

#### **Run Info**

Host Name GXB01275 (localhost)

Position X2
Experiment Name kokia
Sample ID Kc

Run ID **f6a3b4f3-d5f7-47be-beb0-3112461af3bb** 

Acquisition ID(s) 0402f5908d8c99c9b87d2e6f33cf808ab7c43f7b, 5b74c43dc6d838db6414f8be01f05446db224dcf

Flow Cell Id FAU08677
Start Time June 10, 14:15
Run Length 3d 0h 3m

#### **Run Summary**

Reads Generated1.78 MPassed Bases8.64 GbFailed Bases2.01 GbEstimated Bases12.19 GbPercentage Basecalled102%

#### **Run Parameters**

Flow Cell Type FLO-MIN112 Kit SQK-LSK112 Initial bias voltage -200 mV FAST5 output **Enabled** FASTQ output **Enabled** BAM output Disabled Bulk file output Disabled Active channel selection **Enabled** Basecalling **Enabled** Specified run length 72 hours 4000 FAST5 reads per file

FAST5 output options vbz\_compress,fastq,raw

FASTQ reads per file 4000
FASTQ output options compress

Mux scan period 1 hour 30 minutes

Reserved pores 0 %

Basecall model dna\_r10.4\_e8.1\_hac.cfg

Read filtering min\_qscore=9
Read splitting enable=on

#### Versions

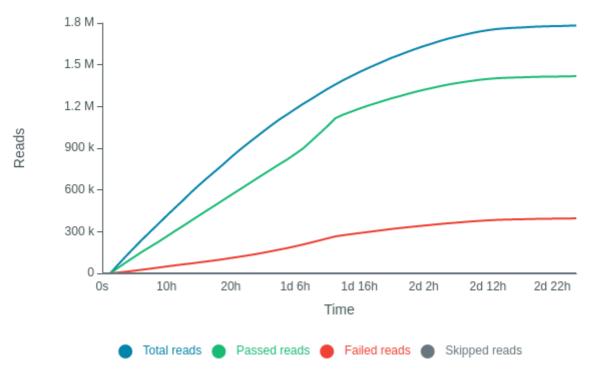
 MinKNOW
 21.11.7

 MinKNOW Core
 4.5.4

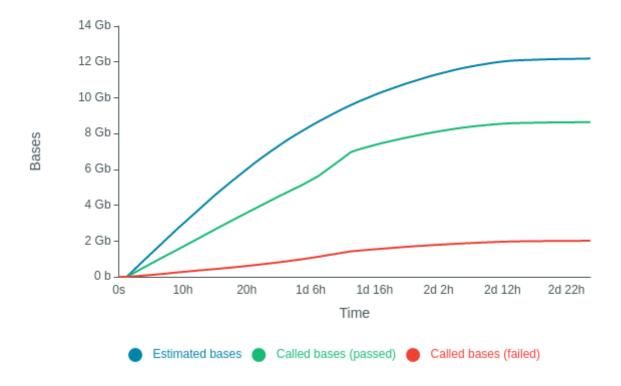
 Bream
 6.3.5

 Guppy
 5.1.13

## **Cumulative Output Reads**

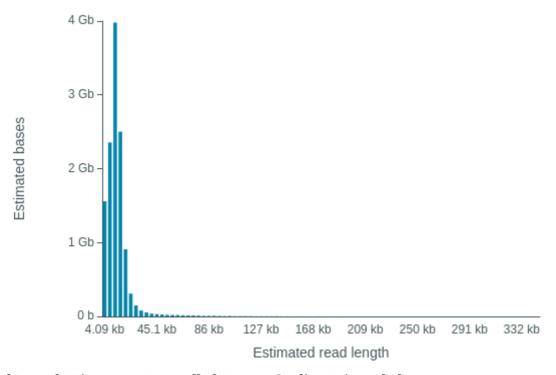


# **Cumulative Output Bases**



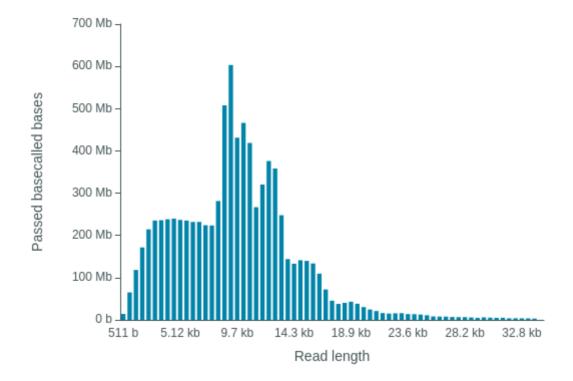
## Read Length Histogram Estimated Bases - Outliers Discarded

