Run Info

Experiment Name	**GS11_pool_06Jan2020_1**
Sample ID	**GS11**
Run ID	**4bbeb7ac-caef-4afe-9d81-3635a325e868**
Flow Cell Id	**FAL40909**
Start Time	**January 6, 16:27**
Run Length	**16h 47m**

Run Summary

Reads Generated	**325.67 K**
Estimated Bases	**505.9 Mb**

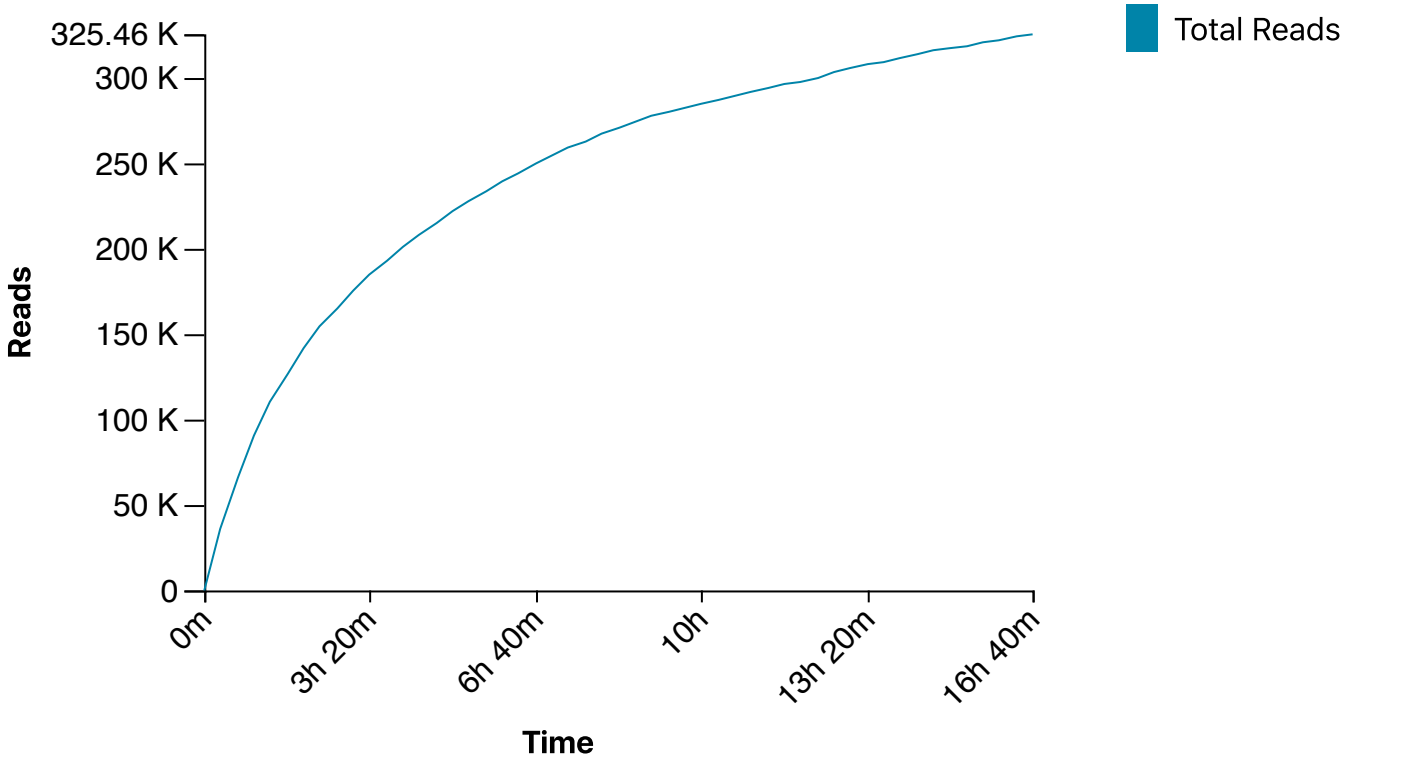
Run Parameters

Flow Cell Type	**FLO-MIN106**
Kit	**SQK-LSK108**
Basecalling	**off**
Specified Run Length	**48 hours**
Initial Bias Voltage	**-180 mV**
FAST5 Output	**Enabled**
FAST5 Output Options	**zlib_compress,raw**
FAST5 Reads per File	**4000**
Active Channel Selection	**Enabled**
Mux Scan Period	**1 hour 30 minutes**
Reserved Pores	**0 %**

