



Run Info

Host Name	MC-110337 (localhost)
Experiment Name	TOTO
Sample ID	no_sample
Run ID	dad08772-6d67-403b-920e-54972270ab71
Flow Cell Id	FAO31058
Start Time	March 15, 16:08
Run Length	39m

Run Summary

Reads Generated	21.36 K
Estimated Bases	40.5 Mb

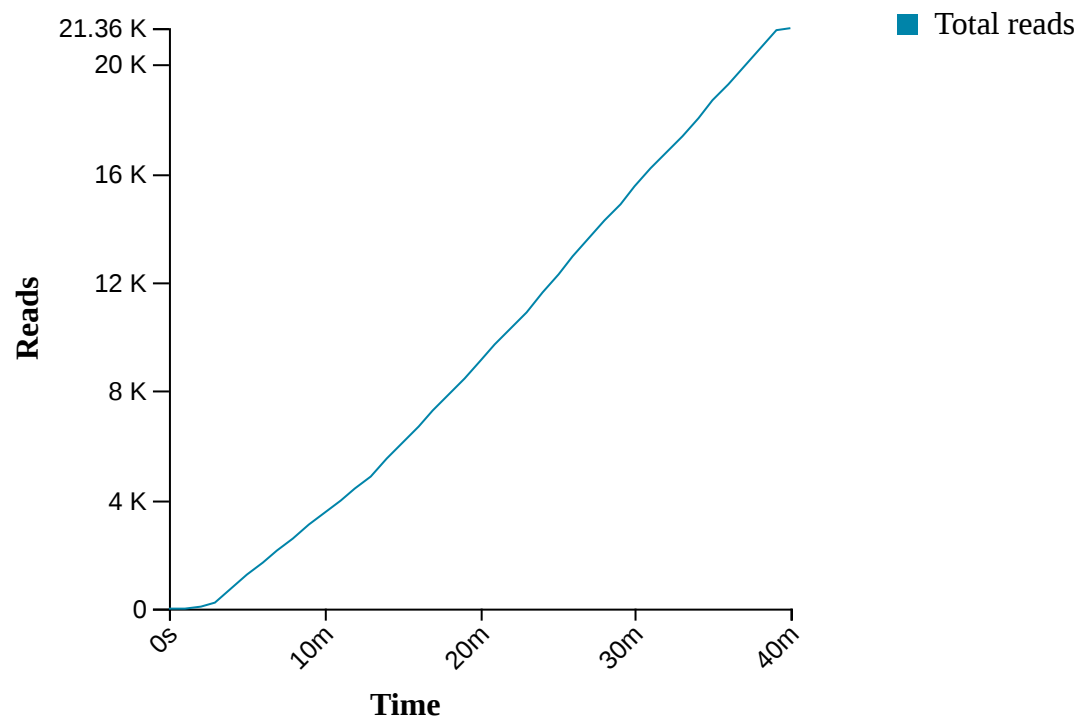
Run Parameters

Flow Cell Type	FLO-MIN106
Kit	SQK-PBK004
Initial Bias Voltage	-210 mV
FAST5 Output	Enabled
FASTQ Output	Disabled
BAM Output	Disabled
Active Channel Selection	Enabled
Basecalling	off
Specified Run Length	72 hours
FAST5 Reads per File	4000
FAST5 Output Options	zlib_compress,fastq,raw
Mux Scan Period	1 hour 30 minutes
Reserved Pores	0 %

