PromethION 2 Solo P2S-01310-B Final report



17 de dez. de 24, 11:23 UTC-3:00 — 20 de dez. de 24, 12:27 UTC-3:00 · Aflosc_genome_ULRnanopore · Aflosc_run1 · P2S-01310-B Protocol run ID: bb70e90f-5b66-4805-b0e2-e28675d2d3fa

Run summary | Run configuration | Sequence output | Run health | Run log

▲ Run summary

Estimated bases 11.75 Gb Reads generated 171.61 k Estimated N50 Fun time 73 hrs 1 min / 120 hrs 0 mins Elapsed time Run limit Run status STOPPED · By user

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DATA OUTPUT RUN DURATION

Estimated bases 11.75 Gb	Run time
Reads generated 171.61 k	73 hrs 1 min / 120 hrs 0 mins Elapsed time Run limit
Estimated N50 596.82 kb	Run status STOPPED · By user

Run configuration

RUN SETUP

Flow cell type FLO-PRO114M
Flow cell type alias FLO-PRO114M
Flow cell ID PAS53393
Kit type SQK-ULK114

RUN SETTINGS

Run limit 120 hrs
Pore scan freq. 1.5 hrs
Reserved pores On
Basecalling Off
Modified basecalling Off

△ DATA OUTPUT SETTINGS

FAST5 output Off FASTQ data output Off

POD5 data output One file per hour, or

500000000 bases per batch

BAM file output Off Bulk file output Off

Data location C:\data\.\Aflosc_genome_UL

Rnanopore/Aflosc_run1/2024 1217_0823_P2S-01310-B_PAS53393_bb70e90f

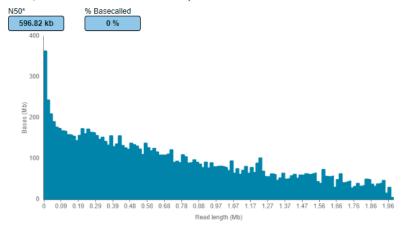
^ SOFTWARE VERSIONS

MinKNOW	24.11.8
Bream	8.2.5
Configuration	6.2.12
Dorado	7.6.7
MinKNOW Core	6.2.6

▲ Sequence output

▲ READ LENGTHS · OUTLIERS REMOVED

The read length graph shows the total number of bases vs the read length. The longest 1% of strands are classified as outliers, and excluded to allow focus on the main body of data.



∧ OUTLIERS

The longest 1% of strands are classified as outliers, and aggregated into groups to show their relative amounts.

Read length (Mb)	Bases (Mb)
1.966 - 2.097	76.4
2.097 - 2.228	36.51
2.228 - 2.327	4.6

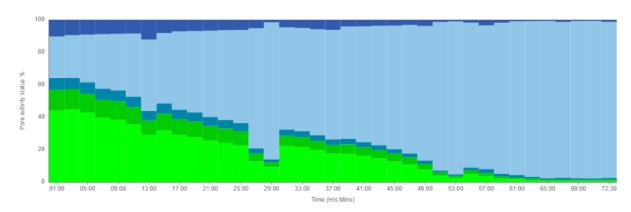
▲ PORE ACTIVITY

The Pore activity graph shows the performance of your sample as it is being sequenced during a run.









Basecalled

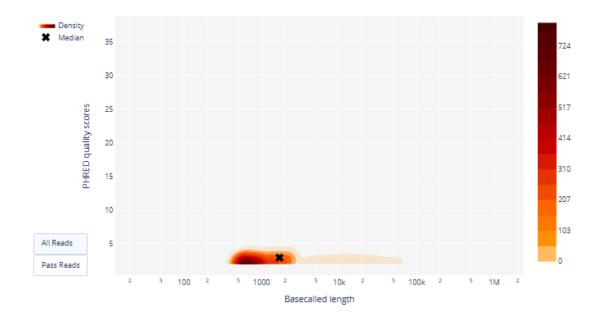
All Reads		
Pass Reads		

Reads	Bases	Med Read Length	N50 Length		Active Channels	Run Duration (h)
172,333	7,072,642,89	1,706.00	380,422.00	2.92	2,602	72.95

All Reads
Pass Reads

Reads	Bases	Med Read Length	N50 Length		Active Channels	Run Duration (h)
15,579	147,898,896	3,287.00	23,124.00	17.87	2,442	72.74

Basecalled reads length vs reads PHRED quality



Basecalled reads length vs reads PHRED quality