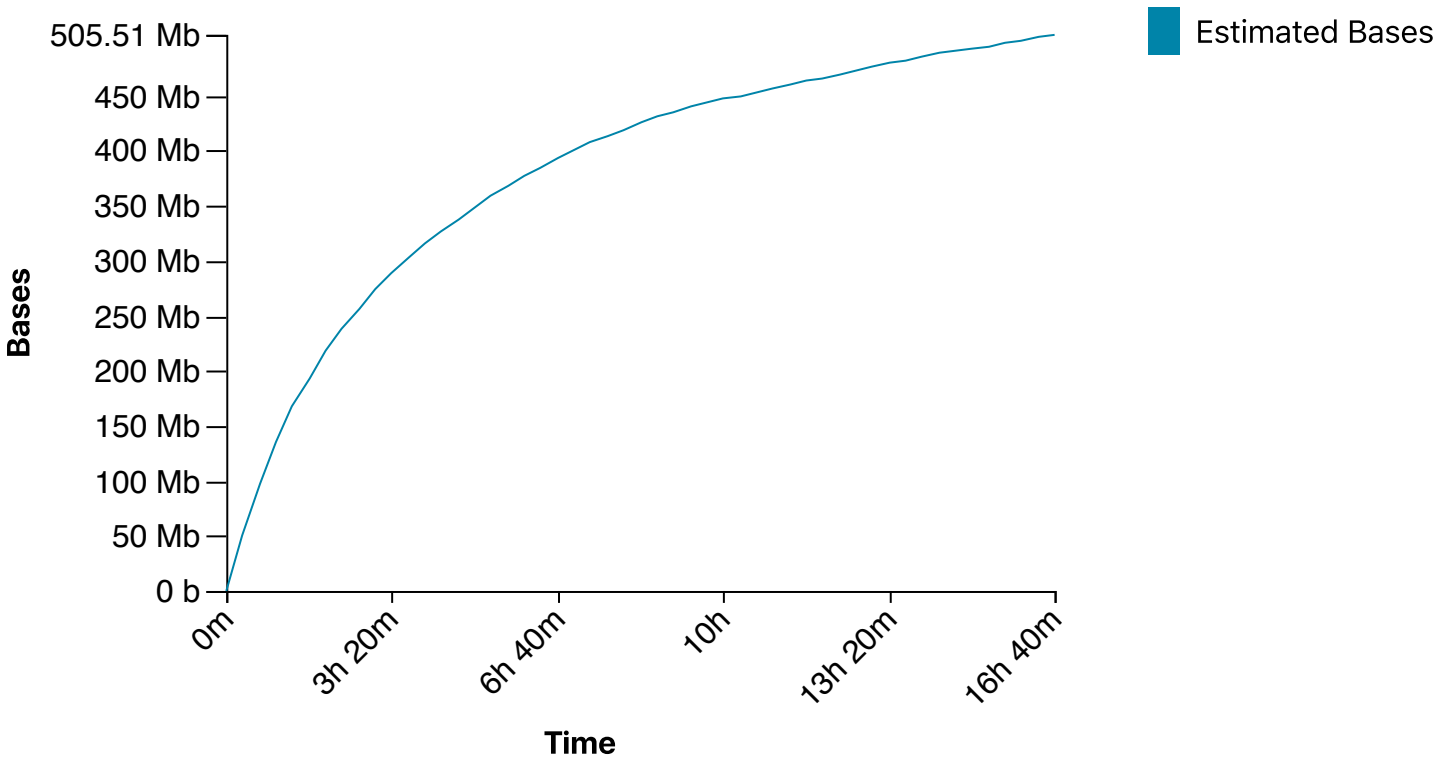
Versions

MinKNOW Core	**3.6.0**
Bream	**4.3.12**
Guppy	**3.2.8**

Cumulative Output Reads

