

Run Info

Host Name GXB02097 (localhost)

Position X5

Experiment Name VX06_H61211d

Sample ID **H61211d**

Run ID **a34a034d-c81a-43b2-a2e6-b519132ec846**

Acquisition ID(s) **0f2ce8ad26cd15706e1d754a661315d4922e13a1,**

672f4a768319fee5321e071ea2b125995f8bc8da

Flow Cell Id FAL78159
Start Time July 22, 10:55
Run Length 3d 0h 4m

Run Summary

Reads Generated14.15 MPassed Bases5.54 GbFailed Bases3.26 GbEstimated Bases9.03 Gb

Run Parameters

FAST5 reads per file

Flow Cell Type FLO-MIN106 **SQK-DCS109** Kit -180 mV Initial bias voltage FAST5 output **Enabled** FASTQ output **Enabled** BAM output Disabled Bulk file output Disabled Active channel selection **Enabled** Basecalling **Enabled** Specified run length 72 hours

FAST5 output options vbz_compress,fastq,raw

4000

FASTQ reads per file 4000
FASTQ output options compress

Mux scan period 1 hour 30 minutes

Reserved pores 0 %

Basecall model High-accuracy basecalling

Read filtering min_qscore=9

Versions

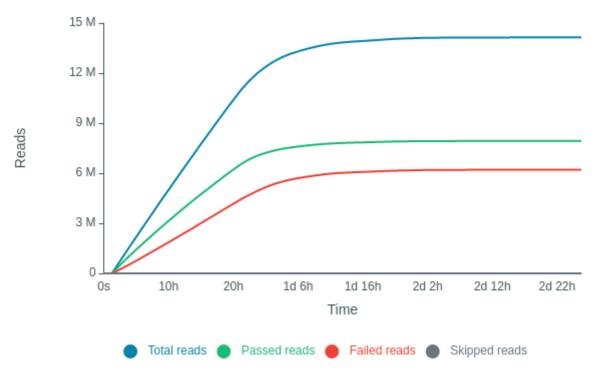
 MinKNOW
 21.05.12

 MinKNOW Core
 4.3.7

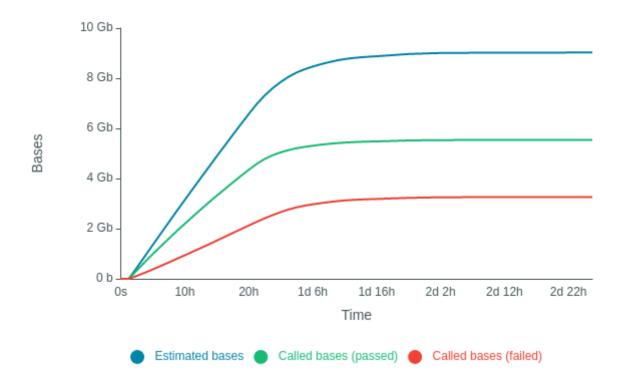
 Bream
 6.2.5

 Guppy
 5.0.12

Cumulative Output Reads

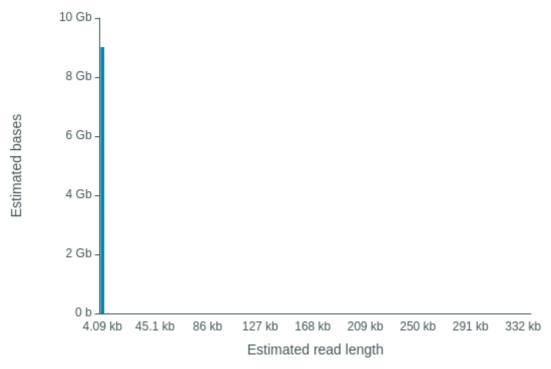


Cumulative Output Bases



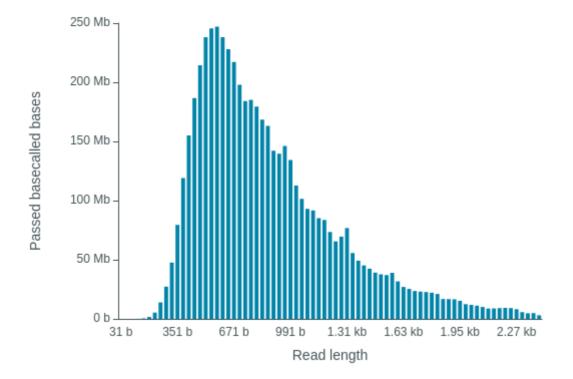
Read Length Histogram Estimated Bases - Outliers Discarded

