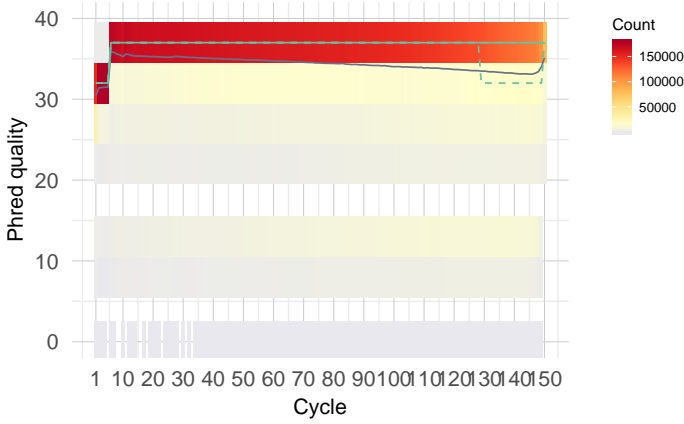
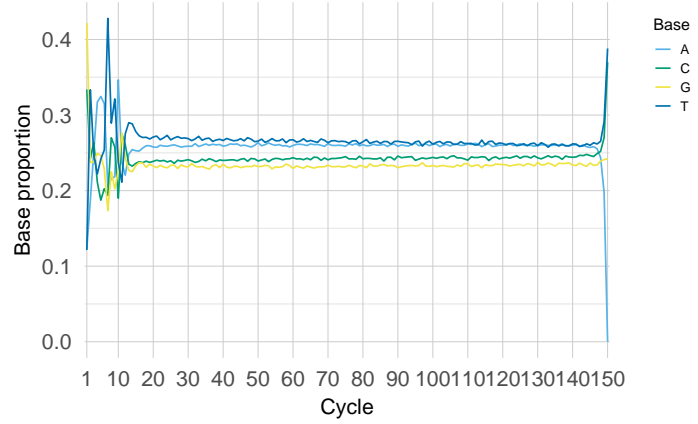


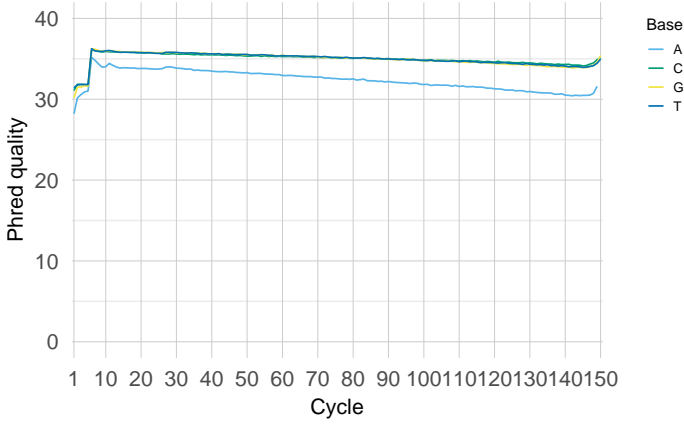
SRR1804235: Per cycle base quality



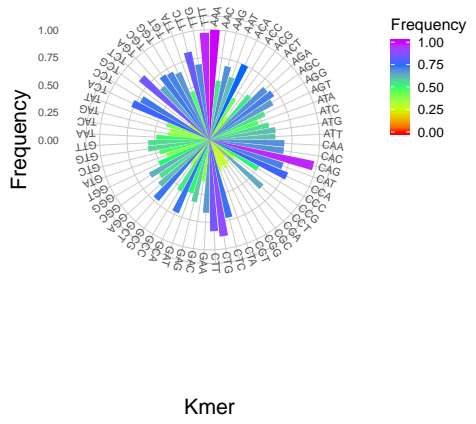
SRR1804235: Per cycle base frequency



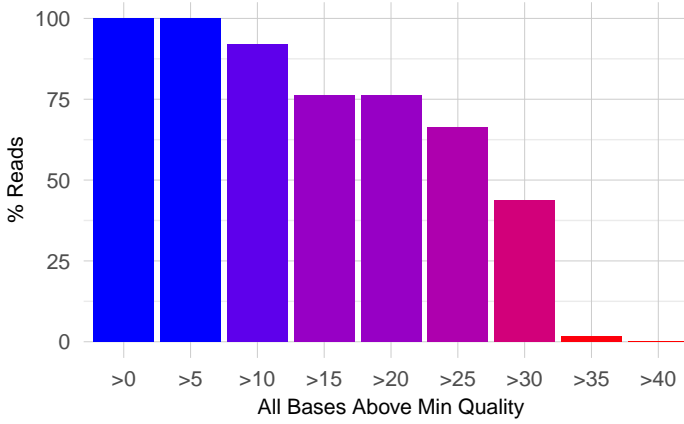
SRR1804235: Per cycle average base quality



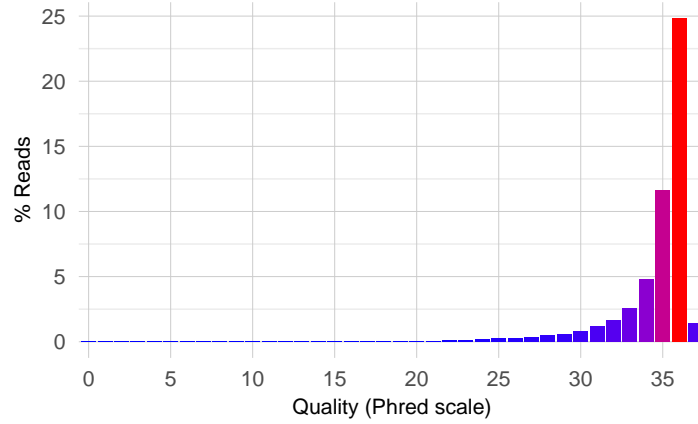
SRR1804235: Trinucleotide frequency



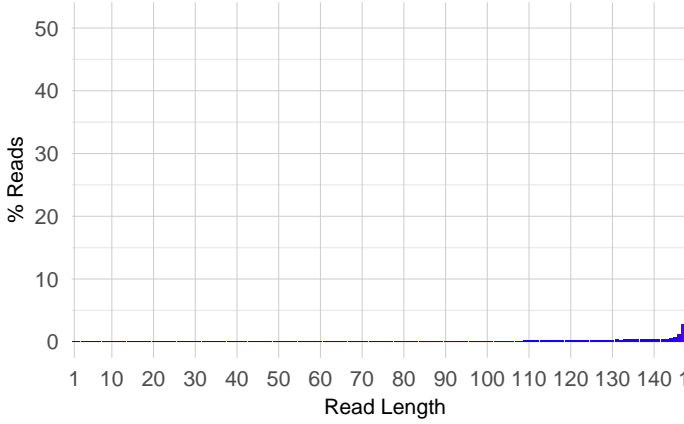
SRR1804235: Phred score quality cutoffs



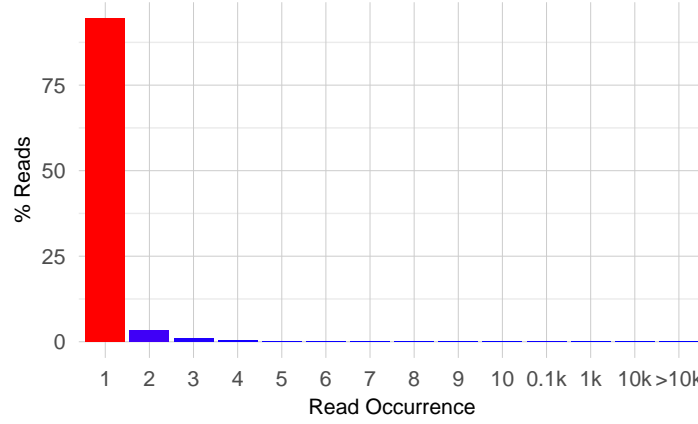
SRR1804235 : read quality for 191,680 of 3,833,604 reads



