Basic Statistics

Filename1: /sbgenomics/Projects/10b38e1d-256a-4288-aa1a-5b5775fc35

Filename2: /sbgenomics/Projects/10b38e1d-256a-4288-aa1a-5b5775fc35
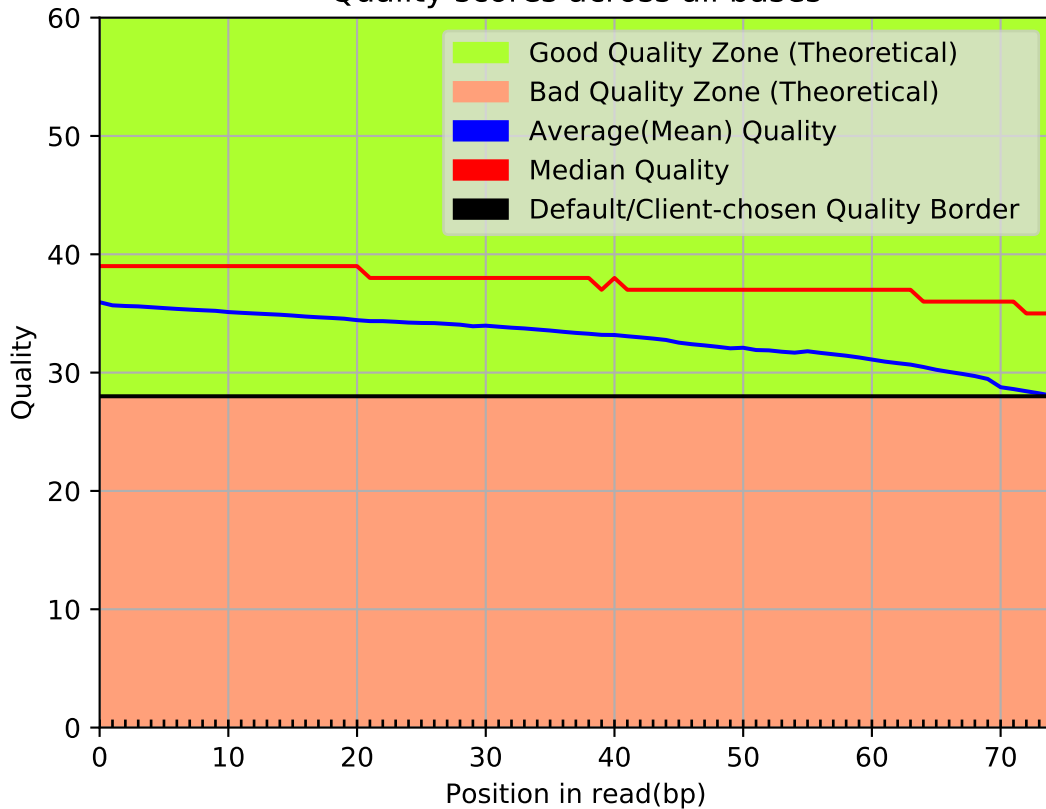
Poor Quality Border: 28

Total Sequences 9885628
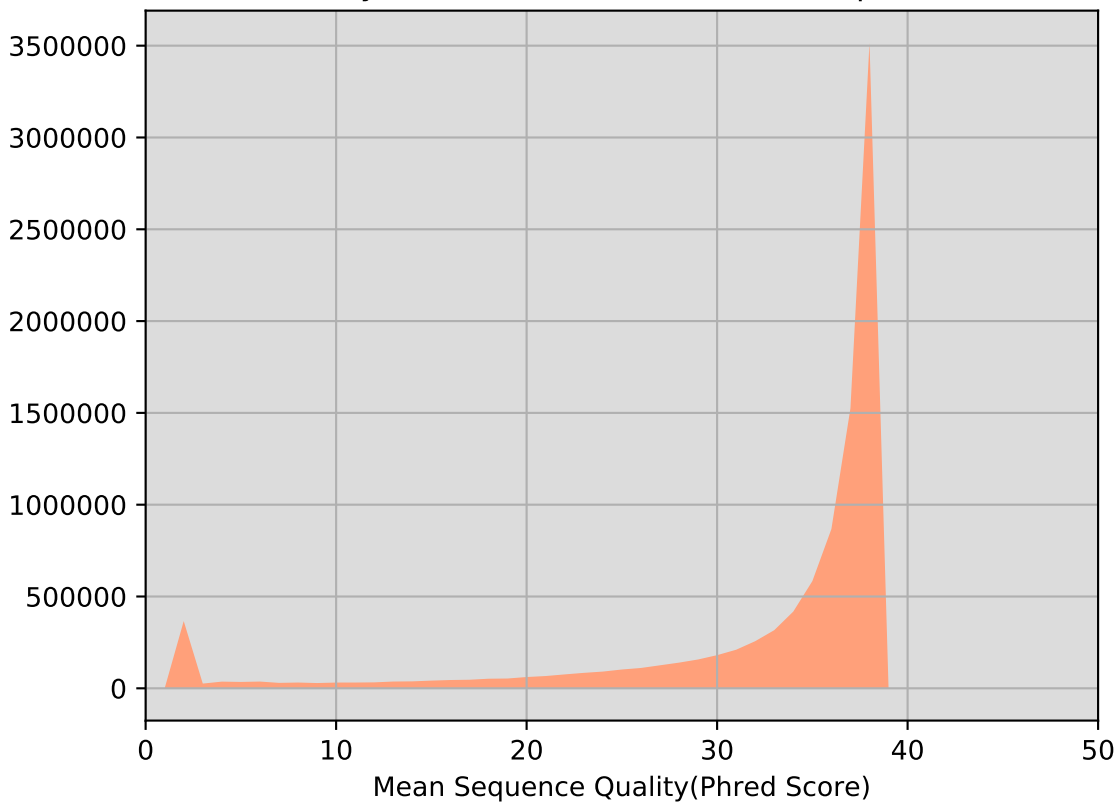
Seq. Flaged as poor Quality: 1715469

