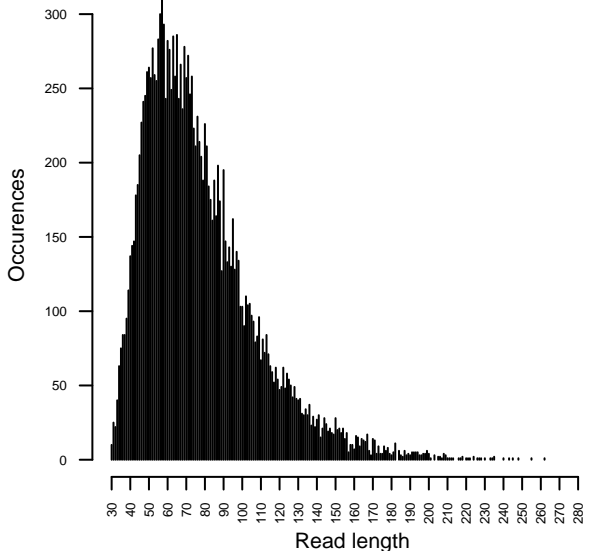
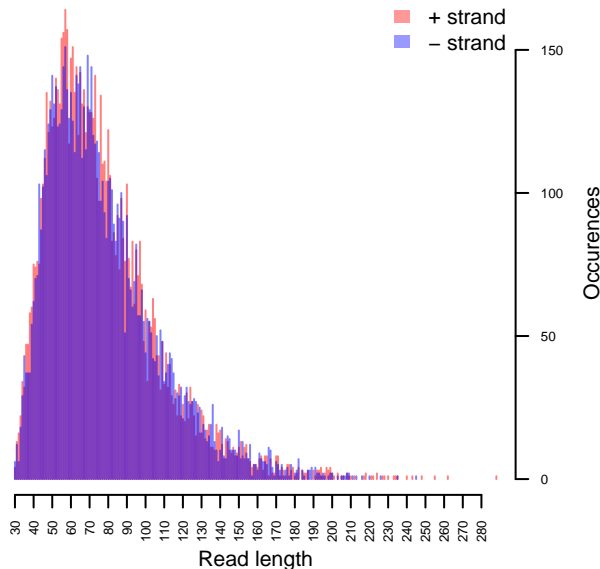


AD-372-H.trimmed.merged.mapped.q30.bwa.sort.rmdup.uniq

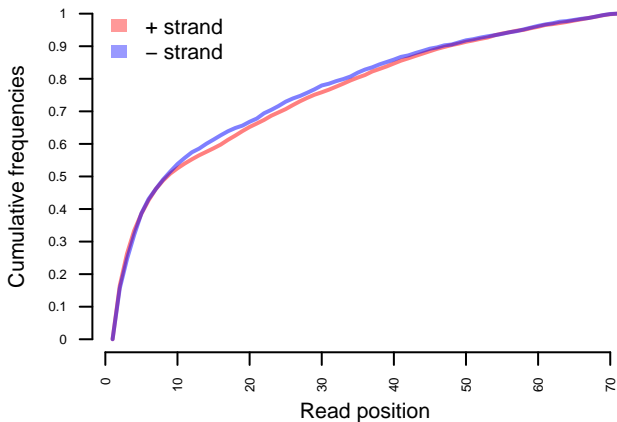
Single-end read length distribution



Single-end read length per strand



C>T



G>A

