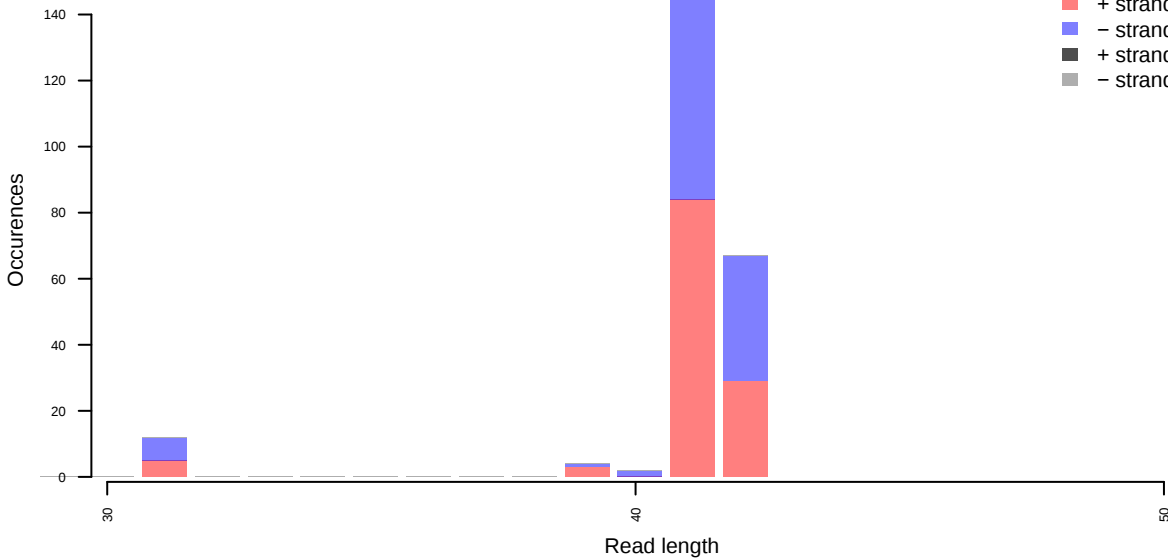


2-AP_24000093367.1

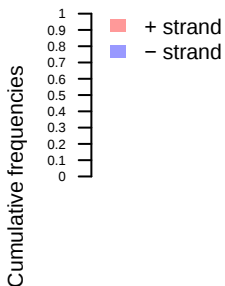
Length distribution

- + strand (SE)
- strand (SE)
- + strand (PE)
- strand (PE)

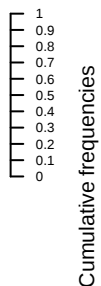


C>T

G>A



- + strand
- strand



Read position

Read position