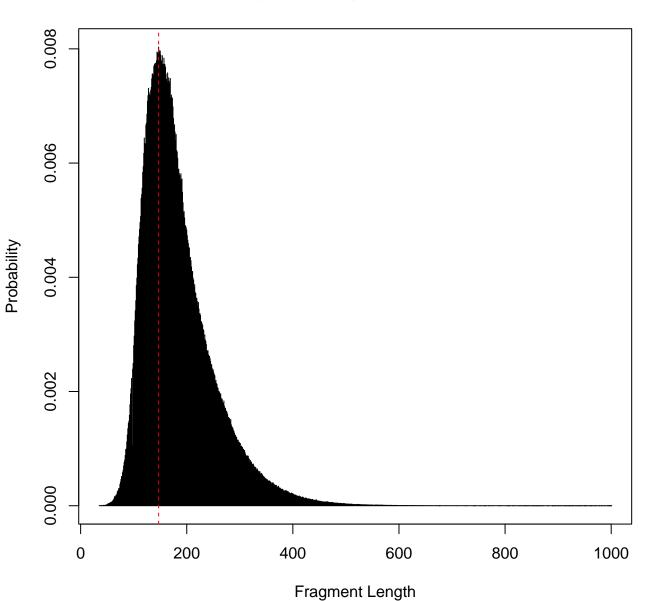
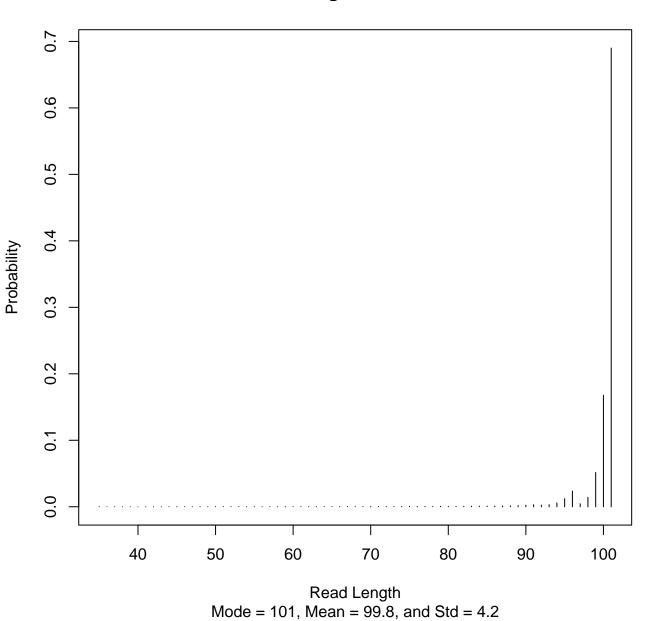
Fragment Length Distribution

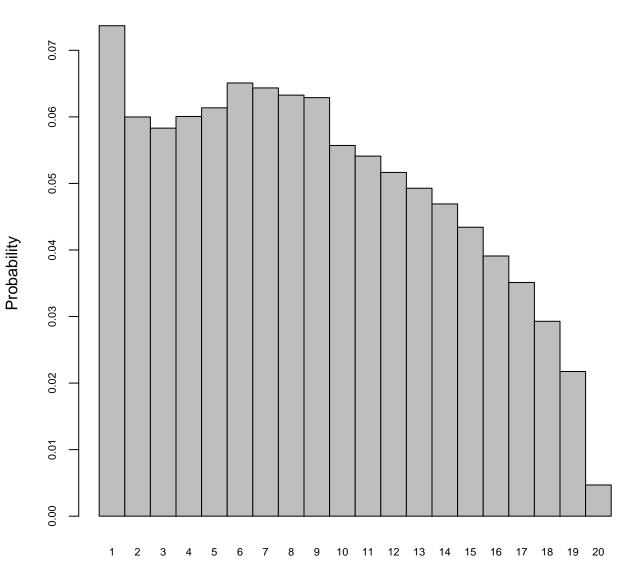


Mode = 147, Mean = 184.3, and Std = 67.3

Read Length Distribution

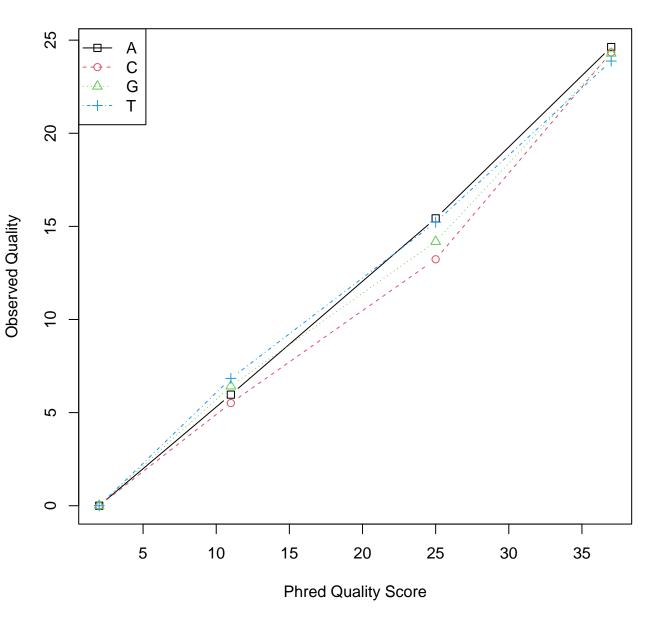


Read Start Position Distribution



Bin#

Observed Quality vs. Phred Quality Score



Alignment statistics

