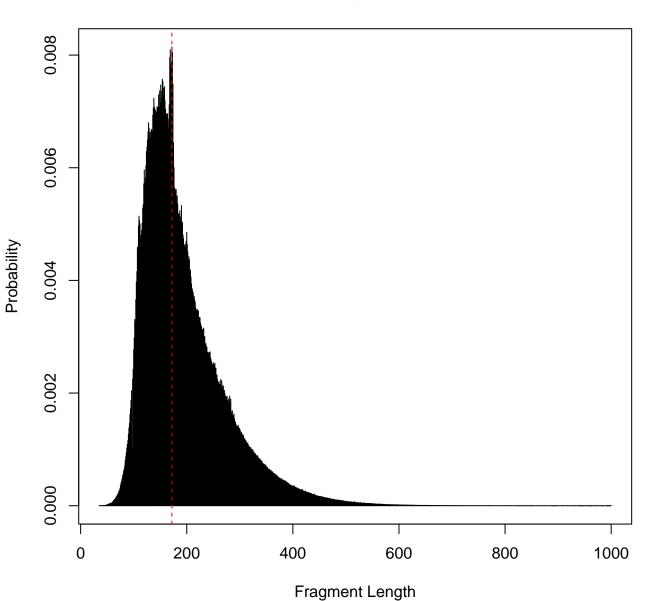
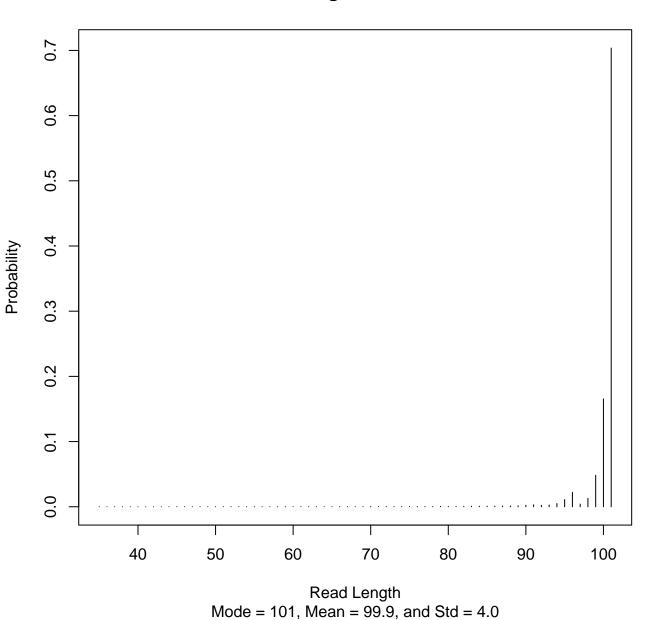
Fragment Length Distribution

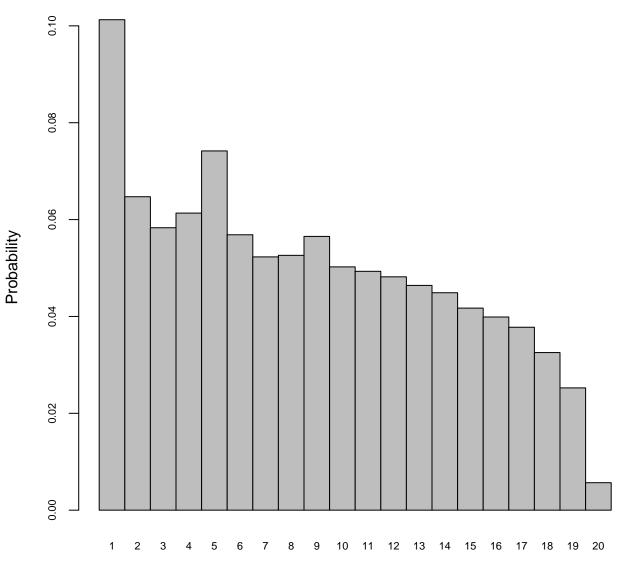


Mode = 172, Mean = 193.8, and Std = 77.7

Read Length Distribution

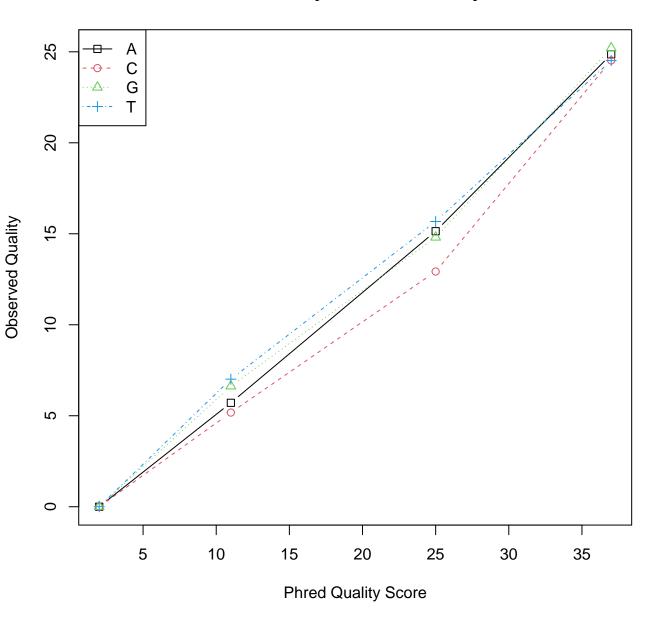


Read Start Position Distribution



Bin#

Observed Quality vs. Phred Quality Score



Alignment statistics

