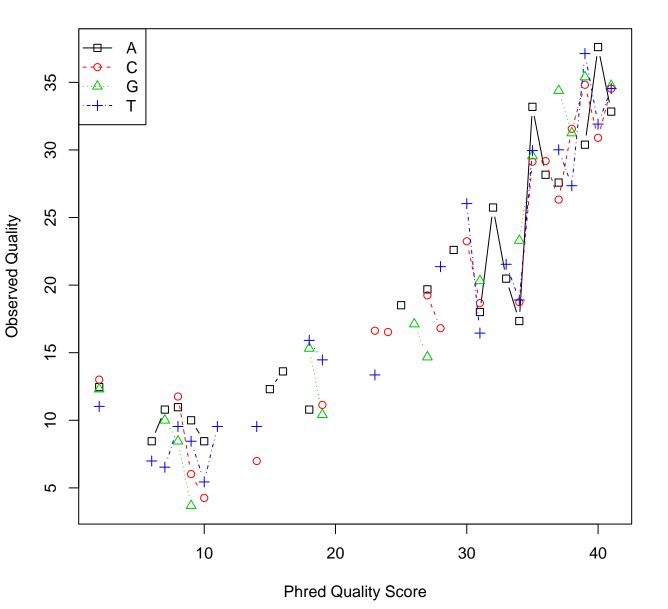


Read Start Position Distribution



Bin#

Observed Quality vs. Phred Quality Score



Histogram of Alignments per Fragment