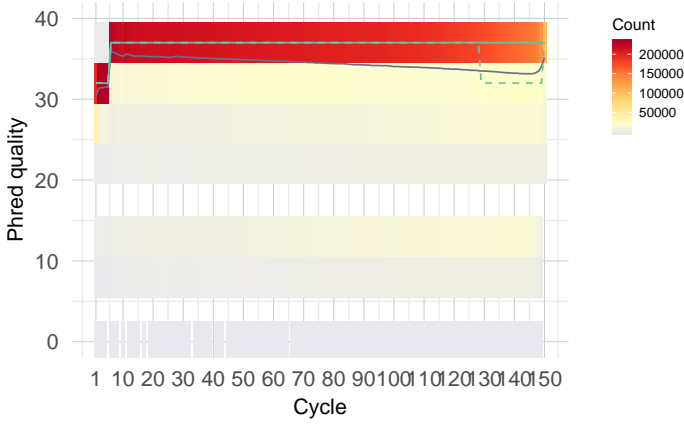
SRR1804235: Read length distribution



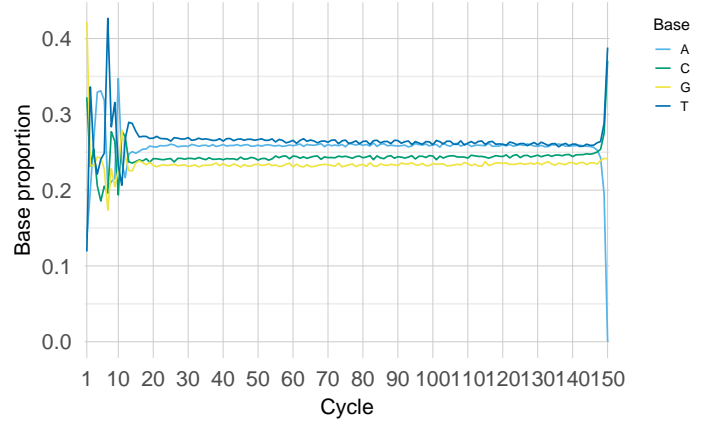
SRR1804235: Read occurrence distribution



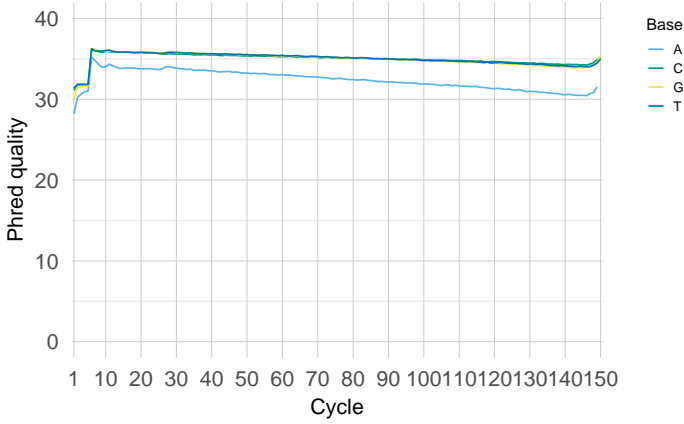
SRR1804236: Per cycle base quality



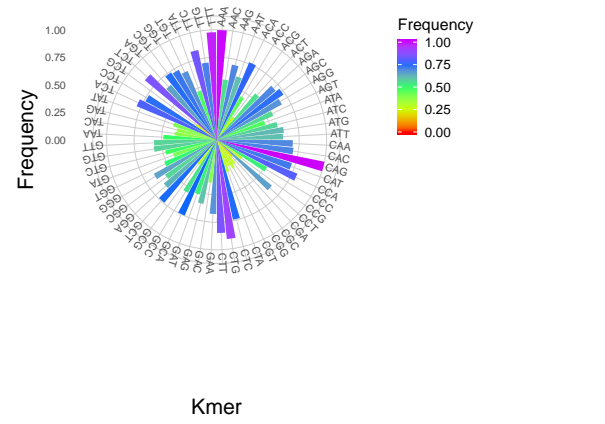
SRR1804236: Per cycle base frequency



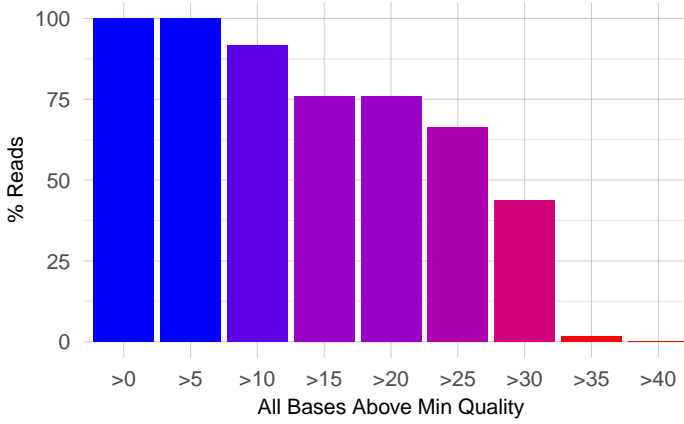
SRR1804236: Per cycle average base quality



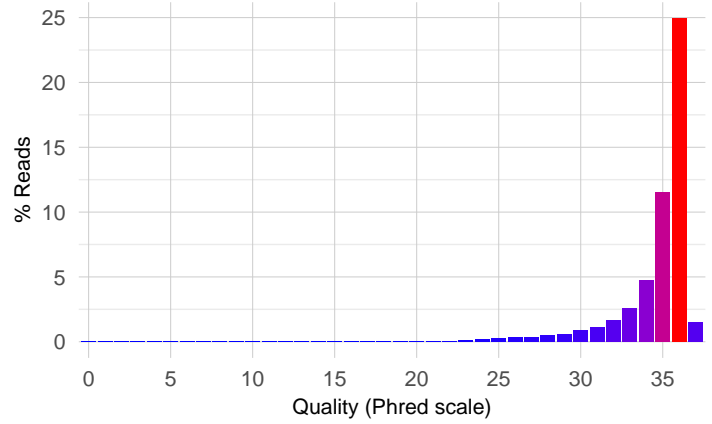
SRR1804236: Trinucleotide frequency



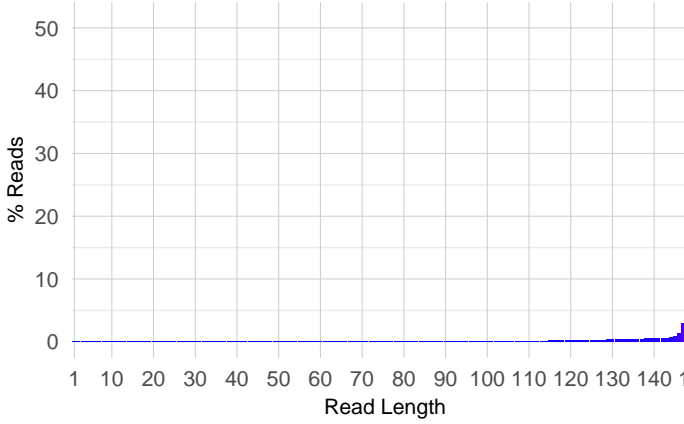
SRR1804236: Phred score quality cutoffs



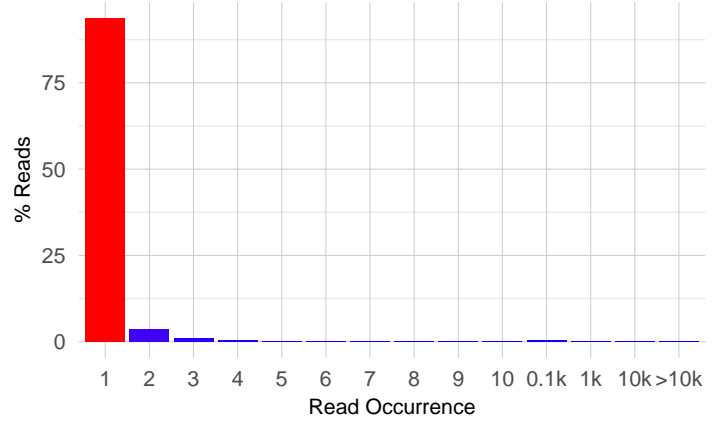
SRR1804236 : read quality for 247,937 of 4,958,739 reads

