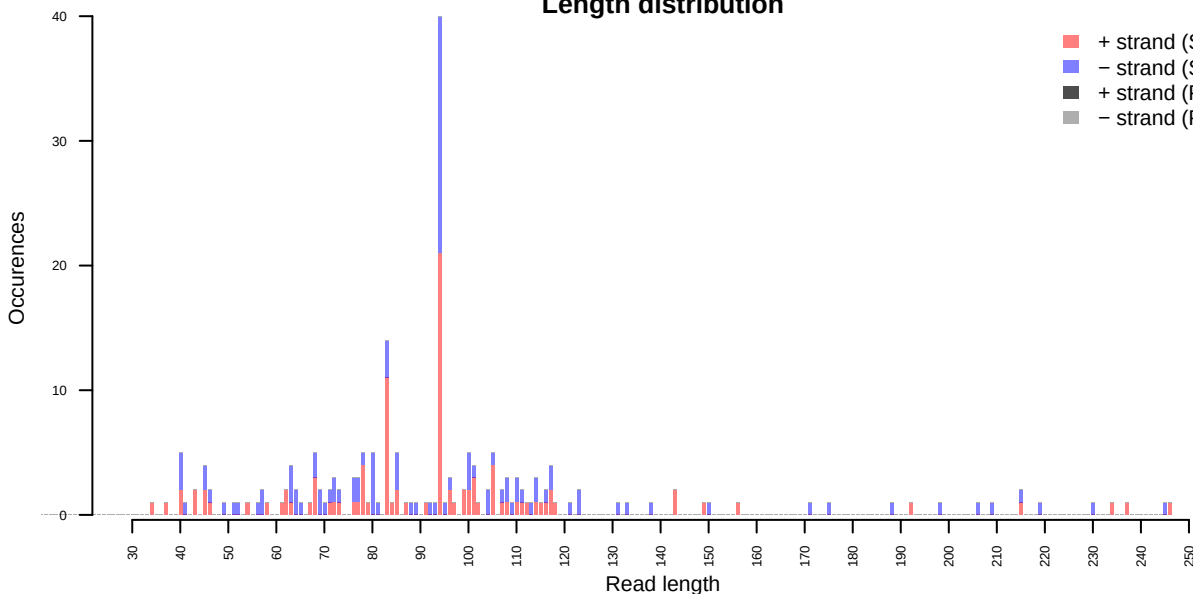


2-AP_32402349385.1

Length distribution

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



C>T

G>A

