

Run Info

Experiment Name	Lmin010720
Sample ID	Lmin
Run ID	634663b8-21af-45fc-a2ec-9ec6f01975fd
Flow Cell Id	FAO01594
Start Time	July 1, 18:21
Run Length	20h 48m

Run Summary

Reads Generated	1.73 M
Bases Generated	4.09 Gb
Estimated Bases	3.75 Gb

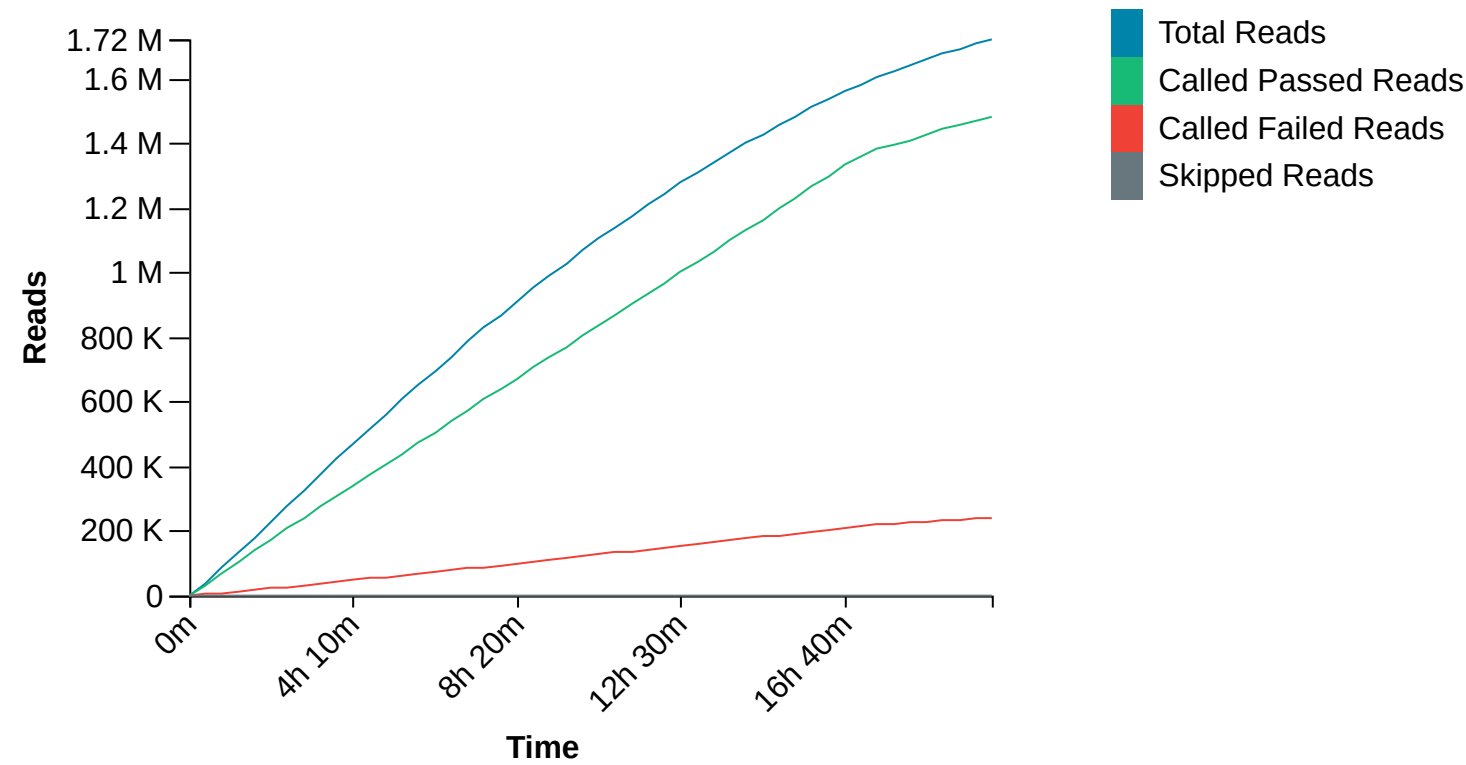
Run Parameters

Flow Cell Type	FLO-MIN106
Kit	SQK-LSK109
Basecalling	on
Specified Run Length	72 hours
Initial Bias Voltage	-185 mV
FAST5 Output	Enabled
FAST5 Output Options	zlib_compress,fastq,raw
FAST5 Reads per File	4000
FASTQ Output	Enabled
FASTQ Reads per File	4000
Active Channel Selection	Enabled
Mux Scan Period	1 hour 30 minutes
Reserved Pores	0 %
Basecall Model	Fast basecalling

