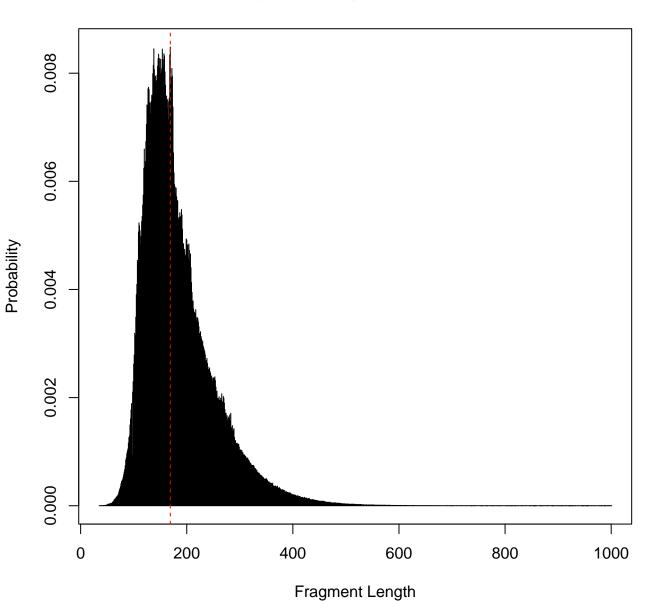
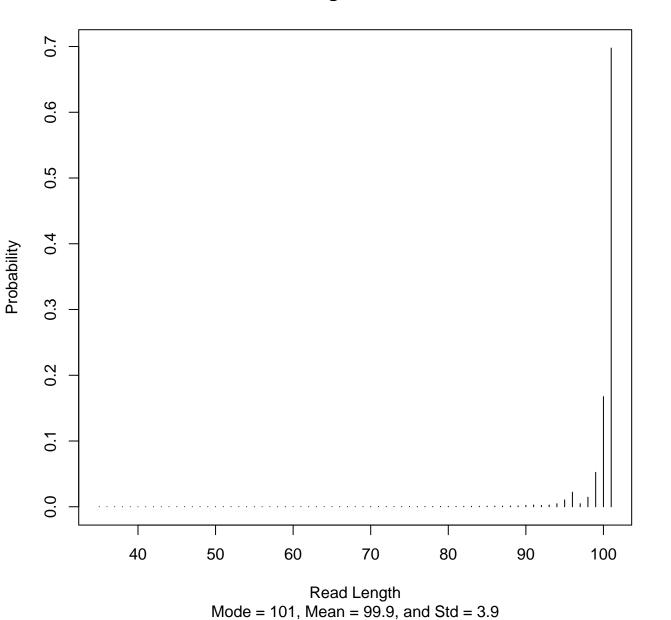
Fragment Length Distribution

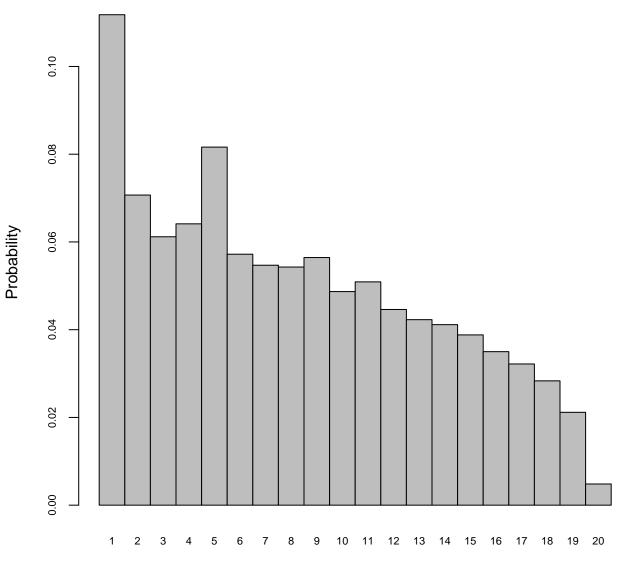


Mode = 169, Mean = 184.4, and Std = 67.5

Read Length Distribution

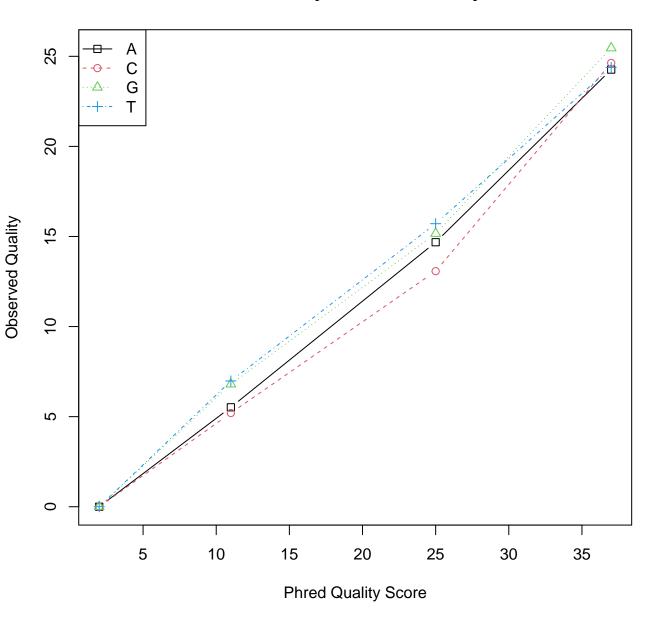


Read Start Position Distribution



Bin#

Observed Quality vs. Phred Quality Score



Alignment statistics

