

QC stats

Reads #: 239400

Total bases: 9576000

Reads Length: 40.00

Processed 239400 reads for quality check only

Reads length < 36 bp: 0 (0.00 %)

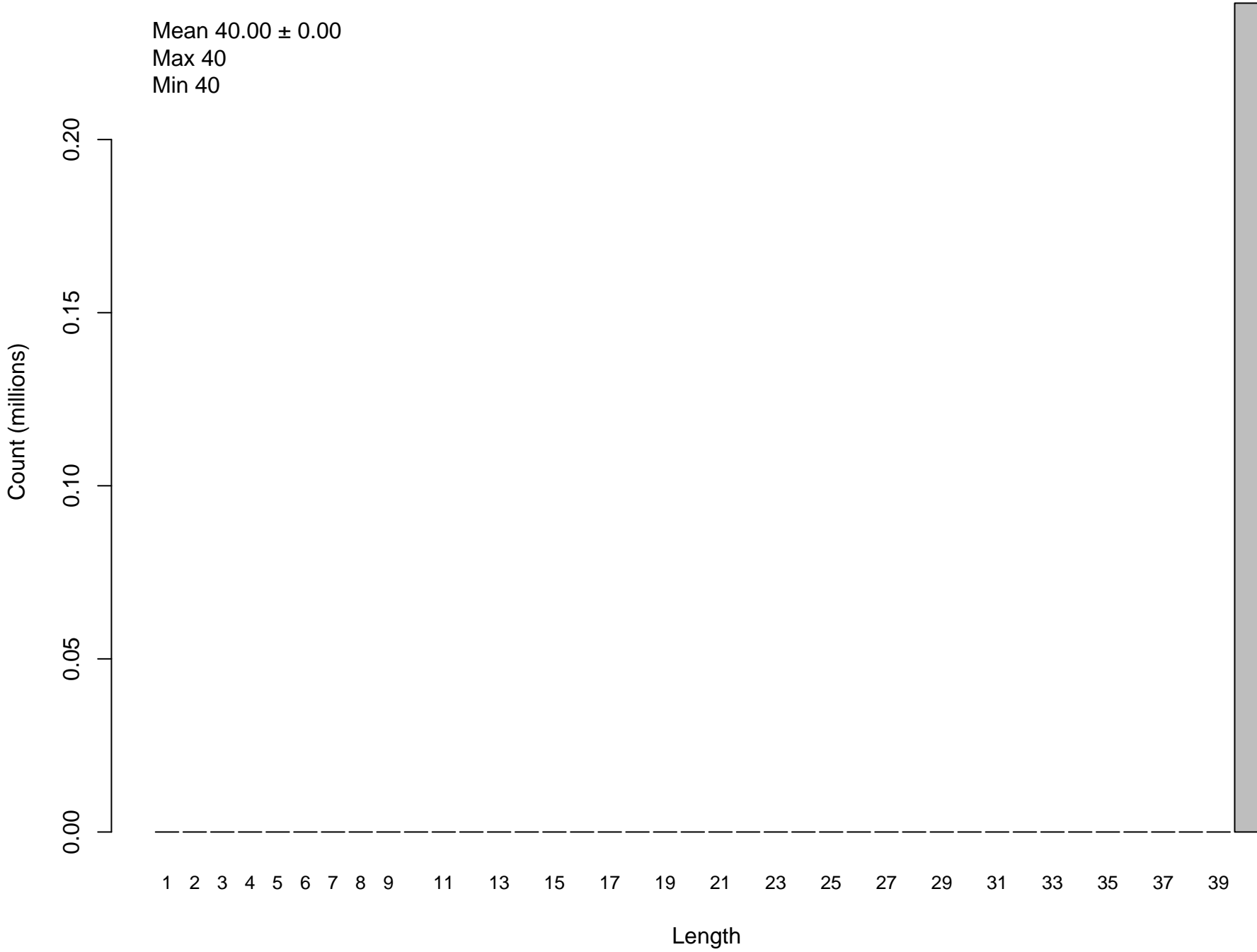
Reads have 2 continuous base "N": 17 (0.01 %)

