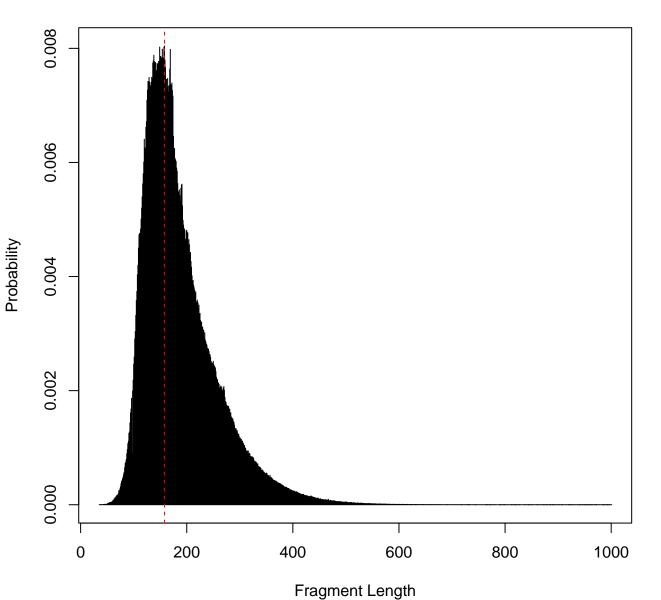
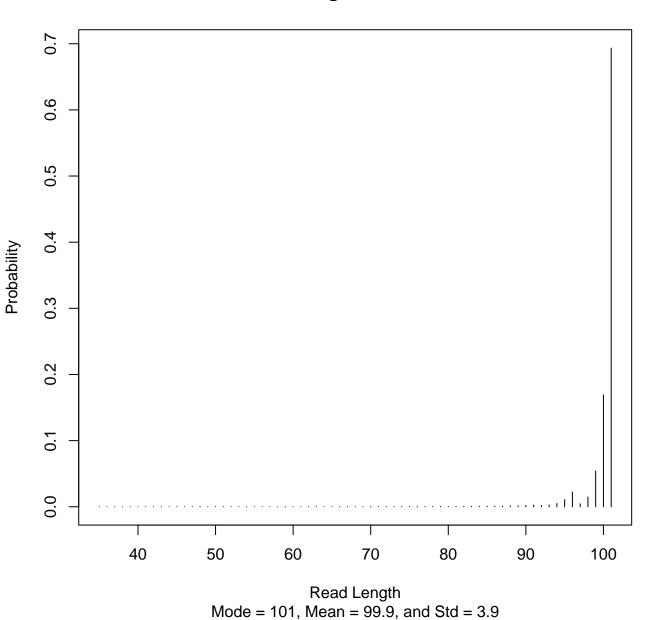
Fragment Length Distribution

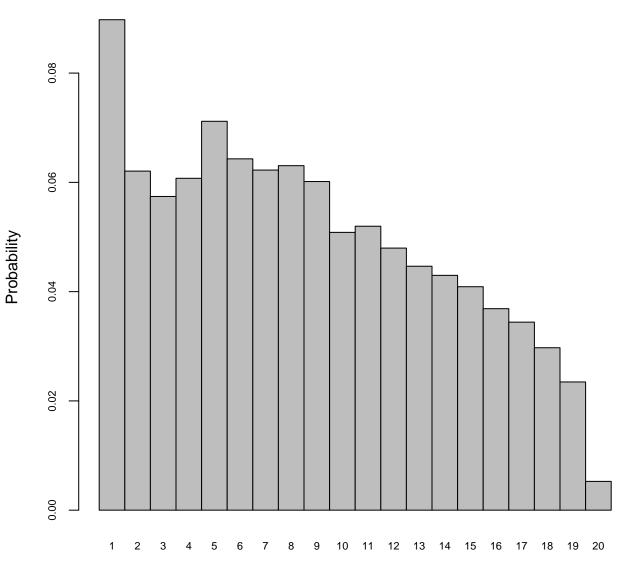


Mode = 158, Mean = 187.7, and Std = 70.0

Read Length Distribution

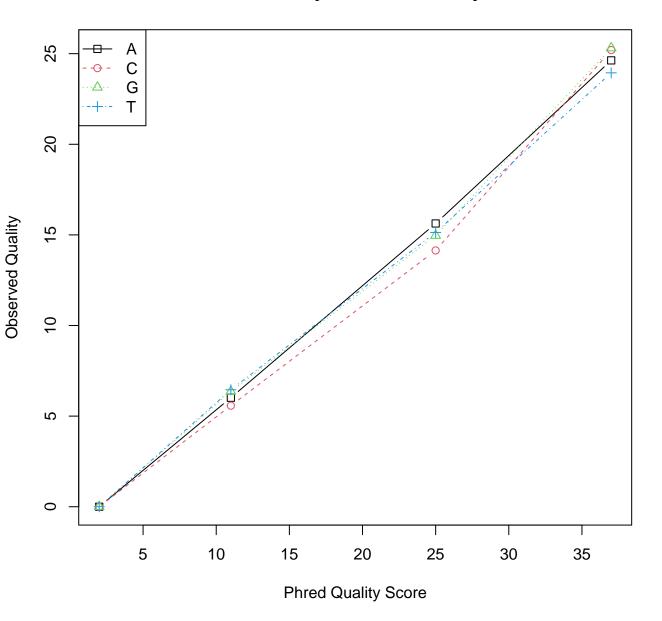


Read Start Position Distribution



Bin#

Observed Quality vs. Phred Quality Score



Alignment statistics

