Peak Overlap-ALL-H3K4ME3

Stephen Kelly

AGK

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CBK

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EV.

FLV

GH

SPI

SRF

ZGI

All Salliple

FastQC Sample 1

All Samples

Session Information

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

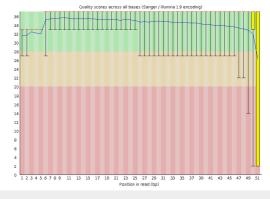
Adapter Content

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



Produced by FastQC (version 0.11.4)