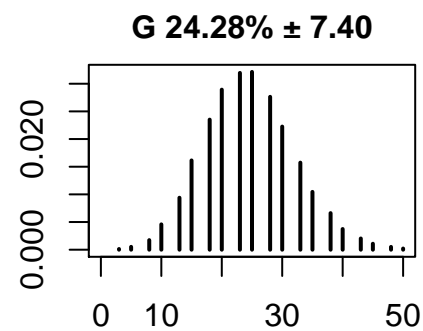
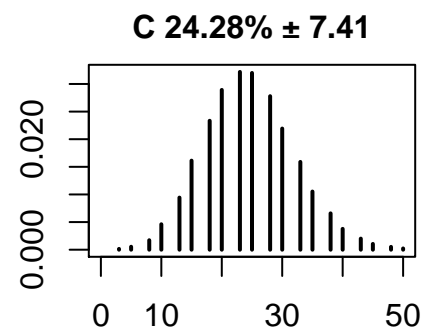
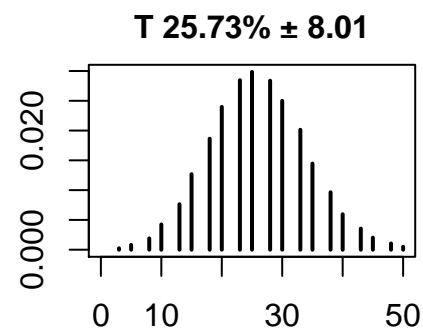
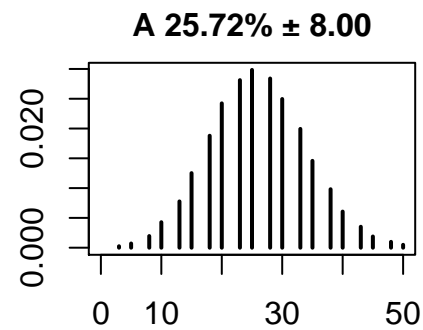
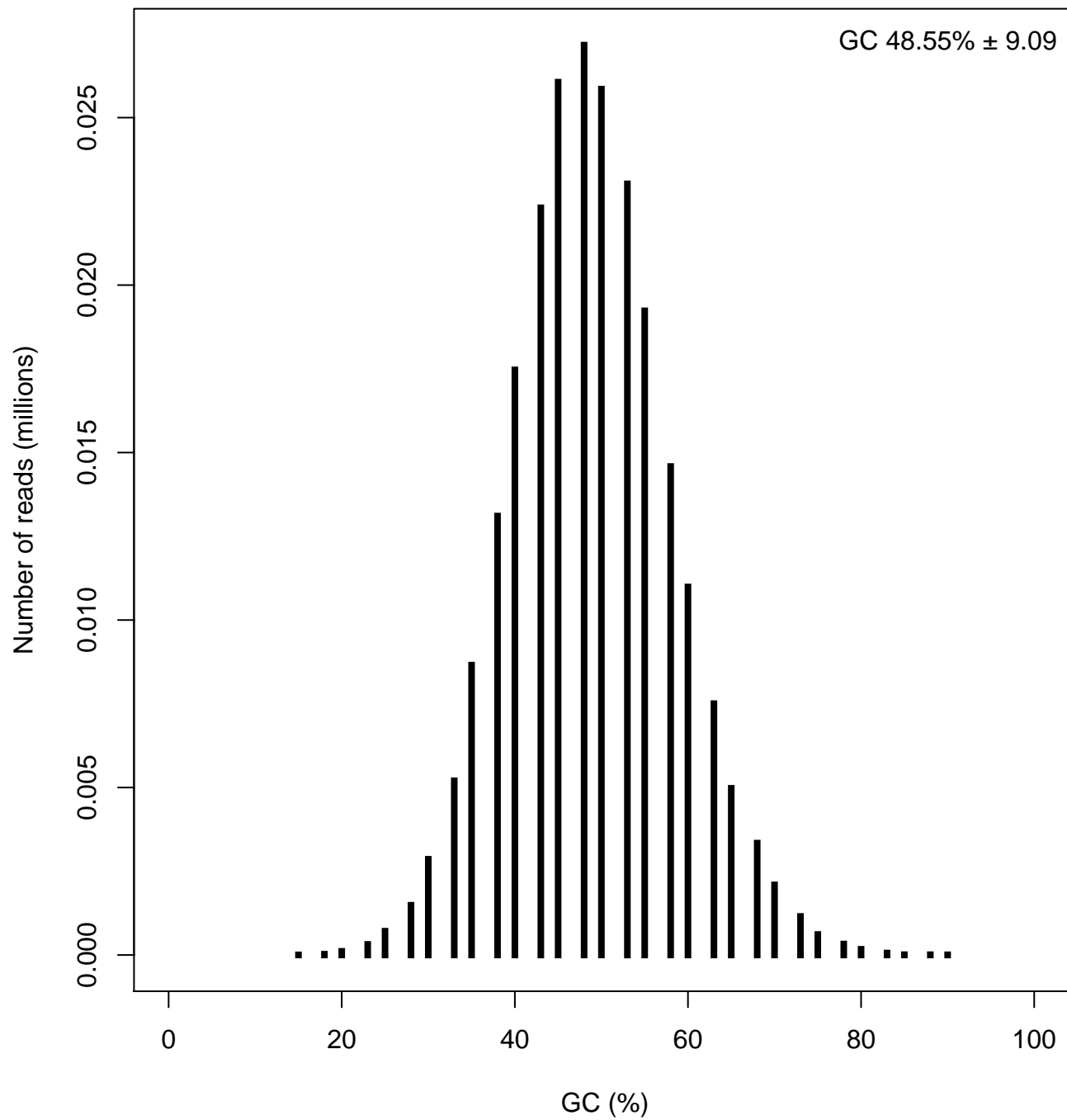
Low complexity Reads (>85.00% mono/di-nucleotides): 0 (0.00 %)