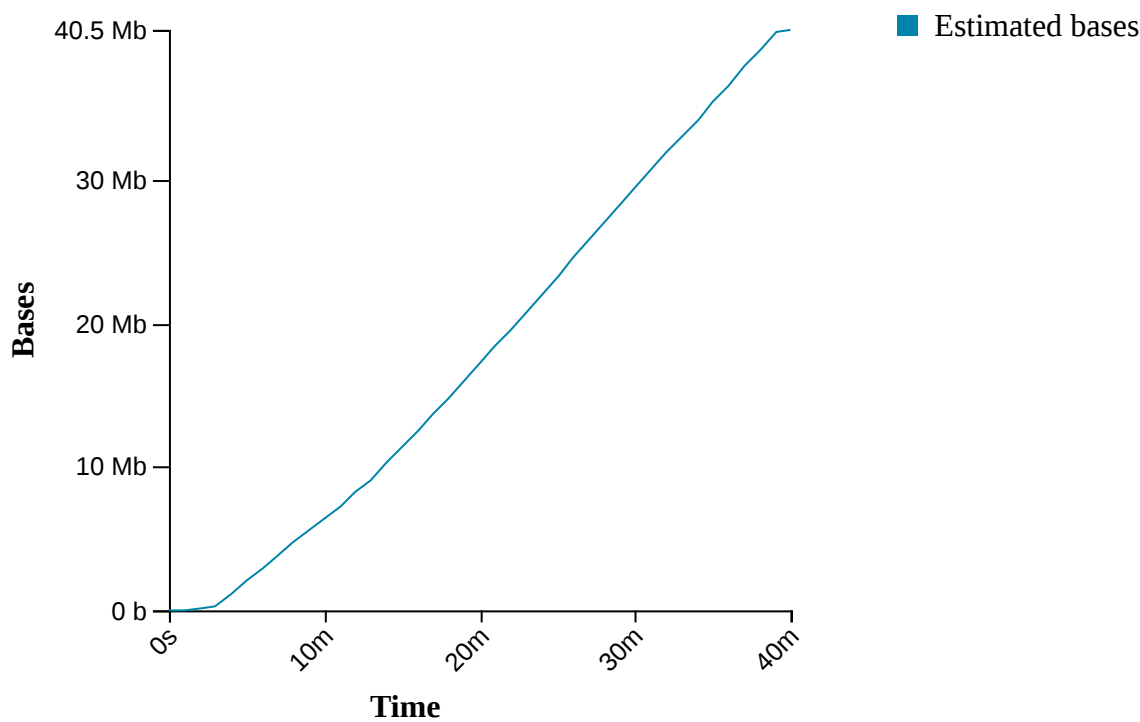
Versions

MinKNOW	21.02.2
MinKNOW Core	4.2.4
Bream	6.1.10
Guppy	4.3.4

Cumulative Output Reads