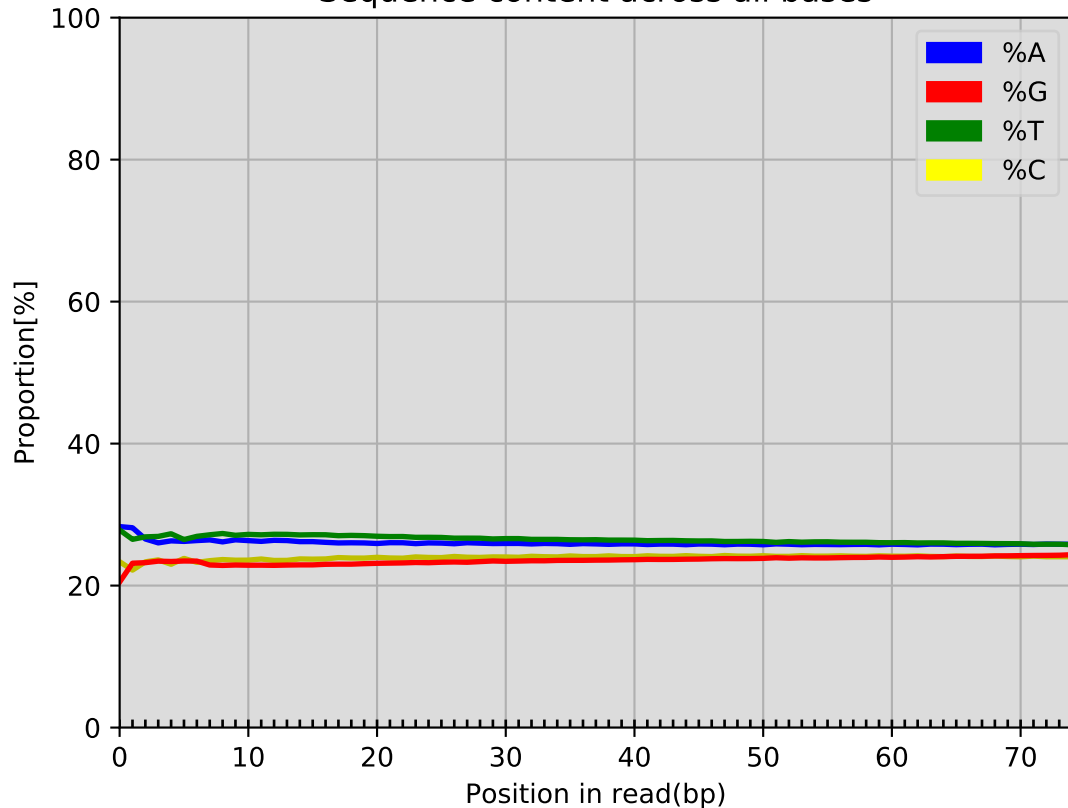**Sequence Length:** 77

%GC: 47.5
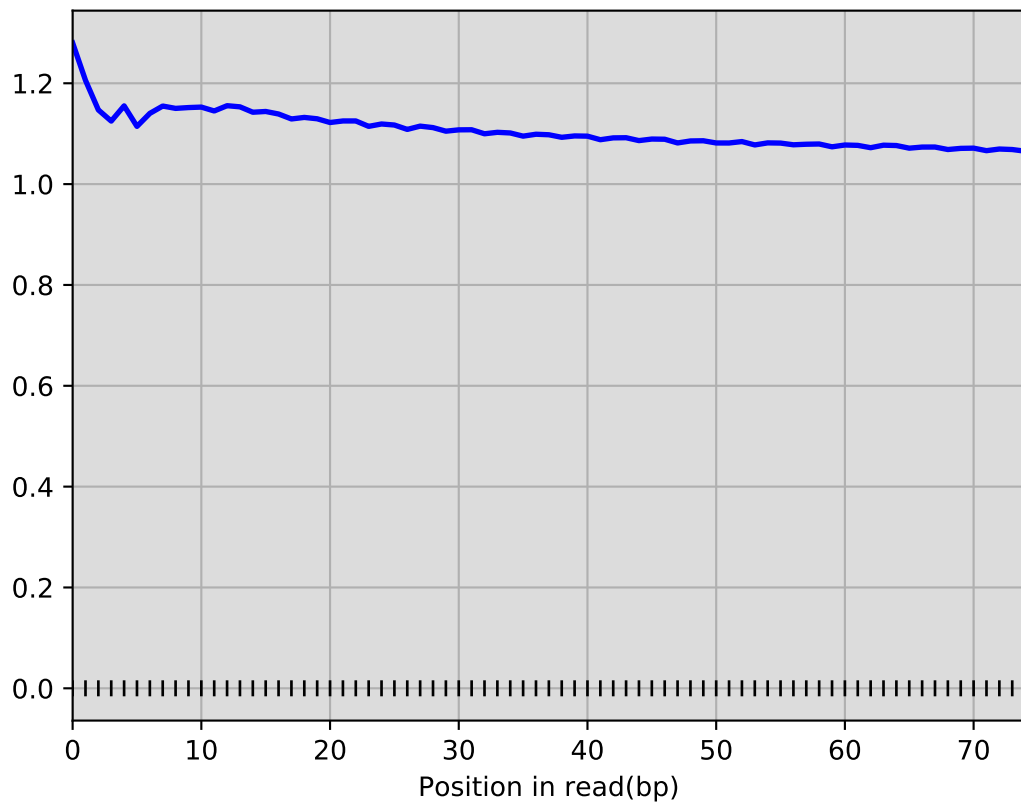
