plot

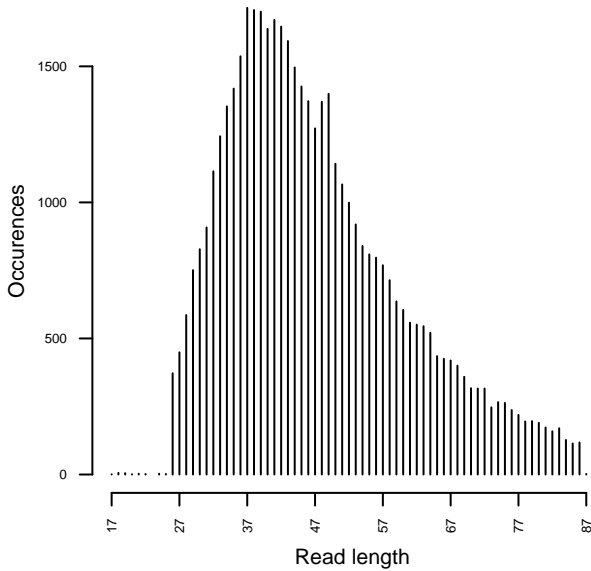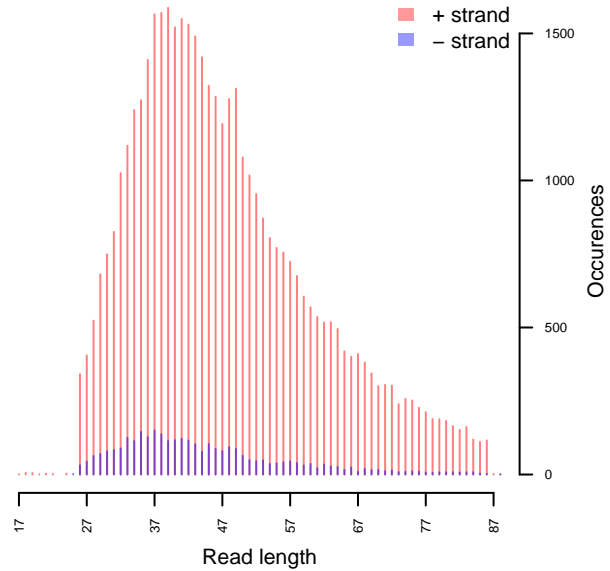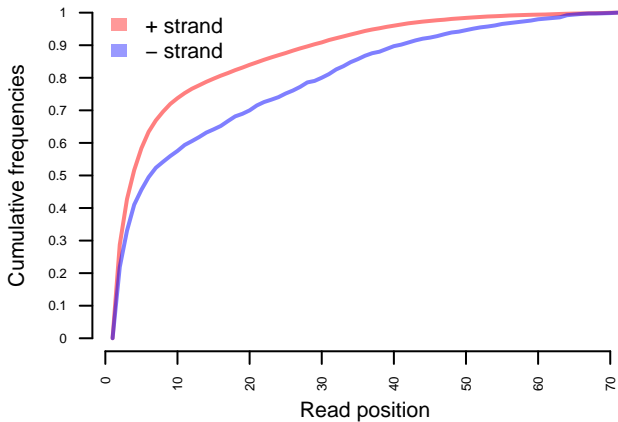**Single−end read length distribution**

Single−end read length per strand

C>T

G>A