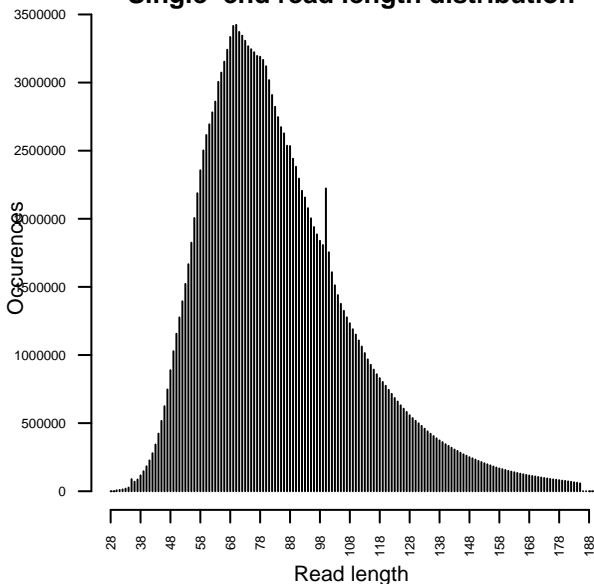
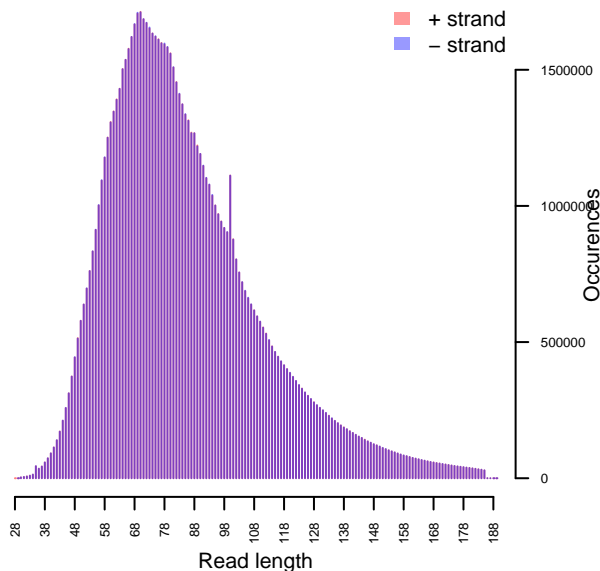


nochr-Chan_hg19_merged_duprem30_realigned

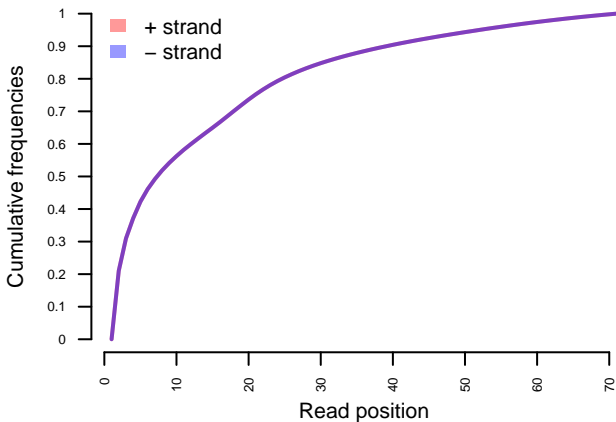
Single-end read length distribution



Single-end read length per strand



C>T



G>A