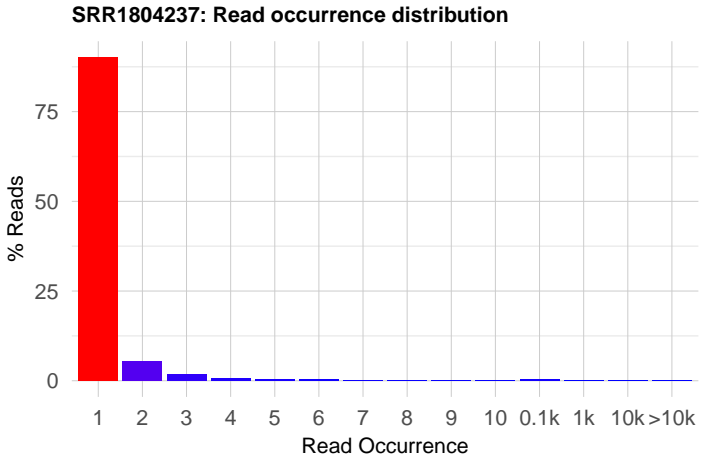
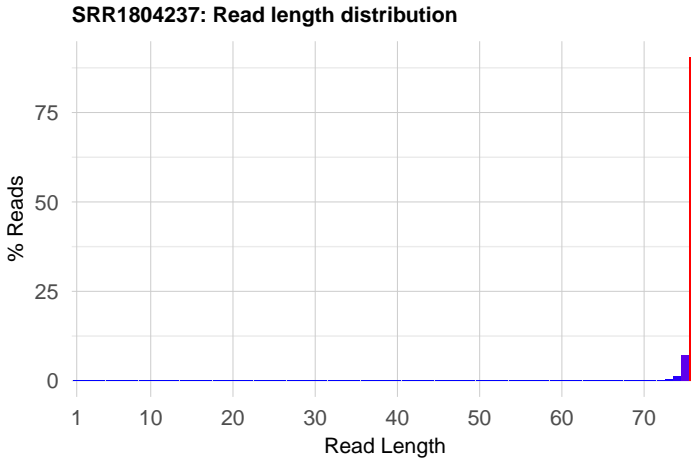
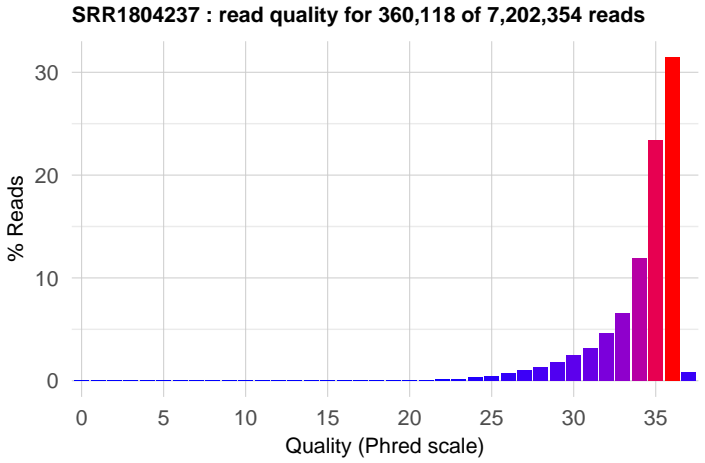
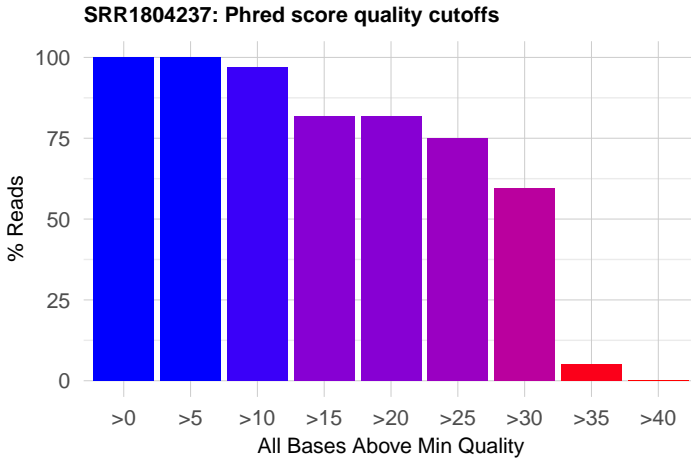
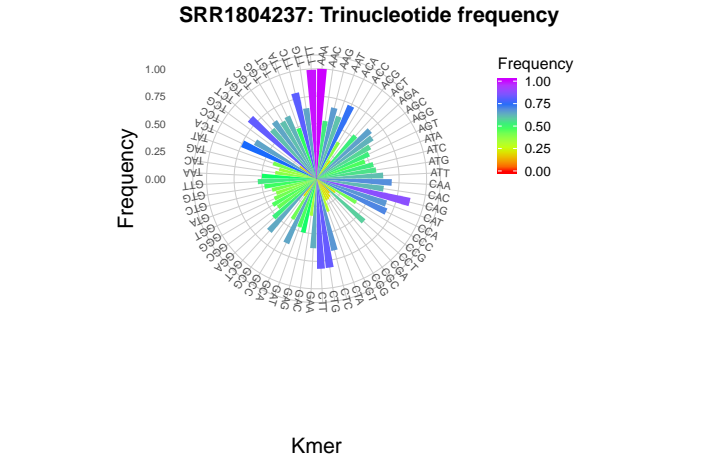
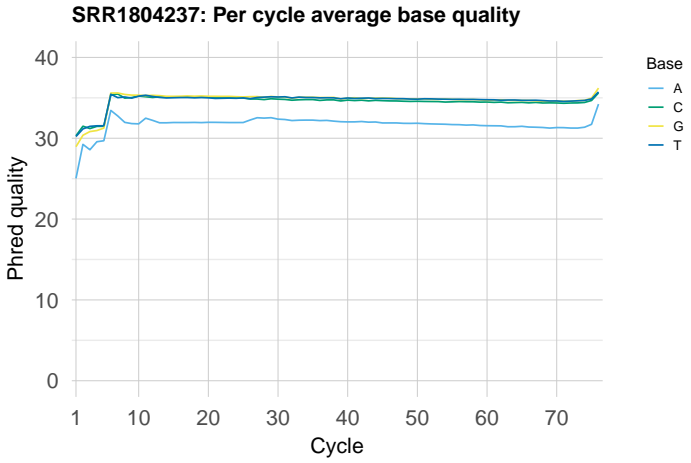
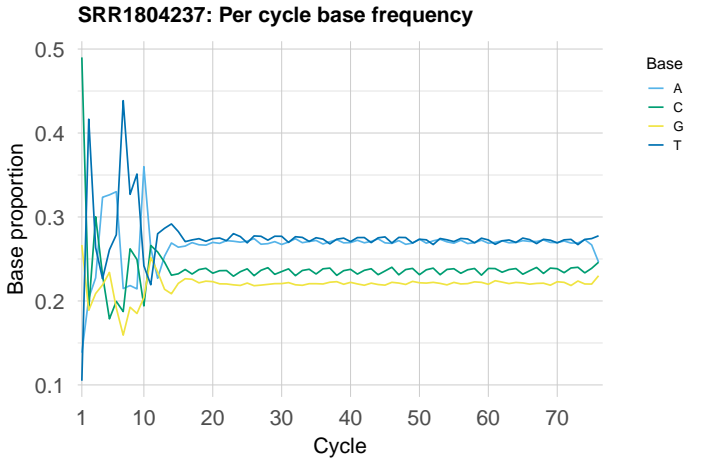
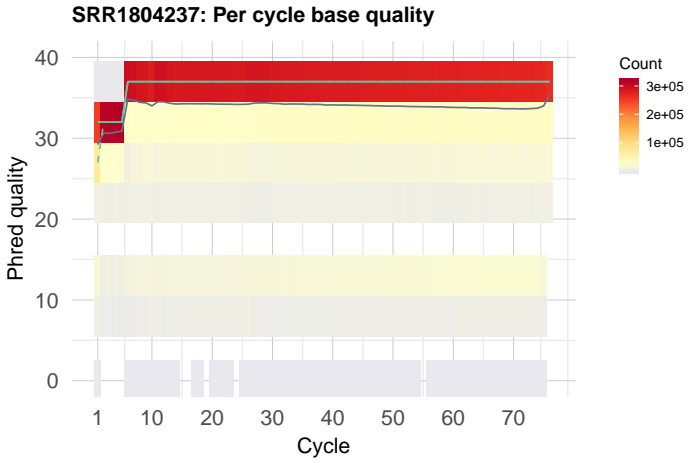


