Basic Statistics

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

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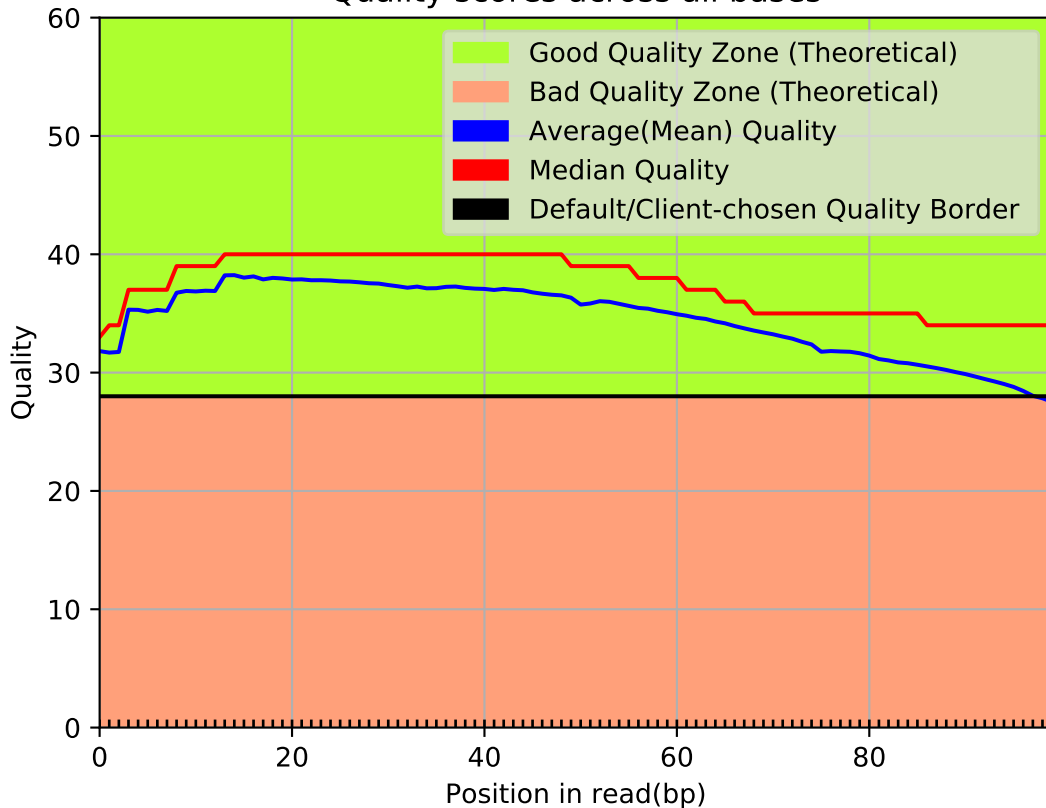
Poor Quality Border: 28

Total Sequences 153213946

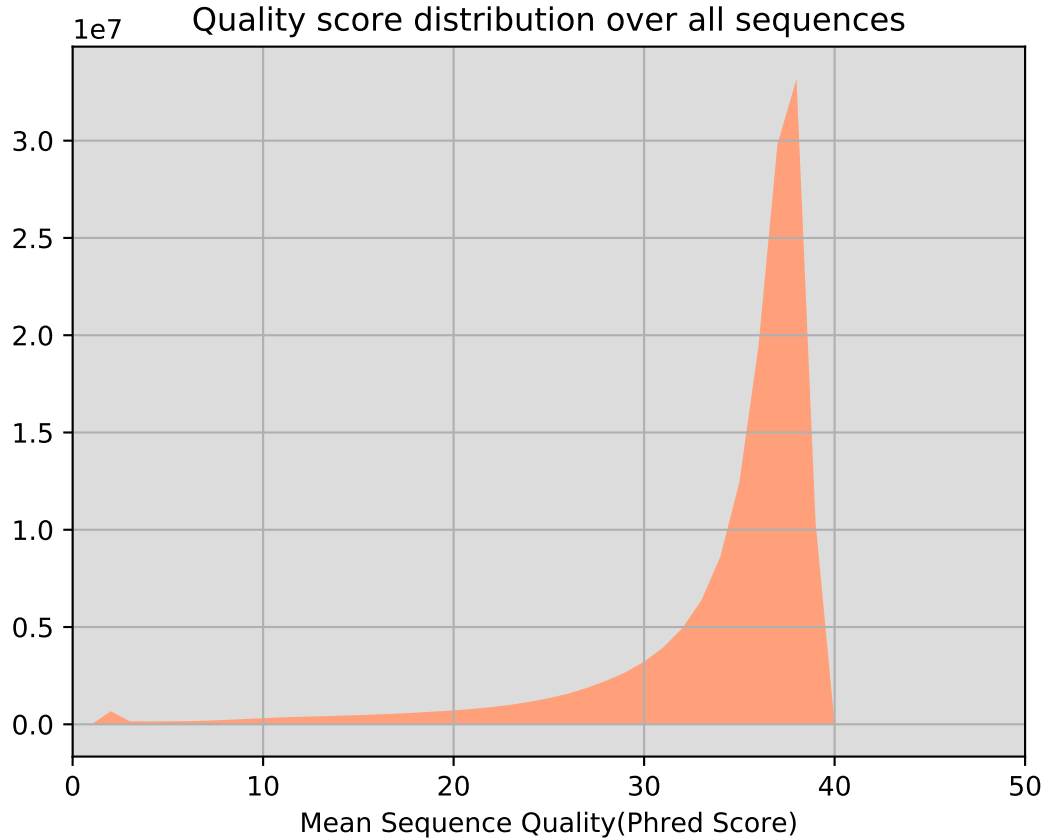**Seq. Flaged as poor Quality:** 15994935