Estimated N50: 10.46 kb



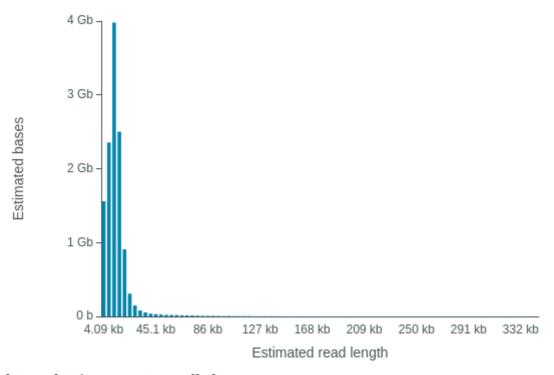
Read Length Histogram Basecalled Bases - Outliers Discarded

Estimated N50: 9.19 kb



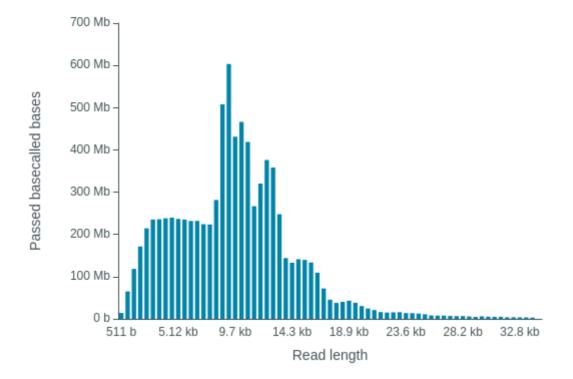
# **Read Length Histogram Estimated Bases**

