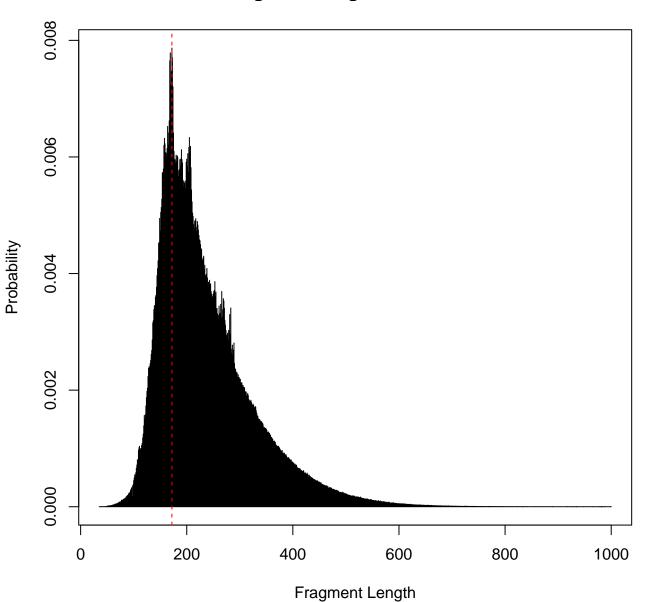
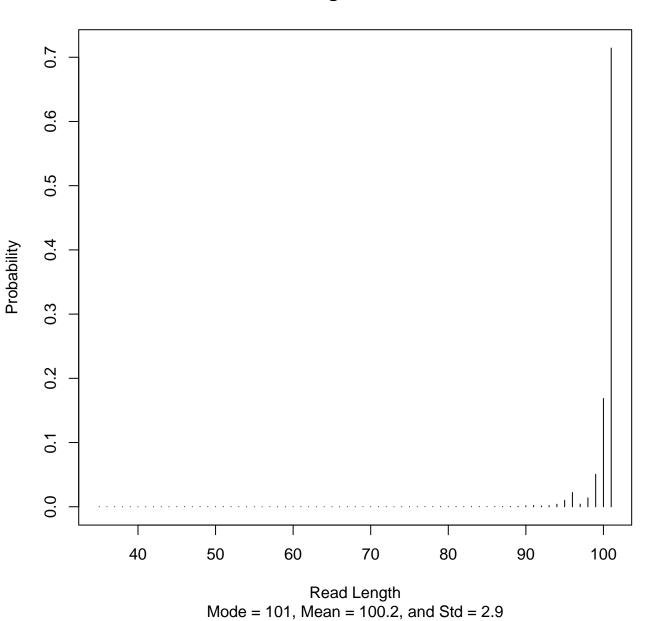
Fragment Length Distribution

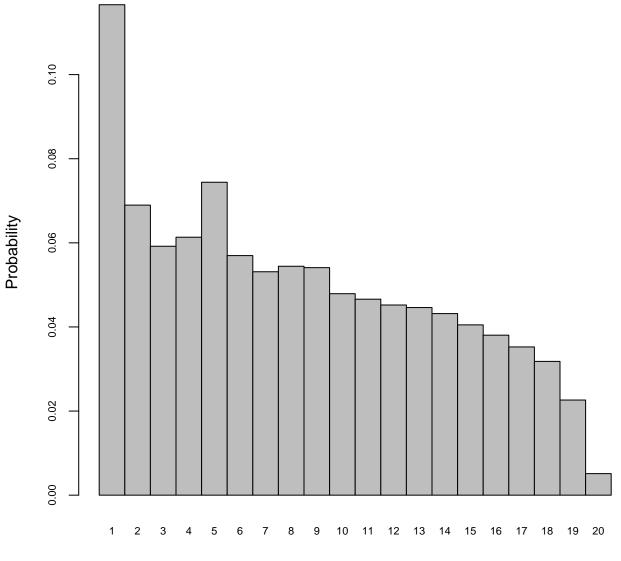


Mode = 172, Mean = 236.1, and Std = 89.5

Read Length Distribution

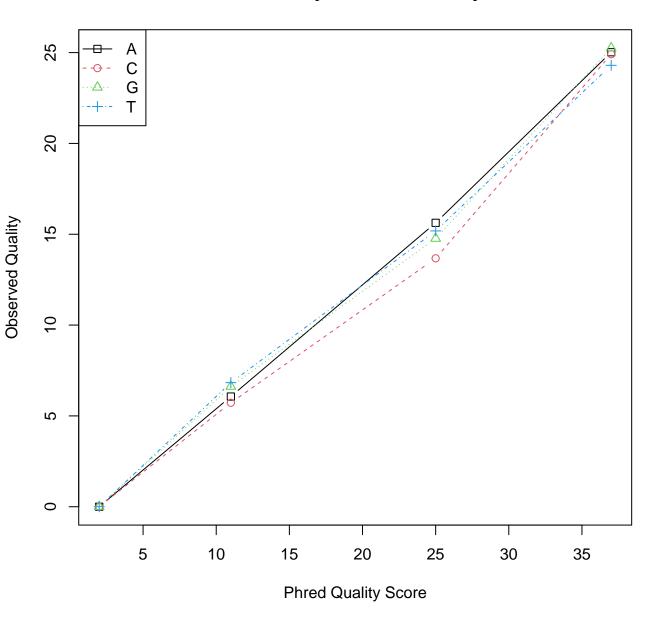


Read Start Position Distribution

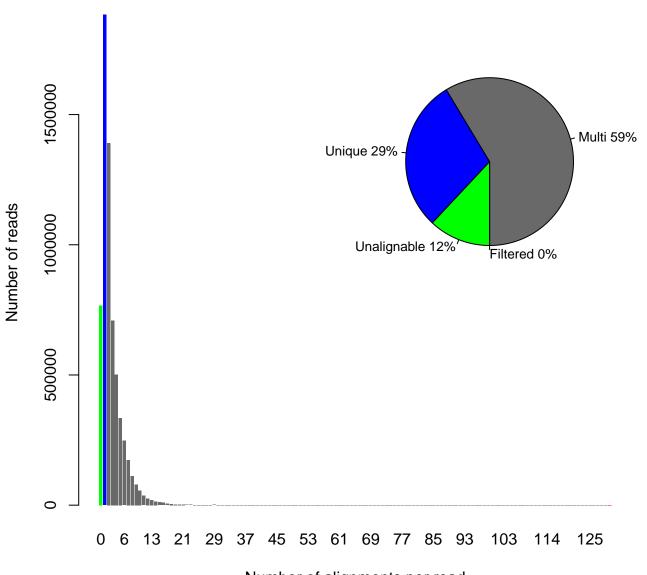


Bin#

Observed Quality vs. Phred Quality Score



Alignment statistics



Number of alignments per read