Estimated N50: 755 b



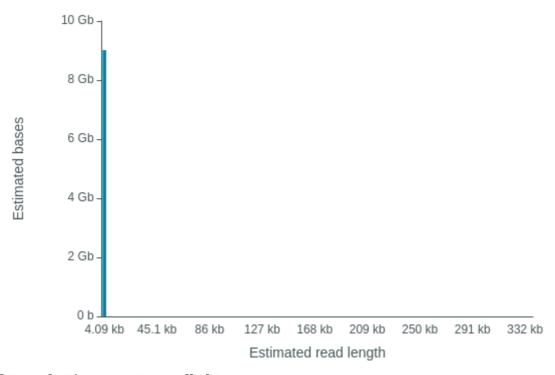
Read Length Histogram Basecalled Bases - Outliers Discarded

Estimated N50: 752 b



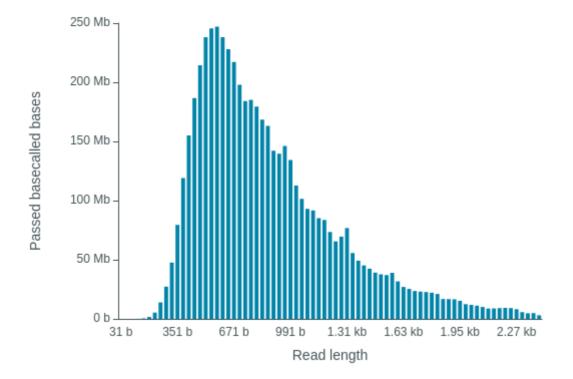
Read Length Histogram Estimated Bases

Estimated N50: 755 b

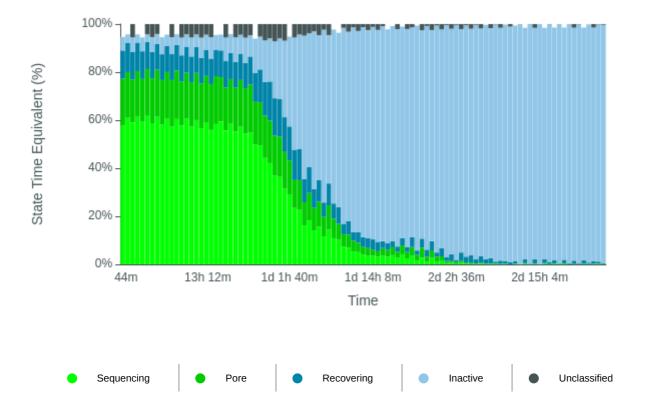


Read Length Histogram Basecalled Bases

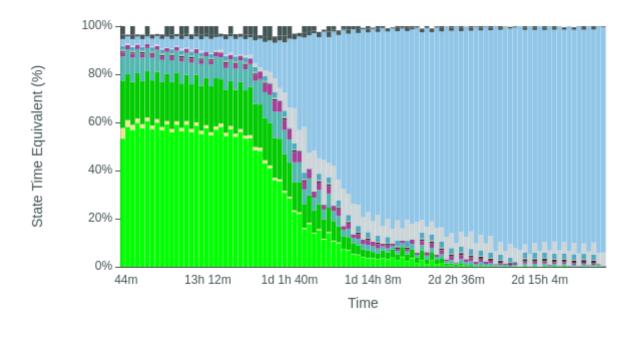
Estimated N50: 752 b



Duty Time Grouped

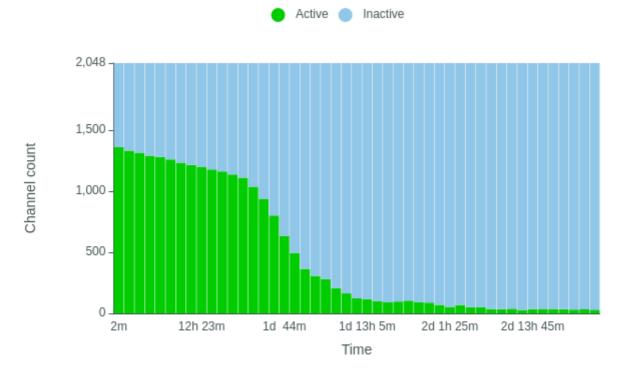


Duty time Categorised

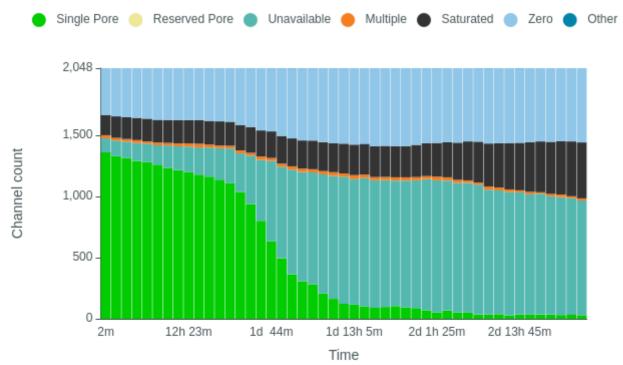




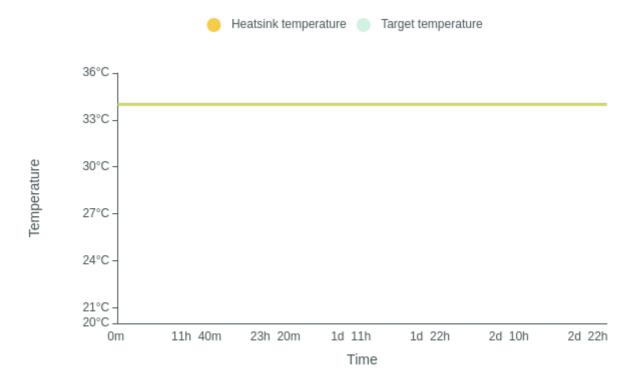
Mux Scan Grouped



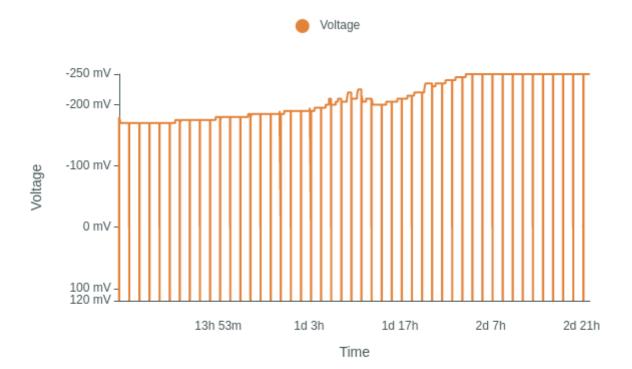
Mux Scan Categorised



Temperature History



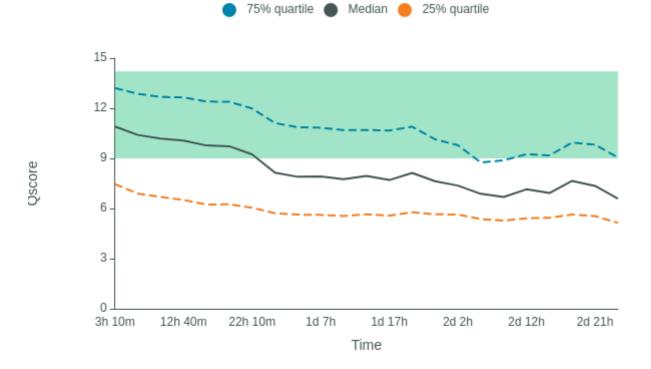
Bias Voltage History



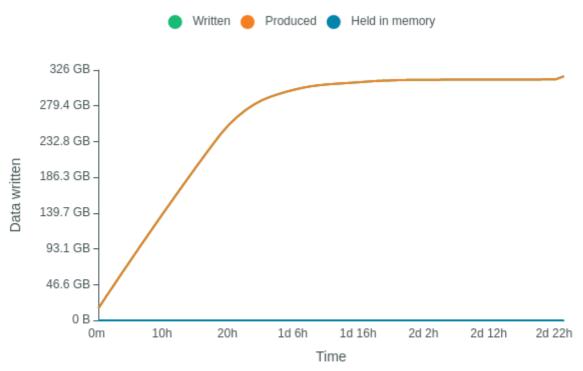
Translocation Speed



QScore



Disk Write Performance



Run Debug Messages

- The sequencing run has finished, but basecalling may continue July 25, 10:59
- Mux scan for flow cell FAL78159 has found a total of 30 pores. 30 pores available for immediate sequencing July 25, 10:00
- Performing Mux Scan July 25, 09:57
- Mux scan for flow cell FAL78159 has found a total of 33 pores. 33 pores available for immediate sequencing July 25, 08:27