Reads < average quality 0.0: 0 (0.00 %)

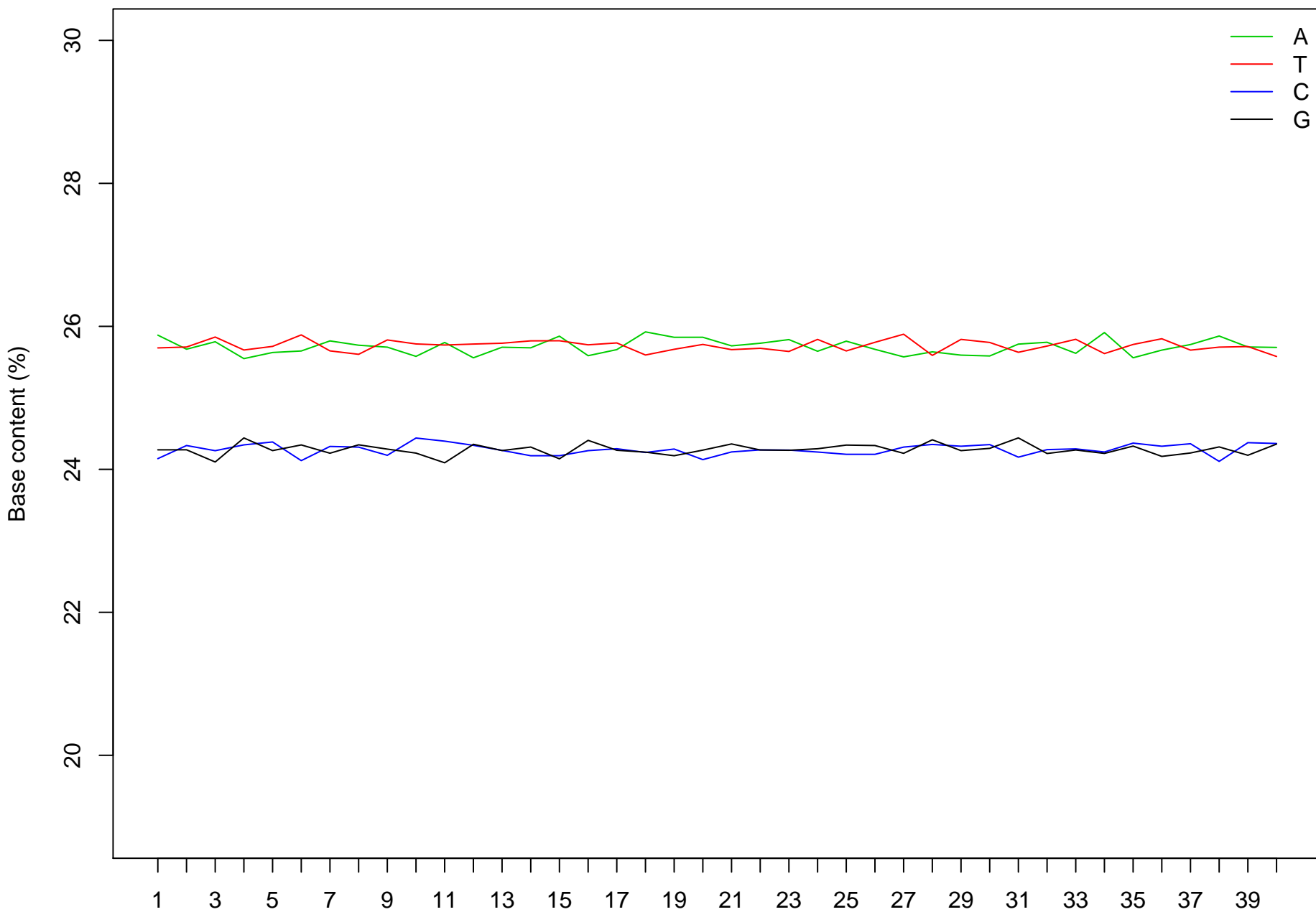
Reads Length Histogram

Mean 40.00 ± 0.00