Estimated N50: 10.46 kb

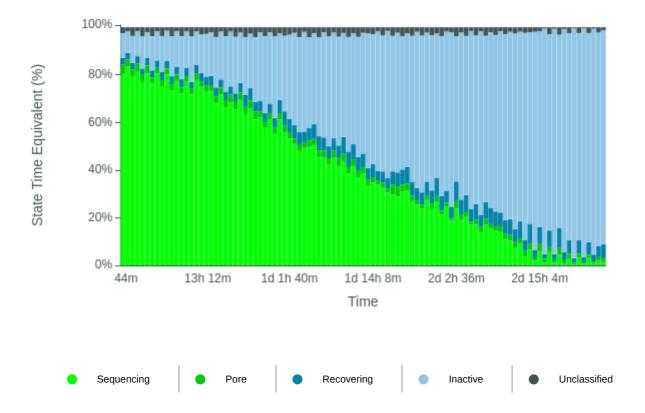


# **Read Length Histogram Basecalled Bases**

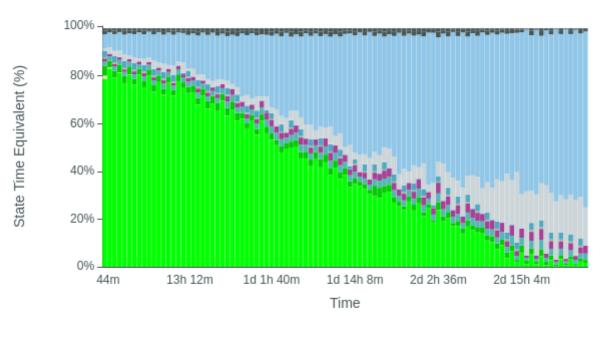
Estimated N50: 9.19 kb



#### **Duty Time Grouped**

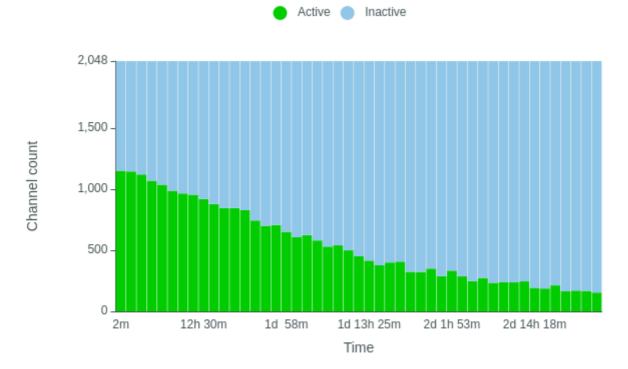


# **Duty time Categorised**

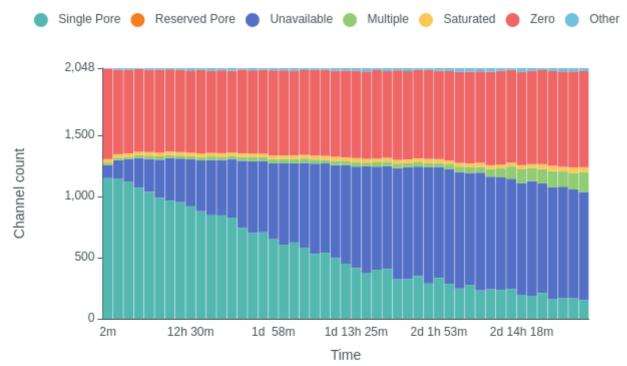




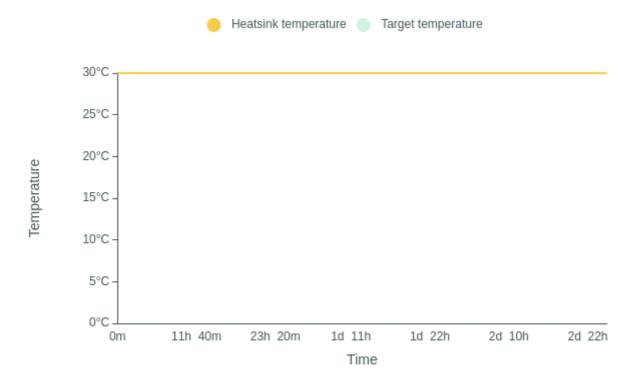
# **Mux Scan Grouped**



## **Mux Scan Categorised**



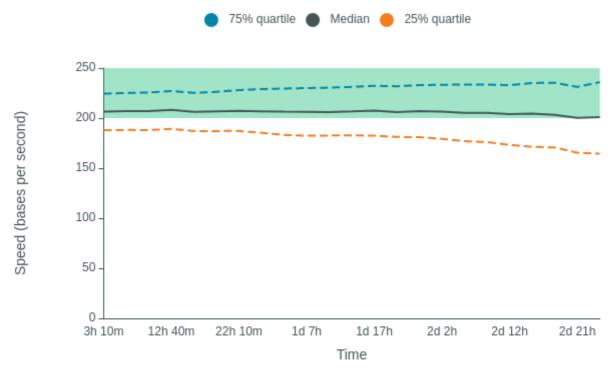
## **Temperature History**



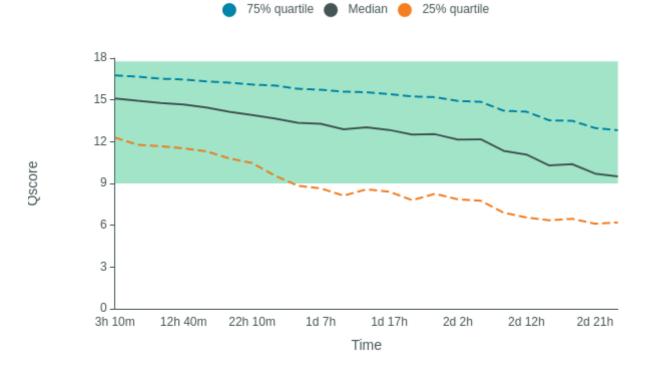
# <u>Bias Voltage History</u>



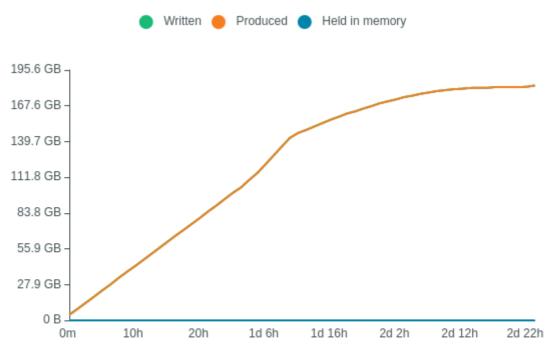
# **Translocation Speed**



## **QScore**



# **Disk Write Performance**



#### **Run Debug Messages**

- Mux scan for flow cell FAU08677 has found a total of 154 pores. 134 pores available for immediate sequencing June 13, 13:52
- Performing Mux Scan June 13, 13:50
- Mux scan for flow cell FAU08677 has found a total of 168 pores. 147 pores available for immediate sequencing June 13, 12:20
- Performing Mux Scan June 13, 12:17
- Mux scan for flow cell FAU08677 has found a total of 170 pores. 147 pores available for immediate sequencing June 13, 10:47
- Performing Mux Scan June 13, 10:45
- Mux scan for flow cell FAU08677 has found a total of 164 pores. 143 pores available for immediate sequencing June 13, 09:15
- Performing Mux Scan June 13, 09:12
- Mux scan for flow cell FAU08677 has found a total of 212 pores. 179 pores available for immediate sequencing June 13, 07:42
- Performing Mux Scan June 13, 07:40
- Mux scan for flow cell FAU08677 has found a total of 187 pores. 159 pores available for immediate sequencing June 13, 06:10
- Performing Mux Scan June 13, 06:07