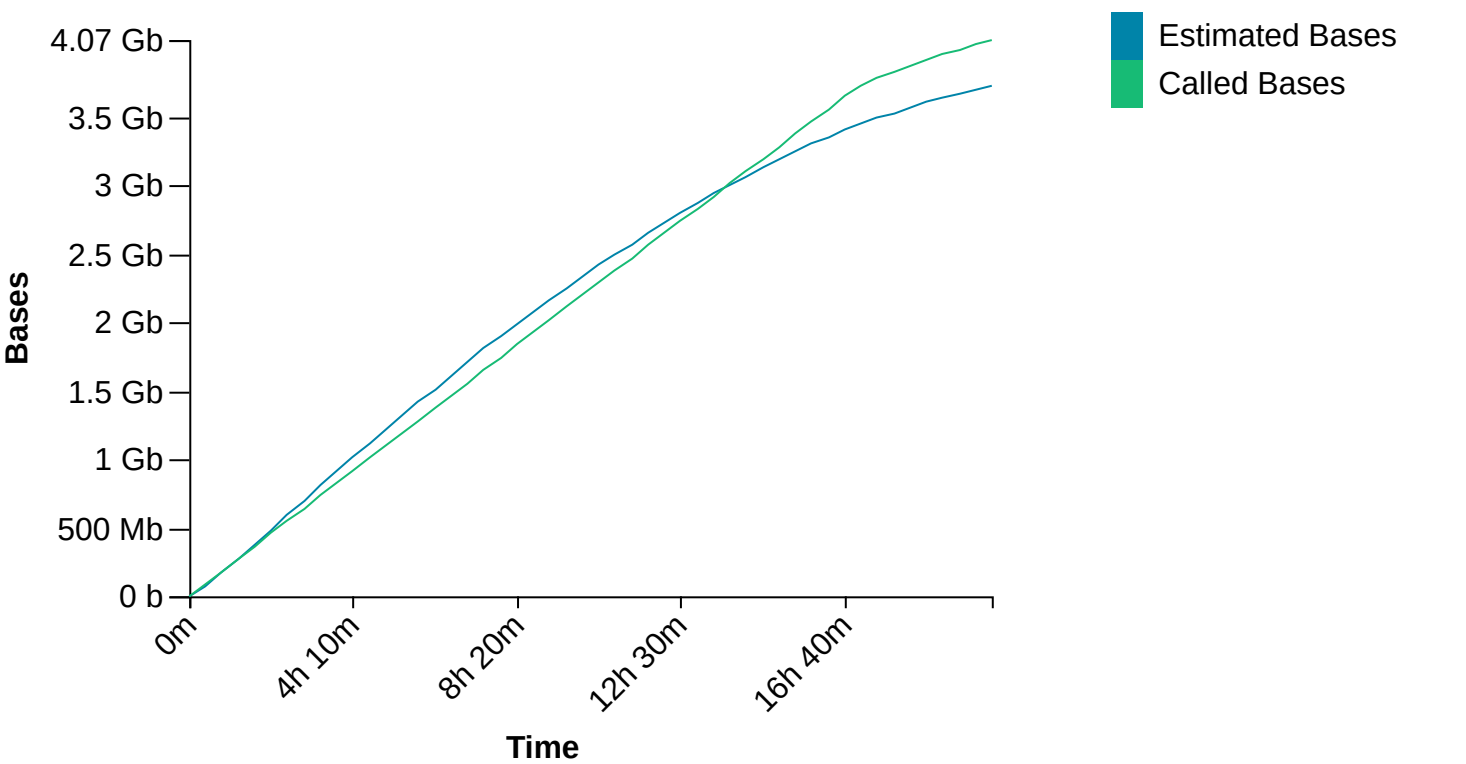
Versions

MinKNOW Core	3.6.5
Bream	4.3.16
Guppy	3.2.10

Cumulative Output Reads

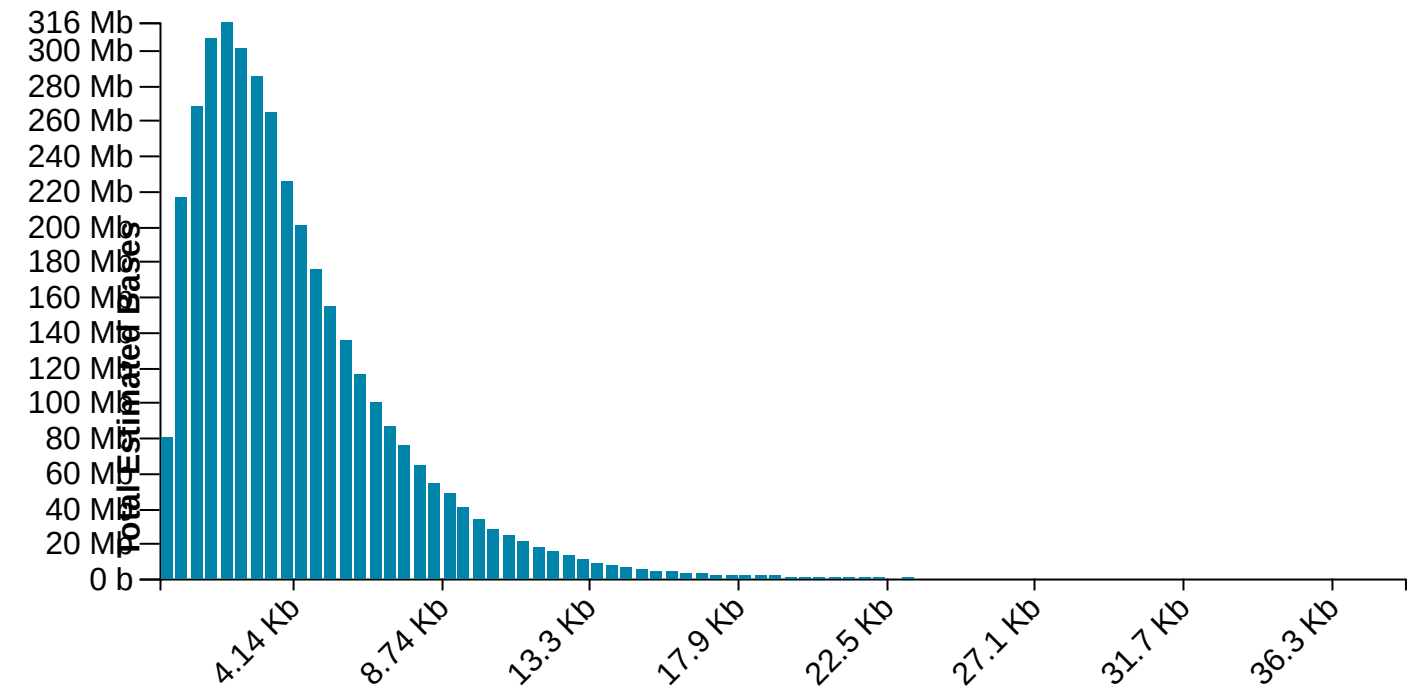


Cumulative Output Bases



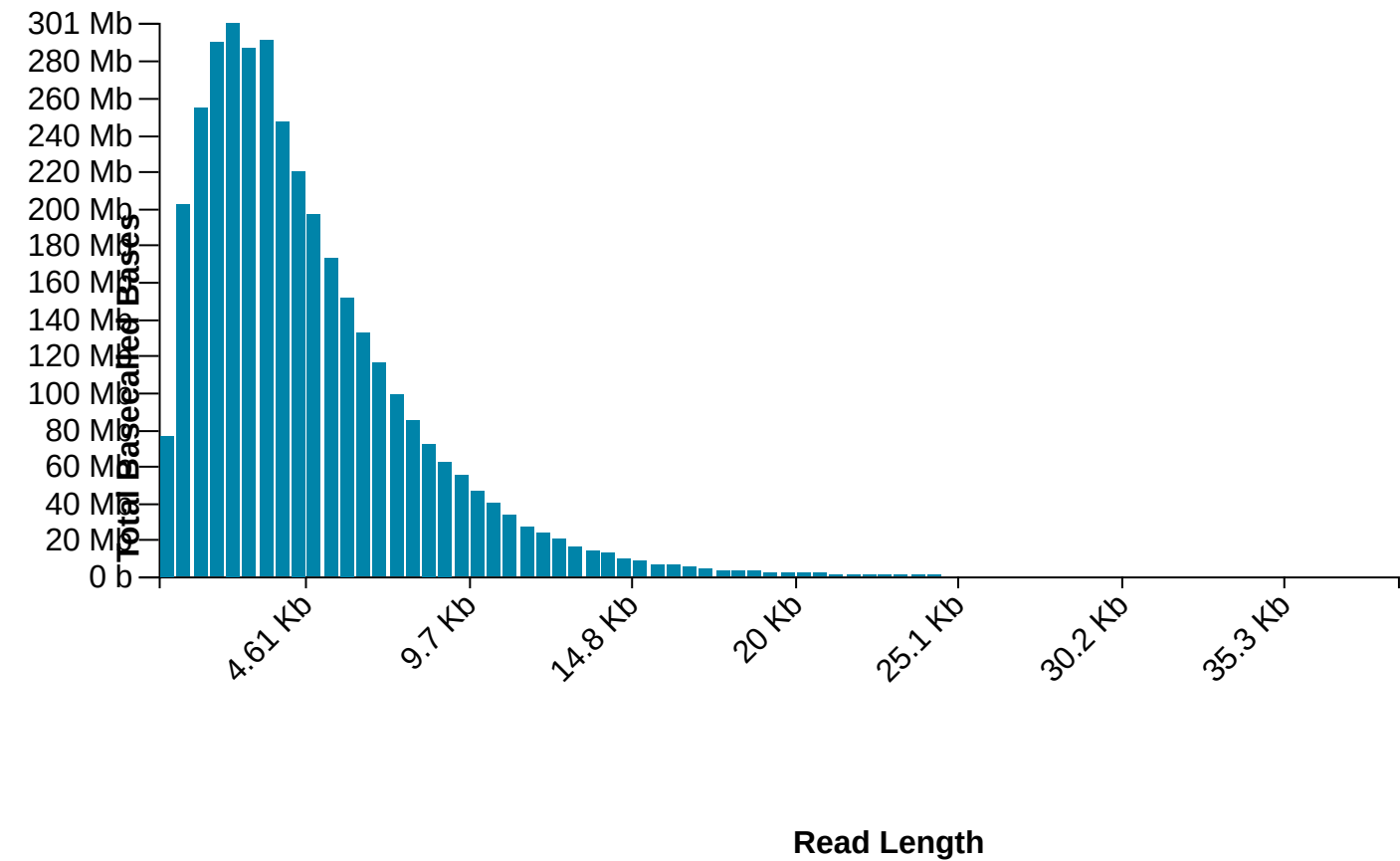
Read Length Histogram Estimated Bases

Estimated N50: 3.39 Kb

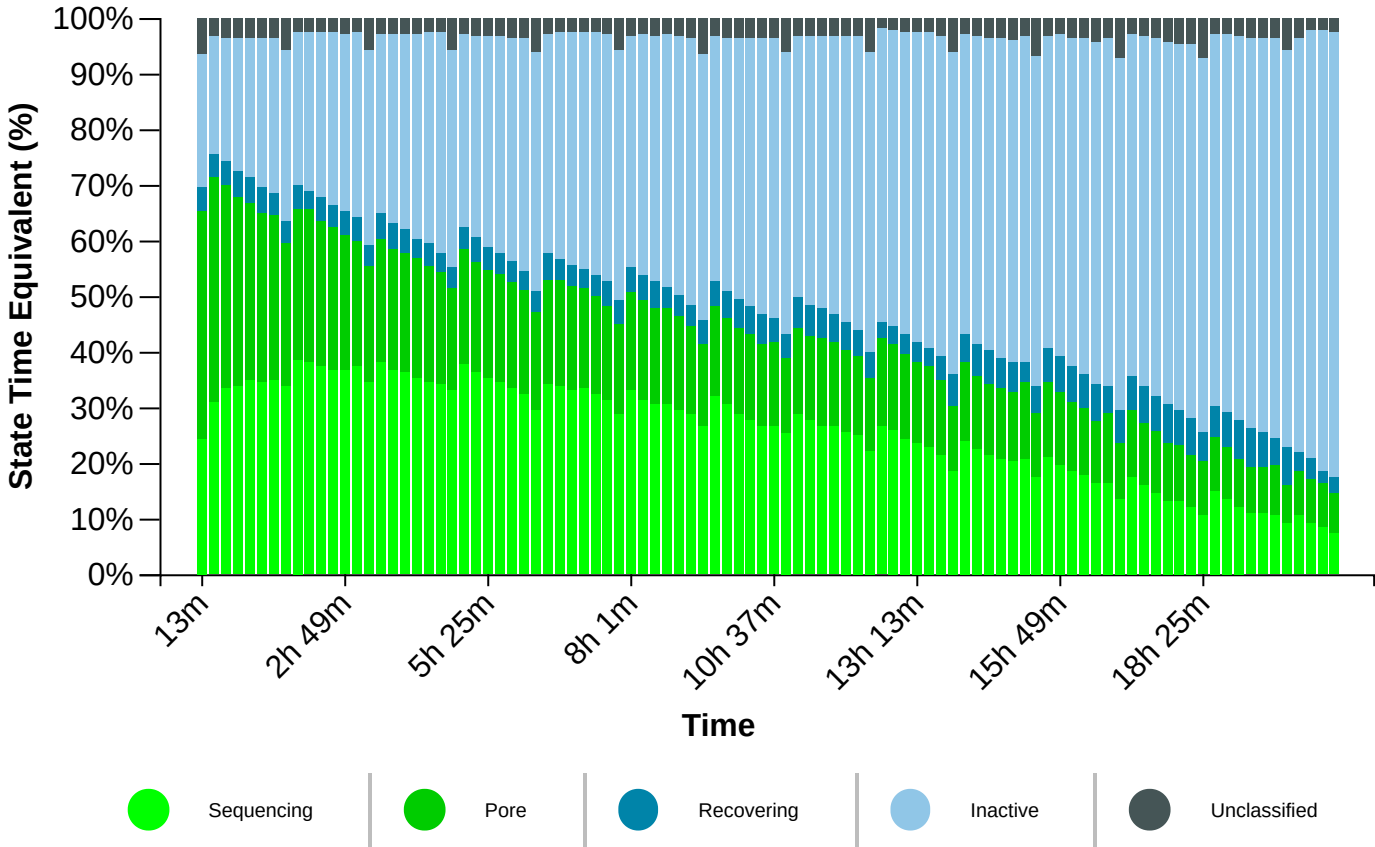


Read Length Histogram Basecalled Bases

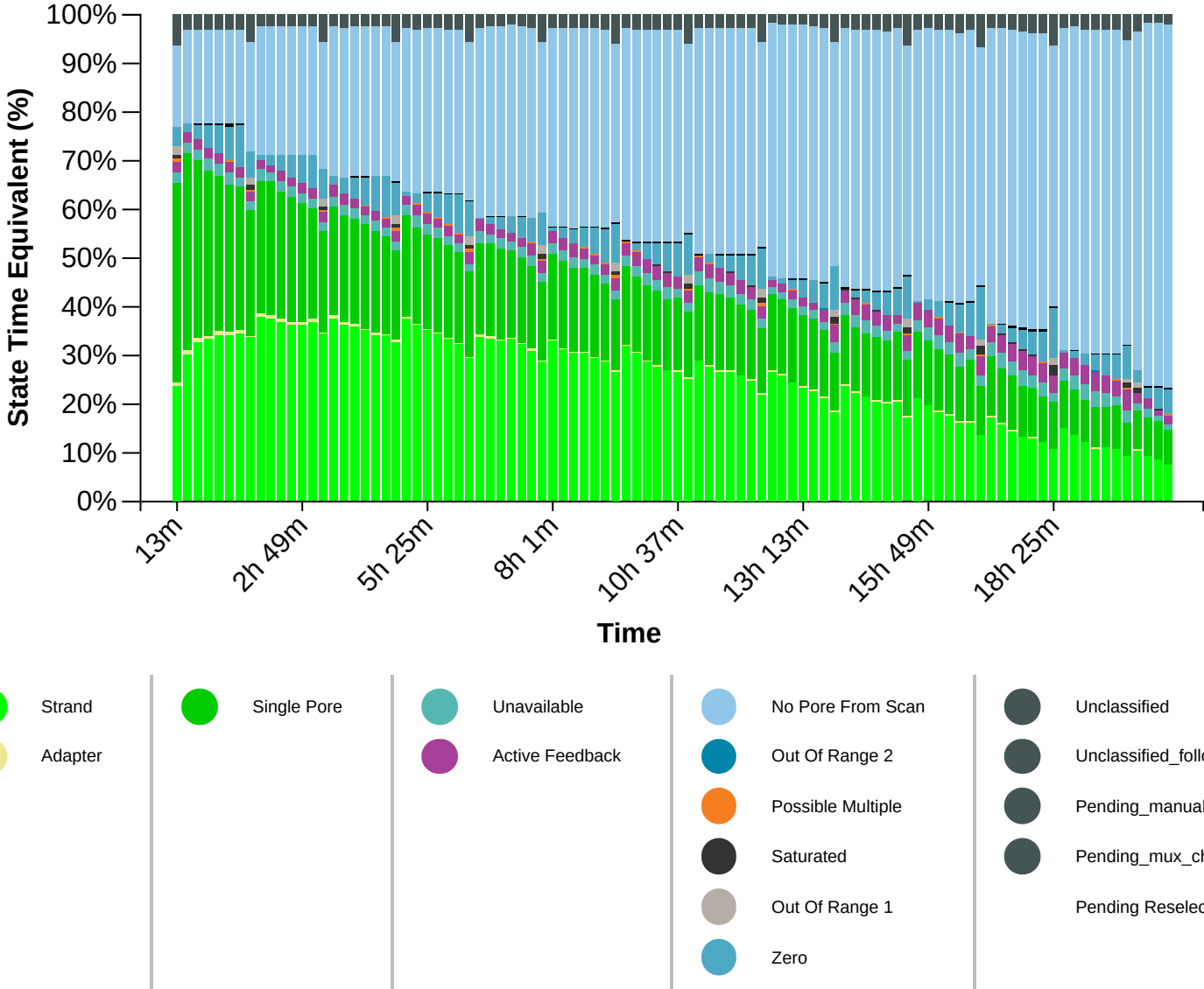
Estimated N50: 3.8 Kb



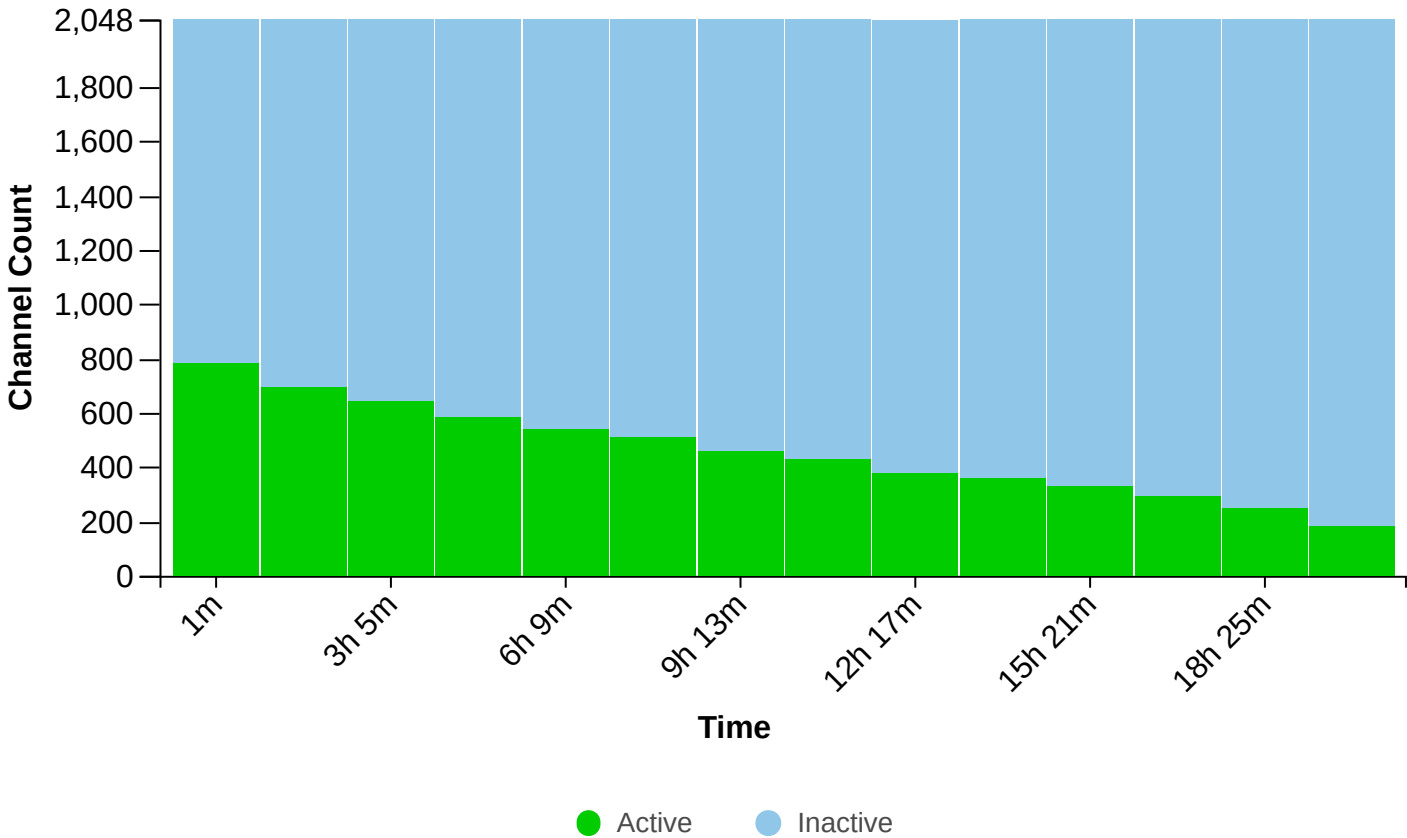
Duty Time Grouped



