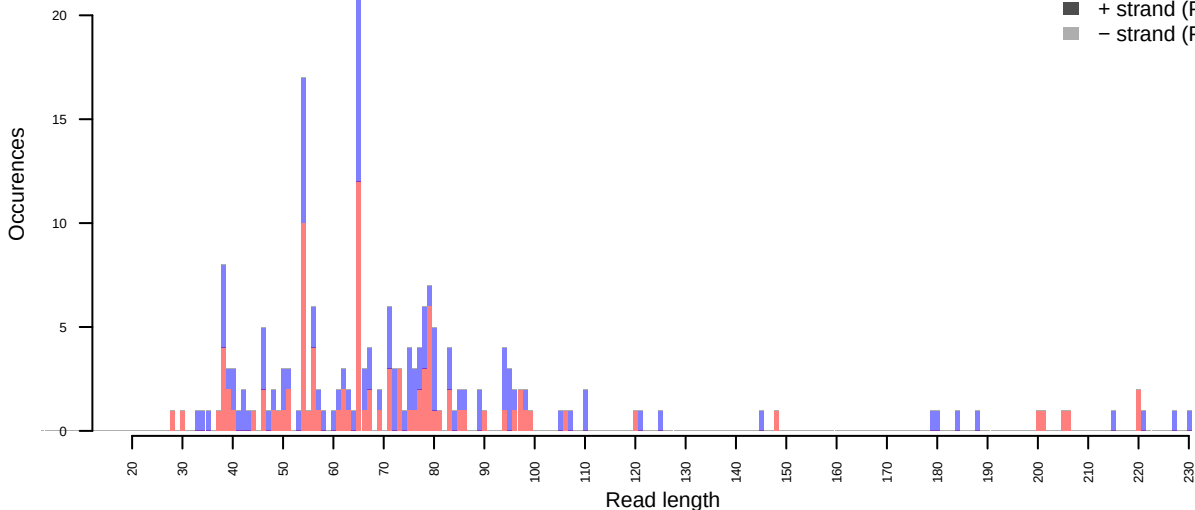


2-AP_183300232499.1

Length distribution

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



C>T

G>A