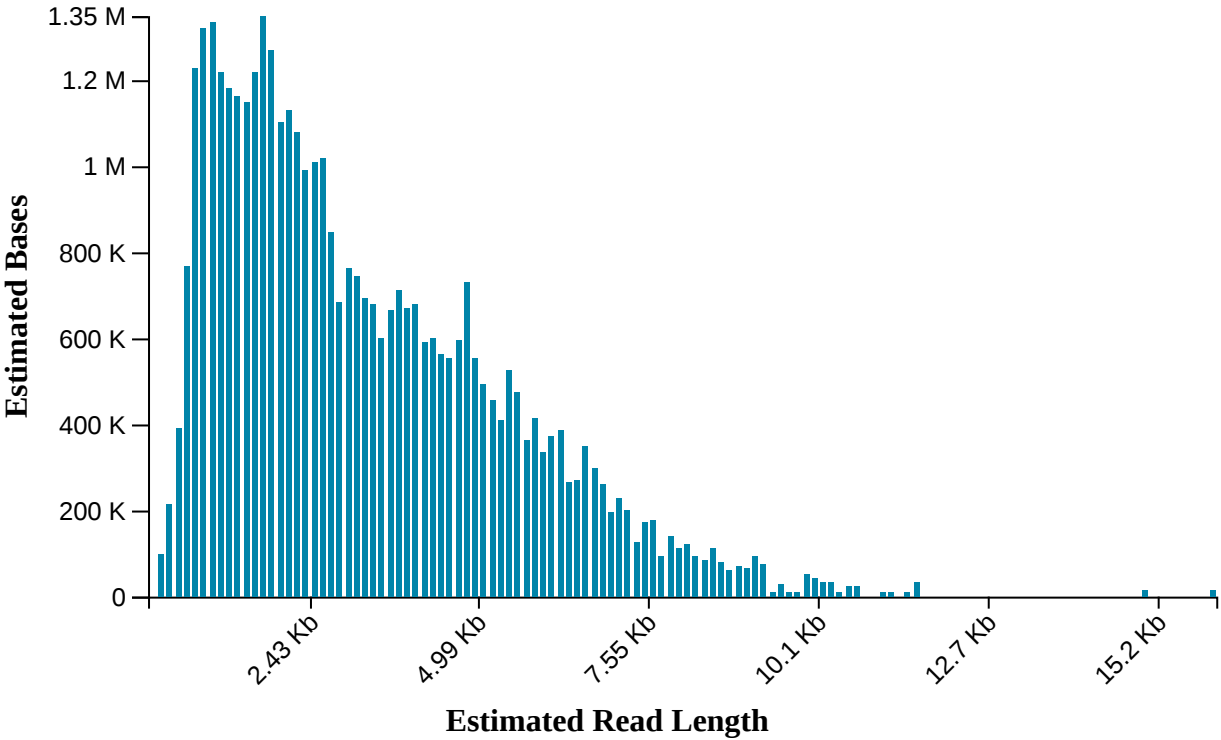


Cumulative Output Bases



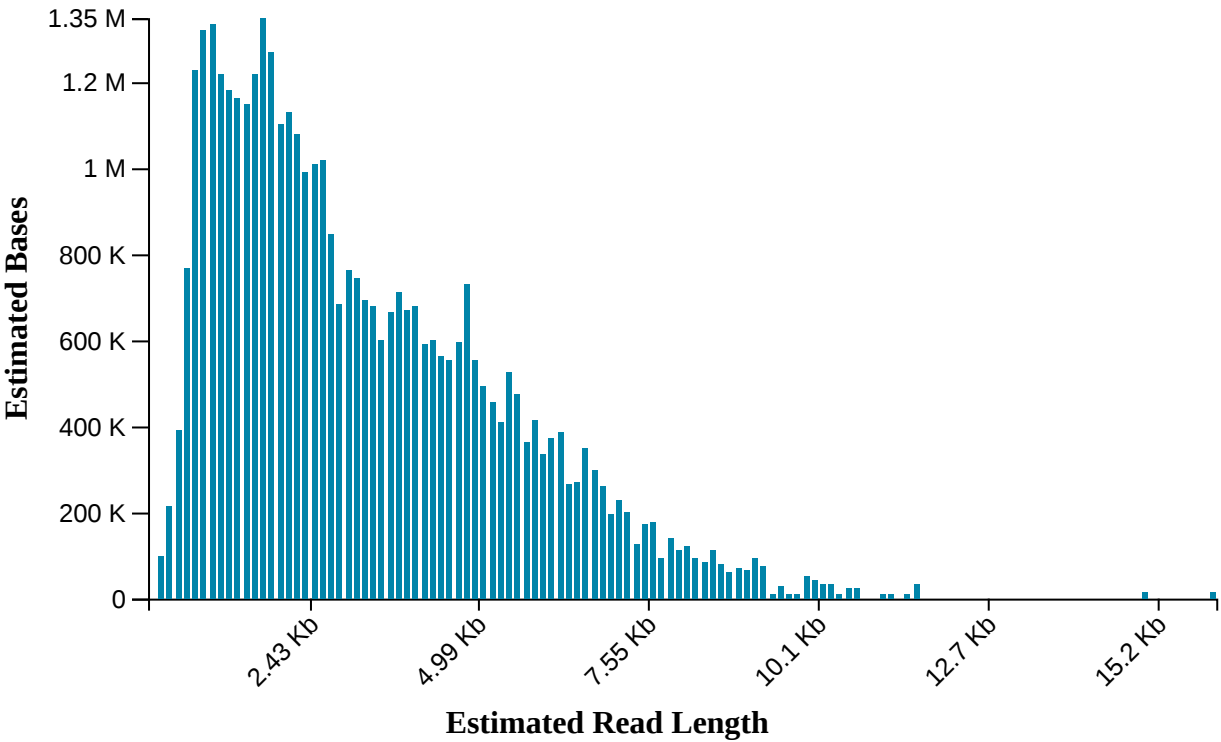
Read Length Histogram Estimated Bases - Outliers Discarded

