

Basic Statistics

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

Filename2: /sbgenomics/Projects/10b38e1d-256a-4288-aa1a-5b5775fc35

Poor Quality Border: 28

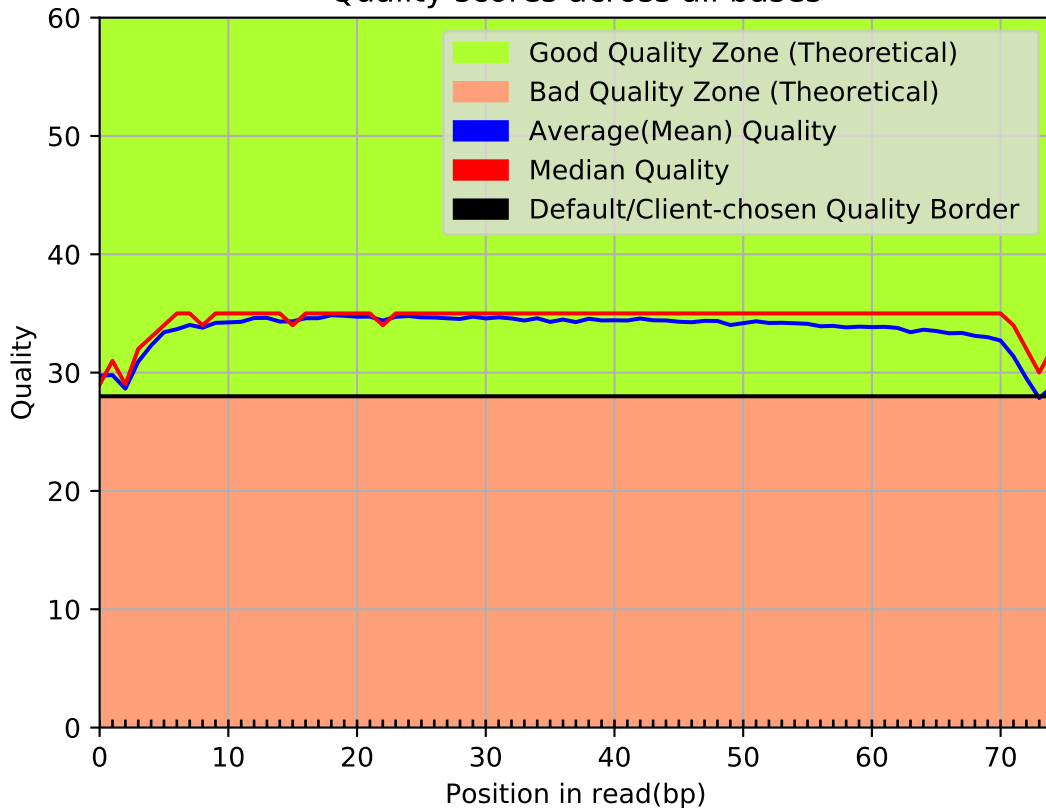
Total Sequences 1632890

Seq. Flaged as poor Quality: 94500

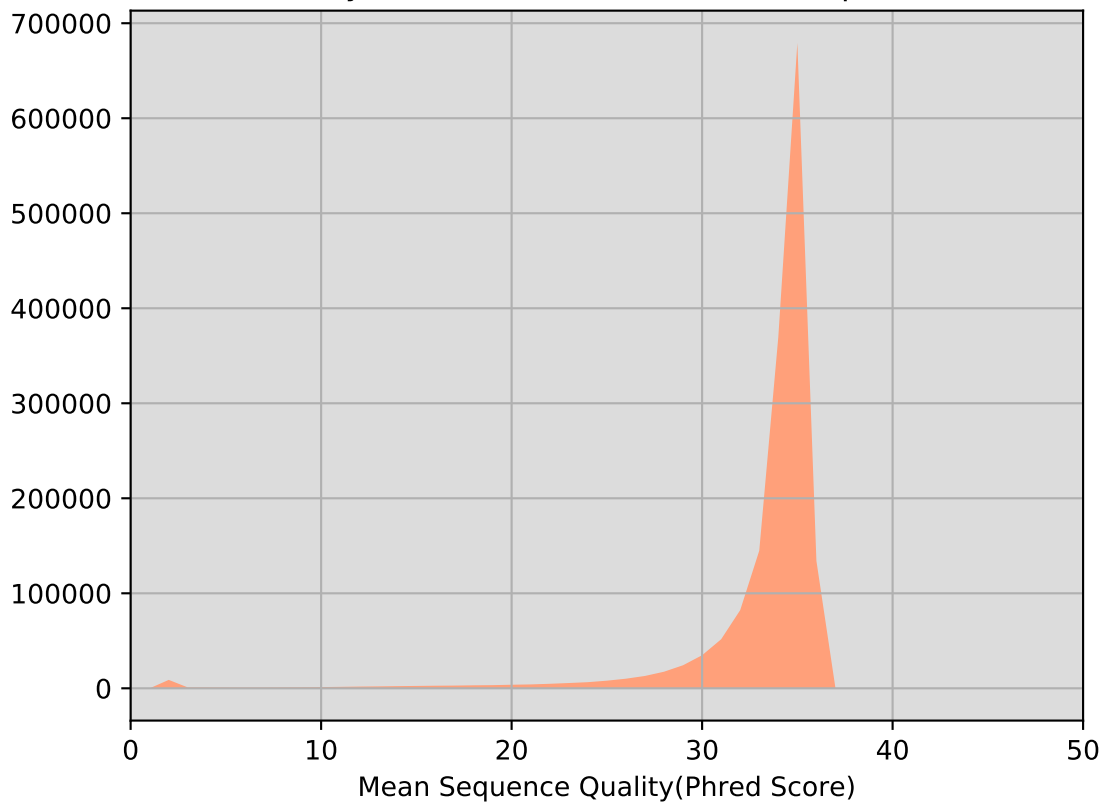
Sequence Length: 77

%GC: 52.4