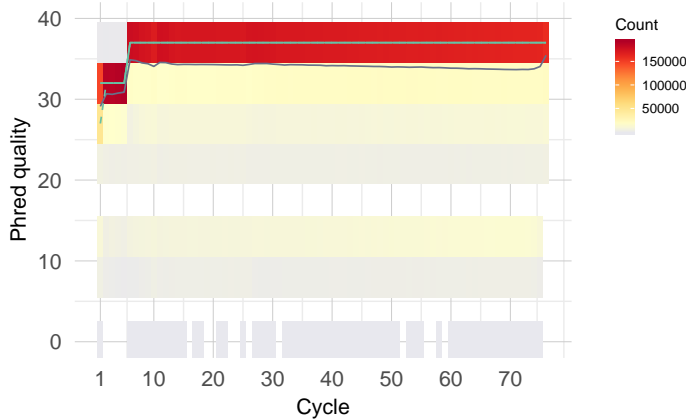
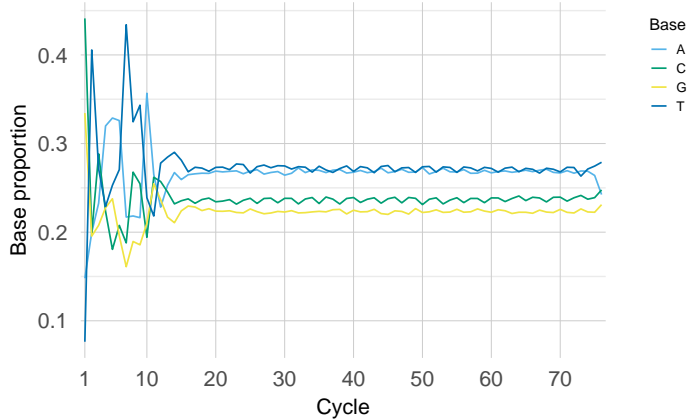
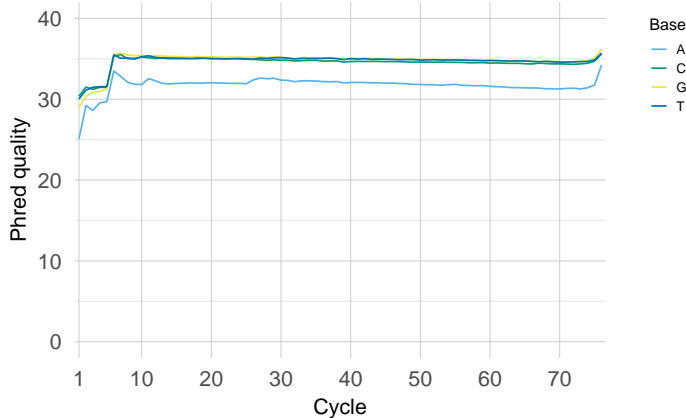
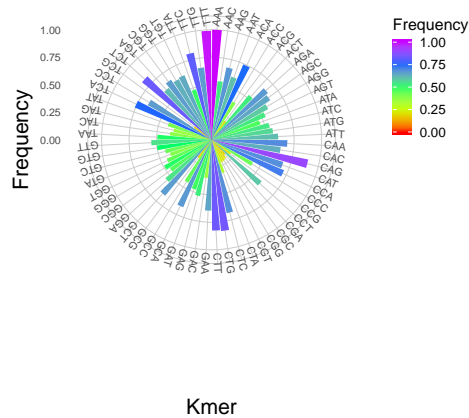
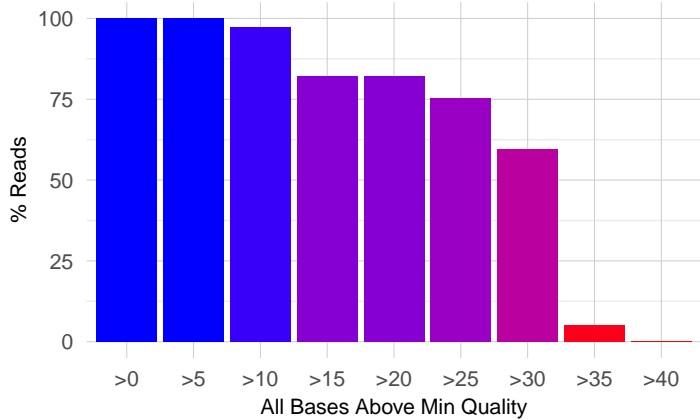
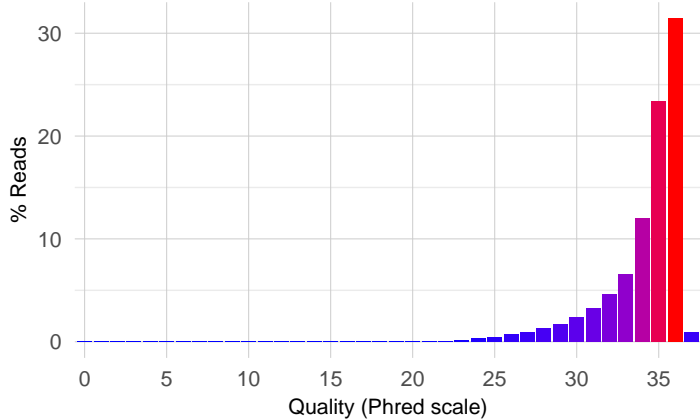
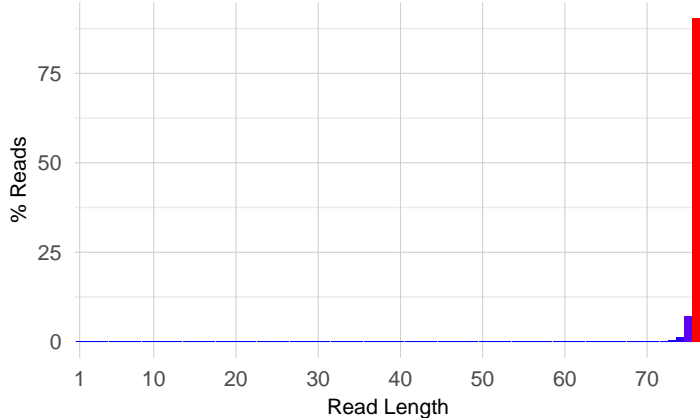
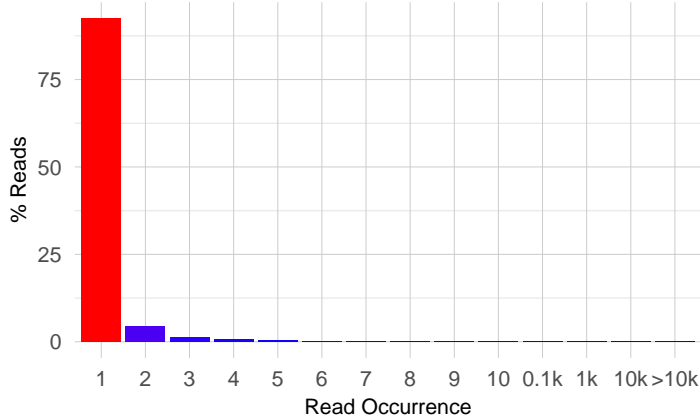
SRR1804236: Read length distribution