Max 40
Min 40

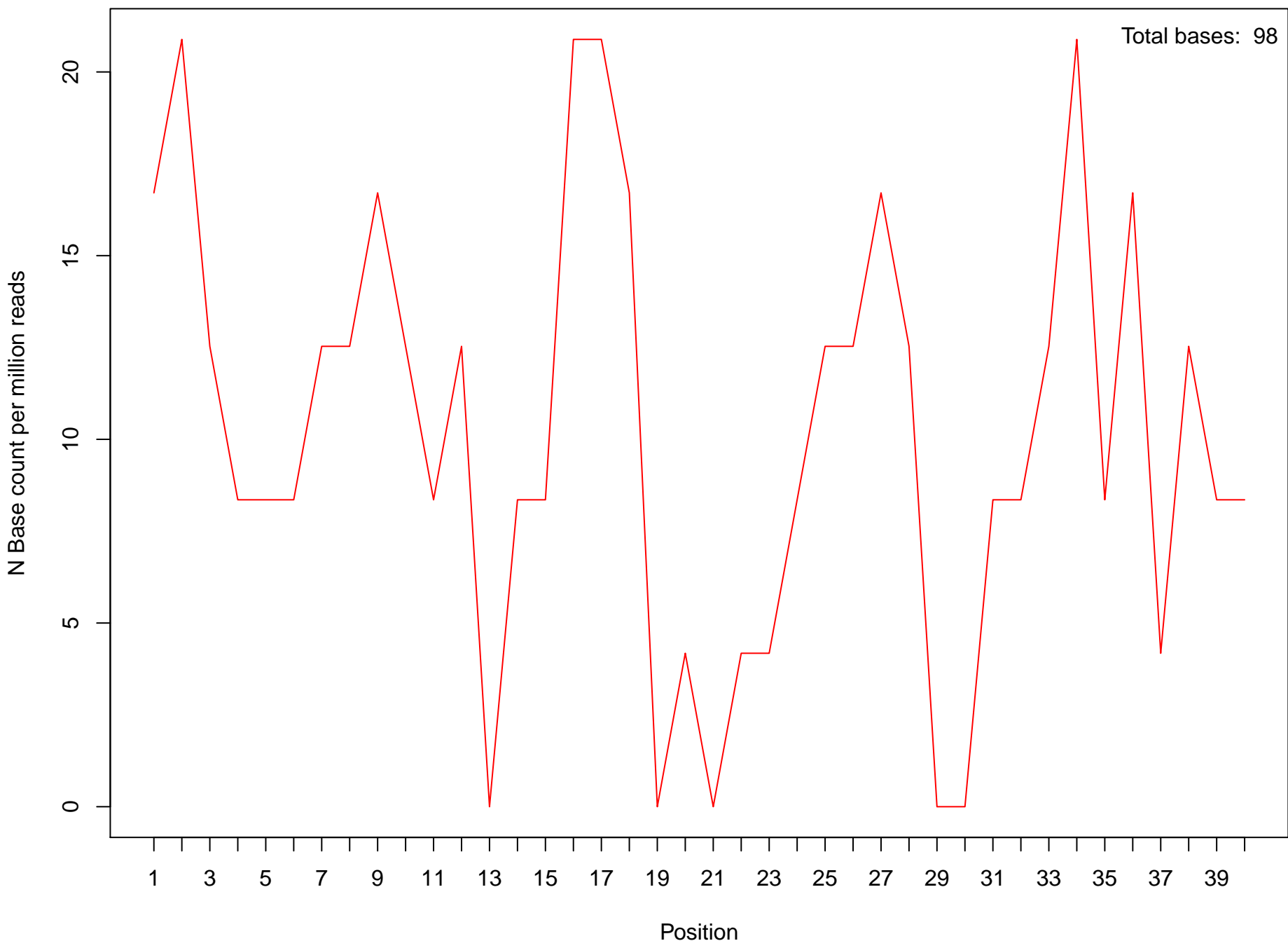




Nucleotide Content Per Cycle

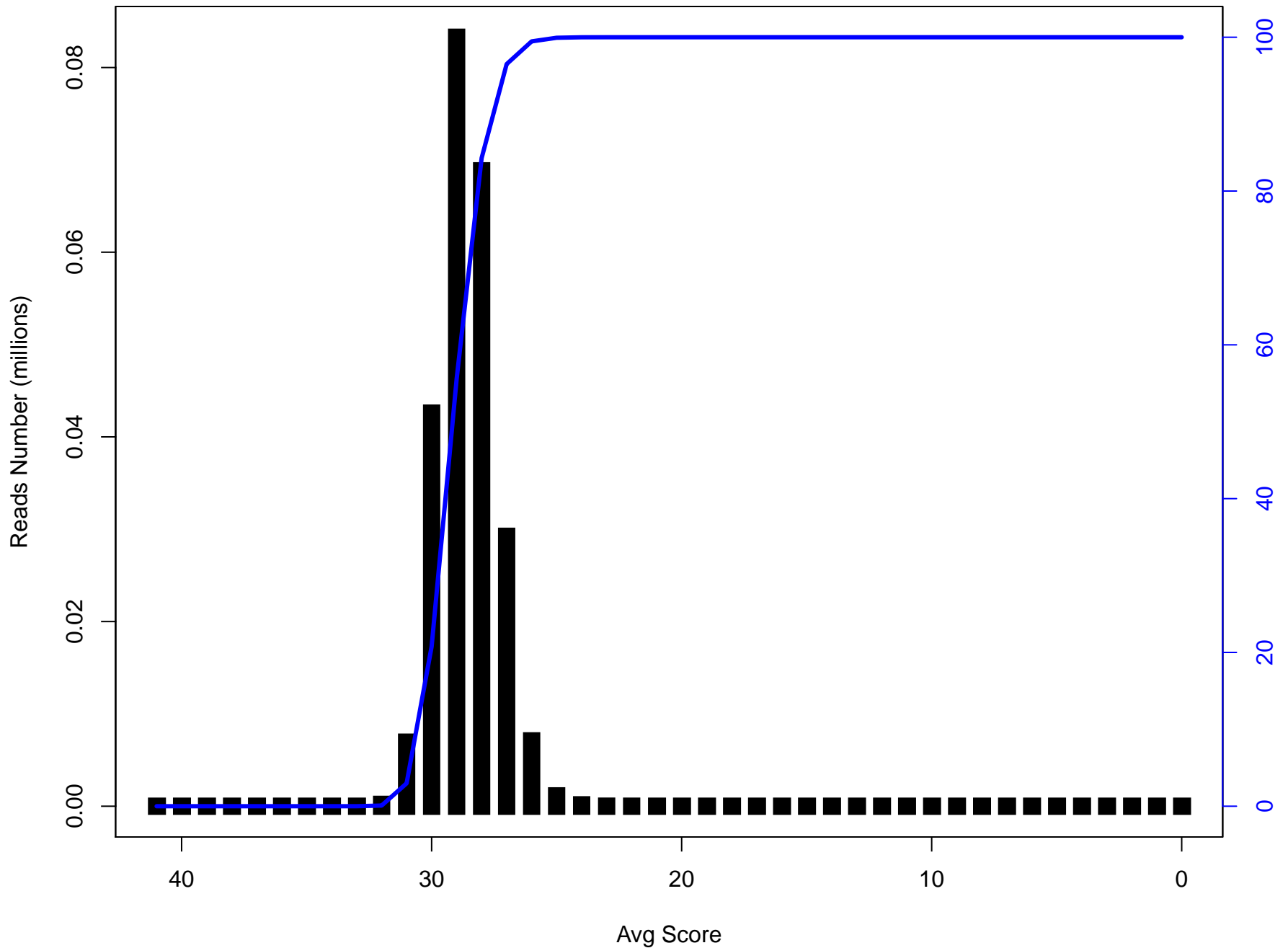


