

Sequencing Report

V65_v2

3	Flow cells
121.1 GB	Total bases
29.0 kB	N50
1763.54 GB	Total disk usage

Nanopype v1.0.1-11-ga124850

Nanopype

1 Flow cells

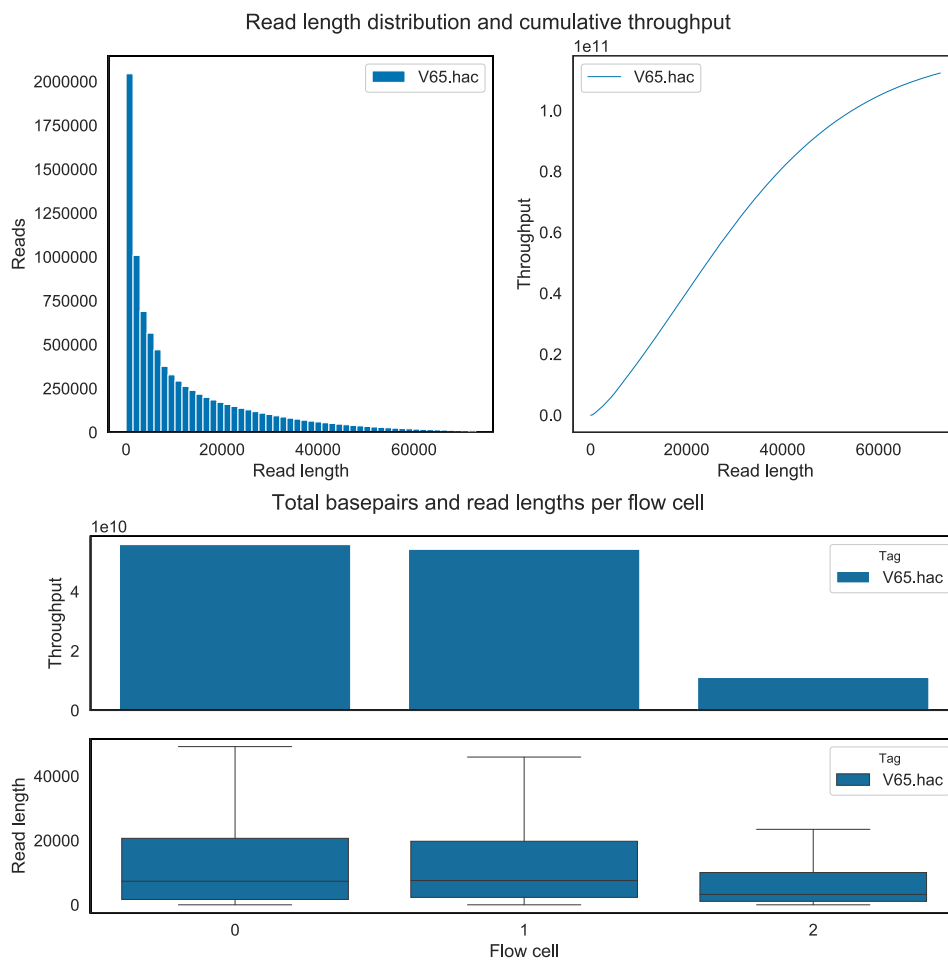
0: 20200708_PAE91419_FLO-PRO002_SQK-LSK109_V65_WT
1: 20200709_PAE86216_FLO-PRO002_SQK-LSK109_V65_WT
2: 20200710_FAN41394_FLO-MIN106_SQK-LSK109_V65_WT

2 Basecalling

2.1 Summary

		Sum	Mean	Median	N50	Maximum
V65.hac	guppy	121110031025	13239	6603	29048	1315417

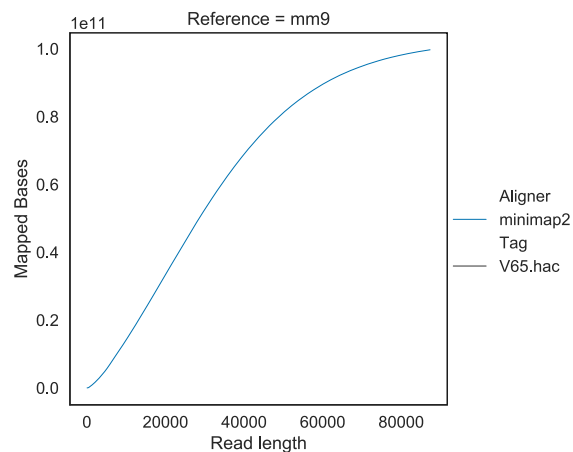
2.2 Guppy:



