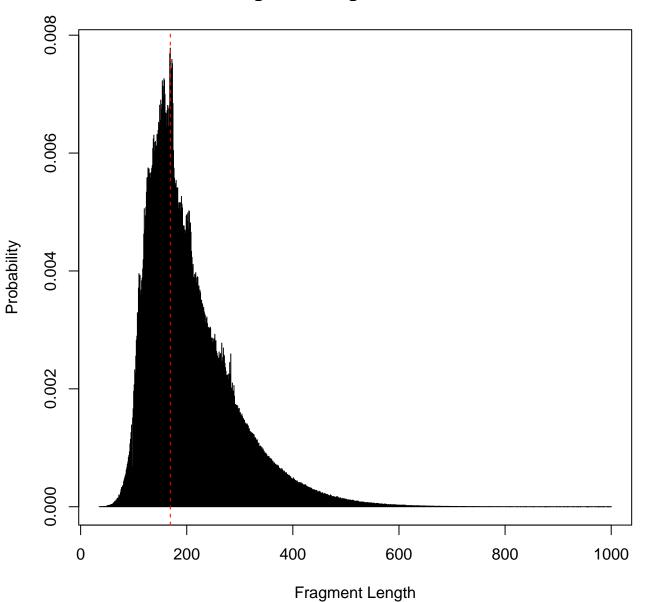
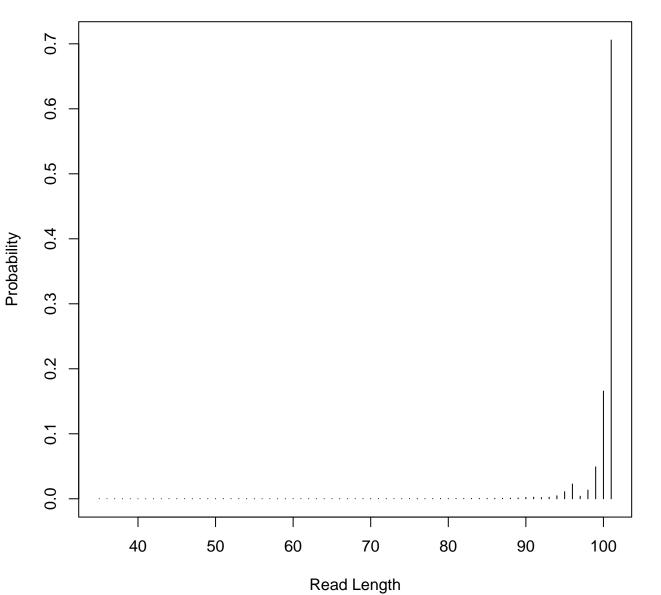
Fragment Length Distribution



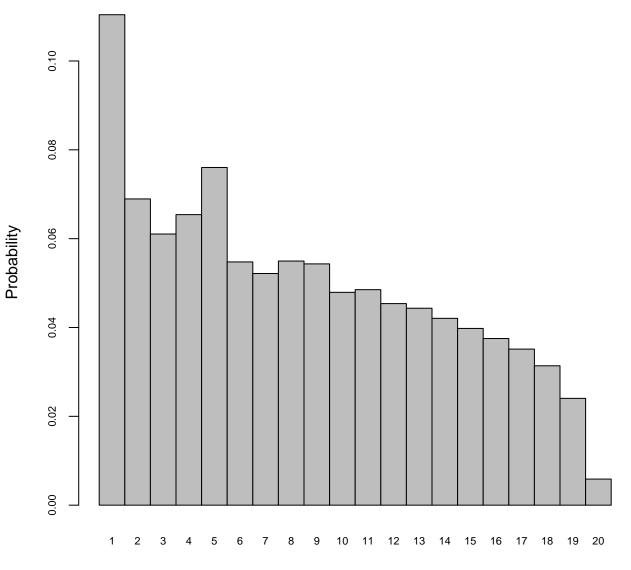
Mode = 169, Mean = 207.1, and Std = 84.5

Read Length Distribution



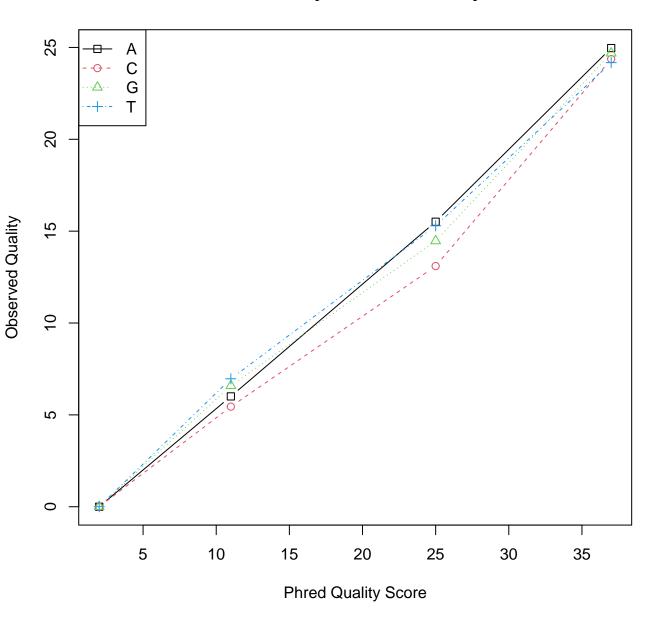
Mode = 101, Mean = 100.0, and Std = 4.2

Read Start Position Distribution

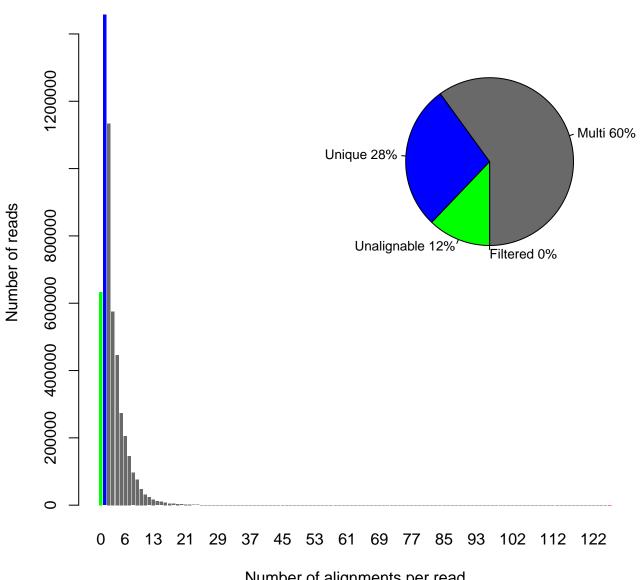


Bin#

Observed Quality vs. Phred Quality Score







Number of alignments per read