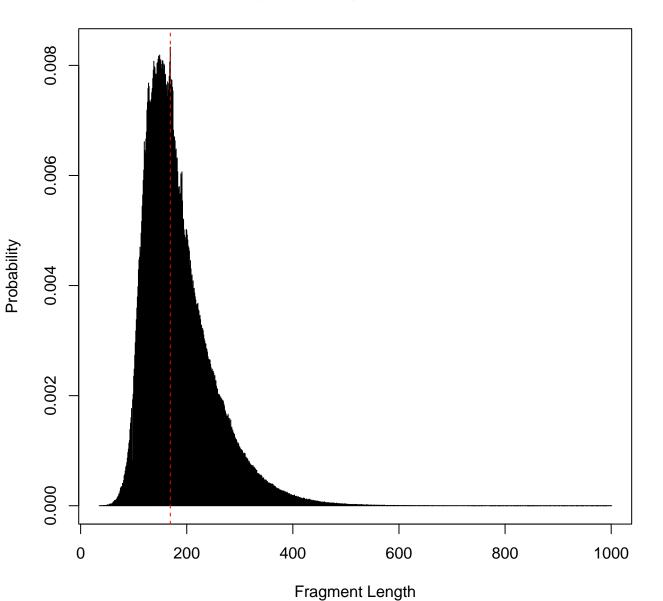
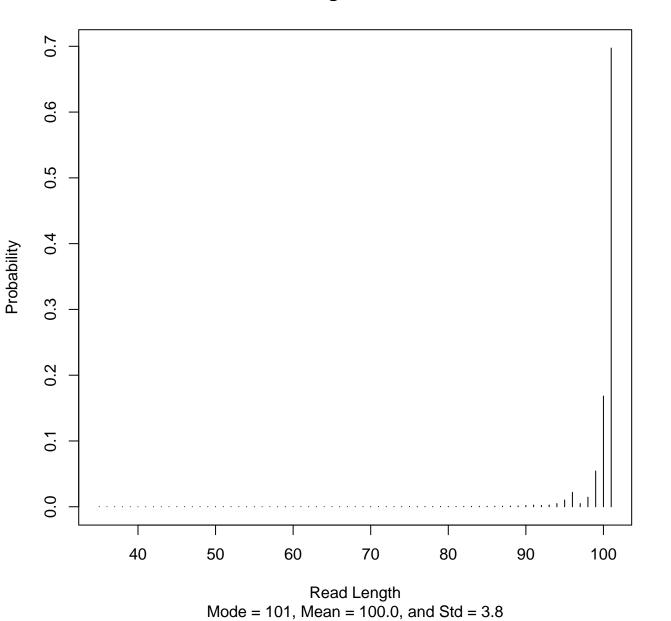
Fragment Length Distribution

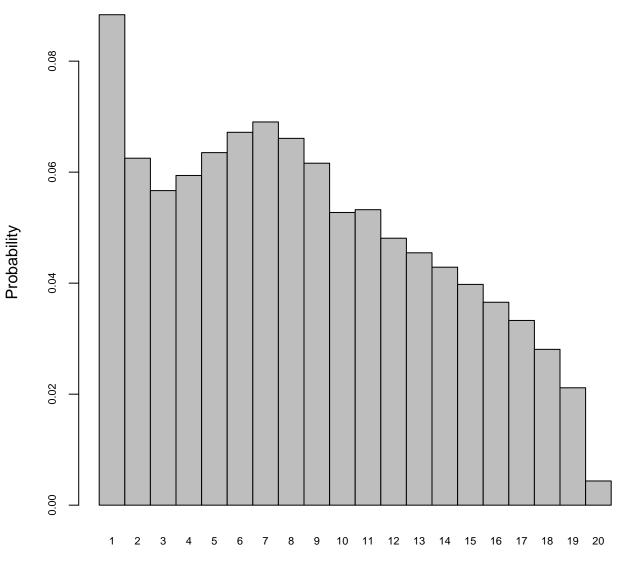


Mode = 169, Mean = 185.0, and Std = 65.8

Read Length Distribution

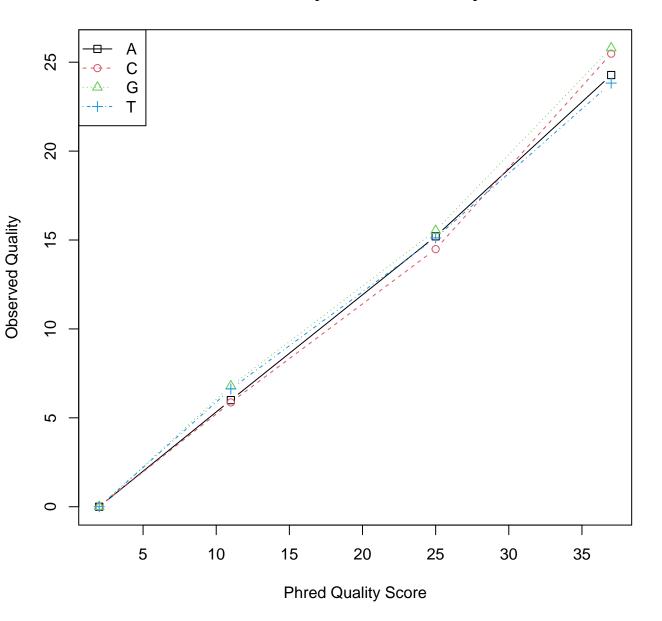


Read Start Position Distribution



Bin#

Observed Quality vs. Phred Quality Score



Alignment statistics

