

Sample 1 Quality Control

FastQC Report

Mon 28 Mar 2016
7Q-SU+UV_CGCTCATT_L005_R1_001.fastq.gz

Summary

- ✓ Basic Statistics
- ✗ Per base sequence quality
- ✗ Per tile sequence quality
- ✓ Per sequence quality scores
- ✗ Per base sequence content
- ✗ Per sequence GC content
- ✓ Per base N content
- ✓ Sequence Length Distribution
- ✗ Sequence Duplication Levels
- ✗ Overrepresented sequences
- ✓ Adapter Content
- ✗ Kmer Content

✓ Basic Statistics

Measure	Value
Filename	7Q-SU+UV_CGCTCATT_L005_R1_001.fastq.gz
File type	Conventional base calls
Encoding	Sanger / Illumina 1.9
Total Sequences	32874428
Sequences flagged as poor quality	0
Sequence length	51
%GC	61

✗ Per base sequence quality