- Performing Mux Scan July 25, 08:25
- Mux scan for flow cell FAL78159 has found a total of 31 pores. 29 pores available for immediate sequencing July 25, 06:55
- Performing Mux Scan July 25, 06:52
- Mux scan for flow cell FAL78159 has found a total of 34 pores. 34 pores available for immediate sequencing July 25, 05:22
- Performing Mux Scan July 25, 05:20
- Mux scan for flow cell FAL78159 has found a total of 36 pores. 35 pores available for immediate sequencing July 25, 03:50
- Performing Mux Scan July 25, 03:47
- Mux scan for flow cell FAL78159 has found a total of 37 pores. 35 pores available for immediate sequencing July 25, 02:17
- Performing Mux Scan July 25, 02:15
- Mux scan for flow cell FAL78159 has found a total of 34 pores. 33 pores available for immediate sequencing July 25, 00:45
- Performing Mux Scan July 25, 00:42
- Mux scan for flow cell FAL78159 has found a total of 28 pores. 28 pores available for immediate sequencing July 24, 23:12
- Performing Mux Scan July 24, 23:10
- Mux scan for flow cell FAL78159 has found a total of 38 pores. 36 pores available for immediate sequencing July 24, 21:40
- Performing Mux Scan July 24, 21:37
- Mux scan for flow cell FAL78159 has found a total of 34 pores. 34 pores available for immediate sequencing July 24, 20:07
- Performing Mux Scan July 24, 20:05
- Mux scan for flow cell FAL78159 has found a total of 36 pores. 34 pores available for immediate sequencing July 24, 18:35
- Performing Mux Scan July 24, 18:32
- Mux scan for flow cell FAL78159 has found a total of 52 pores. 51 pores available for immediate sequencing July 24, 17:02
- Performing Mux Scan July 24, 17:00
- Mux scan for flow cell FAL78159 has found a total of 51 pores. 51 pores available for immediate sequencing July 24, 15:30
- Performing Mux Scan July 24, 15:27
- Mux scan for flow cell FAL78159 has found a total of 69 pores. 66 pores available for immediate sequencing July 24, 13:57
- Performing Mux Scan July 24, 13:55
- Mux scan for flow cell FAL78159 has found a total of 53 pores. 52 pores available for immediate sequencing July 24, 12:25
- Performing Mux Scan July 24, 12:22
- Mux scan for flow cell FAL78159 has found a total of 70 pores. 65 pores available for immediate sequencing July 24, 10:52
- Performing Mux Scan July 24, 10:50
- Mux scan for flow cell FAL 78159 has found a total of 88 pores. 85 pores available for immediate