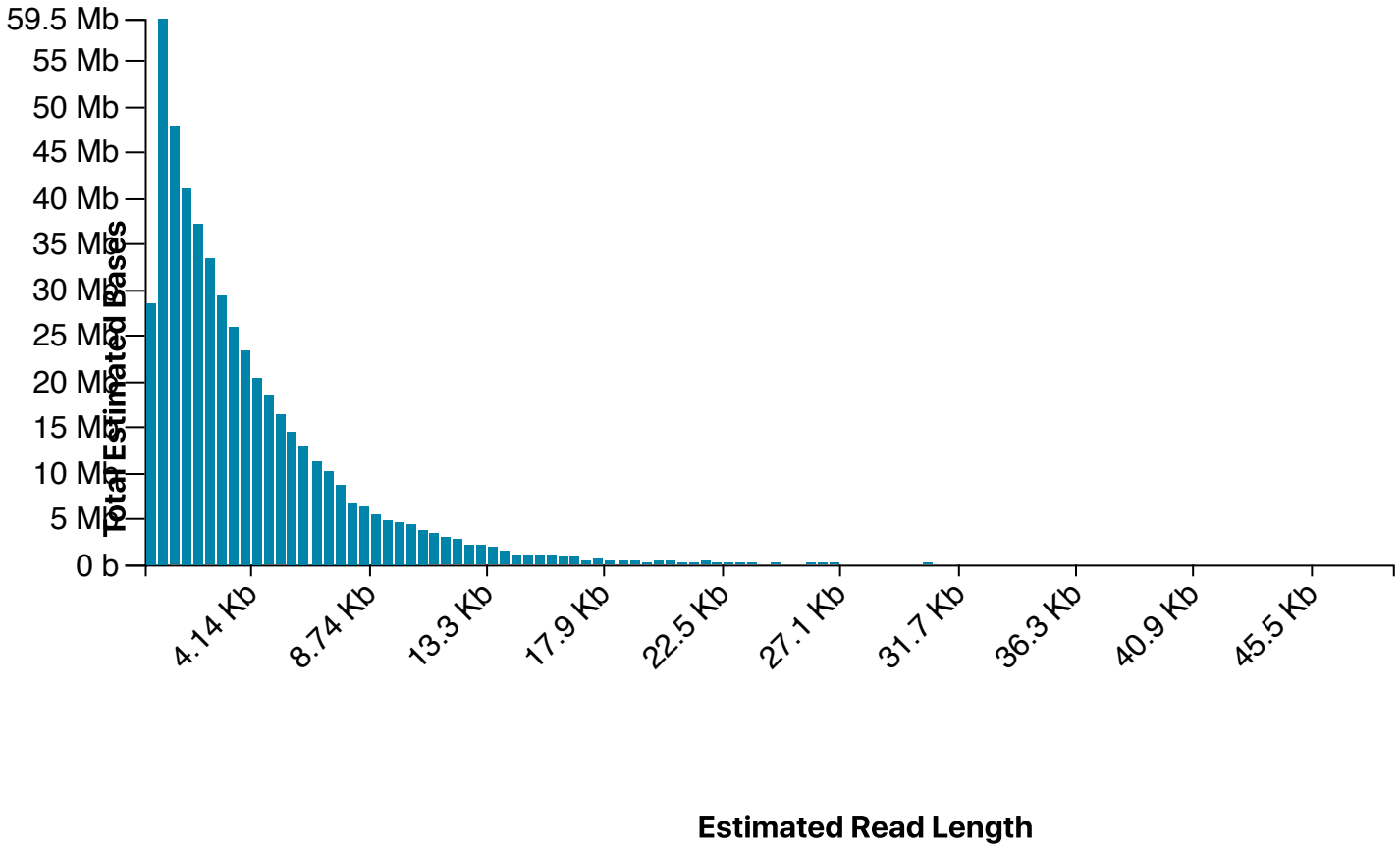


Cumulative Output Bases

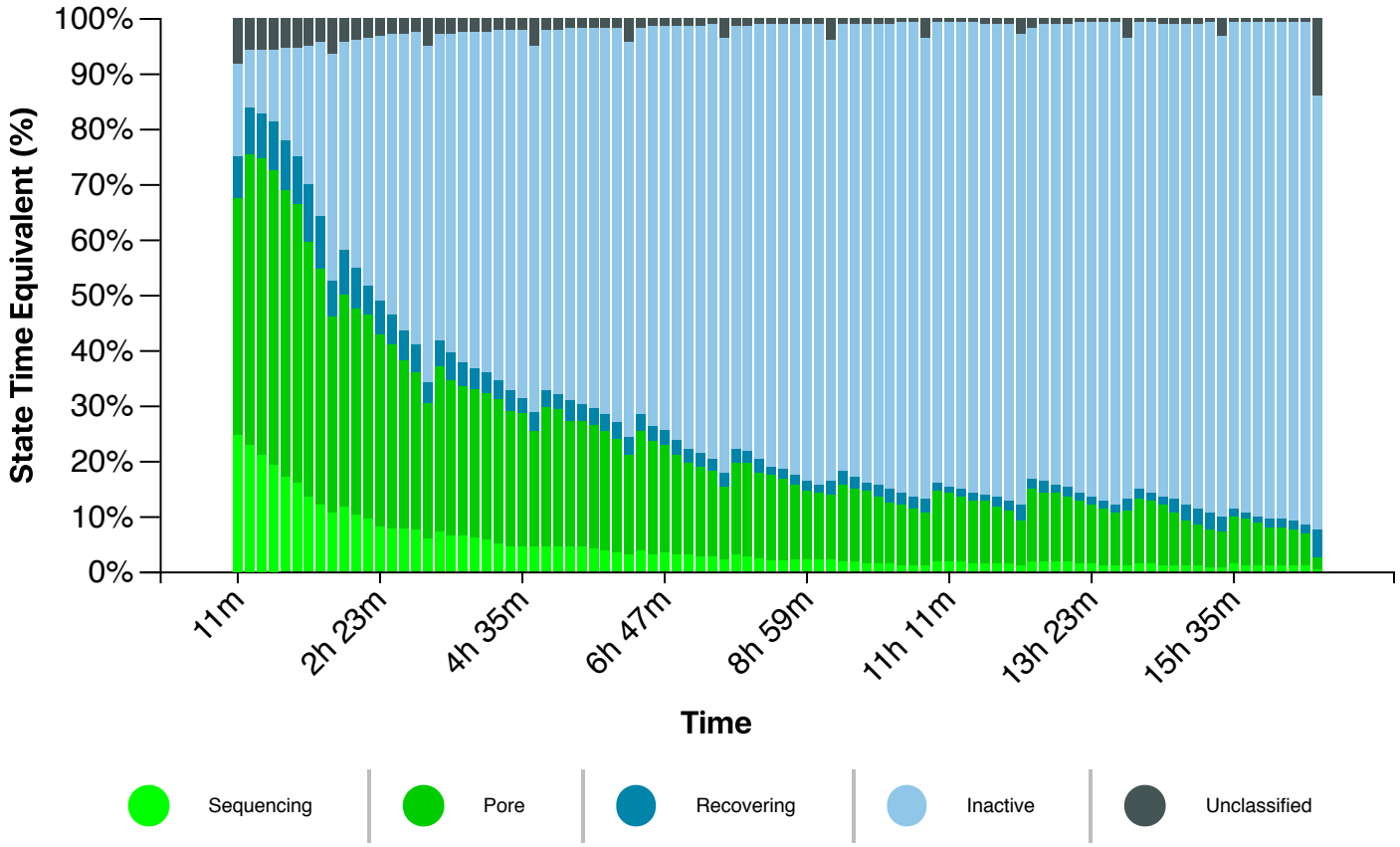


Read Length Histogram Estimated Bases