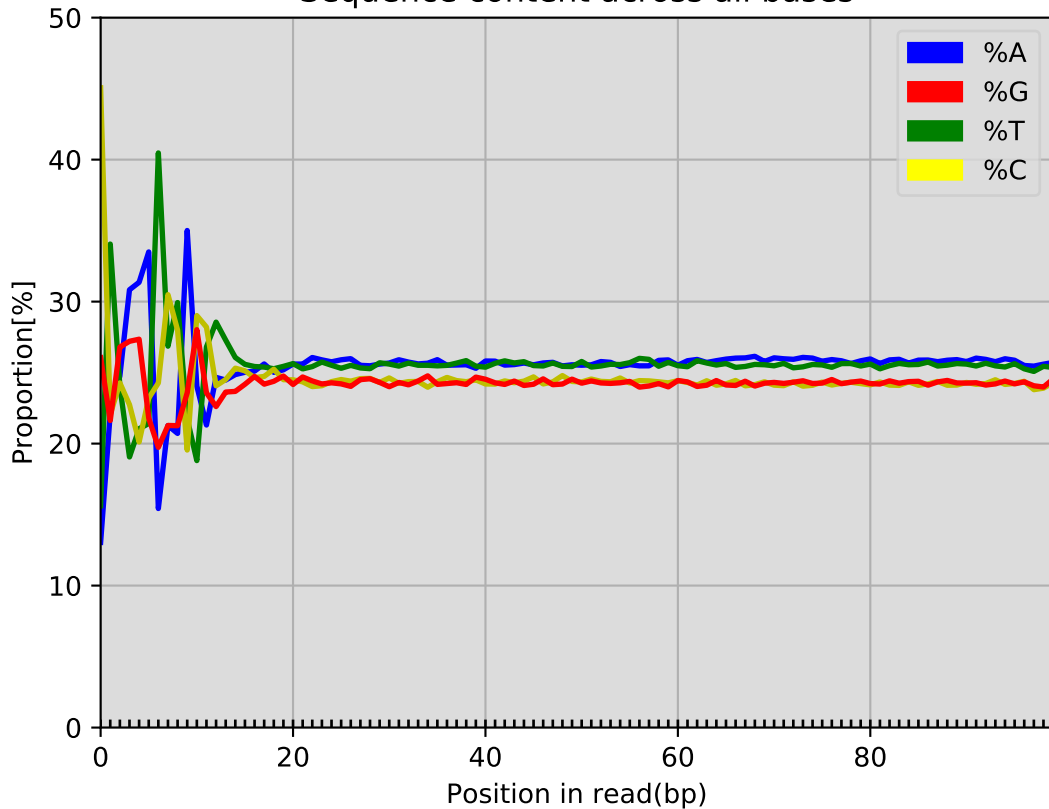**Sequence Length:** 102

%GC: 48.8
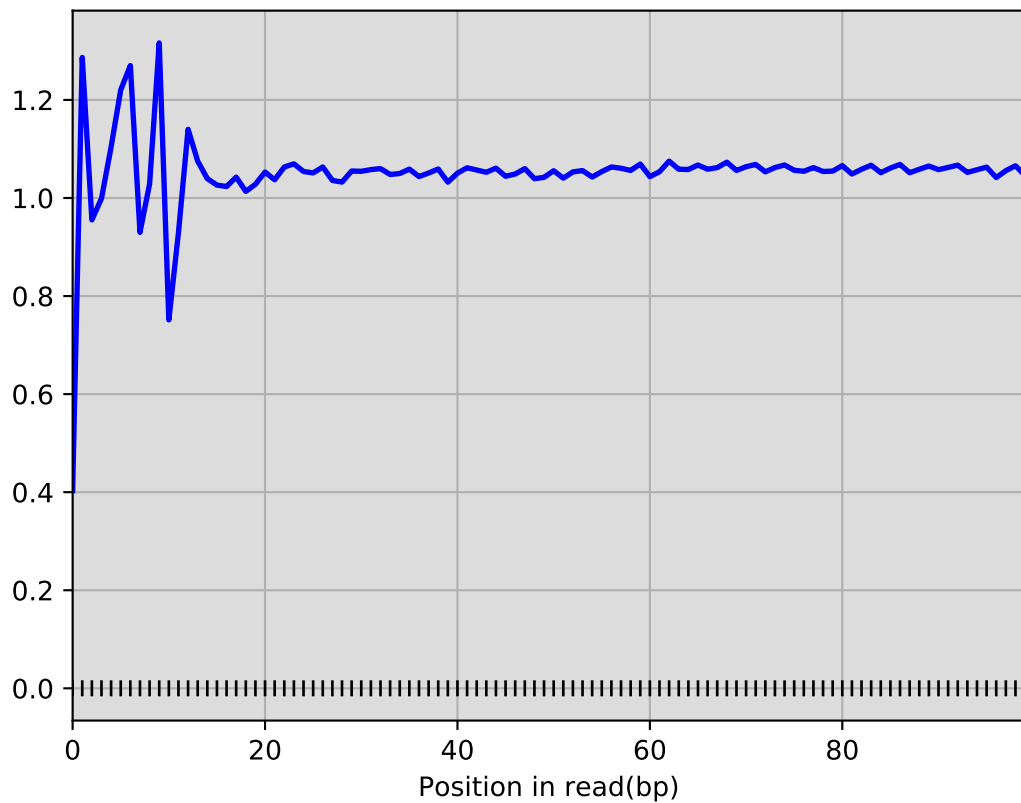