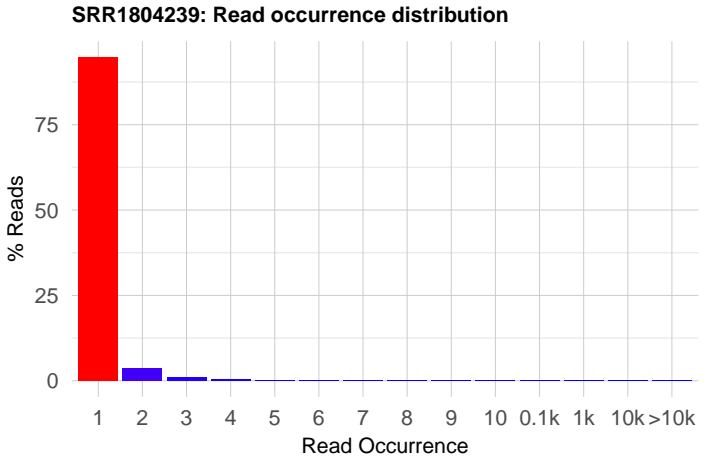
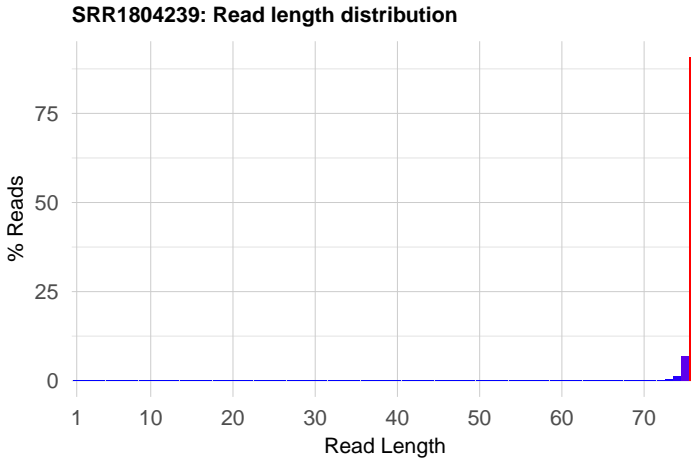
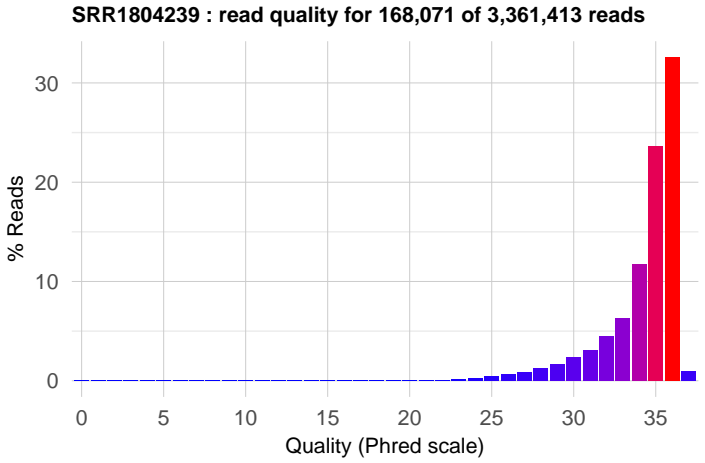
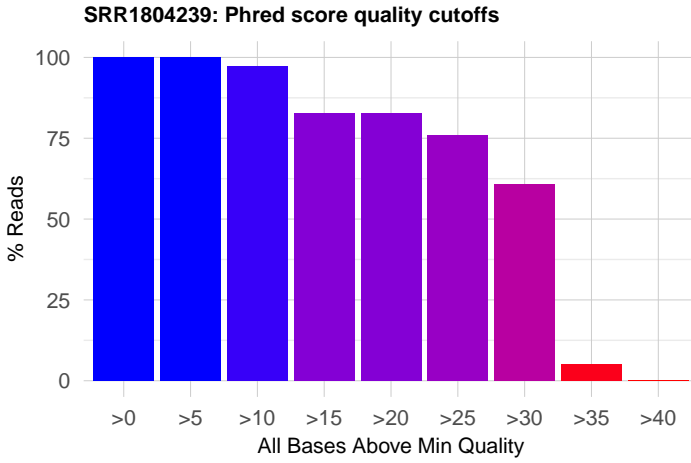
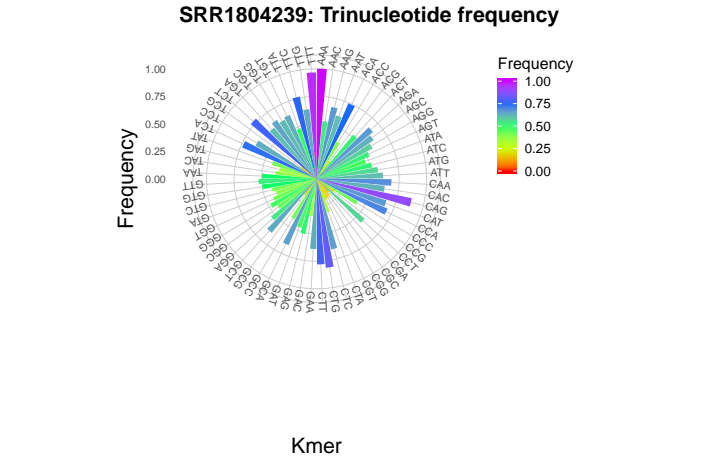
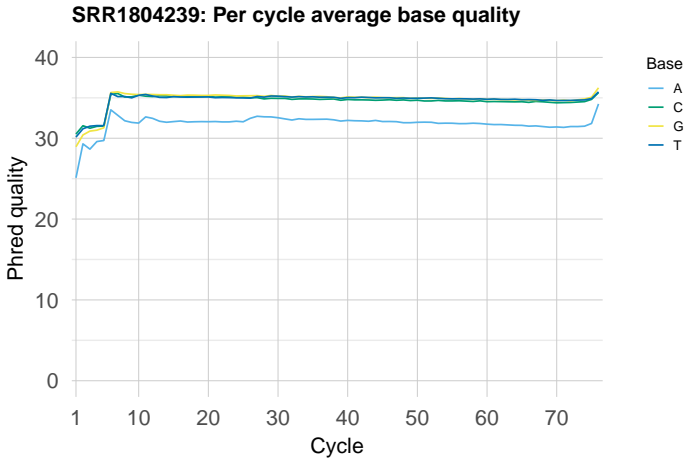
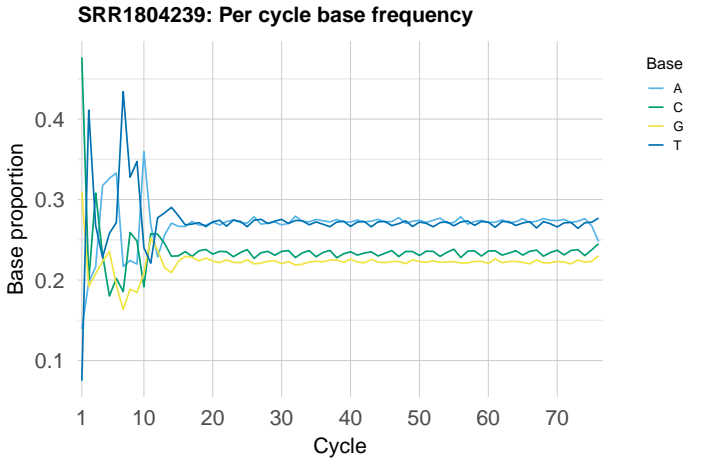
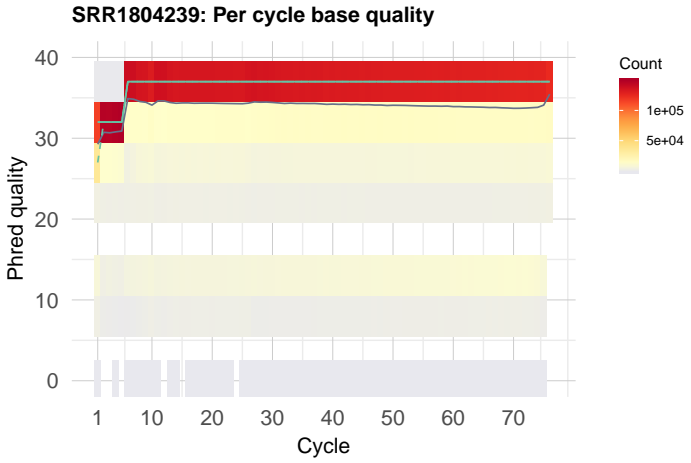


