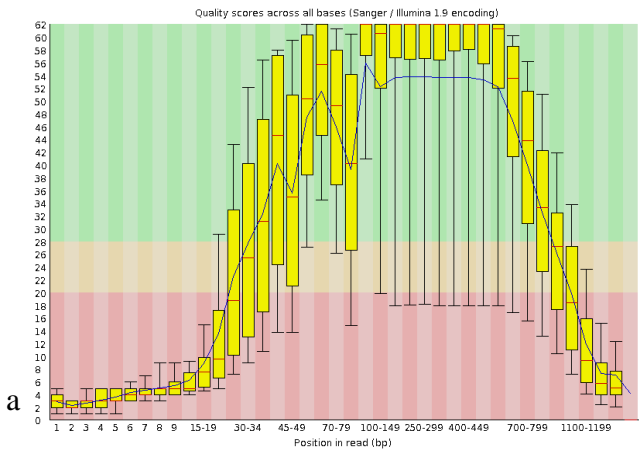
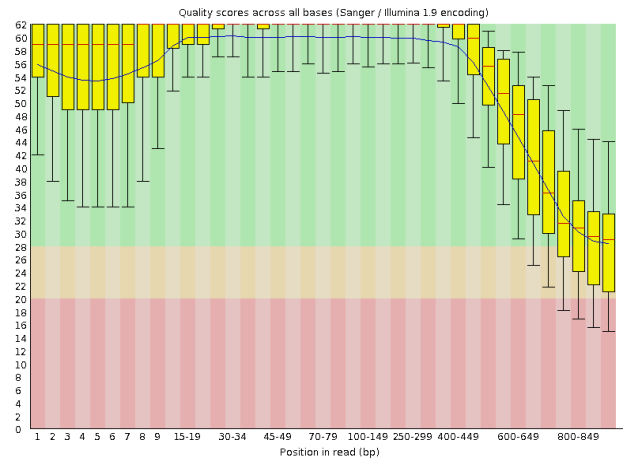


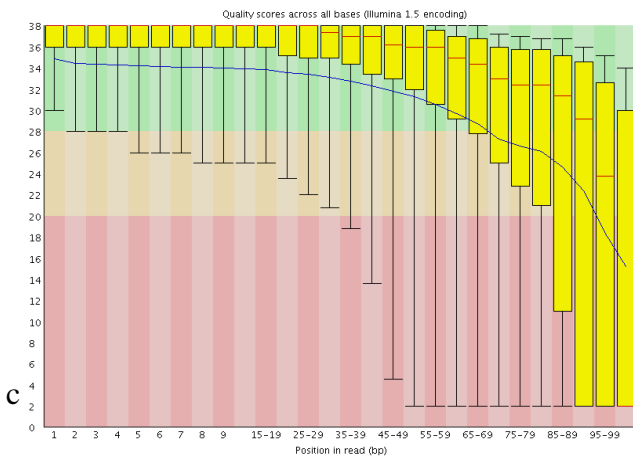
✘ Per base sequence quality



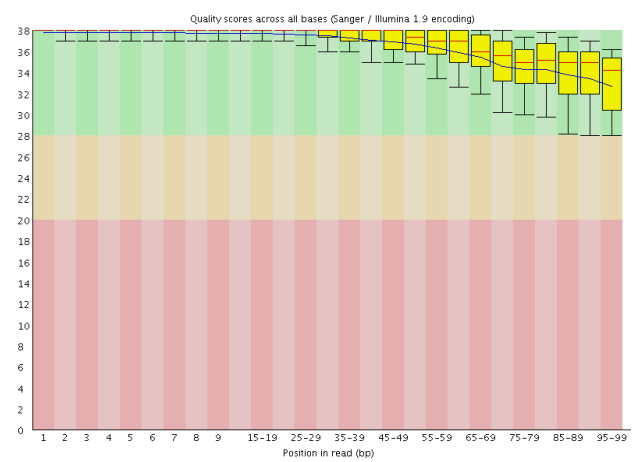
✔ Per base sequence quality



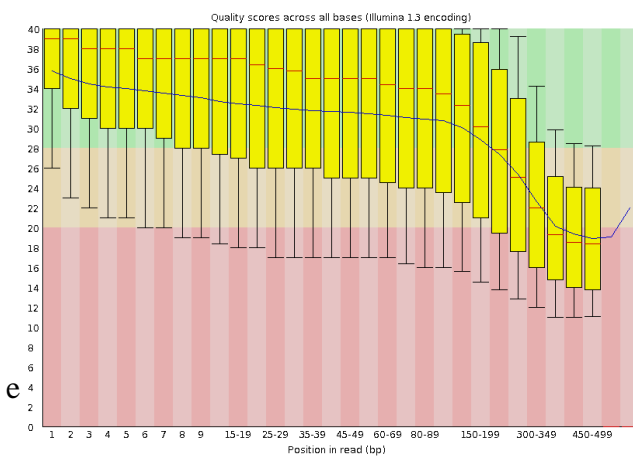
✘ Per base sequence quality



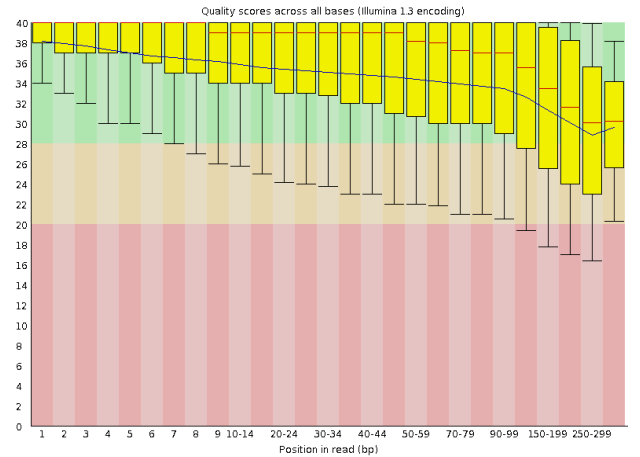
✔ Per base sequence quality



✘ Per base sequence quality



✔ Per base sequence quality



Additional File 1. Quality control of sequencing reads

Sanger (a,b), Illumina (c,d), and Roche 454 (e,f) sequencing data