- sequencing July 24, 09:20
- Performing Mux Scan July 24, 09:17
- Mux scan for flow cell FAL78159 has found a total of 93 pores. 89 pores available for immediate sequencing July 24, 07:47
- Performing Mux Scan July 24, 07:45
- Mux scan for flow cell FAL78159 has found a total of 104 pores. 98 pores available for immediate sequencing July 24, 06:15
- Performing Mux Scan July 24, 06:12
- Mux scan for flow cell FAL78159 has found a total of 97 pores. 87 pores available for immediate sequencing July 24, 04:42
- Performing Mux Scan July 24, 04:40
- Mux scan for flow cell FAL78159 has found a total of 90 pores. 85 pores available for immediate sequencing July 24, 03:10
- Performing Mux Scan July 24, 03:07
- Mux scan for flow cell FAL78159 has found a total of 101 pores. 91 pores available for immediate sequencing July 24, 01:37
- Performing Mux Scan July 24, 01:35
- Mux scan for flow cell FAL78159 has found a total of 117 pores. 103 pores available for immediate sequencing July 24, 00:05
- Performing Mux Scan July 24, 00:02
- Mux scan for flow cell FAL78159 has found a total of 123 pores. 112 pores available for immediate sequencing July 23, 22:32
- Performing Mux Scan July 23, 22:30
- Mux scan for flow cell FAL78159 has found a total of 163 pores. 140 pores available for immediate sequencing July 23, 21:00
- Performing Mux Scan July 23, 20:57
- Mux scan for flow cell FAL78159 has found a total of 207 pores. 174 pores available for immediate sequencing July 23, 19:27
- Performing Mux Scan July 23, 19:25
- Mux scan for flow cell FAL78159 has found a total of 281 pores. 227 pores available for immediate sequencing July 23, 17:54
- Performing Mux Scan July 23, 17:52
- Mux scan for flow cell FAL78159 has found a total of 307 pores. 239 pores available for immediate sequencing July 23, 16:22
- Performing Mux Scan July 23, 16:19
- Mux scan for flow cell FAL78159 has found a total of 364 pores. 262 pores available for immediate sequencing July 23, 14:49
- Performing Mux Scan July 23, 14:47
- Mux scan for flow cell FAL78159 has found a total of 495 pores. 313 pores available for immediate sequencing July 23, 13:17
- Performing Mux Scan July 23, 13:14
- Mux scan for flow cell FAL78159 has found a total of 635 pores. 365 pores available for immediate sequencing July 23, 11:44
- Performing Mux Scan July 23, 11:42
- Mux scan for flow cell FAL78159 has found a total of 801 pores. 411 pores available for immediate sequencing July 23, 10:11
- Performing Mux Scan July 23, 10:09
- Mux scan for flow cell FAL78159 has found a total of 939 pores. 445 pores available for immediate sequencing July 23, 08:39
- Performing Mux Scan July 23, 08:36
- Mux scan for flow cell FAL78159 has found a total of 1036 pores. 451 pores available for immediate sequencing July 23, 07:06

- Performing Mux Scan July 23, 07:04
- Mux scan for flow cell FAL78159 has found a total of 1110 pores. 475 pores available for immediate sequencing July 23, 05:33
- Performing Mux Scan July 23, 05:31
- Mux scan for flow cell FAL78159 has found a total of 1136 pores. 475 pores available for immediate sequencing July 23, 04:01
- Performing Mux Scan July 23, 03:58
- Mux scan for flow cell FAL78159 has found a total of 1162 pores. 477 pores available for immediate sequencing July 23, 02:28
- Performing Mux Scan July 23, 02:26
- Mux scan for flow cell FAL78159 has found a total of 1178 pores. 484 pores available for immediate sequencing July 23, 00:55
- Performing Mux Scan July 23, 00:53
- Mux scan for flow cell FAL78159 has found a total of 1200 pores. 480 pores available for immediate sequencing July 22, 23:23
- Performing Mux Scan July 22, 23:20
- Mux scan for flow cell FAL78159 has found a total of 1215 pores. 487 pores available for immediate sequencing July 22, 21:50
- Performing Mux Scan July 22, 21:48
- Mux scan for flow cell FAL78159 has found a total of 1232 pores. 489 pores available for immediate sequencing July 22, 20:17
- Performing Mux Scan July 22, 20:15
- Mux scan for flow cell FAL78159 has found a total of 1260 pores. 492 pores available for immediate sequencing July 22, 18:45
- Performing Mux Scan July 22, 18:42
- Mux scan for flow cell FAL78159 has found a total of 1280 pores. 489 pores available for immediate sequencing July 22, 17:12
- Performing Mux Scan July 22, 17:10
- Mux scan for flow cell FAL78159 has found a total of 1291 pores. 494 pores available for immediate sequencing July 22, 15:40
- Performing Mux Scan July 22, 15:37
- Mux scan for flow cell FAL78159 has found a total of 1315 pores. 496 pores available for immediate sequencing July 22, 14:07
- Performing Mux Scan July 22, 14:04
- Mux scan for flow cell FAL78159 has found a total of 1331 pores. 497 pores available for immediate sequencing July 22, 12:34
- Performing Mux Scan July 22, 12:32
- Mux scan for flow cell FAL78159 has found a total of 1359 pores. 497 pores available for immediate sequencing July 22, 11:02
- Performing Mux Scan July 22, 10:59
- Starting sequencing procedure July 22, 10:59
- Waiting up to 300 seconds for temperature to stabilise at 34.0°C July 22, 10:55