Quality scores across all bases
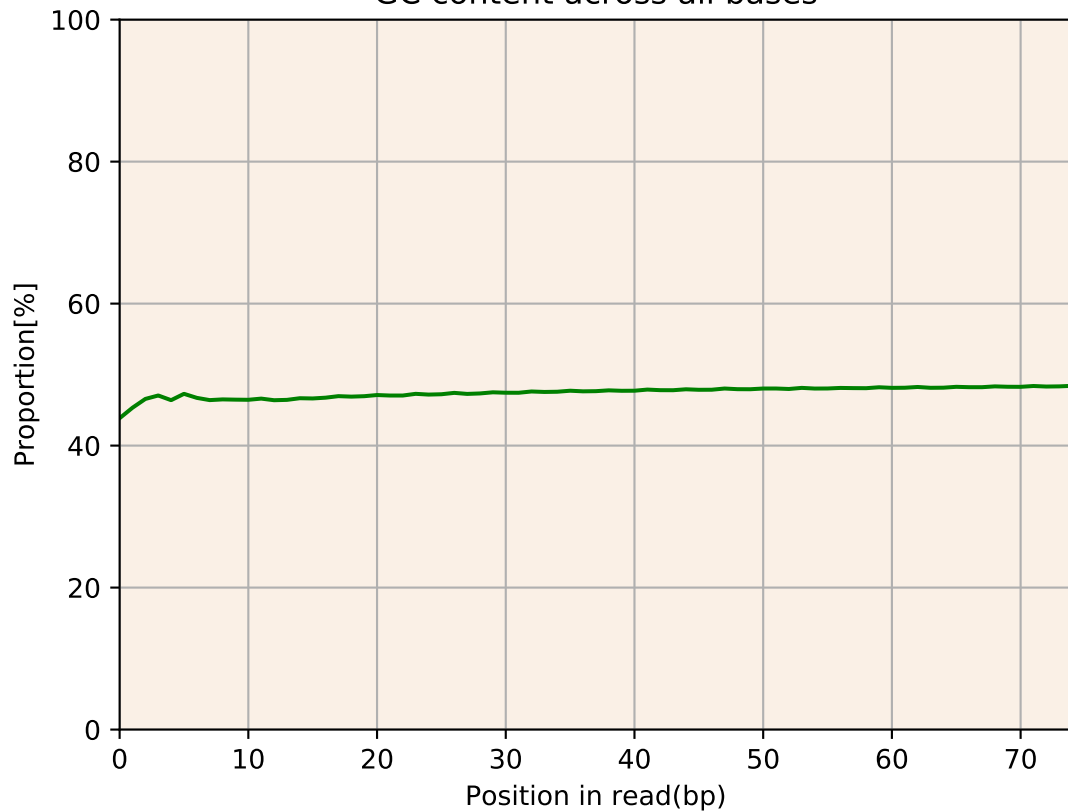
Quality score distribution over all sequences