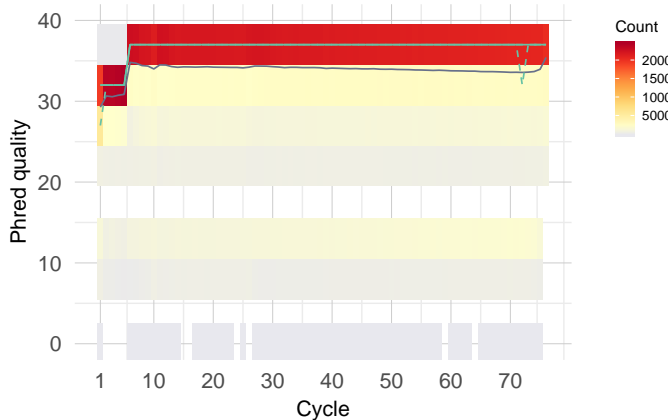
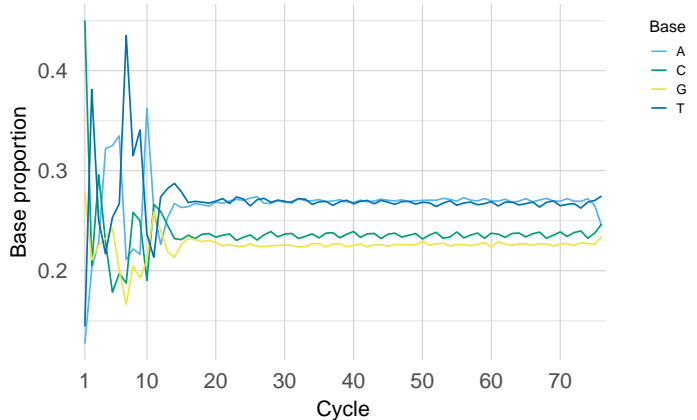
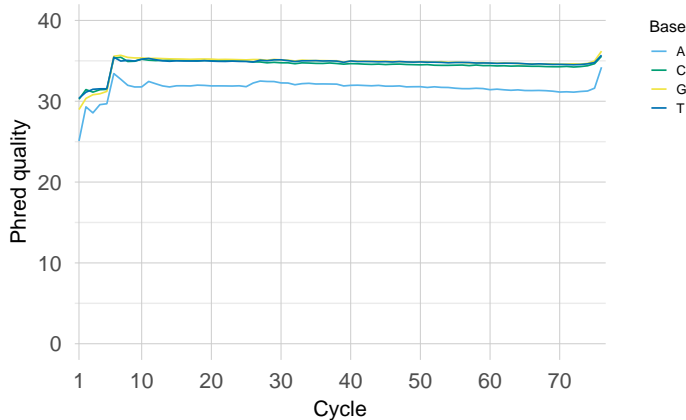
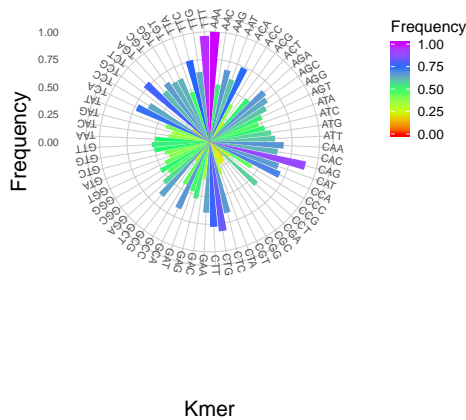
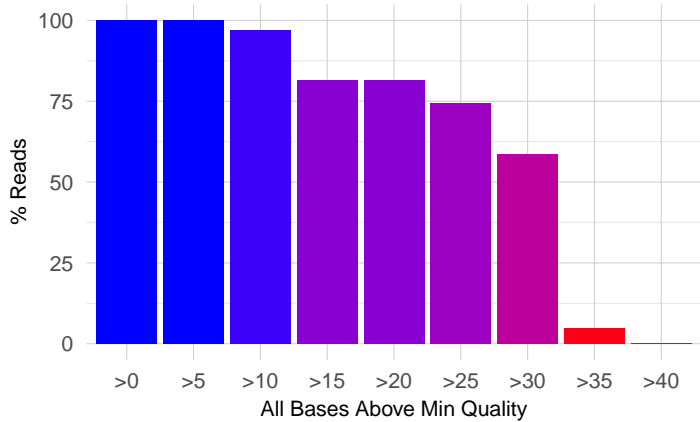
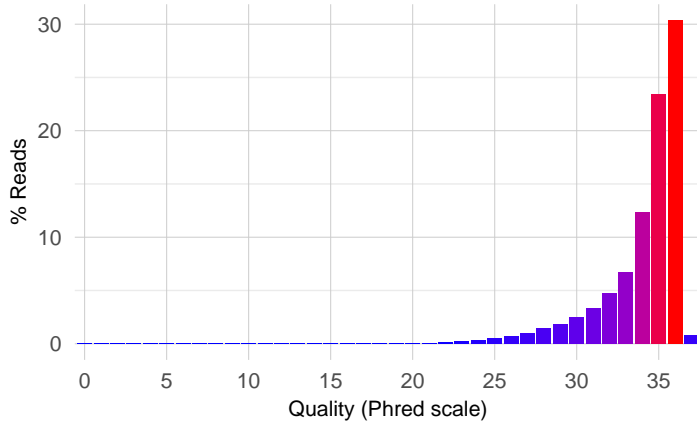
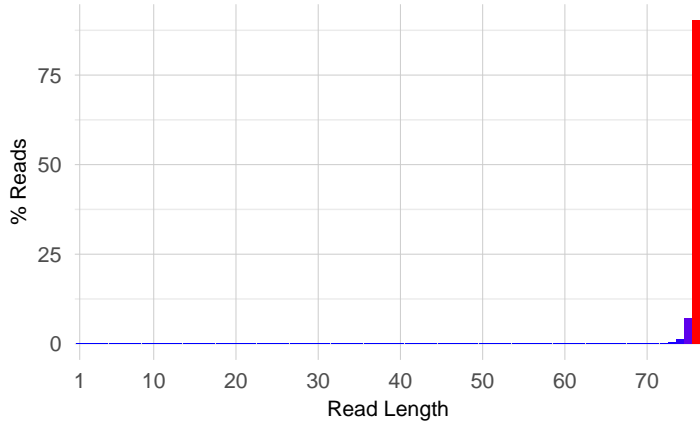
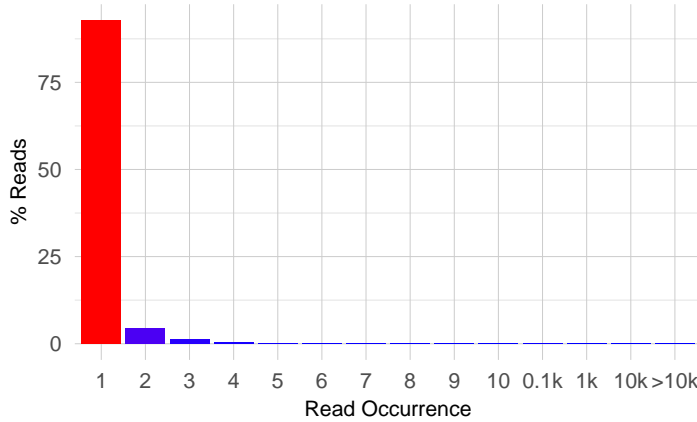
SRR1804236: Read occurrence distribution



