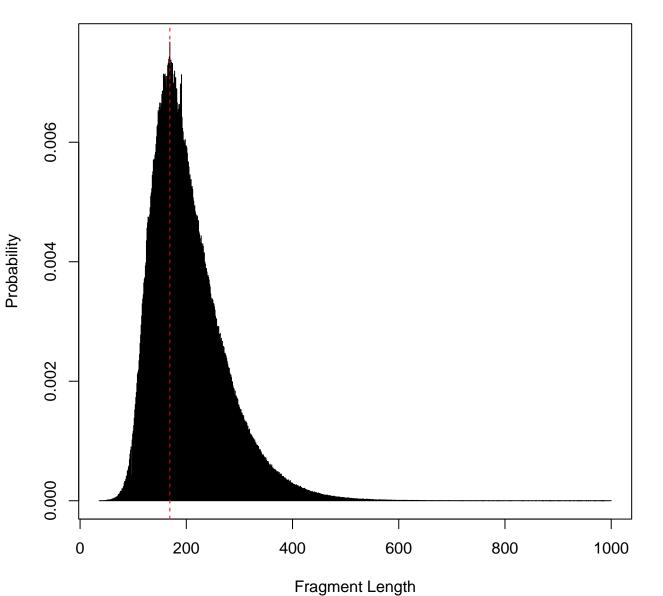
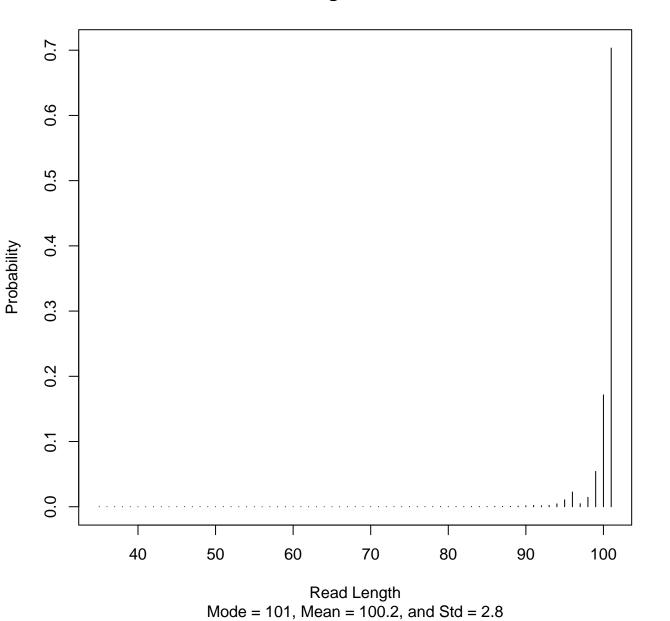
Fragment Length Distribution

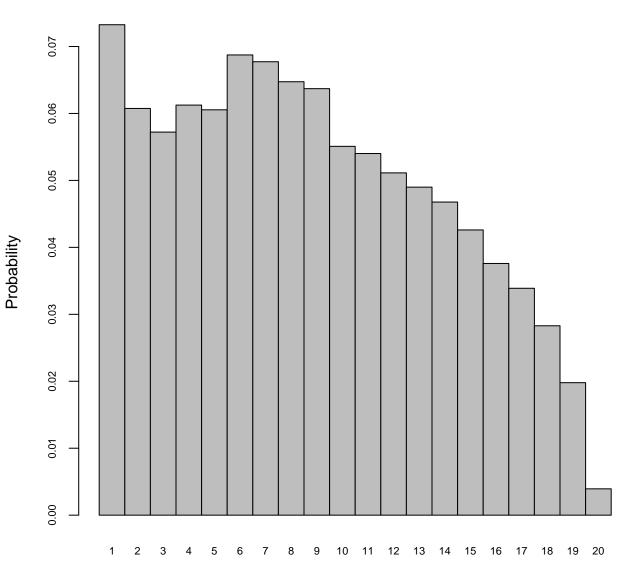


Mode = 169, Mean = 204.3, and Std = 69.9

Read Length Distribution

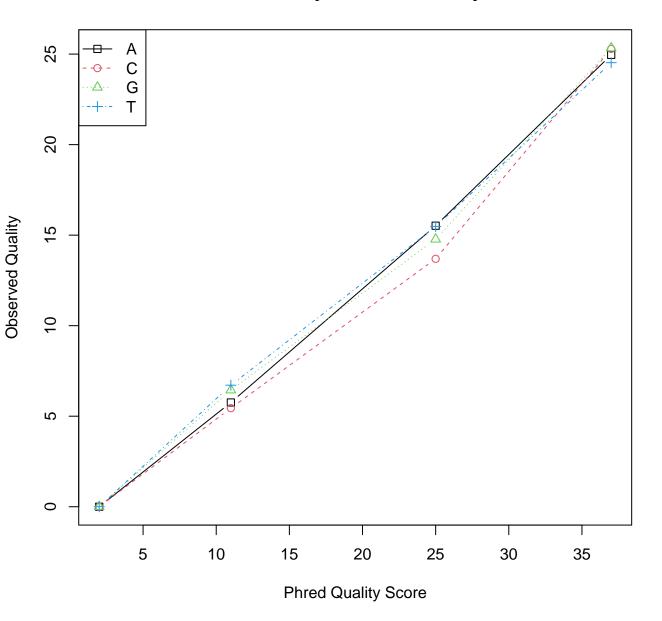


Read Start Position Distribution

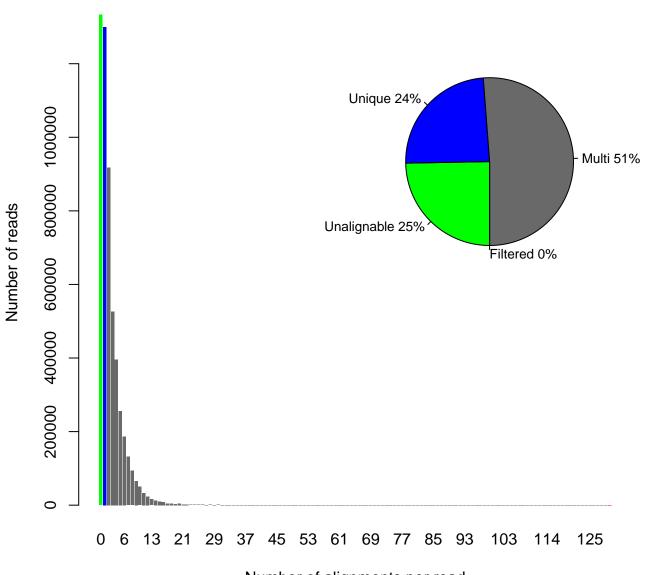


Bin#

Observed Quality vs. Phred Quality Score



Alignment statistics



Number of alignments per read