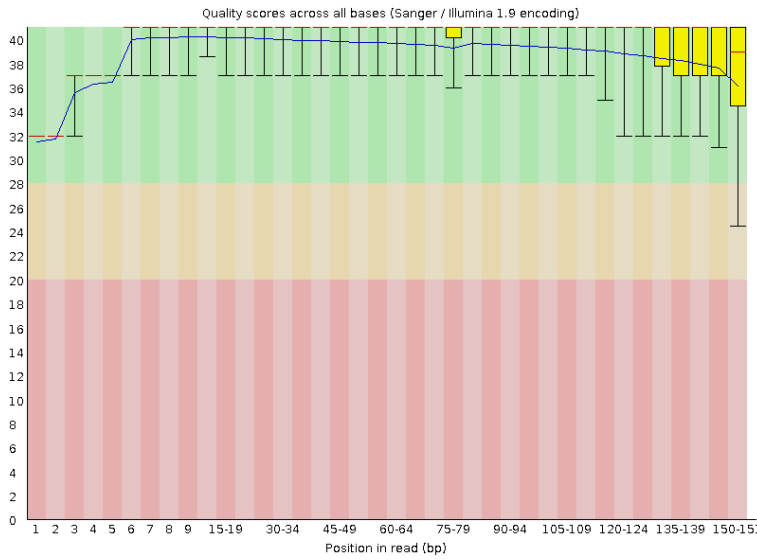
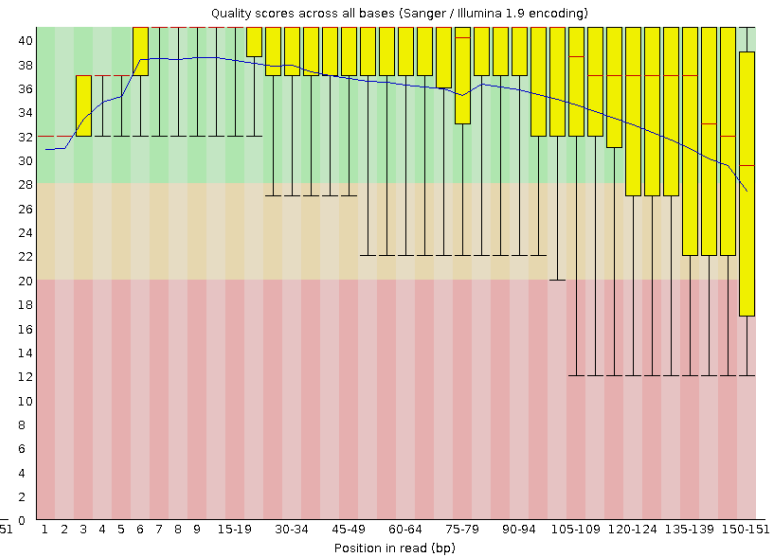
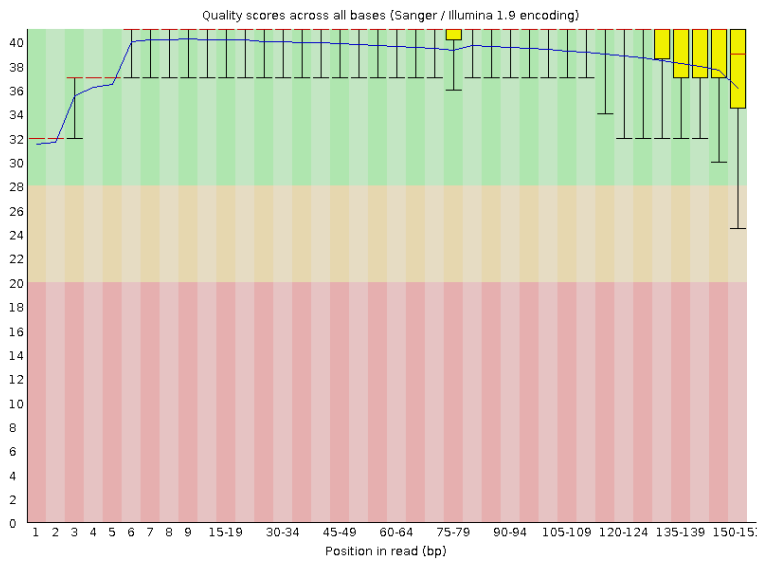
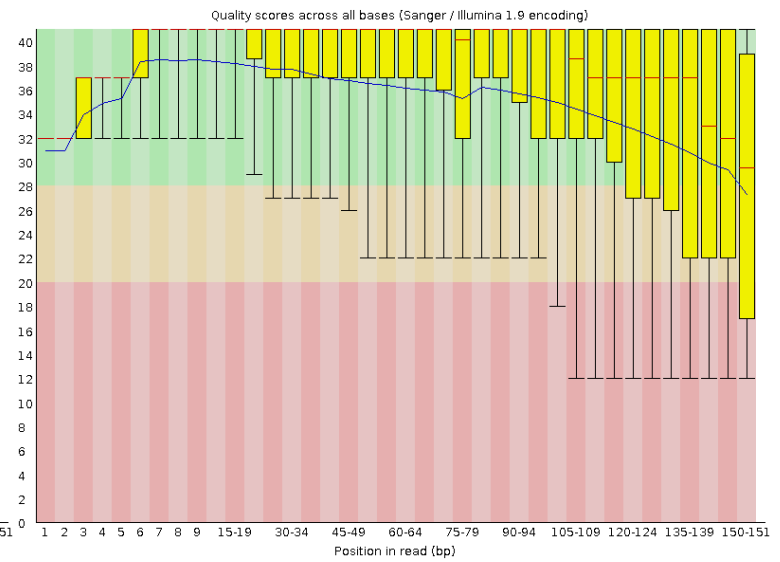
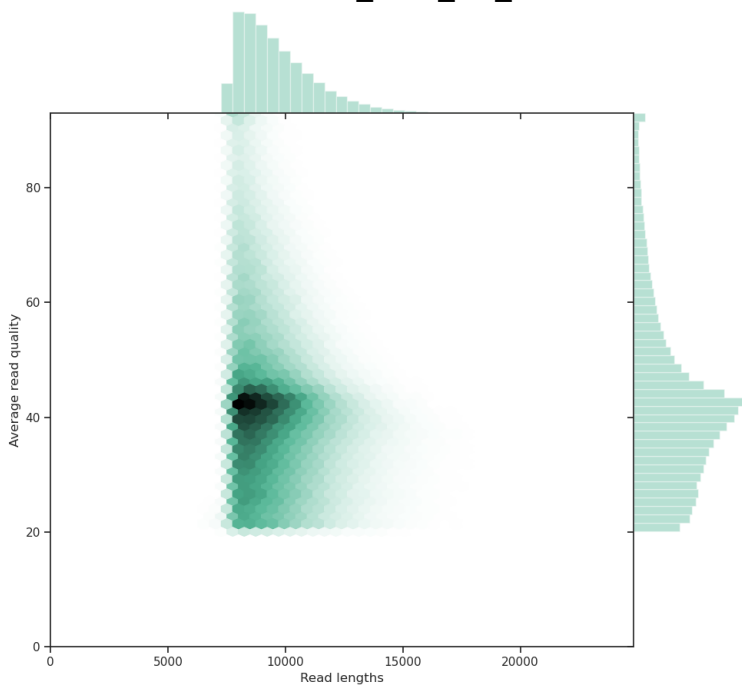
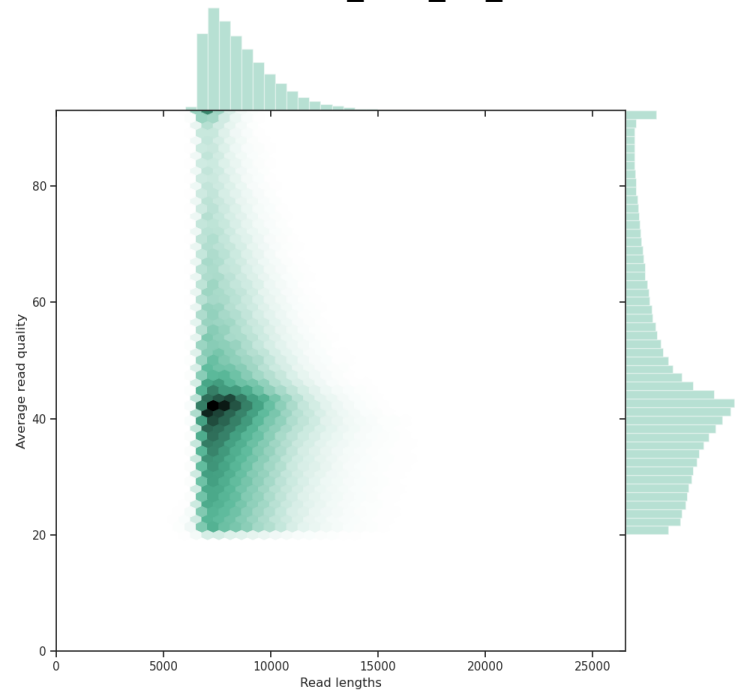


FRAM18_4502_E_R1**FRAM18_4502_E_R2****FRAM18_4502_G_R1****FRAM18_4502_G_R2****FRAM18_4571_PB_I****FRAM18_4571_PB_K**

Supplementary Figure S1. Visualisation of sequence over length. Upper four plots are example outputs of FastQC that summarises the per base sequence quality of Illumina reads. The bottom two plots are example outputs of NanoPlot that summarises the average read quality for different read lengths.