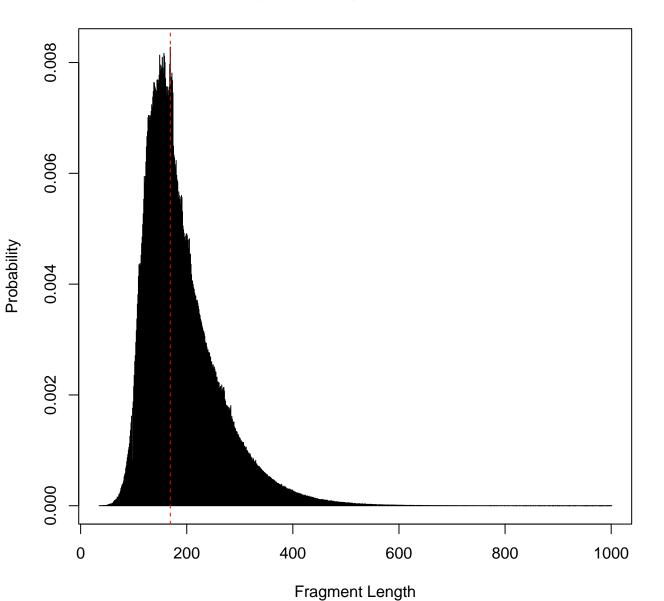
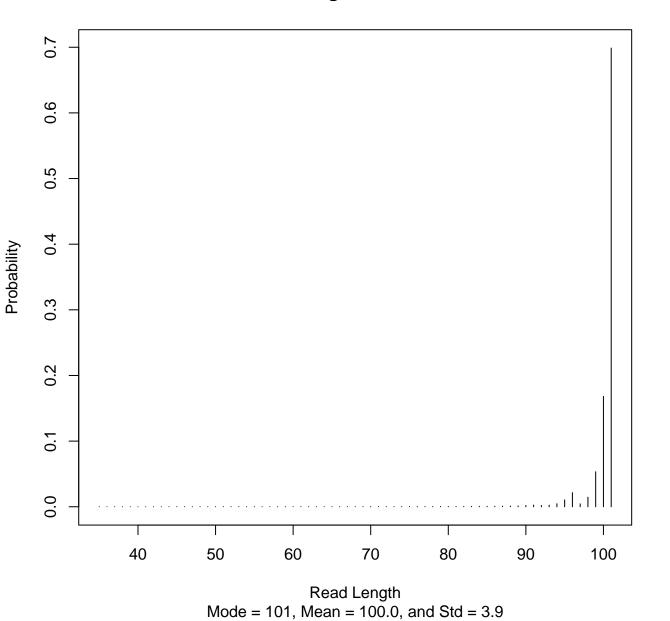
Fragment Length Distribution

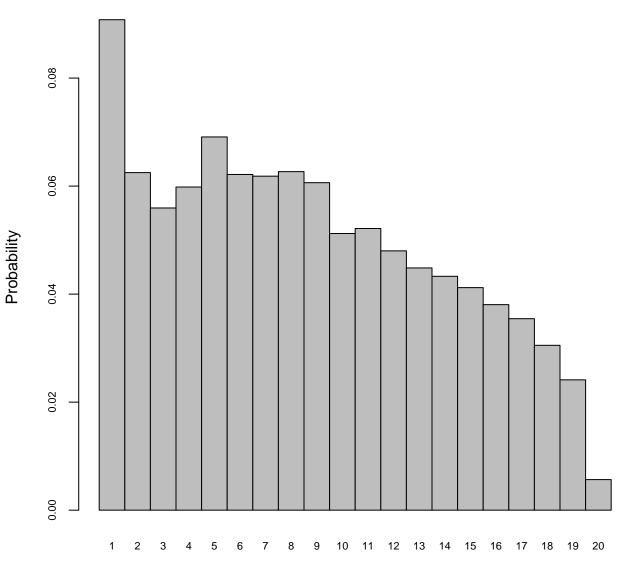


Mode = 169, Mean = 190.7, and Std = 71.4

Read Length Distribution

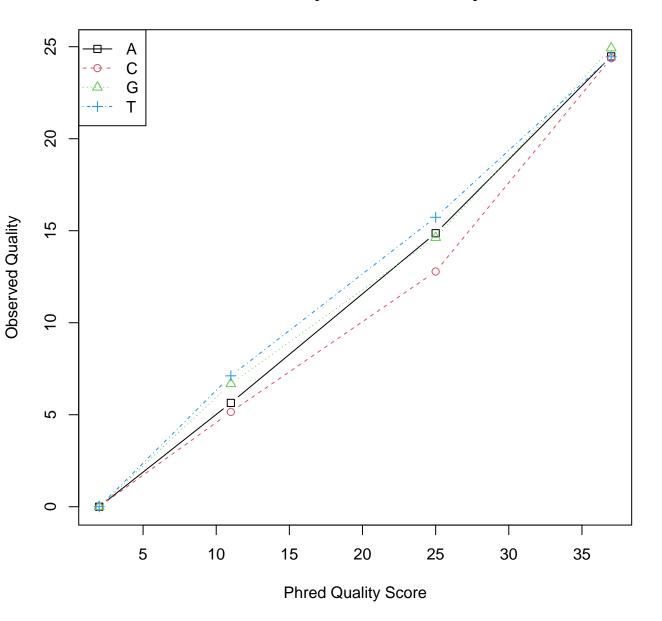


Read Start Position Distribution



Bin#

Observed Quality vs. Phred Quality Score



Alignment statistics

