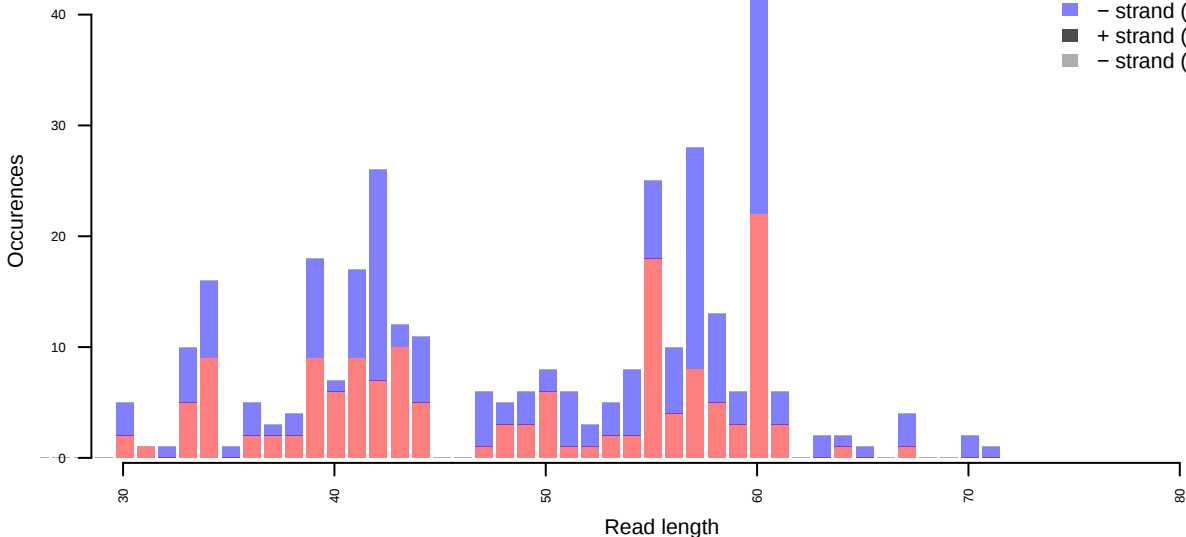


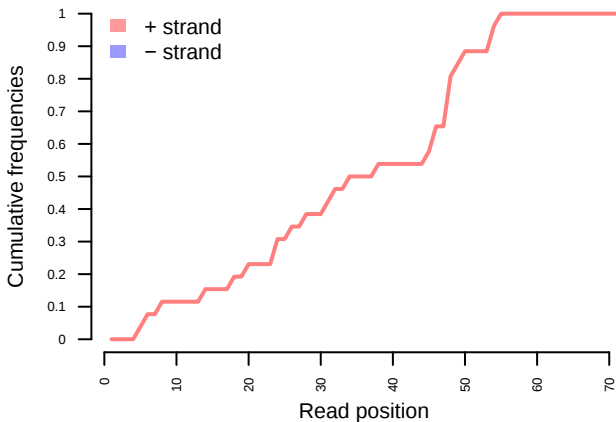
2-AP_85100093986.1

Length distribution

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



C>T



G>A