Estimated N50: 2.69 K

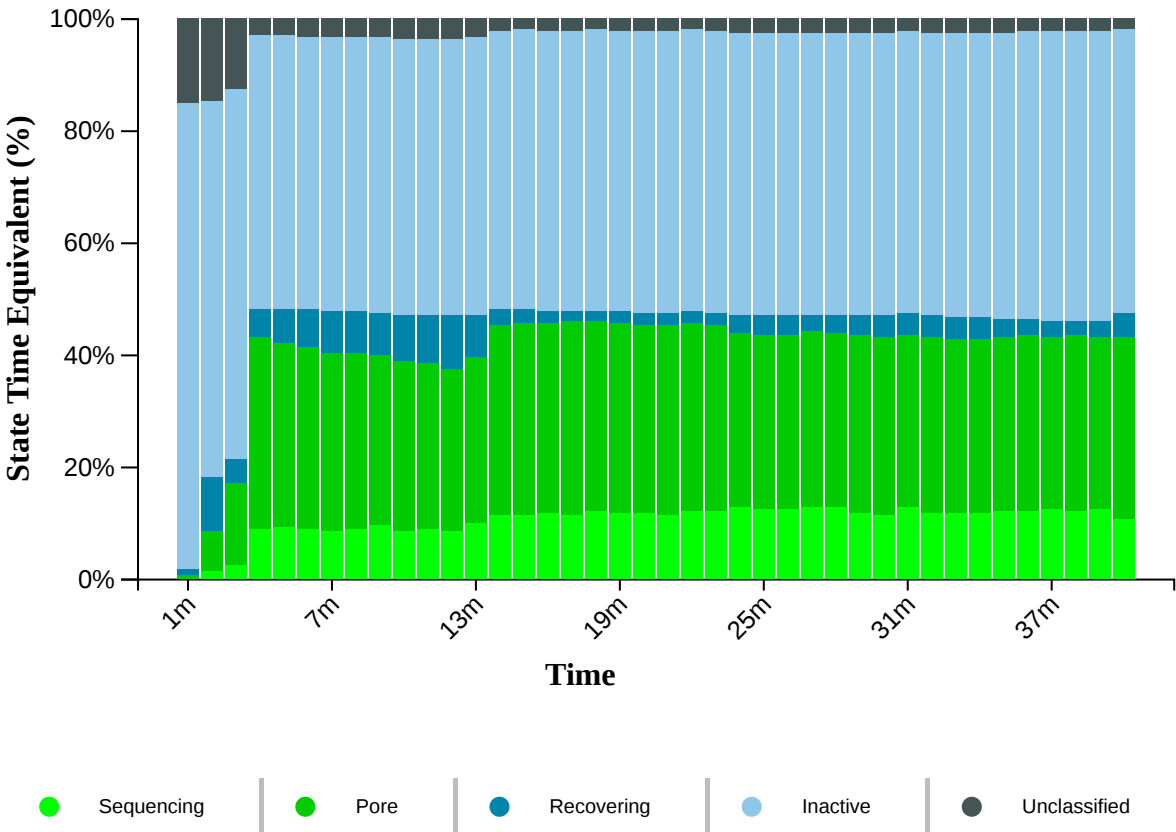


Read Length Histogram Estimated Bases

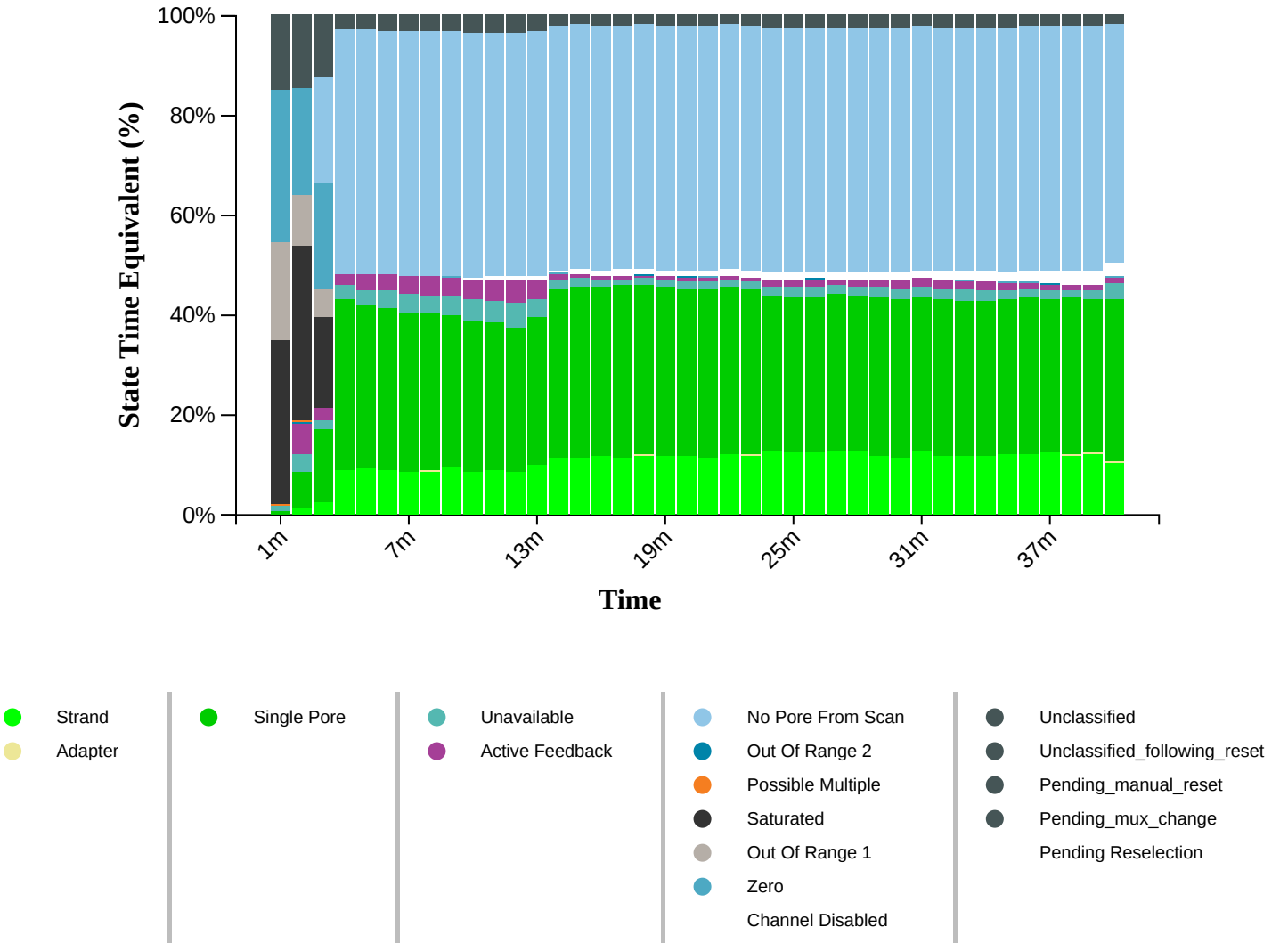
Estimated N50: 2.69 K



