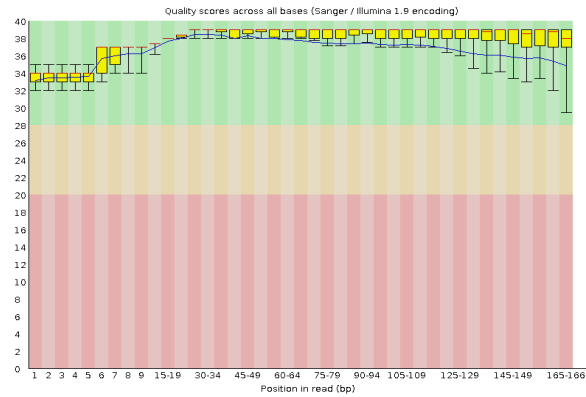
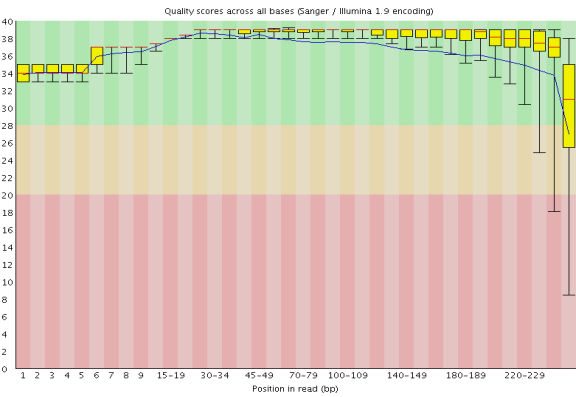
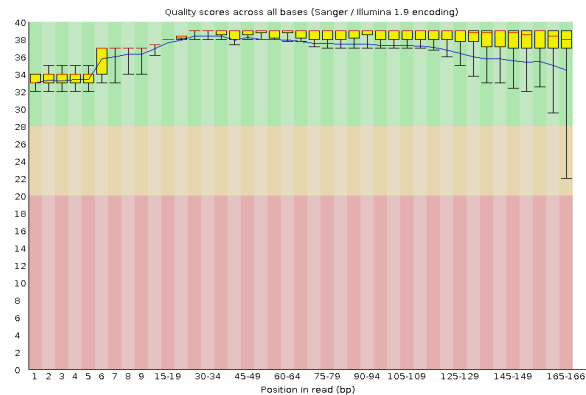
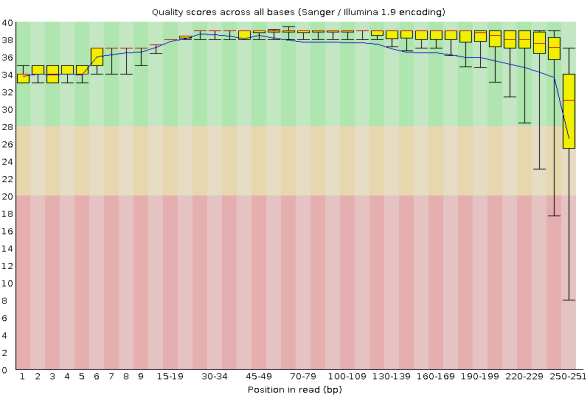


WT sample



Total number of paired reads	217,245
Uncalled bases (%)	0.0
Read length (bp)	250 + 250
GC content (%)	43.5

DLD1-mixed sample



Total number of paired reads	234,803
Uncalled bases (%)	0.0
Read length (bp)	250 + 166
GC content (%)	43

Figure S1