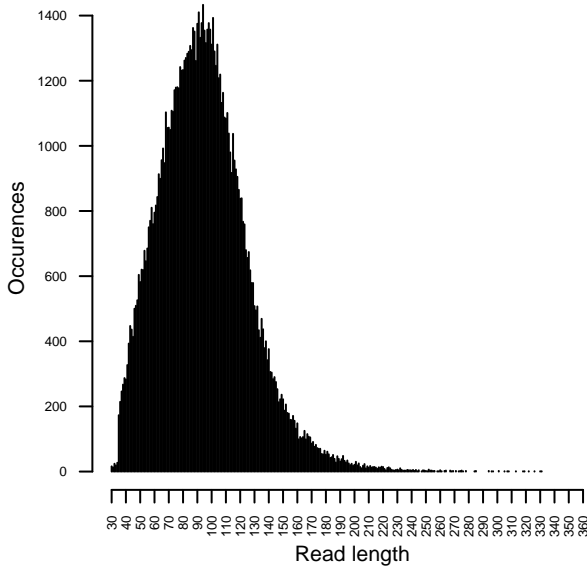
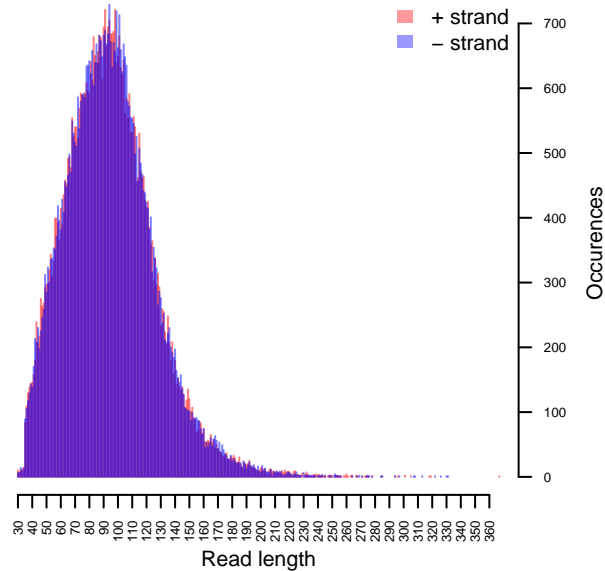


Nodule1_S1

Single-end read length distribution



Single-end read length per strand



C>T

G>A