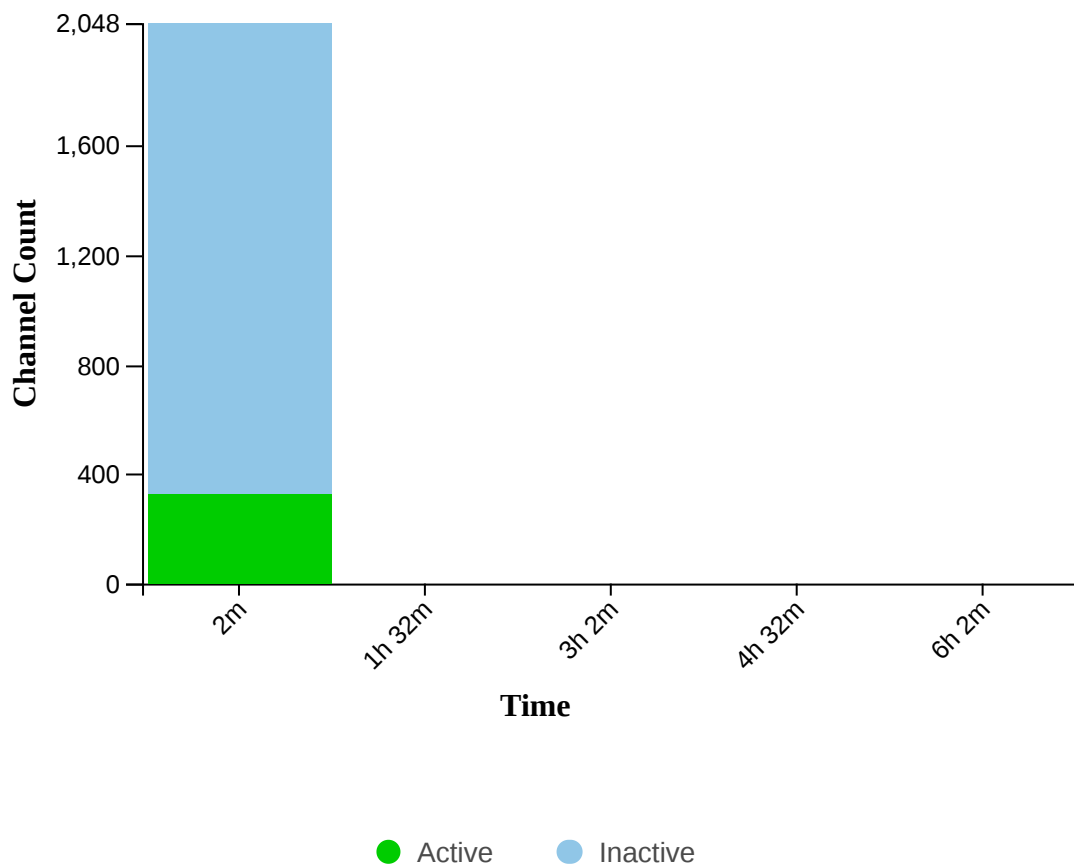
Duty Time Grouped



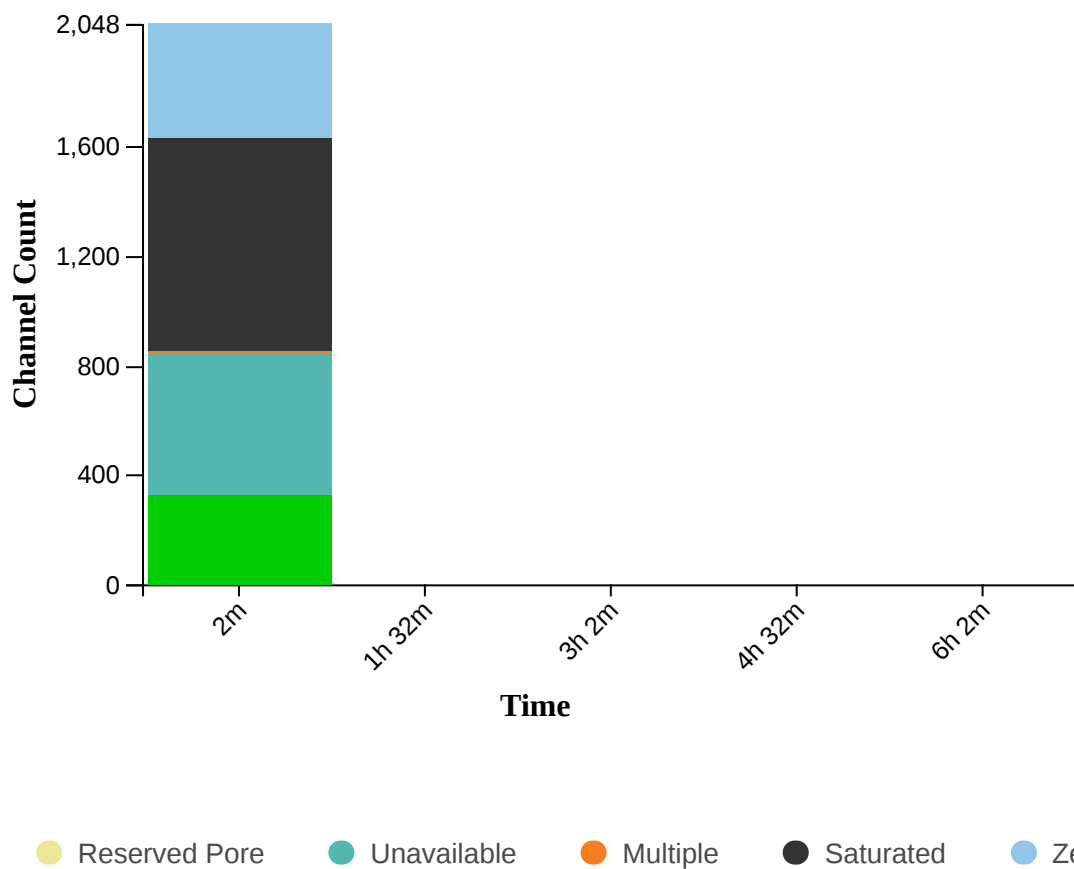
Duty time Categorised



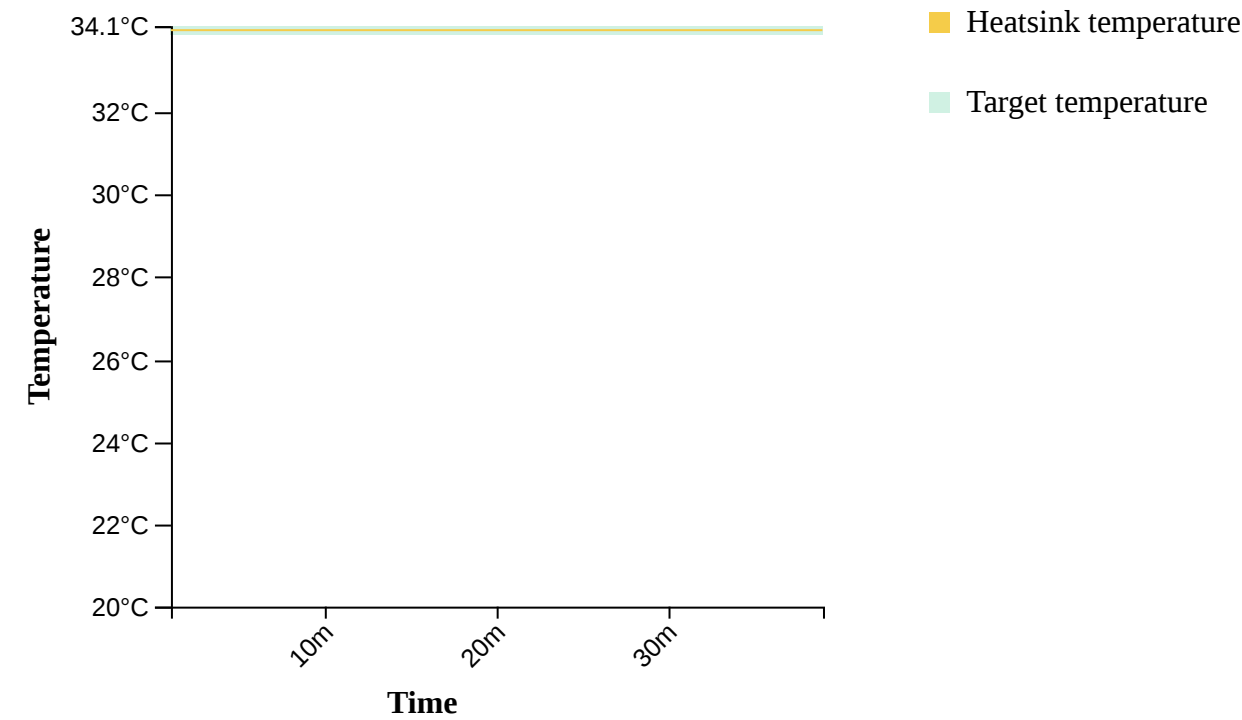
