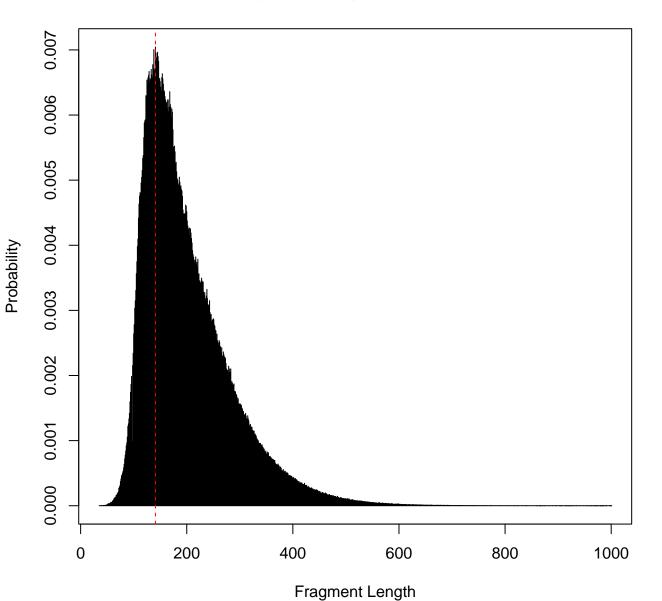
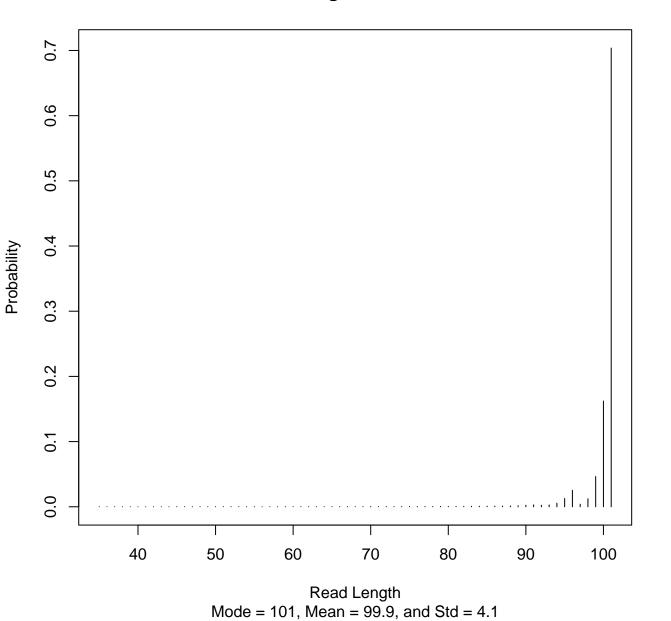
Fragment Length Distribution

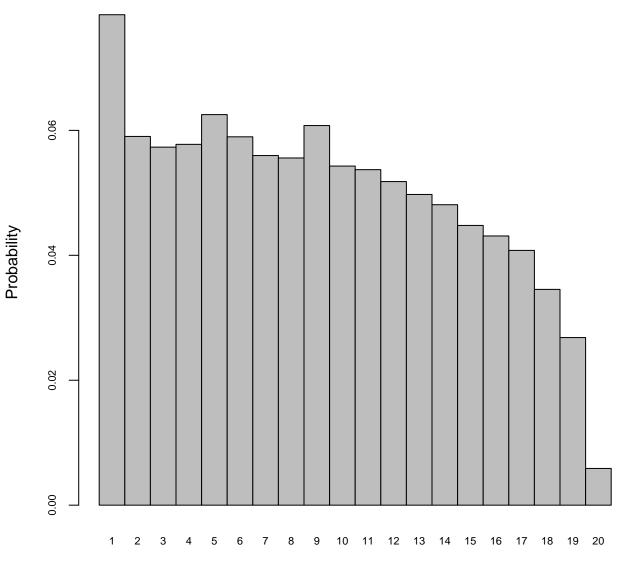


Mode = 141, Mean = 200.8, and Std = 83.4

Read Length Distribution

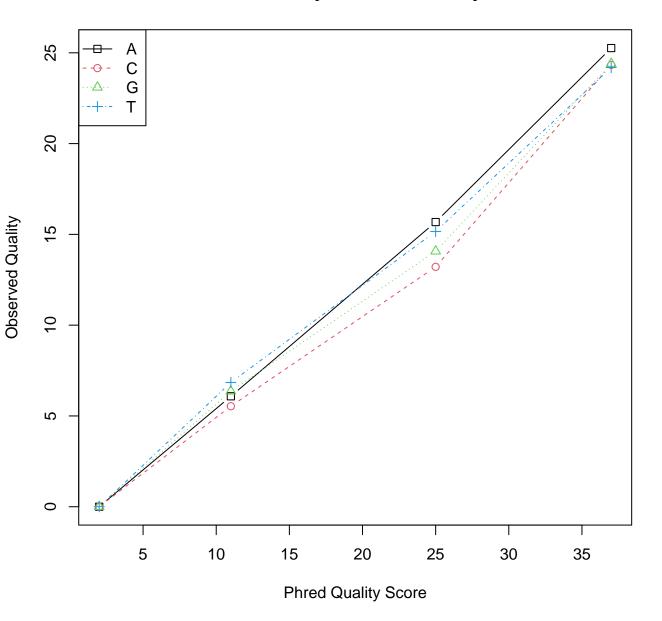


Read Start Position Distribution



Bin#

Observed Quality vs. Phred Quality Score



Alignment statistics

