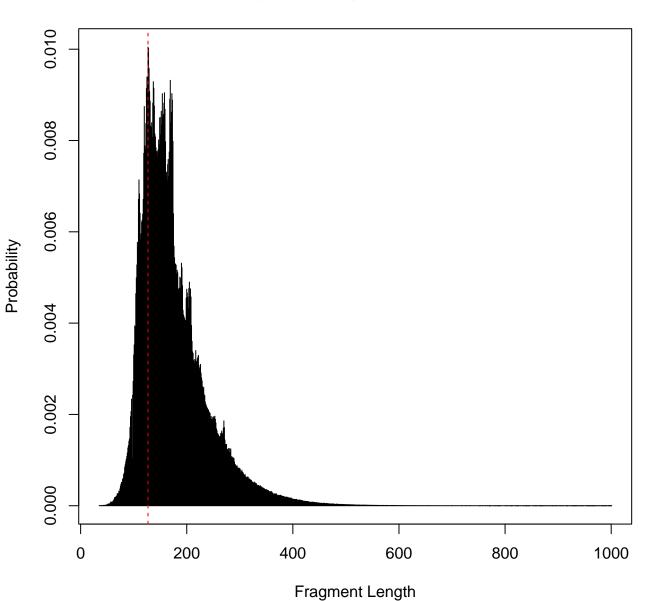
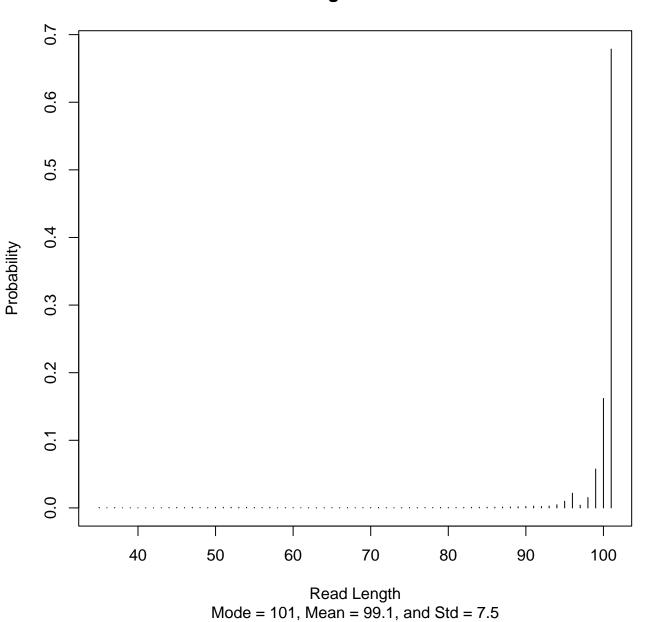
Fragment Length Distribution

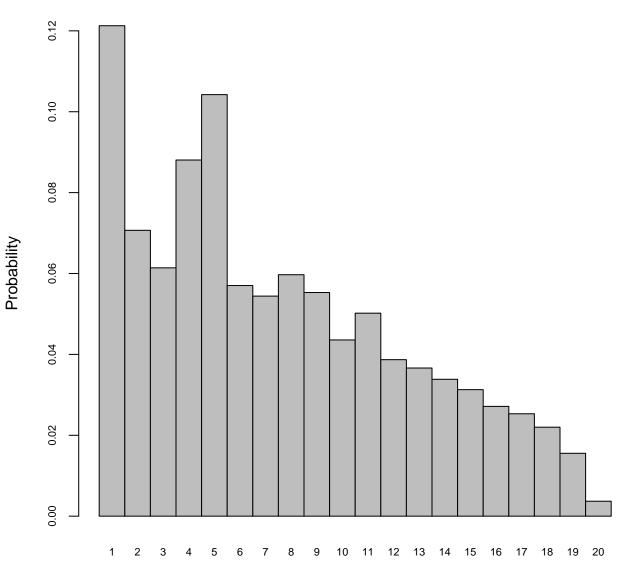


Mode = 127, Mean = 174.9, and Std = 62.8

Read Length Distribution

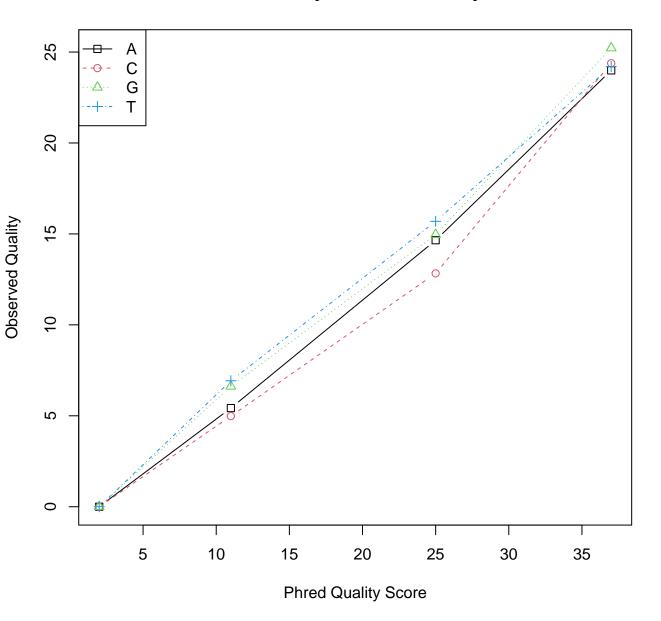


Read Start Position Distribution



Bin#

Observed Quality vs. Phred Quality Score



Alignment statistics