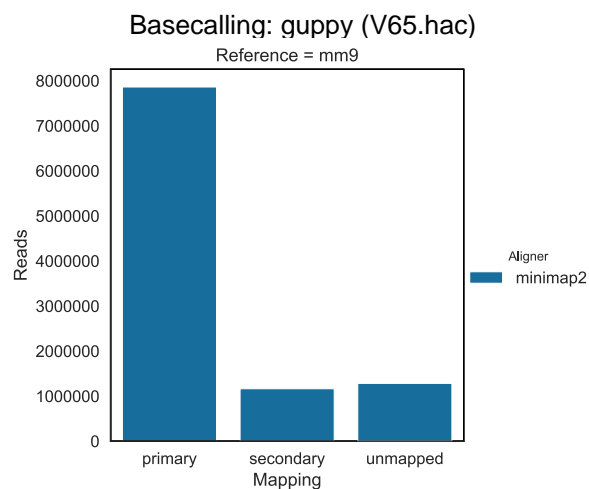
3 Alignments

3.1 Mapped bases (primary alignments)

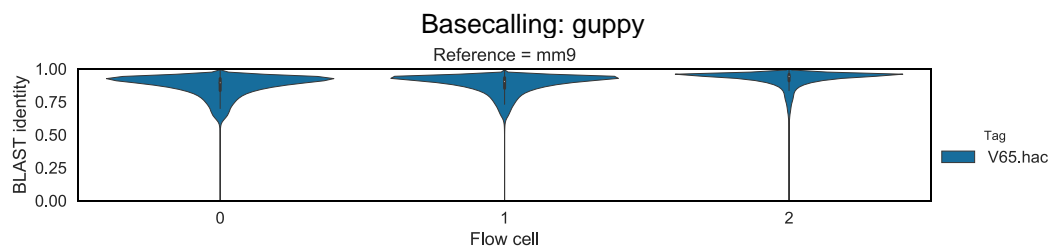
Basecalling: guppy



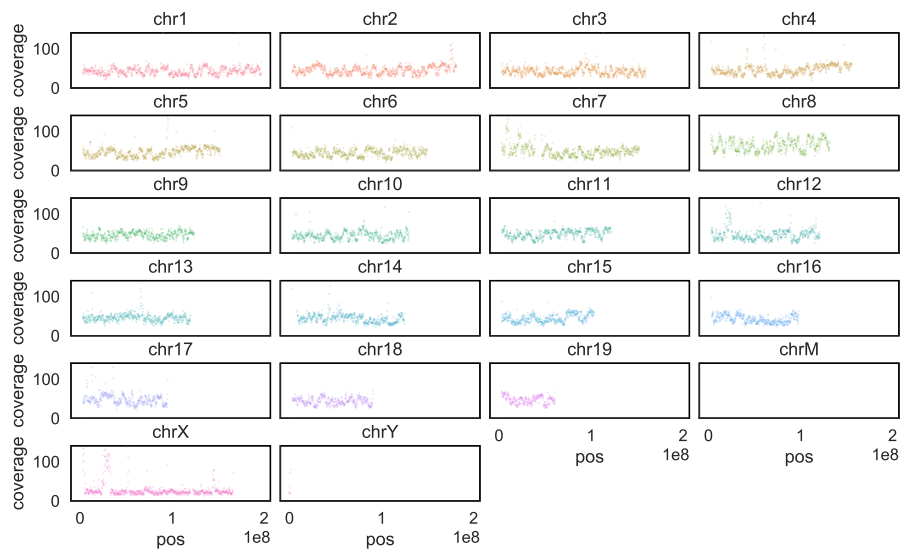
3.2 Mapped reads



3.3 Read identity (primary alignments, all aligners)



3.4 Coverage



4 Methylation

4.1 CpG coverage

