

FastQC: Sequence Counts

16 bars

Unique Reads
Duplicate Reads

5503_clay_control_1
5503_clay_control_2
5503_clay_inoc_1
5503_clay_inoc_2
5503_sandy_control_1
5503_sandy_control_2
5503_sandy_inoc_1
5503_sandy_inoc_2
6007_clay_control_1
6007_clay_control_2
6007_clay_inoc_1
6007_clay_inoc_2
6007_sandy_control_1
6007_sandy_control_2
6007_sandy_inoc_1
6007_sandy_inoc_2

0 5M 10M 15M 20M 25M 30M 35M 40M 45M

Number of reads