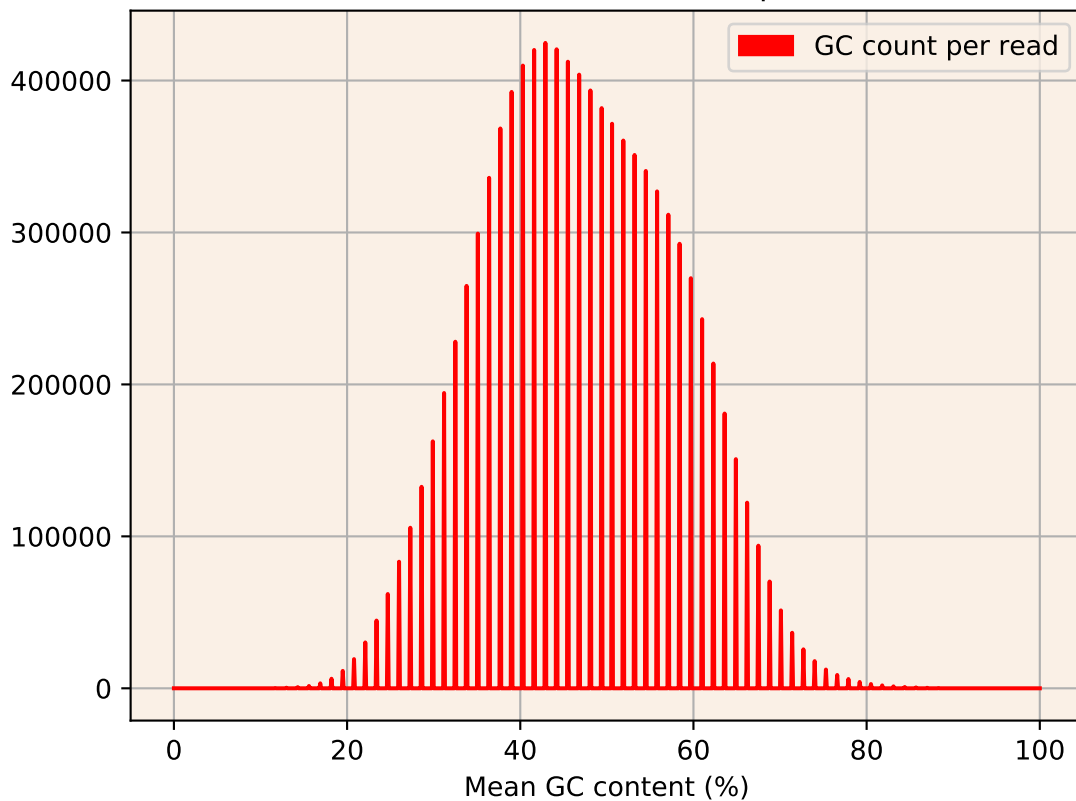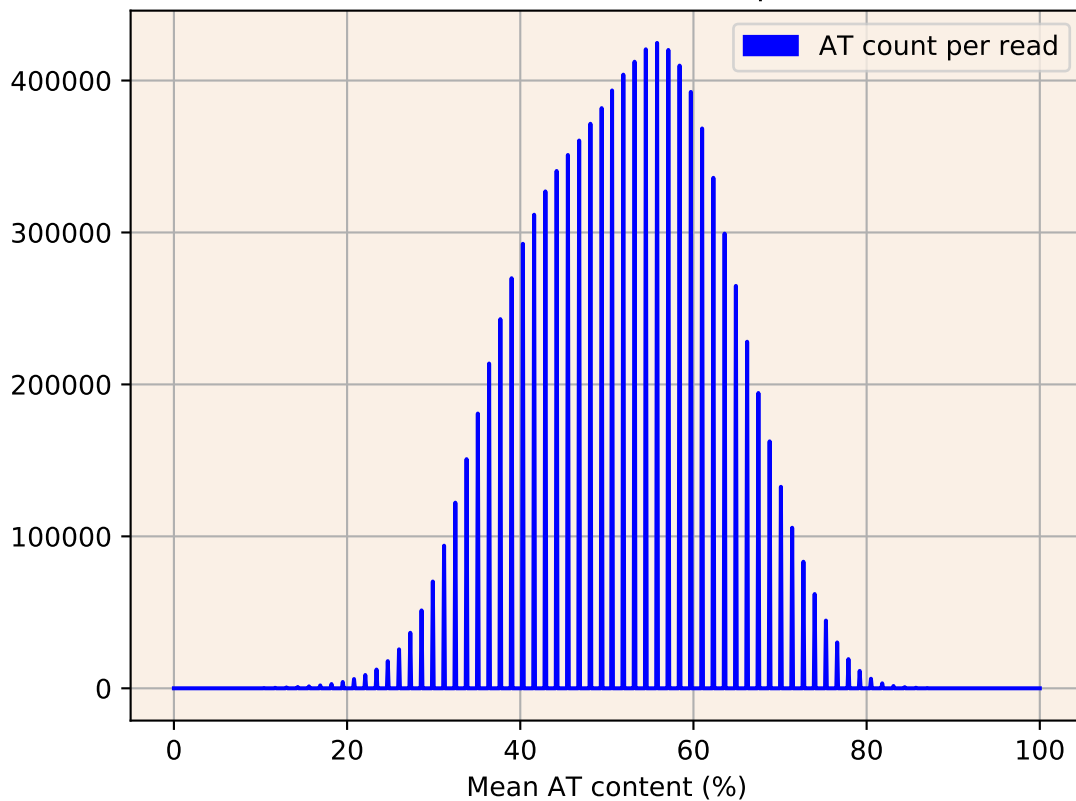Sequence content across all bases