- Mux scan for flow cell FAU08677 has found a total of 193 pores. 165 pores available for immediate sequencing June 13, 04:37
- Performing Mux Scan June 13, 04:35
- Mux scan for flow cell FAU08677 has found a total of 245 pores. 197 pores available for immediate sequencing June 13, 03:05
- Performing Mux Scan June 13, 03:02
- Mux scan for flow cell FAU08677 has found a total of 237 pores. 193 pores available for immediate sequencing June 13, 01:32
- Performing Mux Scan June 13, 01:30
- Mux scan for flow cell FAU08677 has found a total of 241 pores. 179 pores available for immediate sequencing June 13, 00:00
- Performing Mux Scan June 12, 23:57
- Mux scan for flow cell FAU08677 has found a total of 234 pores. 177 pores available for immediate sequencing June 12, 22:26
- Performing Mux Scan June 12, 22:24
- Mux scan for flow cell FAU08677 has found a total of 274 pores. 207 pores available for immediate sequencing June 12, 20:53
- Performing Mux Scan June 12, 20:50
- Mux scan for flow cell FAU08677 has found a total of 248 pores. 178 pores available for immediate sequencing June 12, 19:20
- Performing Mux Scan June 12, 19:17
- Mux scan for flow cell FAU08677 has found a total of 286 pores. 199 pores available for immediate sequencing June 12, 17:46
- Performing Mux Scan June 12, 17:44
- Mux scan for flow cell FAU08677 has found a total of 333 pores. 233 pores available for immediate sequencing June 12, 16:13
- Performing Mux Scan June 12, 16:10
- Mux scan for flow cell FAU08677 has found a total of 289 pores. 185 pores available for immediate sequencing June 12, 14:39
- Performing Mux Scan June 12, 14:37
- Mux scan for flow cell FAU08677 has found a total of 350 pores. 227 pores available for immediate sequencing June 12, 13:06

