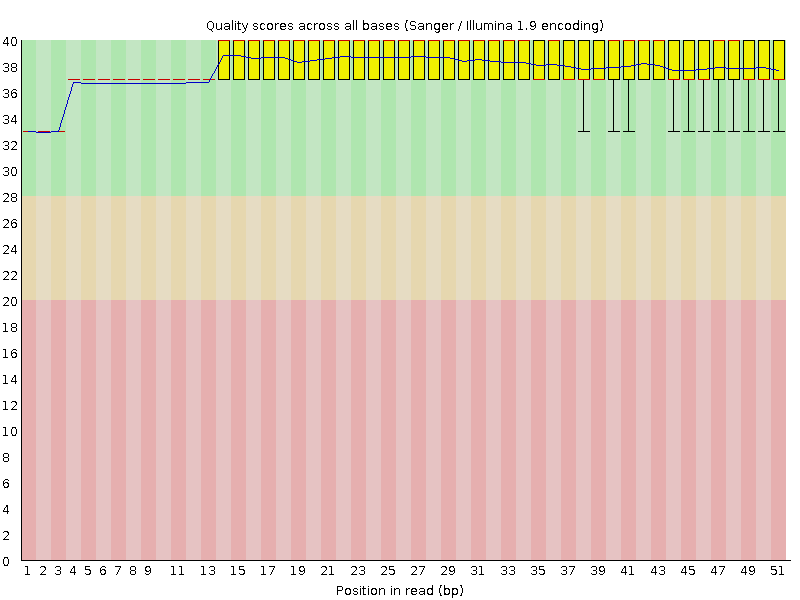
**S2 File**. **Quality control analysis** **after the removal of the sequencing adapter**. Quality scores across all bases (**Figure A**). Quality score distribution over all sequences (**Figure B**). Percentage of adapter sequence (**Figure C**).



A

B

C