Quality scores across all bases

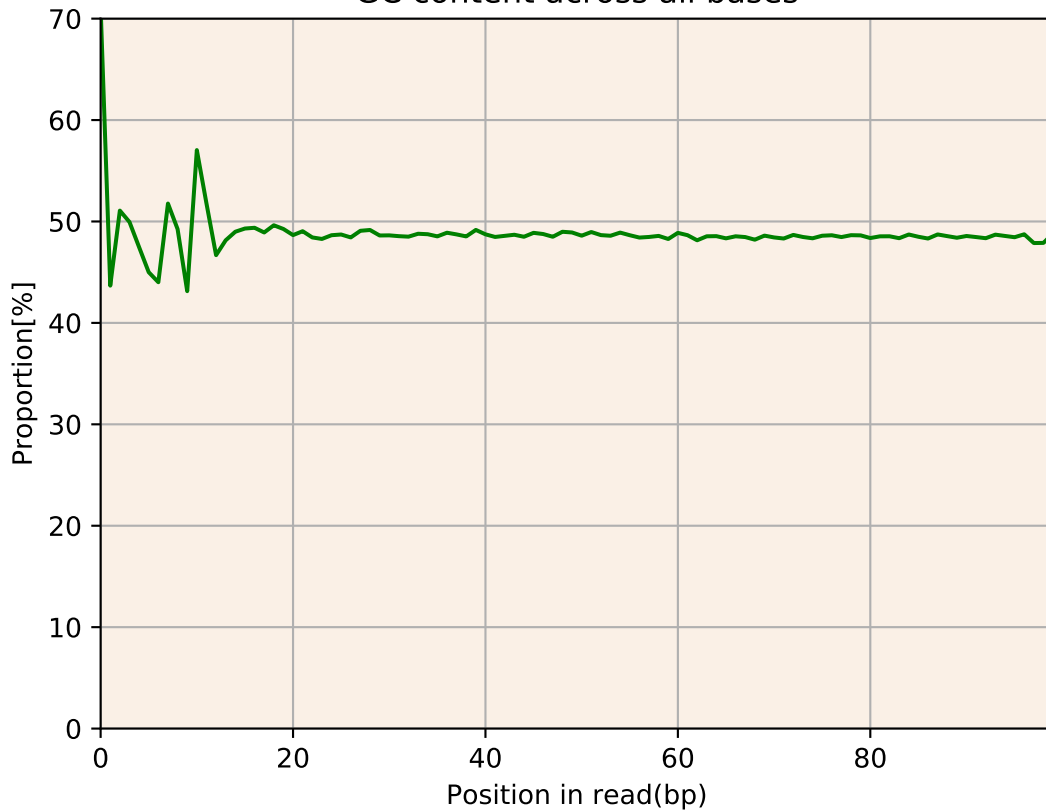Quality score distribution over all sequences