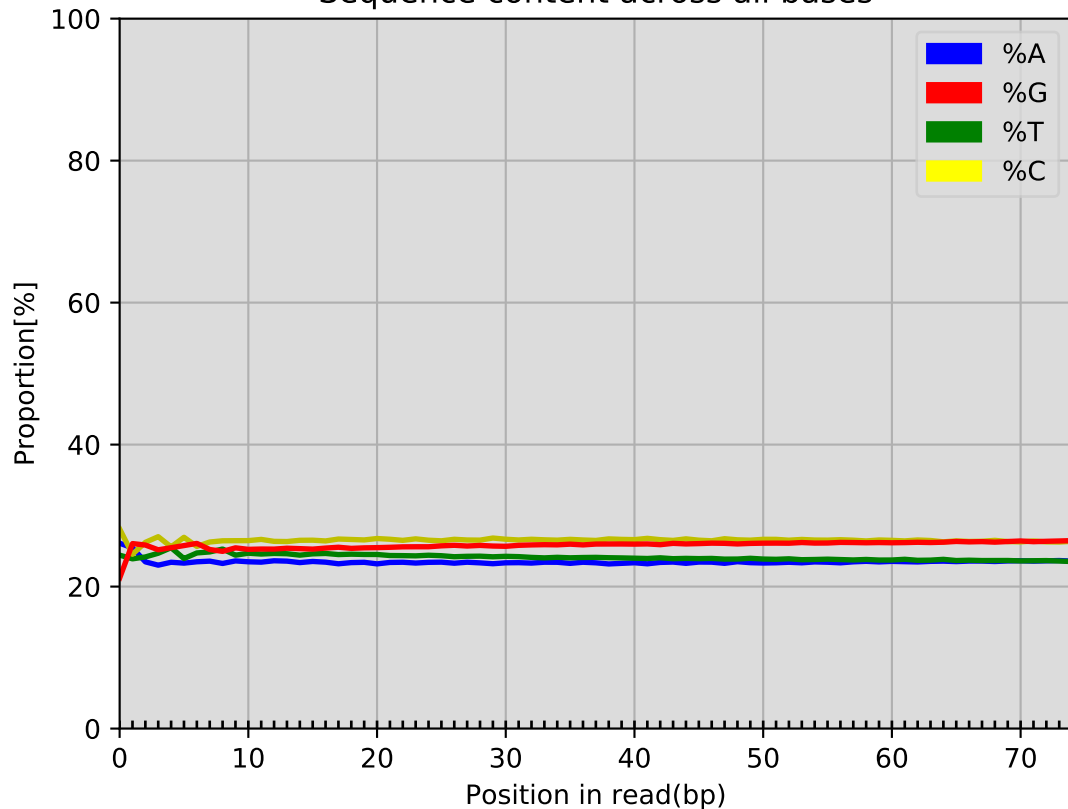
Quality scores across all bases



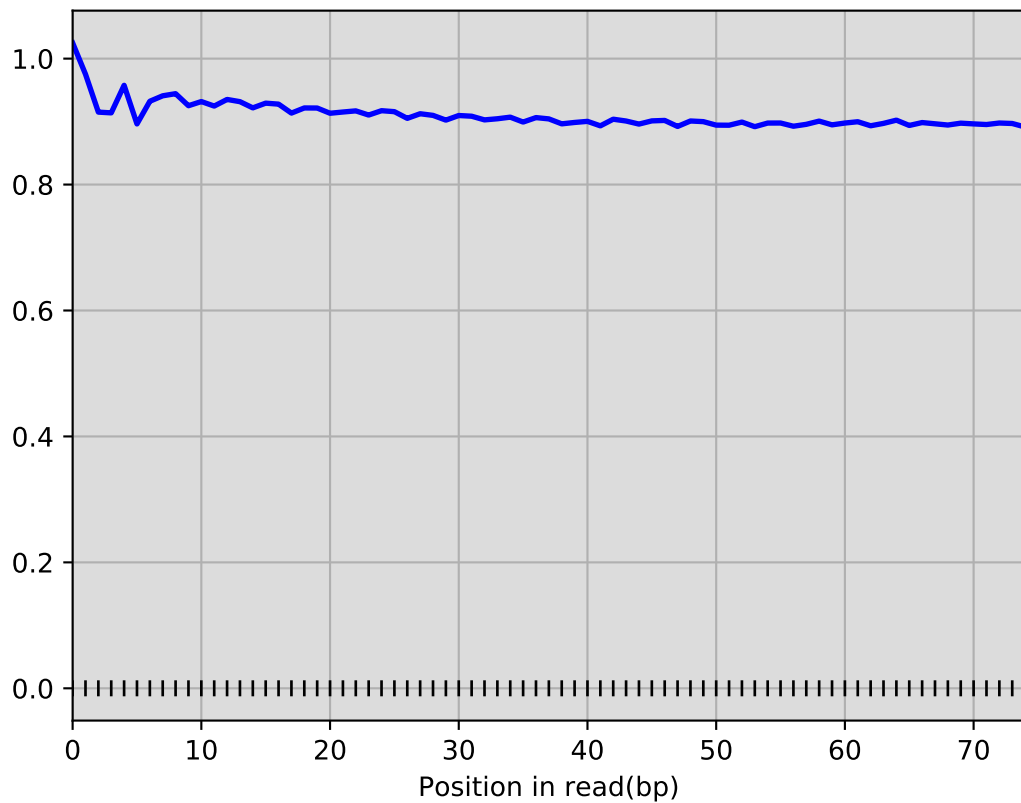
Quality score distribution over all sequences