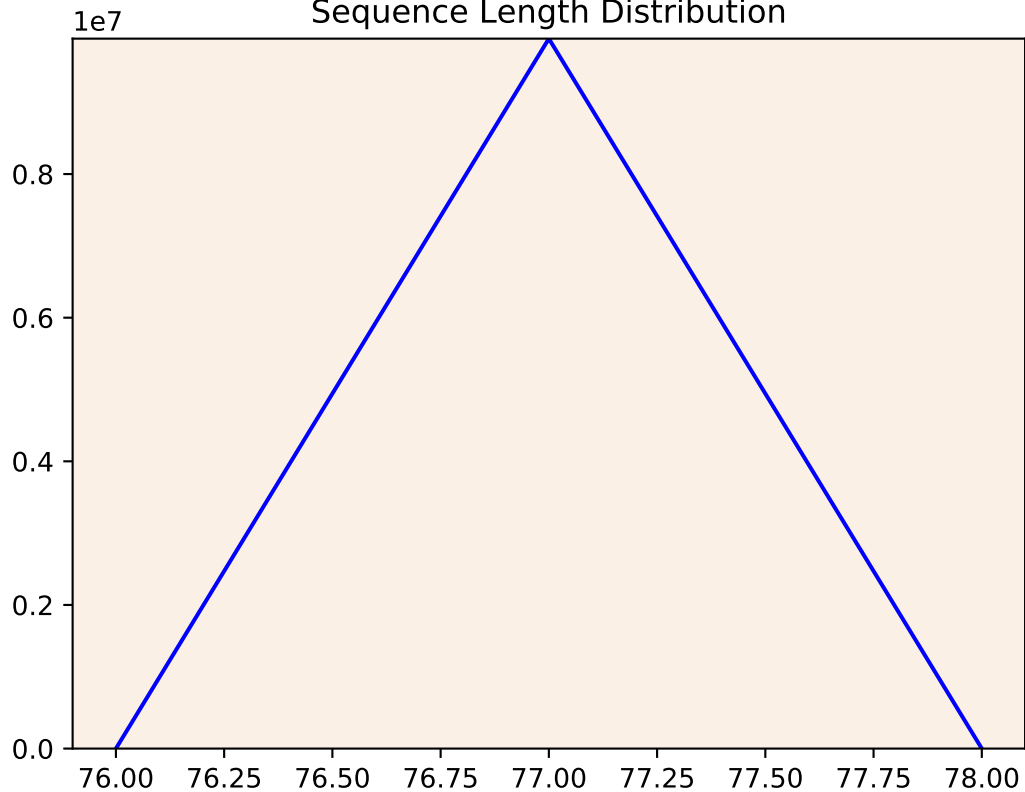AT/GC ratio

GC content across all bases

GC distribution over all sequences

AT distribution over all sequences

Sequence Length Distribution

N content across all bases