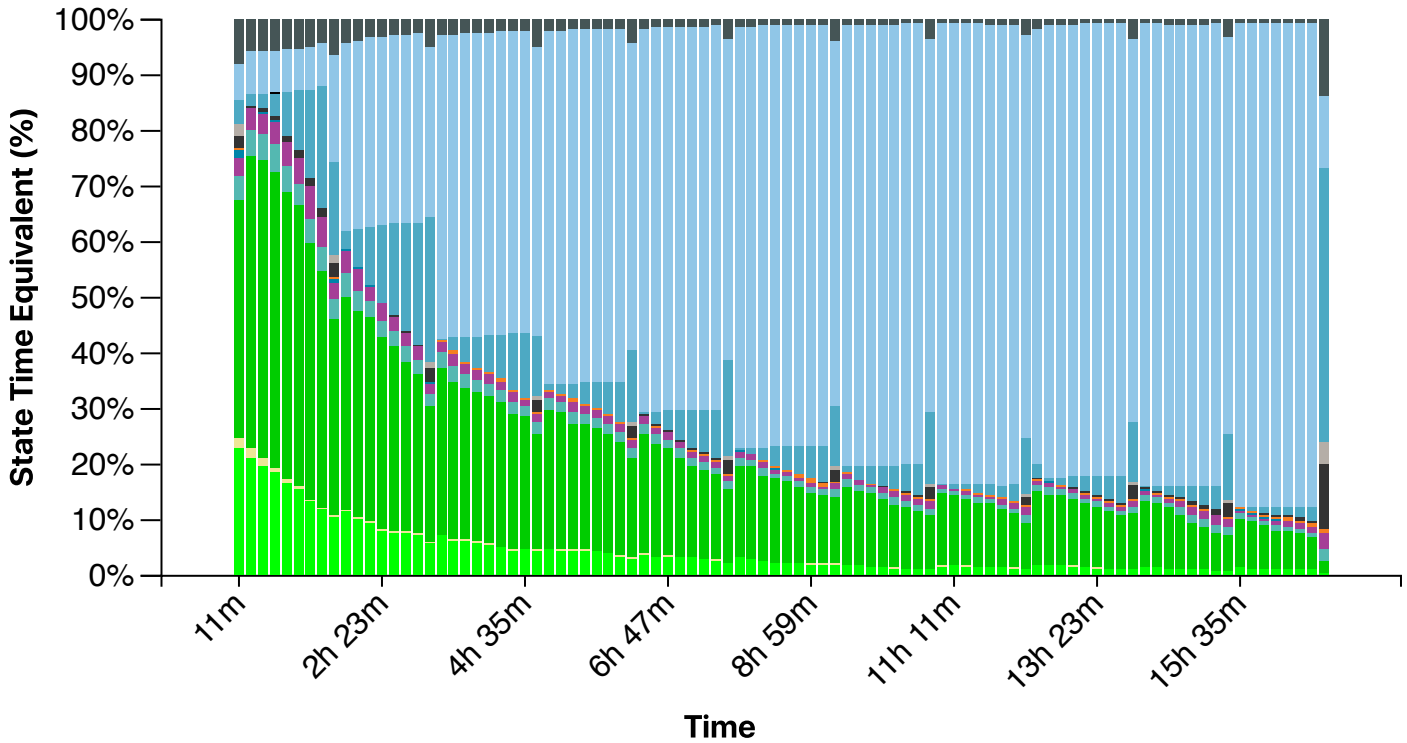
Estimated N50: 2.85 Kb



Duty Time Grouped

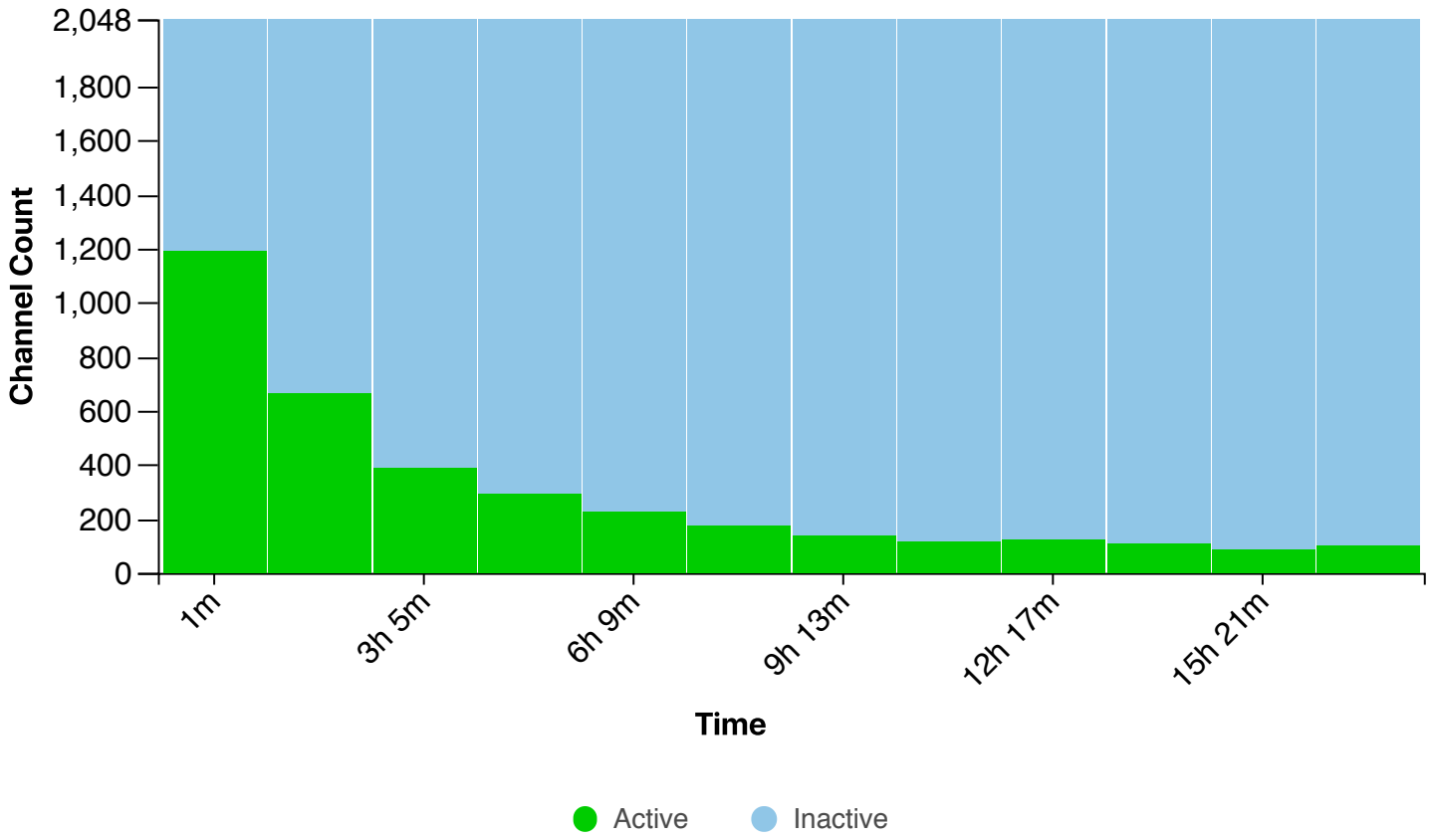


Duty time Categorised