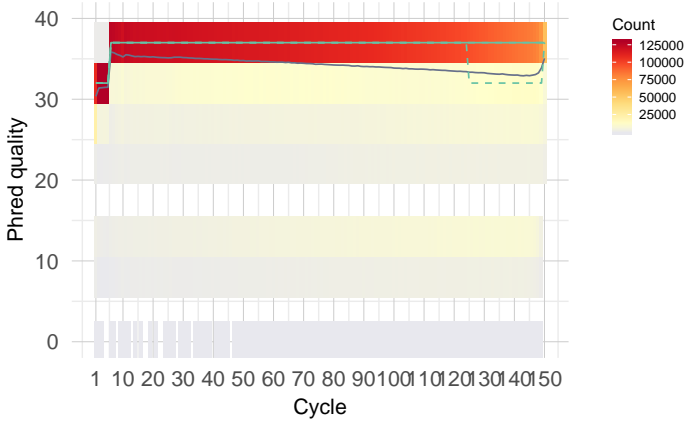


**SRR1804238: Per cycle base quality****SRR1804238: Per cycle base frequency****SRR1804238: Per cycle average base quality****SRR1804238: Trinucleotide frequency****SRR1804238: Phred score quality cutoffs****SRR1804238 : read quality for 217,242 of 4,344,843 reads****SRR1804238: Read length distribution****SRR1804238: Read occurrence distribution**

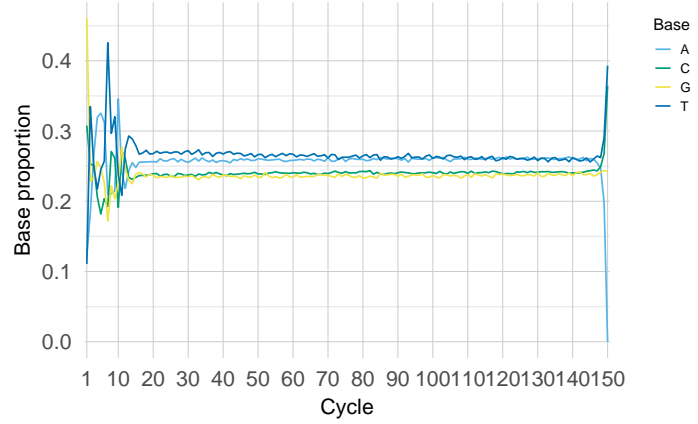


**SRR1804240: Per cycle base quality****SRR1804240: Per cycle base frequency****SRR1804240: Per cycle average base quality****SRR1804240: Trinucleotide frequency****SRR1804240: Phred score quality cutoffs****SRR1804240 : read quality for 276,515 of 5,530,294 reads****SRR1804240: Read length distribution****SRR1804240: Read occurrence distribution**

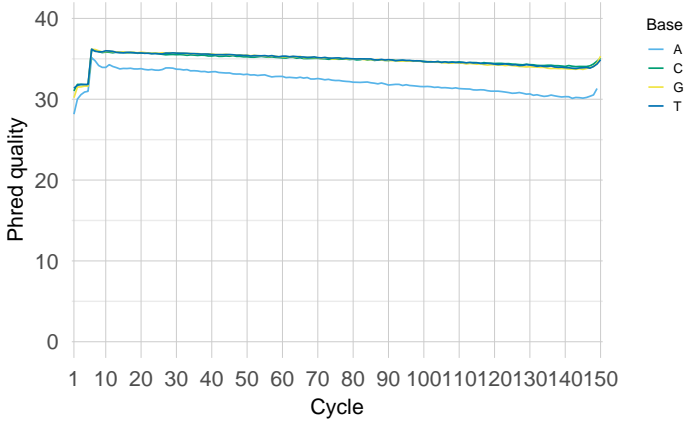
SRR1804241: Per cycle base quality



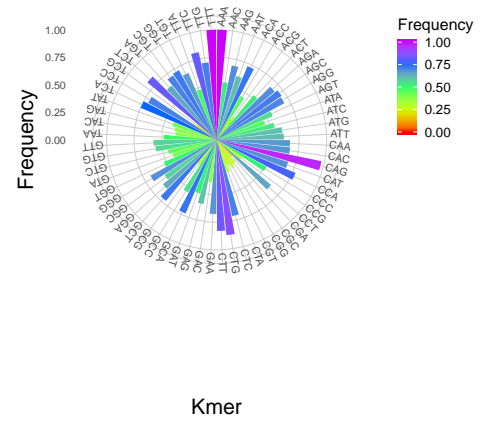
SRR1804241: Per cycle base frequency



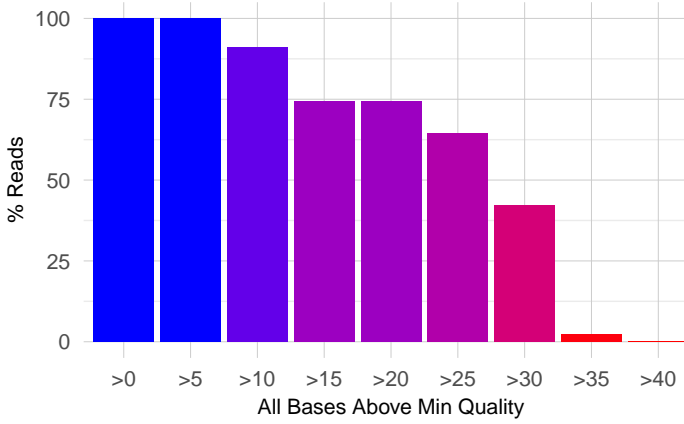
SRR1804241: Per cycle average base quality



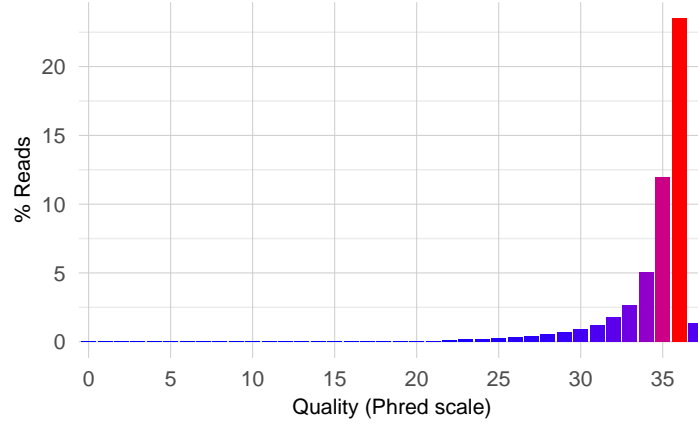
SRR1804241: Trinucleotide frequency



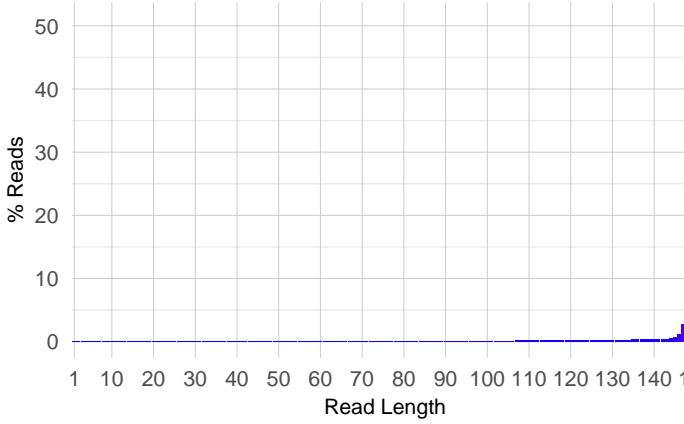
SRR1804241: Phred score quality cutoffs



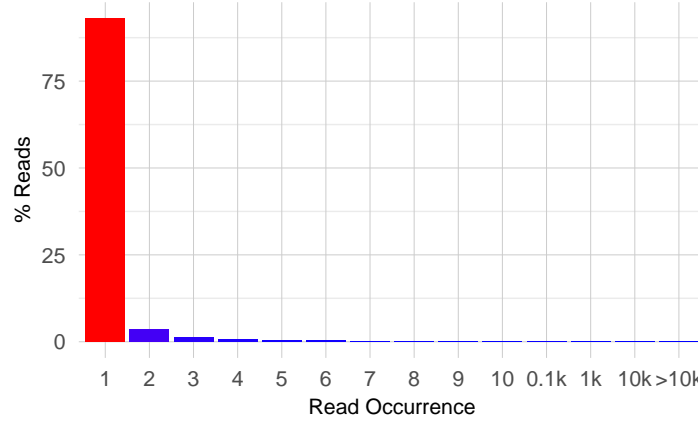
SRR1804241 : read quality for 138,869 of 2,777,383 reads