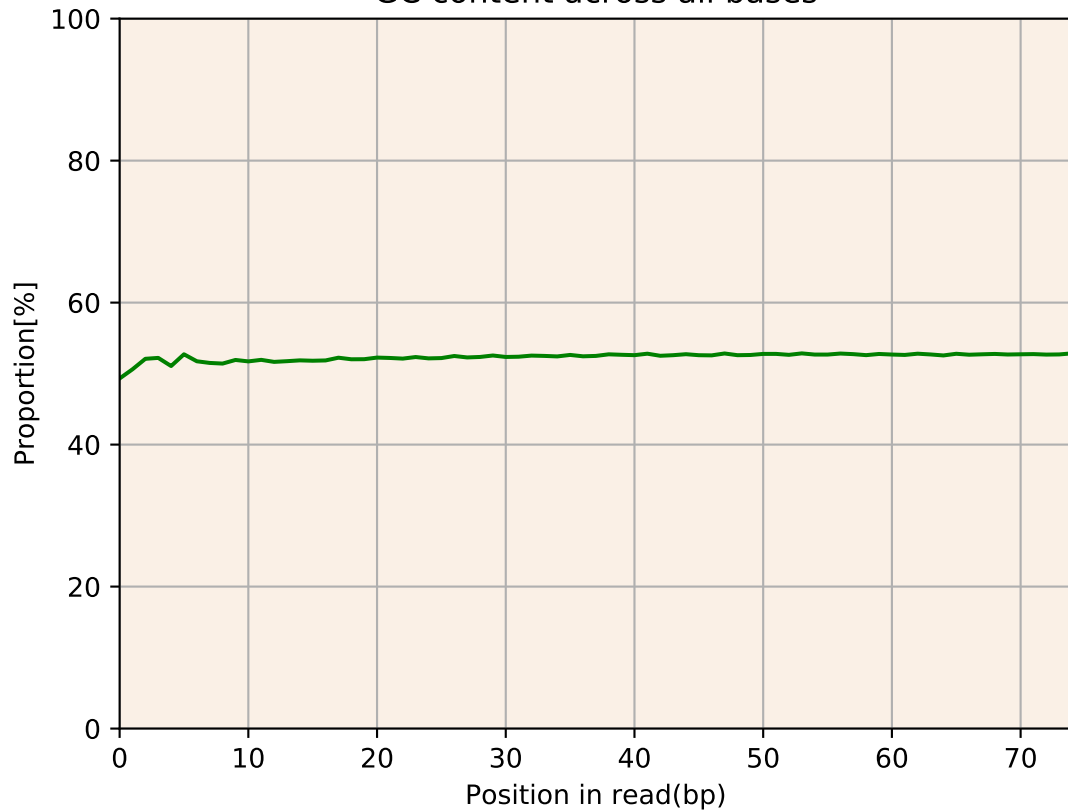
Sequence content across all bases



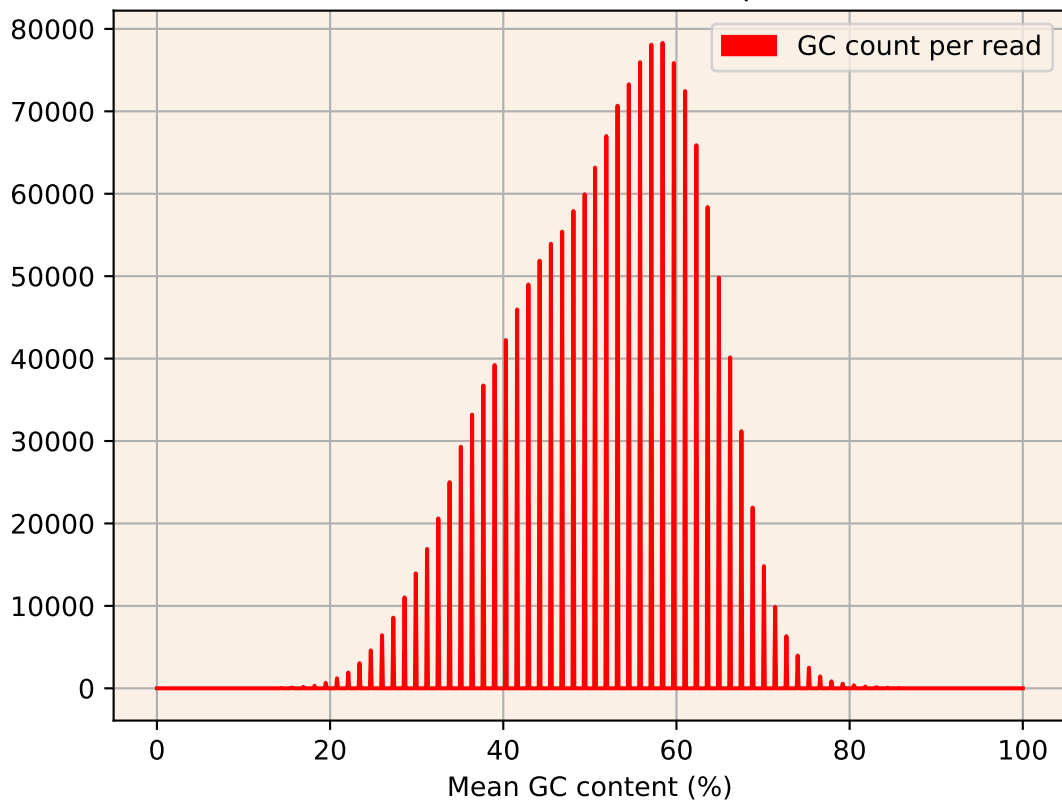
AT/GC ratio



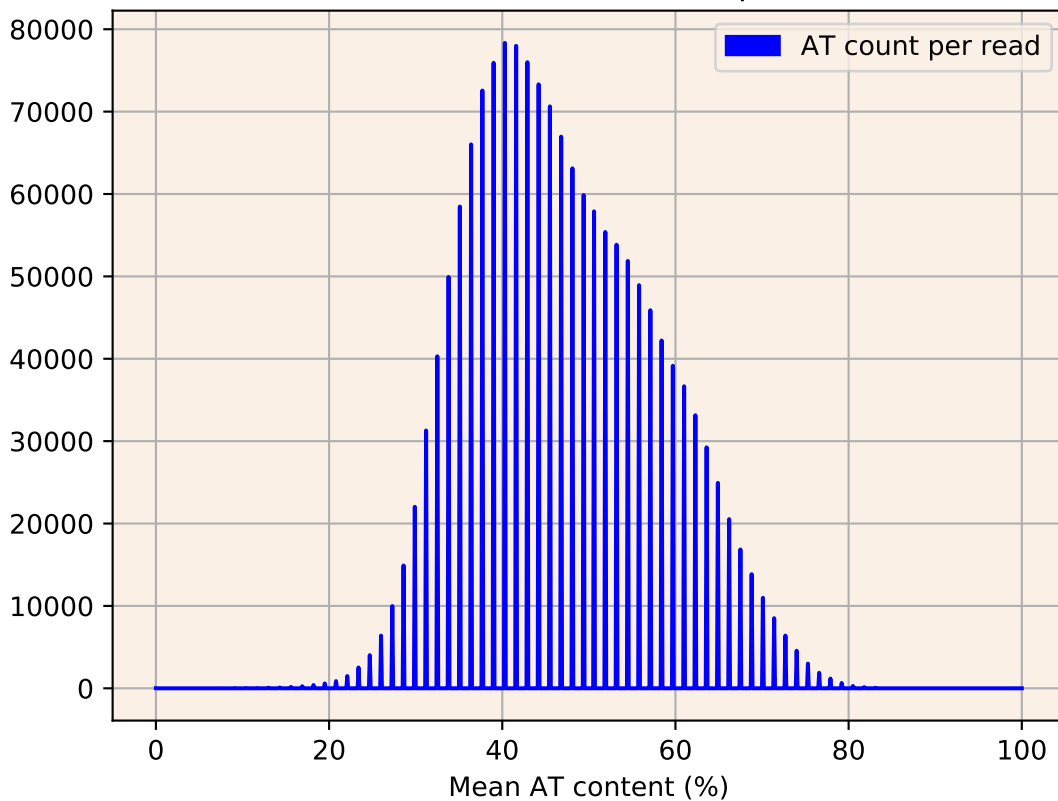
GC content across all bases