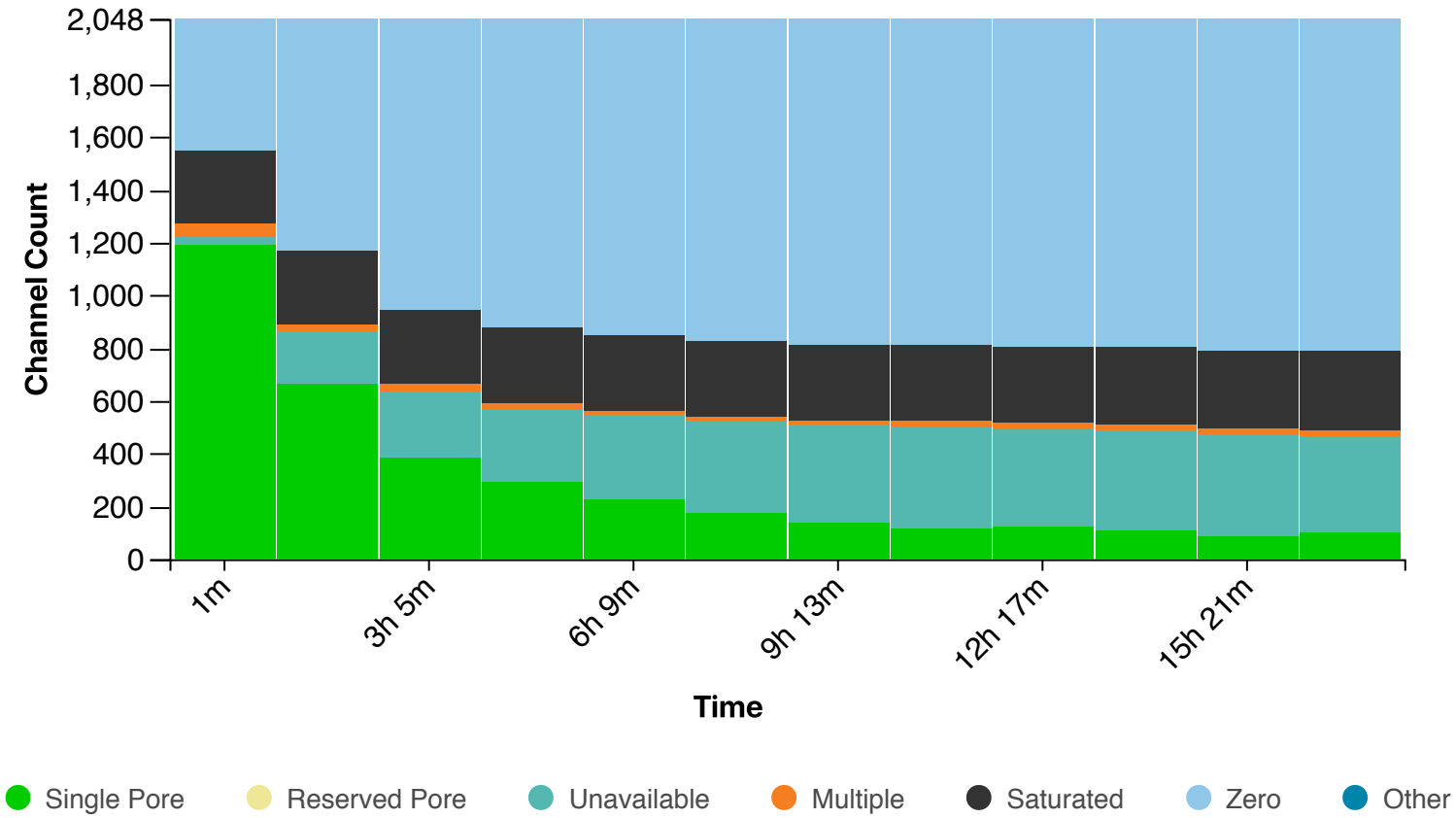


Strand
Adapter
Single Pore
Unavailable
Active Feedback
No Pore From Scan
Out Of Range 2
Possible Multiple
Saturated
Out Of Range 1
Zero
Unclassified
Unclassified_following_reset
Pending_manual_reset
Pending_mux_change
Pending Reselection