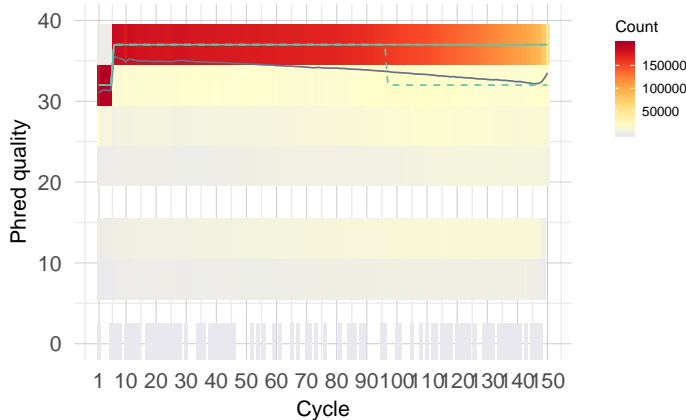
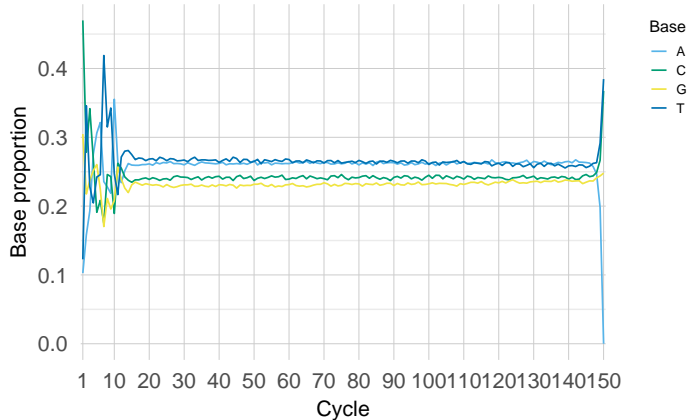
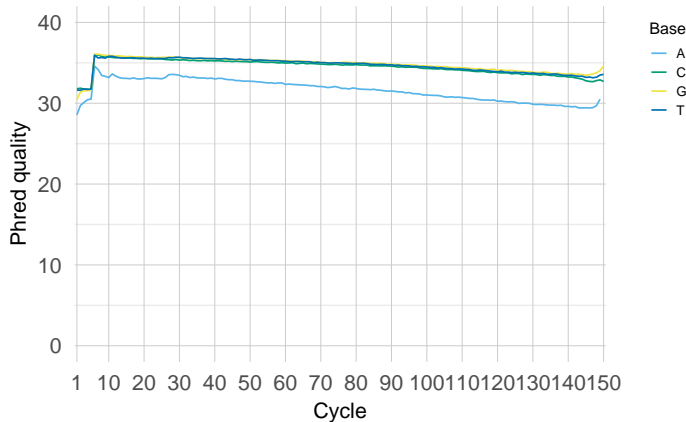
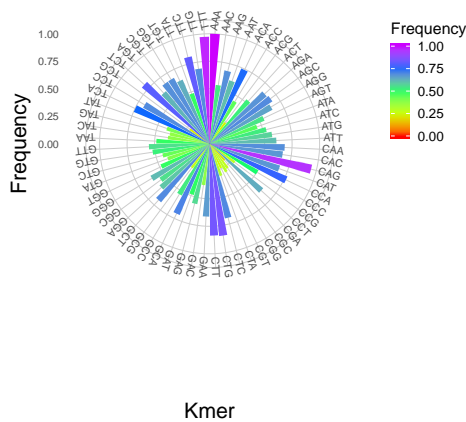
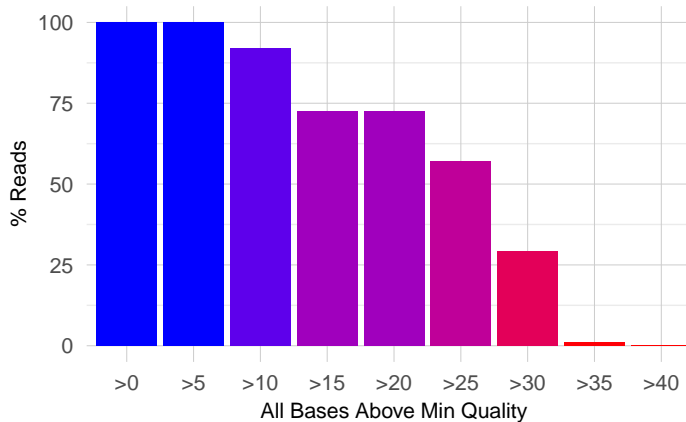
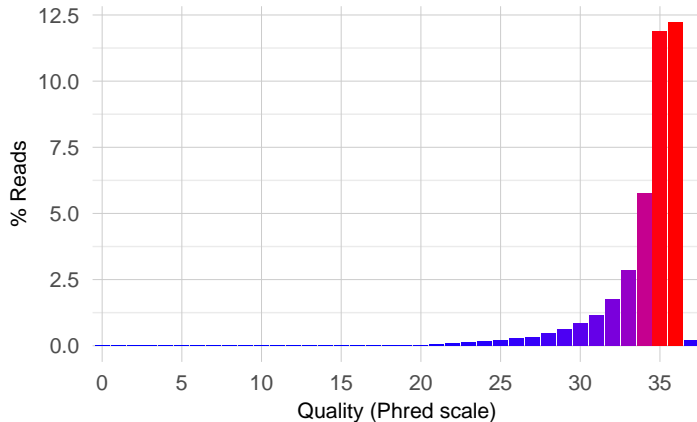
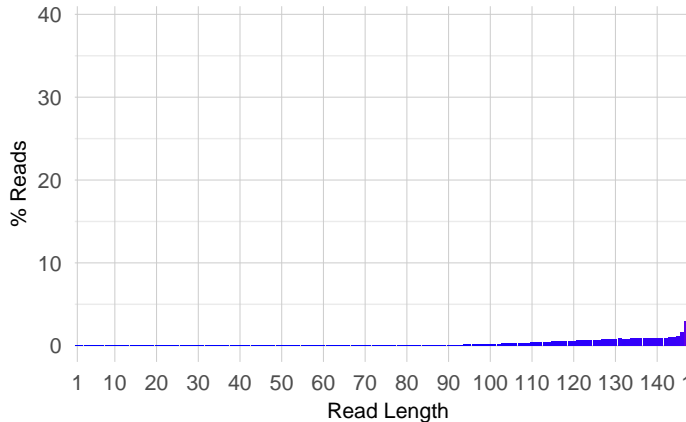


SRR1804241: Read length distribution



SRR1804241: Read occurrence distribution



**SRR1804242: Per cycle base quality****SRR1804242: Per cycle base frequency****SRR1804242: Per cycle average base quality****SRR1804242: Trinucleotide frequency****SRR1804242: Phred score quality cutoffs****SRR1804242 : read quality for 212,385 of 4,247,705 reads****SRR1804242: Read length distribution****SRR1804242: Read occurrence distribution**