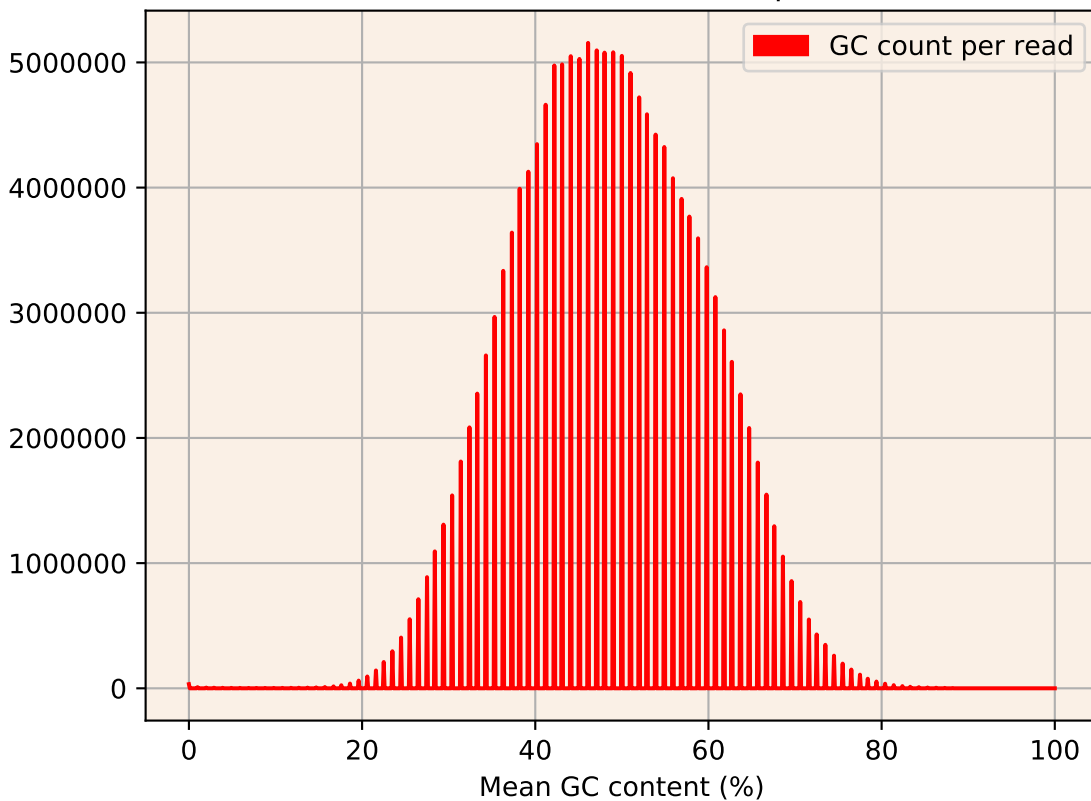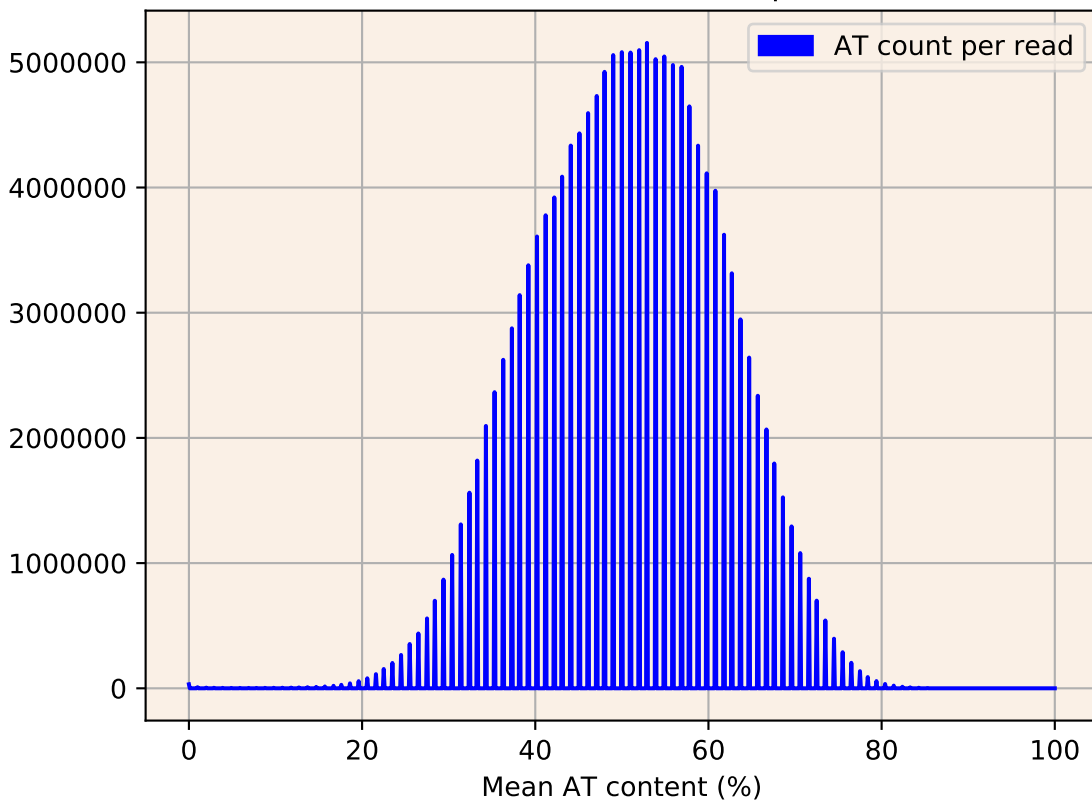Sequence content across all bases