N Nucleotide Content Per Cycle

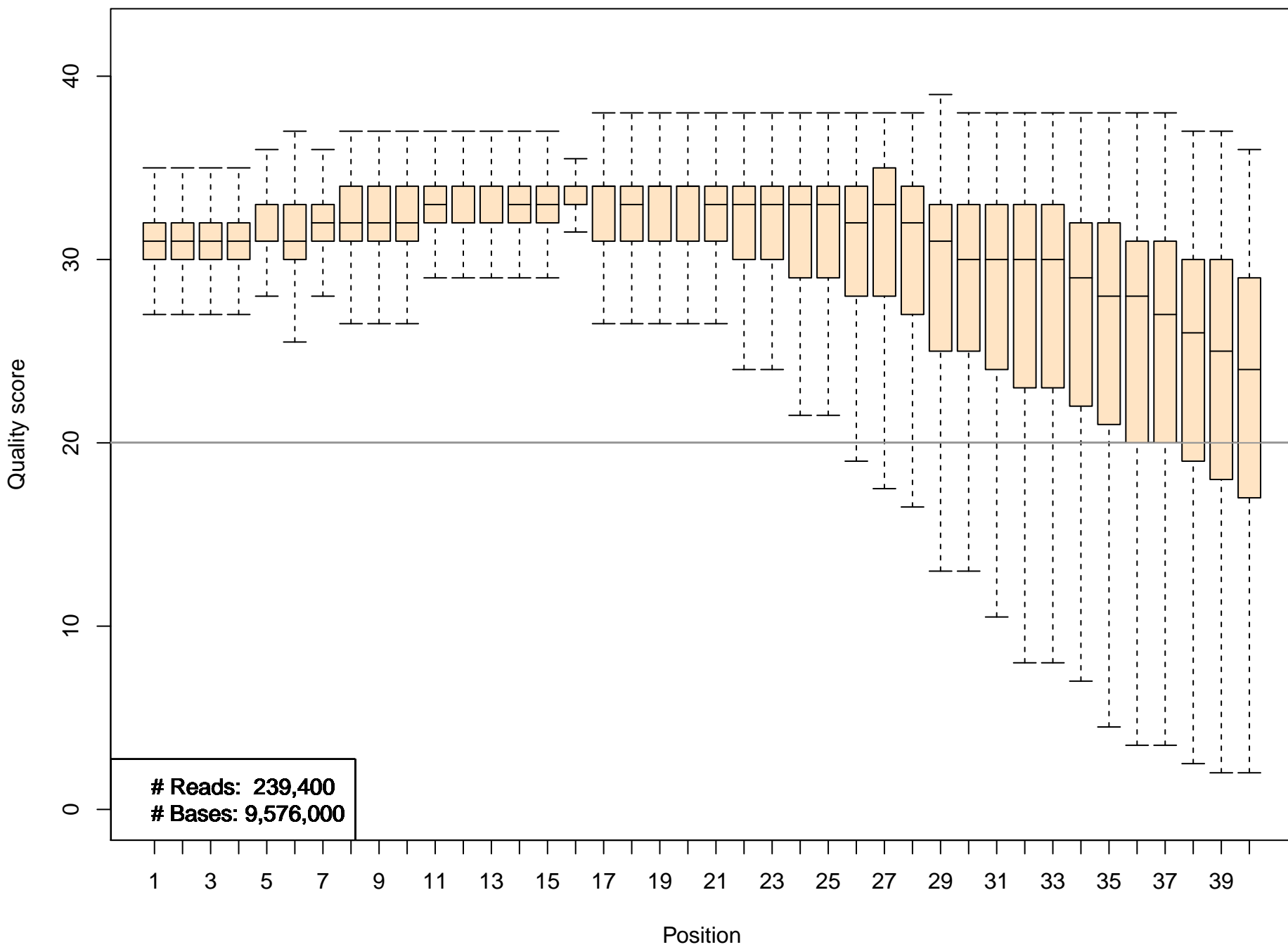


Reads Average Quality Histogram