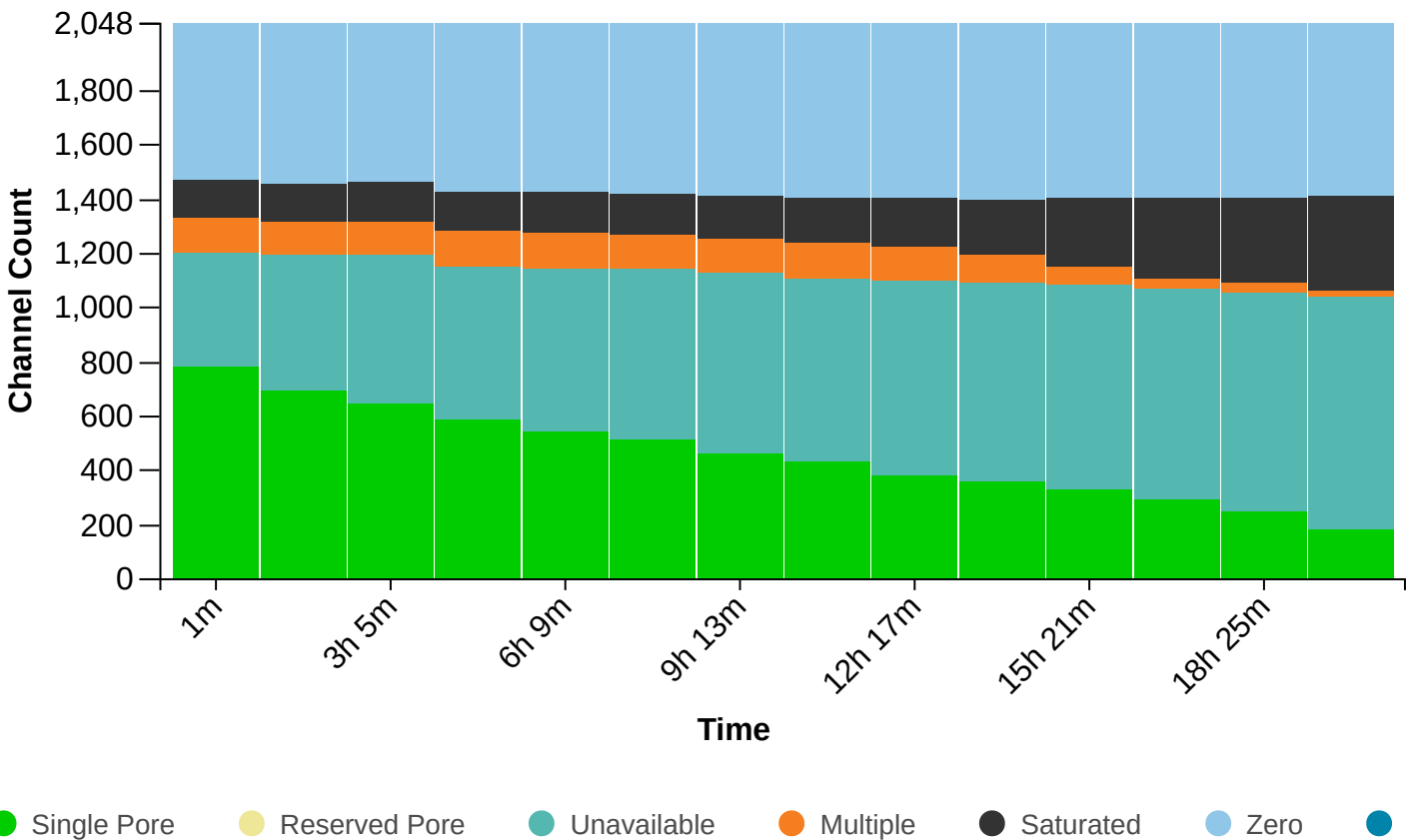
Duty time Categorised



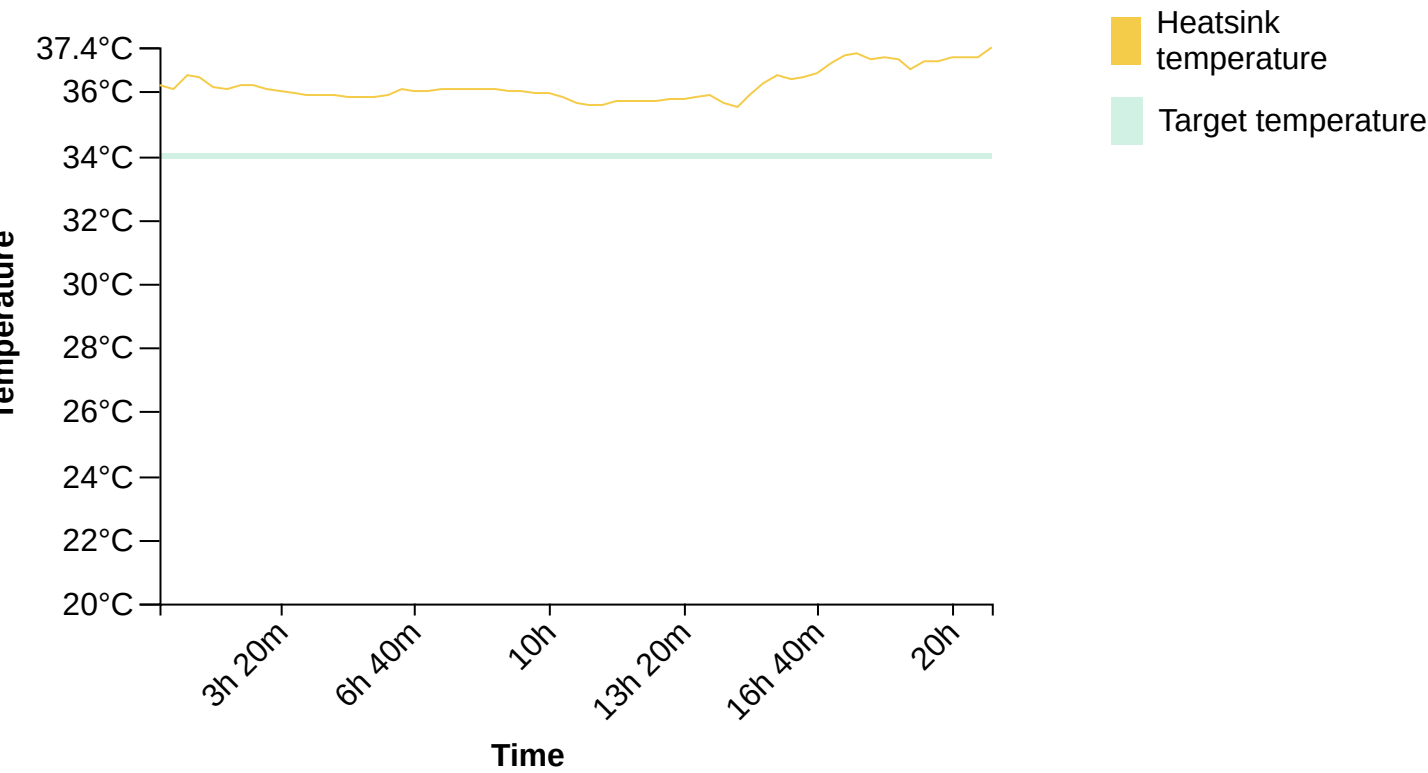
Mux Scan Grouped



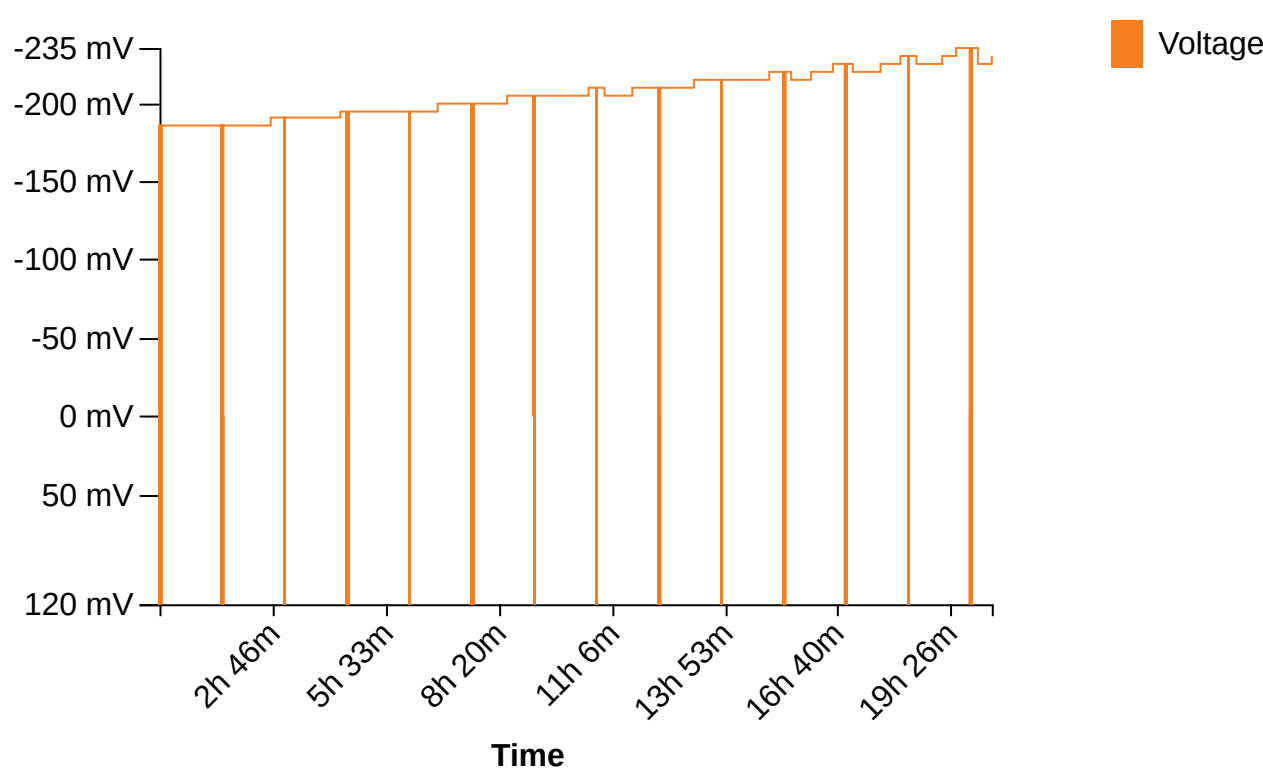
Mux Scan Categorised



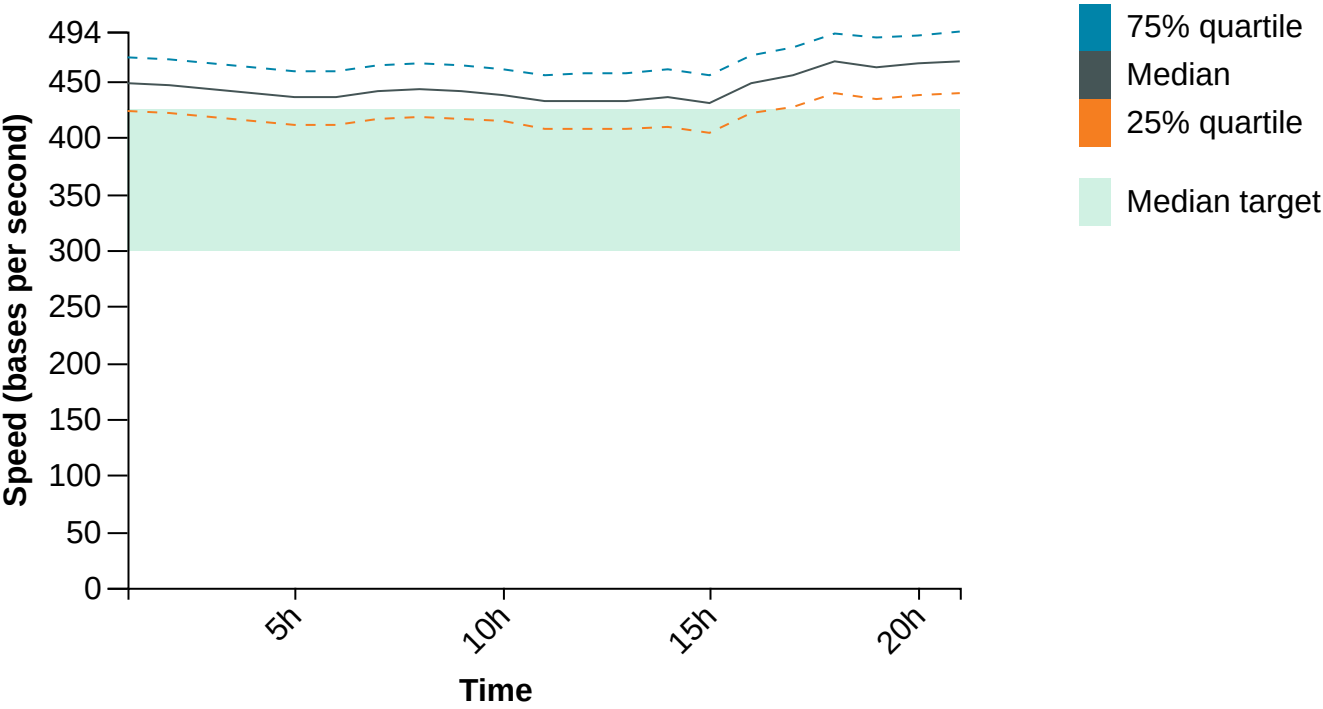
Temperature History



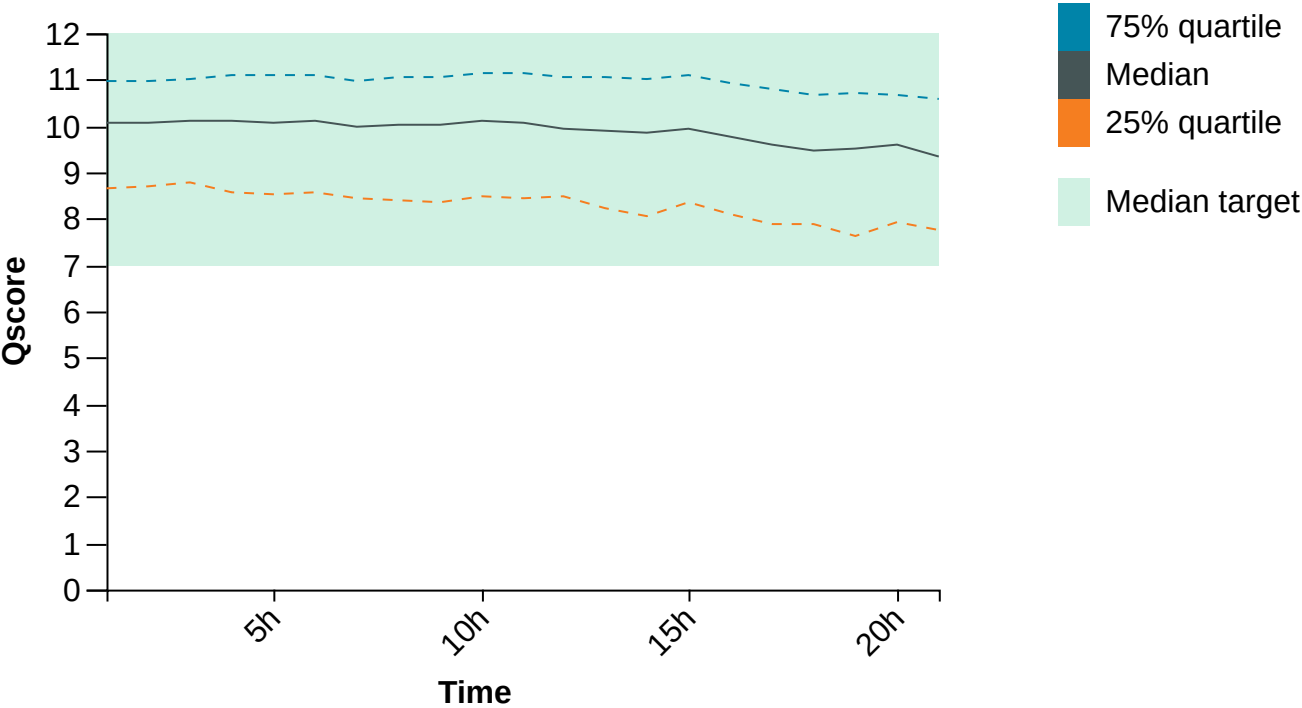
Bias Voltage History



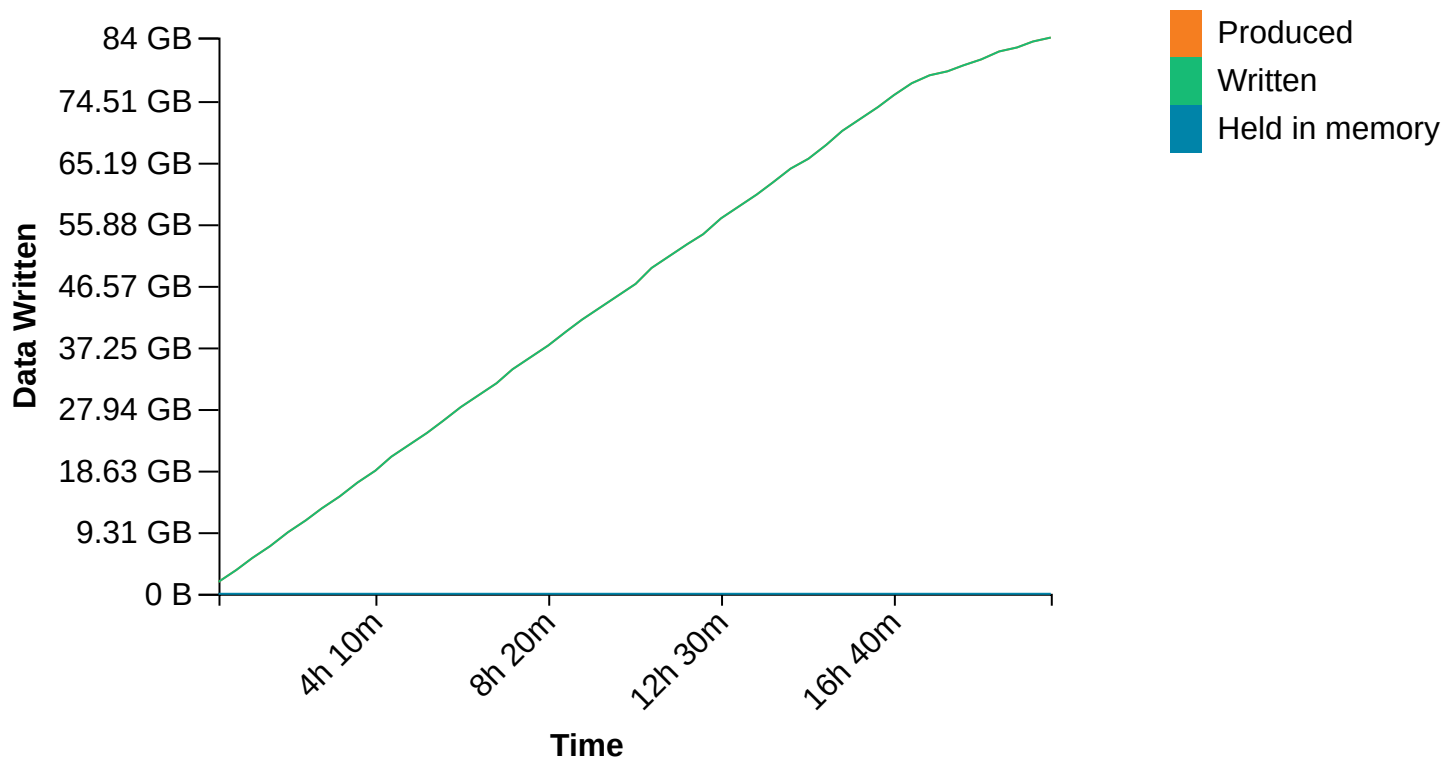
Translocation Speed



QScore



Disk Write Performance



Run Debug Messages