- Performing Mux Scan June 12, 13:03
- Mux scan for flow cell FAU08677 has found a total of 323 pores. 215 pores available for immediate sequencing June 12, 11:32
- Performing Mux Scan June 12, 11:30
- Mux scan for flow cell FAU08677 has found a total of 324 pores. 208 pores available for immediate sequencing June 12, 09:59
- Performing Mux Scan June 12, 09:56
- Mux scan for flow cell FAU08677 has found a total of 407 pores. 266 pores available for immediate sequencing June 12, 08:25
- Performing Mux Scan June 12, 08:23
- Mux scan for flow cell FAU08677 has found a total of 400 pores. 251 pores available for immediate sequencing June 12, 06:52
- Performing Mux Scan June 12, 06:49
- Mux scan for flow cell FAU08677 has found a total of 377 pores. 241 pores available for immediate sequencing June 12, 05:18
- Performing Mux Scan June 12, 05:16
- Mux scan for flow cell FAU08677 has found a total of 415 pores. 249 pores available for immediate sequencing June 12, 03:45
- Performing Mux Scan June 12, 03:42
- Mux scan for flow cell FAU08677 has found a total of 451 pores. 269 pores available for immediate sequencing June 12, 02:11
- Performing Mux Scan June 12, 02:09
- Mux scan for flow cell FAU08677 has found a total of 499 pores. 294 pores available for immediate sequencing June 12, 00:38
- Performing Mux Scan June 12, 00:35
- Mux scan for flow cell FAU08677 has found a total of 540 pores. 311 pores available for immediate sequencing June 11, 23:04
- Performing Mux Scan June 11, 23:02
- Mux scan for flow cell FAU08677 has found a total of 530 pores. 304 pores available for immediate sequencing June 11, 21:31
- Performing Mux Scan June 11, 21:28
- Mux scan for flow cell FAU08677 has found a total of 580 pores. 316 pores available for immediate sequencing June 11, 19:57
- Performing Mux Scan June 11, 19:55
- Mux scan for flow cell FAU08677 has found a total of 624 pores. 348 pores available for immediate sequencing June 11, 18:24
- Performing Mux Scan June 11, 18:21
- Mux scan for flow cell FAU08677 has found a total of 606 pores. 329 pores available for immediate sequencing June 11, 16:50
- Performing Mux Scan June 11, 16:48
- Mux scan for flow cell FAU08677 has found a total of 651 pores. 350 pores available for immediate sequencing June 11, 15:17
- Performing Mux Scan June 11, 15:14
- Mux scan for flow cell FAU08677 has found a total of 707 pores. 378 pores available for immediate sequencing June 11, 13:43
- Performing Mux Scan June 11, 13:41
- Mux scan for flow cell FAU08677 has found a total of 701 pores. 372 pores available for immediate sequencing June 11, 12:10
- Performing Mux Scan June 11, 12:07
- Mux scan for flow cell FAU08677 has found a total of 743 pores. 377 pores available for immediate sequencing June 11, 10:36
- Performing Mux Scan June 11, 10:34

- Mux scan for flow cell FAU08677 has found a total of 827 pores. 406 pores available for immediate sequencing June 11, 09:03
- Performing Mux Scan June 11, 09:00
- Mux scan for flow cell FAU08677 has found a total of 845 pores. 414 pores available for immediate sequencing June 11, 07:29
- Performing Mux Scan June 11, 07:27
- Mux scan for flow cell FAU08677 has found a total of 846 pores. 408 pores available for immediate sequencing June 11, 05:56
- Performing Mux Scan June 11, 05:53
- Mux scan for flow cell FAU08677 has found a total of 879 pores. 422 pores available for immediate sequencing June 11, 04:22
- Performing Mux Scan June 11, 04:20
- Mux scan for flow cell FAU08677 has found a total of 919 pores. 436 pores available for immediate sequencing June 11, 02:49
- Performing Mux Scan June 11, 02:47
- Mux scan for flow cell FAU08677 has found a total of 954 pores. 449 pores available for immediate sequencing June 11, 01:15
- Performing Mux Scan June 11, 01:13
- Mux scan for flow cell FAU08677 has found a total of 965 pores. 442 pores available for immediate sequencing June 10, 23:42
- Performing Mux Scan June 10, 23:40
- Mux scan for flow cell FAU08677 has found a total of 988 pores. 447 pores available for immediate sequencing June 10, 22:09
- Performing Mux Scan June 10, 22:06
- Mux scan for flow cell FAU08677 has found a total of 1037 pores. 456 pores available for immediate sequencing June 10, 20:35
- Performing Mux Scan June 10, 20:33
- Mux scan for flow cell FAU08677 has found a total of 1070 pores. 456 pores available for immediate sequencing June 10, 19:02
- Performing Mux Scan June 10, 18:59
- Mux scan for flow cell FAU08677 has found a total of 1120 pores. 464 pores available for immediate sequencing June 10, 17:28
- Performing Mux Scan June 10, 17:26
- Mux scan for flow cell FAU08677 has found a total of 1144 pores. 473 pores available for immediate sequencing June 10, 15:55
- Performing Mux Scan June 10, 15:52
- Mux scan for flow cell FAU08677 has found a total of 1150 pores. 480 pores available for immediate sequencing June 10, 14:21
- Performing Mux Scan June 10, 14:19
- Starting sequencing procedure June 10, 14:19
- Waiting up to 300 seconds for temperature to stabilise at 30.0°C June 10, 14:15