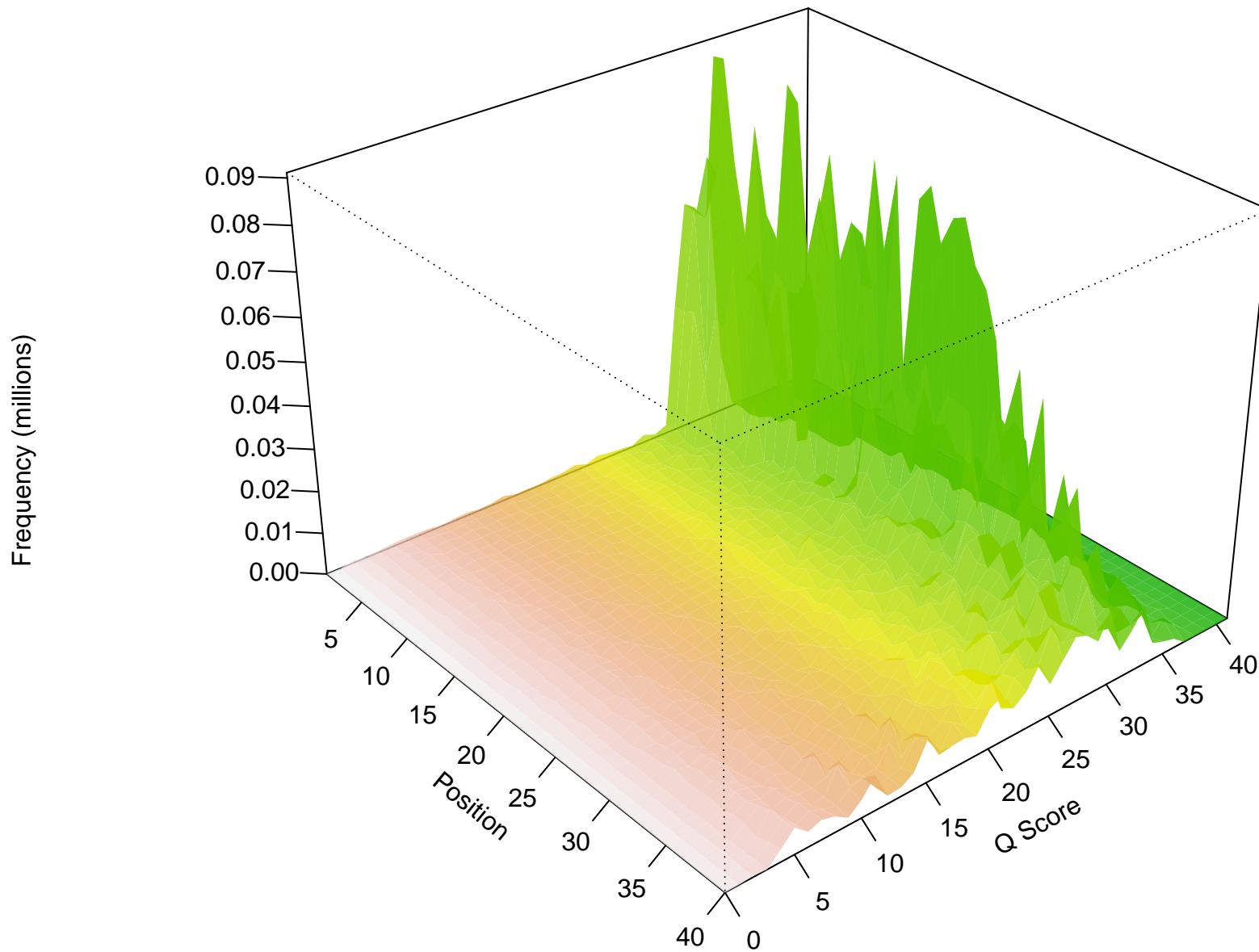
Number of Q>=20 reads: 239,400 (100.00%) , mean Length: 40.00



Quality Boxplot Per Cycle



Quality 3D plot. (Position vs. Score vs. Frequency)



Quality report

Average: 29.08