AT/GC ratio

GC content across all bases
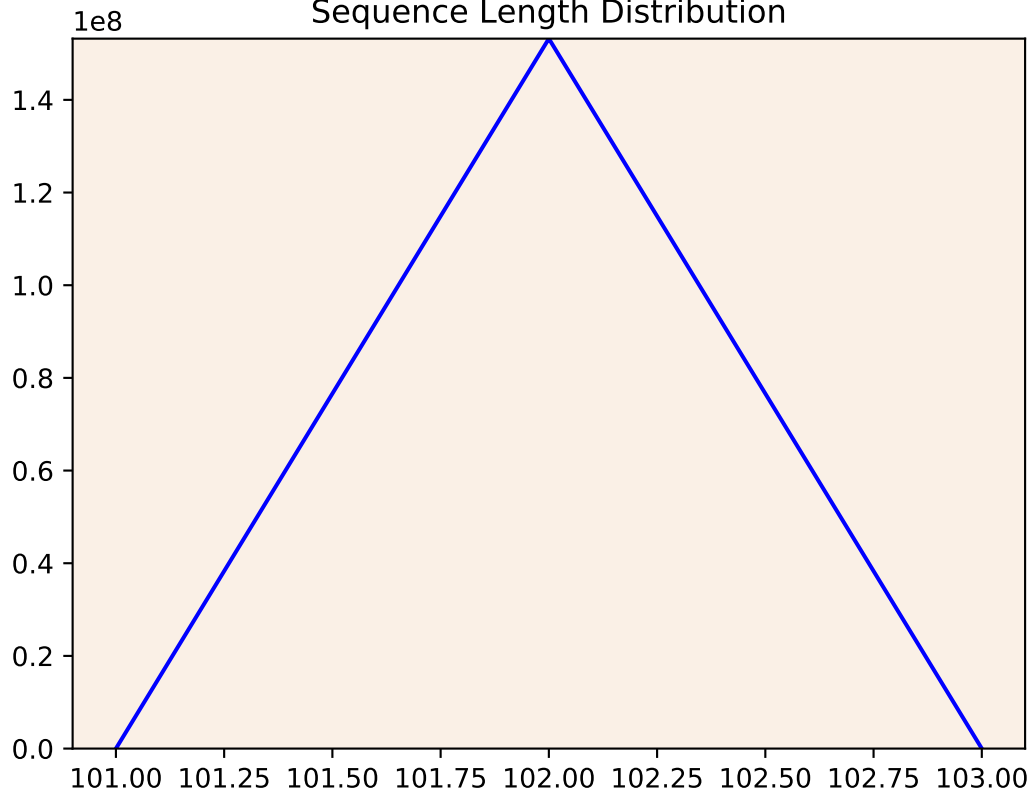
GC distribution over all sequences

AT distribution over all sequences

Sequence Length Distribution

N content across all bases