Mux Scan Grouped



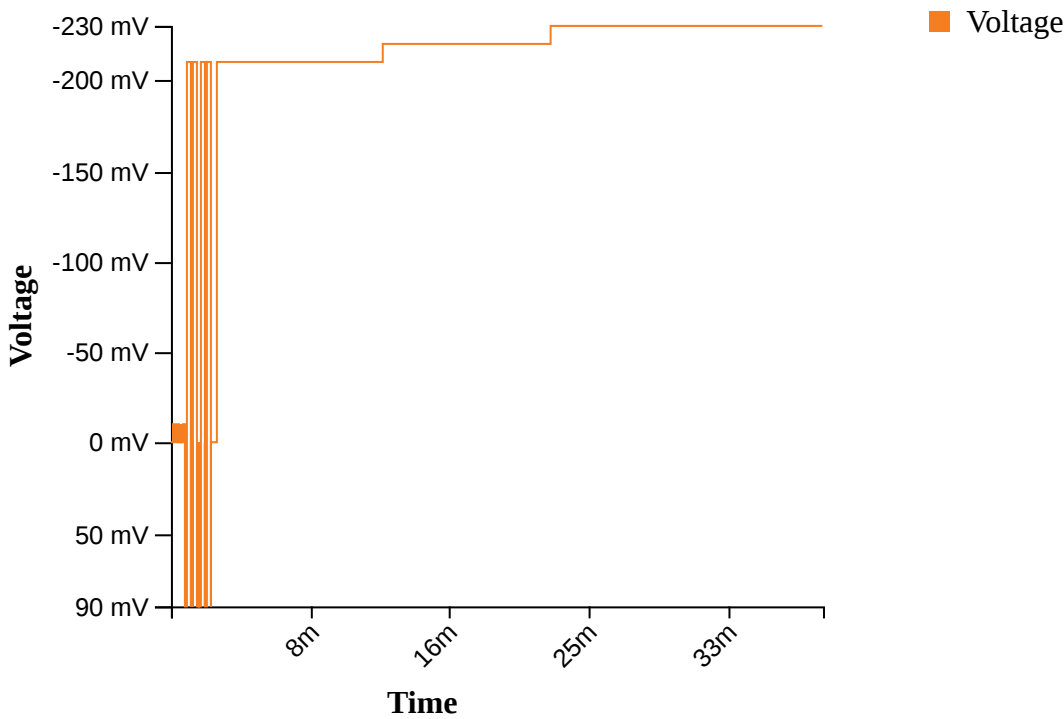
Mux Scan Categorized



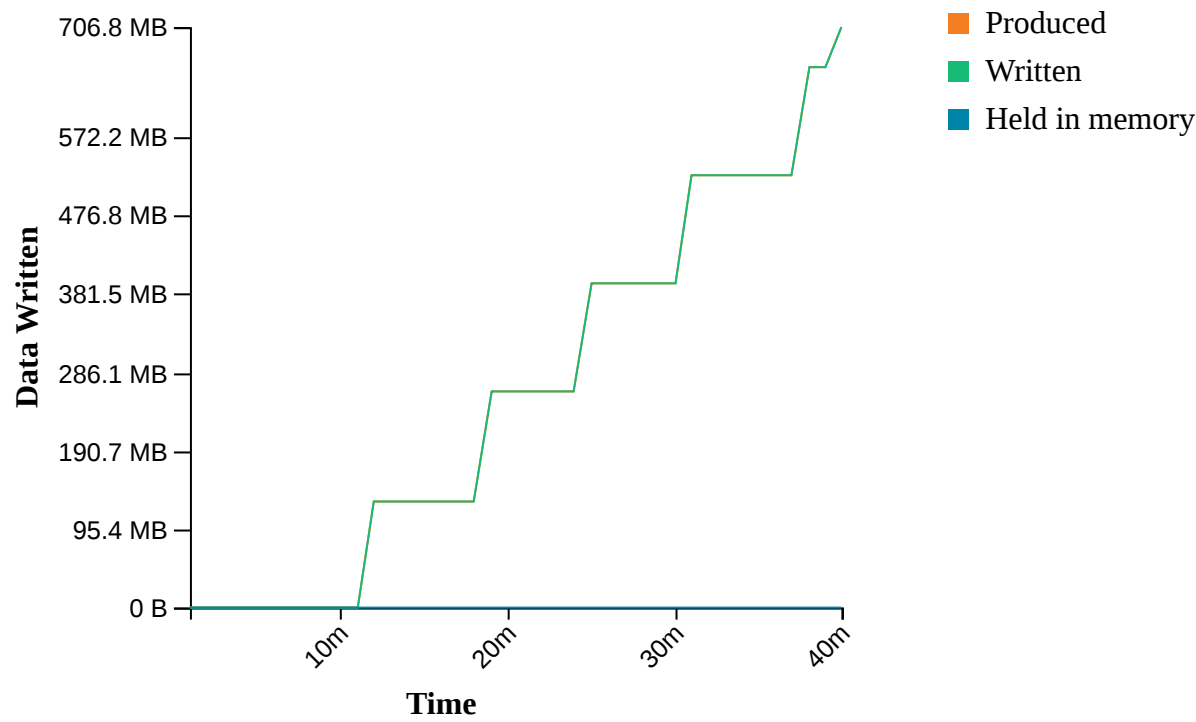
Temperature History.



Bias Voltage History.



Disk Write Performance



Run Debug Messages

- Mux scan for flow cell FAO31058 has found a total of 329 pores. 262 pores available for immediate sequencing March 15, 16:11
- Performing Mux Scan March 15, 16:09
- Starting sequencing procedure March 15, 16:09
- Waiting up to 300 seconds for temperature to stabilise at 34.0°C March 15, 16:08