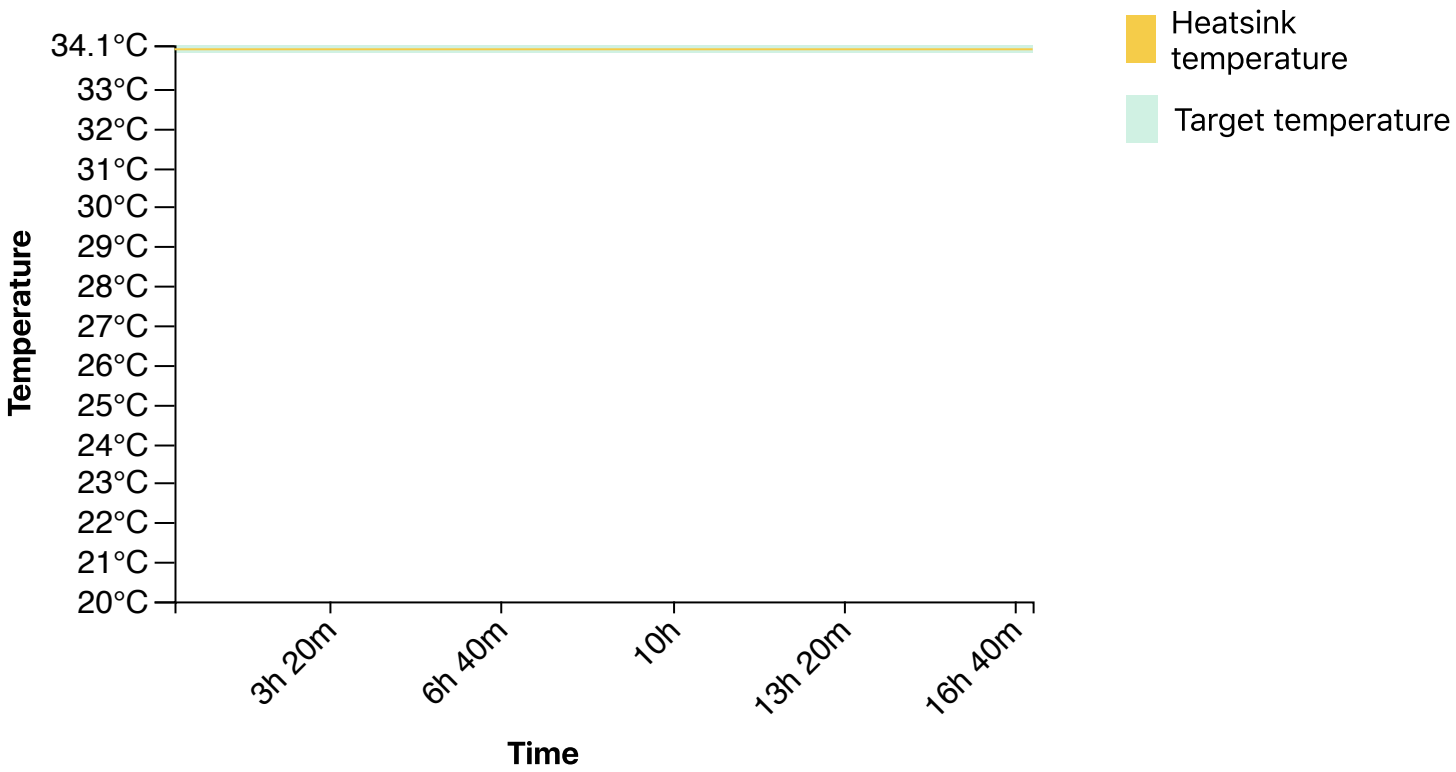
Mux Scan Grouped



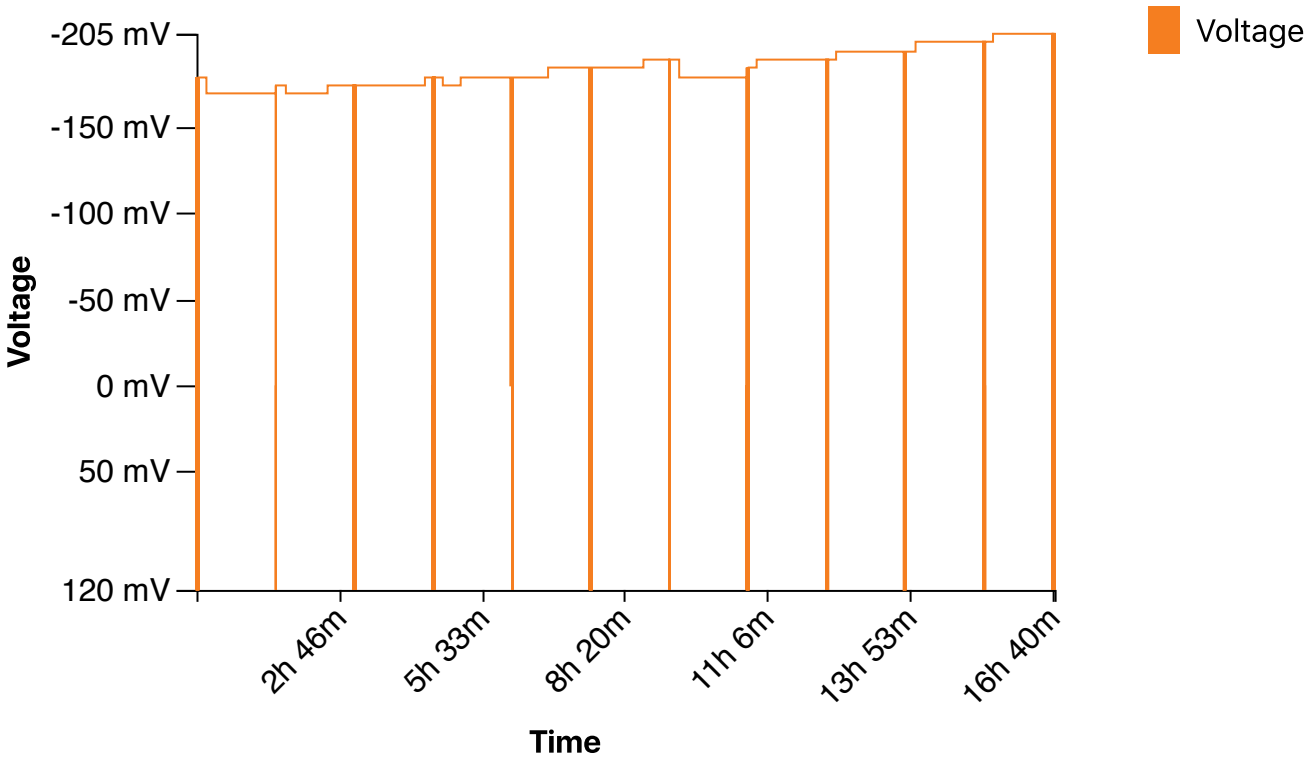
Mux Scan Categorised



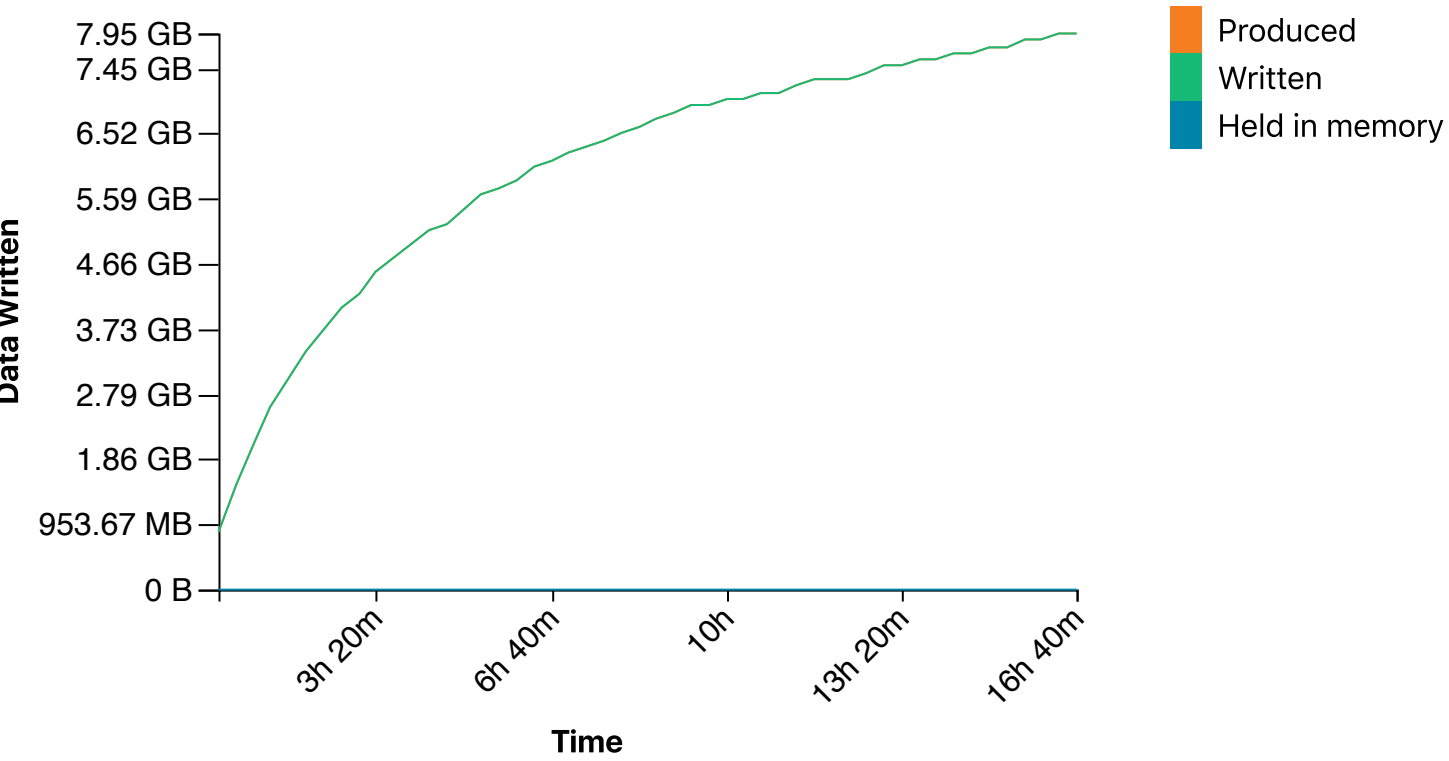
Temperature History



Bias Voltage History



Disk Write Performance



Run Debug Messages

- Flow cell FAL40909 has 100 pores available for sequencing. Starting sequencing with 81 pores January 7, 09:13
- Performing Mux Scan January 7, 09:11
- Flow cell FAL40909 has 85 pores available for sequencing. Starting sequencing with 66 pores January 7, 07:51
- Performing Mux Scan January 7, 07:50
- Flow cell FAL40909 has 108 pores available for sequencing. Starting sequencing with 87 pores January 7, 06:20
- Performing Mux Scan January 7, 06:18
- Flow cell FAL40909 has 127 pores available for sequencing. Starting sequencing with 96 pores January 7, 04:48
- Performing Mux Scan January 7, 04:46
- Flow cell FAL40909 has 120 pores available for sequencing. Starting sequencing with 88 pores January 7, 03:16
- Performing Mux Scan January 7, 03:14
- Flow cell FAL40909 has 141 pores available for sequencing. Starting sequencing with 106 pores January 7, 01:44
- Performing Mux Scan January 7, 01:42
- Flow cell FAL40909 has 176 pores available for sequencing. Starting sequencing with 124 pores January 7, 00:12
- Performing Mux Scan January 7, 00:10
- Flow cell FAL40909 has 228 pores available for sequencing. Starting sequencing with 159 pores January 6, 22:40
- Performing Mux Scan January 6, 22:38
- Flow cell FAL40909 has 289 pores available for sequencing. Starting sequencing with 186 pores January 6, 21:08
- Performing Mux Scan January 6, 21:06
- Flow cell FAL40909 has 387 pores available for sequencing. Starting sequencing with 233 pores January 6, 19:36
- Performing Mux Scan January 6, 19:34
- Flow cell FAL40909 has 667 pores available for sequencing. Starting sequencing with 338 pores January 6, 18:04
- Performing Mux Scan January 6, 18:02
- Flow cell FAL40909 has 1191 pores available for sequencing. Starting sequencing with 472 pores January 6, 16:32
- Performing Mux Scan January 6, 16:30
- Starting sequencing procedure January 6, 16:30
- Waiting for temperature to stabilise at 34.0°C January 6, 16:27
- Disk / has 2735 GB space remaining January 6, 16:27