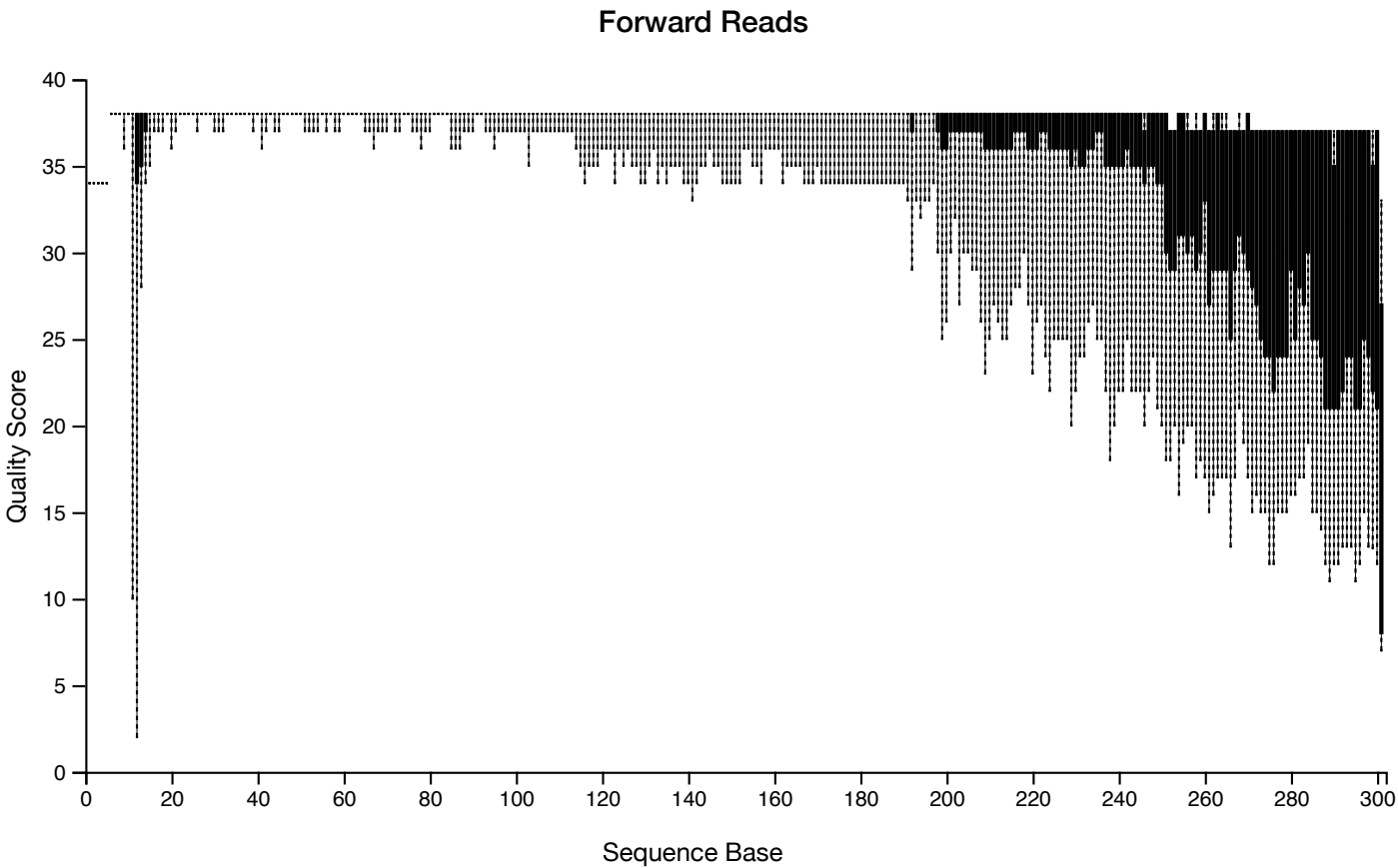


Click and drag on plot to zoom in. Double click to zoom back out to full size. Hover over a box to see the parametric seven-number summary of the quality scores at the corresponding position.