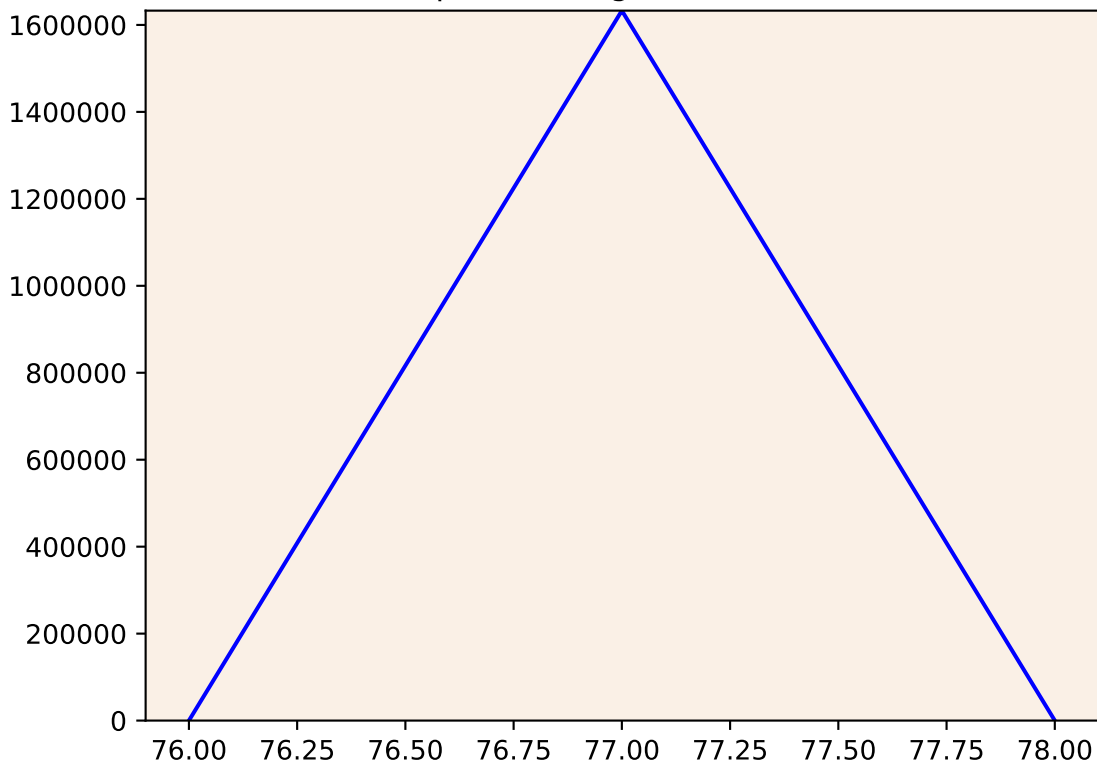
GC distribution over all sequences



AT distribution over all sequences



Sequence Length Distribution



N content across all bases