- Flow cell FAO01594 has 184 pores available for sequencing. Starting sequencing with 129 pores July 2, 14:23
- Performing Mux Scan July 2, 14:21
- Flow cell FAO01594 has 250 pores available for sequencing. Starting sequencing with 172 pores July 2, 12:51
- Performing Mux Scan July 2, 12:49
- Flow cell FAO01594 has 293 pores available for sequencing. Starting sequencing with 201 pores July 2, 11:19
- Performing Mux Scan July 2, 11:17
- Flow cell FAO01594 has 329 pores available for sequencing. Starting sequencing with 227 pores July 2, 09:47
- Performing Mux Scan July 2, 09:46
- Flow cell FAO01594 has 358 pores available for sequencing. Starting sequencing with 239 pores July 2, 08:15
- Performing Mux Scan July 2, 08:14
- Flow cell FAO01594 has 384 pores available for sequencing. Starting sequencing with 245 pores July 2, 06:44
- Performing Mux Scan July 2, 06:42
- Flow cell FAO01594 has 435 pores available for sequencing. Starting sequencing with 275 pores July 2, 05:12
- Performing Mux Scan July 2, 05:10
- Flow cell FAO01594 has 461 pores available for sequencing. Starting sequencing with 289 pores July 2, 03:40
- Performing Mux Scan July 2, 03:38
- Flow cell FAO01594 has 514 pores available for sequencing. Starting sequencing with 303 pores July 2, 02:08
- Performing Mux Scan July 2, 02:06
- Flow cell FAO01594 has 540 pores available for sequencing. Starting sequencing with 312 pores July 2, 00:36
- Performing Mux Scan July 2, 00:34
- Flow cell FAO01594 has 582 pores available for sequencing. Starting sequencing with 340 pores July 1, 23:04
- Performing Mux Scan July 1, 23:02
- Flow cell FAO01594 has 641 pores available for sequencing. Starting sequencing with 355 pores July 1, 21:32
- Performing Mux Scan July 1, 21:30
- Flow cell FAO01594 has 697 pores available for sequencing. Starting sequencing with 377 pores July 1, 20:00
- Performing Mux Scan July 1, 19:58
- Disk usage alert – you only have 392 GB of space free, which is insufficient for the run. Please free up some space, otherwise your run will stop in approximately 1d 21h 11m. July 1, 18:39
- Flow cell FAO01594 has 780 pores available for sequencing. Starting sequencing with 413 pores July 1, 18:28
- Performing Mux Scan July 1, 18:26
- Starting sequencing procedure July 1, 18:26
- Failed to reach 34.0°C within 300 seconds(with 0.1 tolerance). The experiment will continue anyway. July 1, 18:26
- Waiting up to 300 seconds for temperature to stabilise at 34.0°C July 1, 18:21

● Disk / has 394 GB space remaining July 1, 18:21