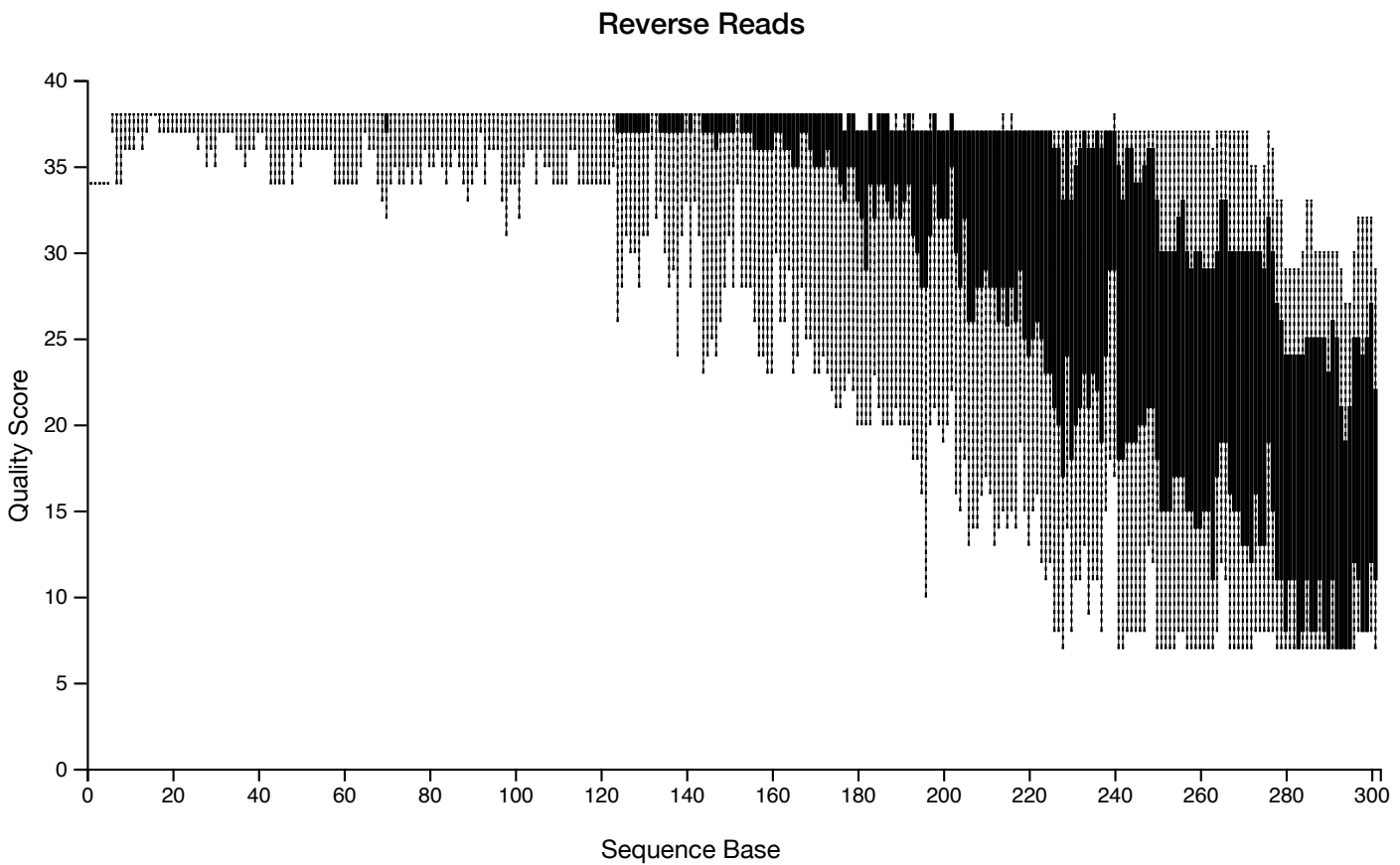


The plot at position 118 was generated using a random sampling of 10000 out of 1473873 sequences without replacement. The minimum sequence length identified during subsampling was 301 bases. Outlier quality scores are not shown in box plots for clarity.

Parametric seven-number summary for position 118		
Box plot feature	Percentile	Quality score
(Not shown in box plot)	2nd	18
Lower Whisker	9th	35
Bottom of Box	25th	38
Middle of Box	50th (Median)	38
Top of Box	75th	38
Upper Whisker	91st	38

(Not shown in box plot)	98th	38
-------------------------	------	----

Download forward parametric seven-number summaries as TSV (forward-seven-number-summaries.tsv)



The plot at position 11 was generated using a random sampling of 10000 out of 1473873 sequences without replacement. The minimum sequence length identified during subsampling was 301 bases. Outlier quality scores are not shown in box plots for clarity.

Parametric seven-number summary for position 11		
Box plot feature	Percentile	Quality score
(Not shown in box plot)	2nd	21
Lower Whisker	9th	36
Bottom of Box	25th	38
Middle of Box	50th (Median)	38
Top of Box	75th	38

Upper Whisker	91st	38
(Not shown in box plot)	98th	38

Download reverse parametric seven-number summaries as TSV ([reverse-seven-number-summaries.tsv](#))

Demultiplexed sequence length summary

Forward Reads

Total Sequences Sampled	10000.0
2%	301 nts
9%	301 nts
25%	301 nts
50% (Median)	301 nts
75%	301 nts
91%	301 nts
98%	301 nts

Reverse Reads

Total Sequences Sampled	10000.0
2%	301 nts
9%	301 nts
25%	301 nts
50% (Median)	301 nts
75%	301 nts
91%	301 nts

98%

301 nts