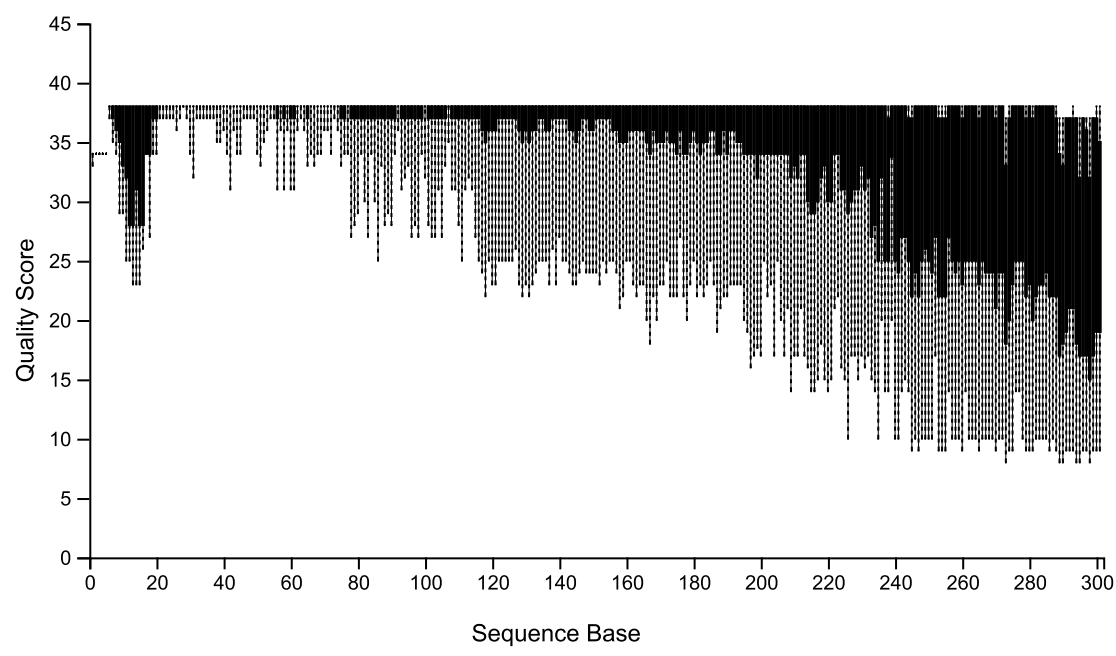
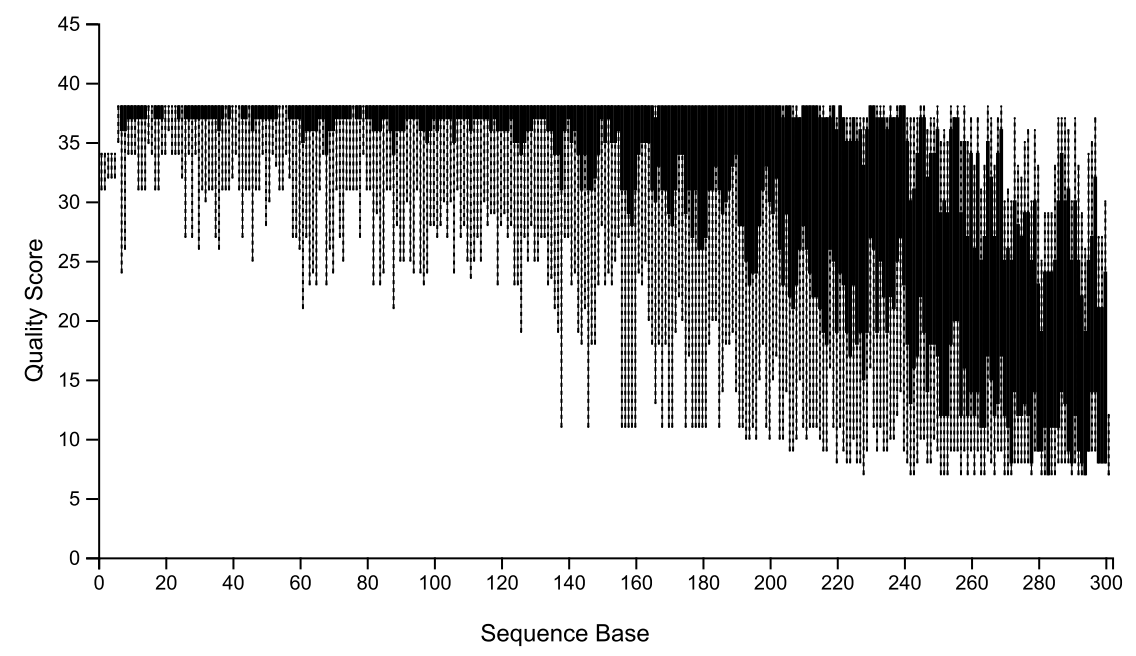


Click and drag on plot to zoom in. Double click to zoom back out to full size. Hover over a box to see the parametric seven-number summary of the quality scores at the corresponding position.

Forward Reads



Reverse Reads



These plots were generated using a random sampling of 10000 out of 8458890 sequences without replacement. The minimum sequence length identified during subsampling was 158 bases. Outlier quality scores are not shown in box plots for clarity.

These plots were generated using a random sampling of 10000 out of 8458890 sequences without replacement. The minimum sequence length identified during subsampling was 79 bases. Outlier quality scores are not shown in box plots for clarity.

Parametric seven-number summary		
Box plot feature	Percentile	Quality score
(Not shown in box plot)	2nd	...
Lower Whisker	9th	...
Bottom of Box	25th	...
Middle of Box	50th (Median)	...
Top of Box	75th	...
Upper Whisker	91st	...
(Not shown in box plot)	98th	...

Parametric seven-number summary		
Box plot feature	Percentile	Quality score
(Not shown in box plot)	2nd	...
Lower Whisker	9th	...
Bottom of Box	25th	...
Middle of Box	50th (Median)	...
Top of Box	75th	...
Upper Whisker	91st	...
(Not shown in box plot)	98th	...

[Download forward parametric seven-number summaries as CSV \(forward-seven-number-summaries.csv\)](#)

[Download reverse parametric seven-number summaries as CSV \(reverse-seven-number-summaries.csv\)](#)

## Demultiplexed sequence length summary

Forward Reads

Percentile	Sequence Length (nts)
<b>Total Sequences Sampled</b>	10000
<b>2%</b>	297 nts
<b>9%</b>	301 nts
<b>25%</b>	301 nts
<b>50% (Median)</b>	301 nts
<b>75%</b>	301 nts
<b>91%</b>	301 nts
<b>98%</b>	301 nts

Reverse Reads

Percentile	Sequence Length (nts)
<b>Total Sequences Sampled</b>	10000
<b>2%</b>	300 nts
<b>9%</b>	300 nts
<b>25%</b>	300 nts
<b>50% (Median)</b>	300 nts
<b>75%</b>	300 nts
<b>91%</b>	301 nts
<b>98%</b>	301 nts