Sequencing Report

J1

3	Flow cells
60.7 GB	Total bases
17.6 kB	N50
1276.65 GB	Total disk usage

Nanopype v0.11.0
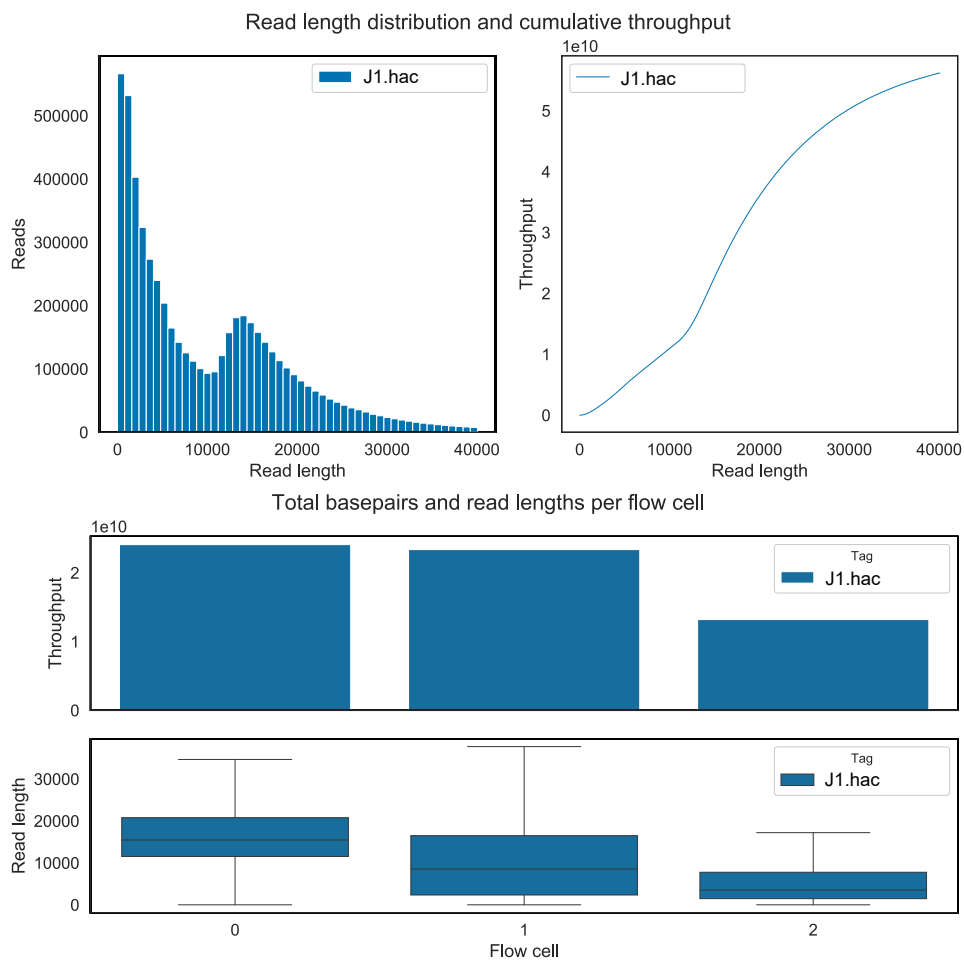
Nanopype

1 Flow cells

0:     20190620_FAK57924_FLO-MIN106_SQK-LSK109_J1
1:     20190620_FAK61976_FLO-MIN106_SQK-LSK109_J1
2:     20200506_FAL08409_FLO-MIN106_SQK-LSK109_J1

2 Basecalling

2.1 Summary

		Sum	Mean	Median	N50	Maximum
J1t.hac	guppy	60702273756	10500	7469	17636	237004

2.2 Guppy:
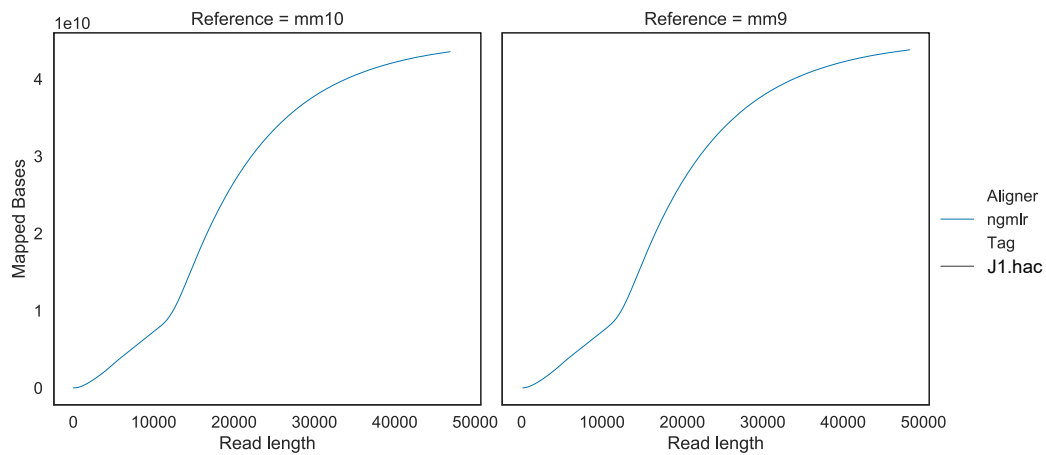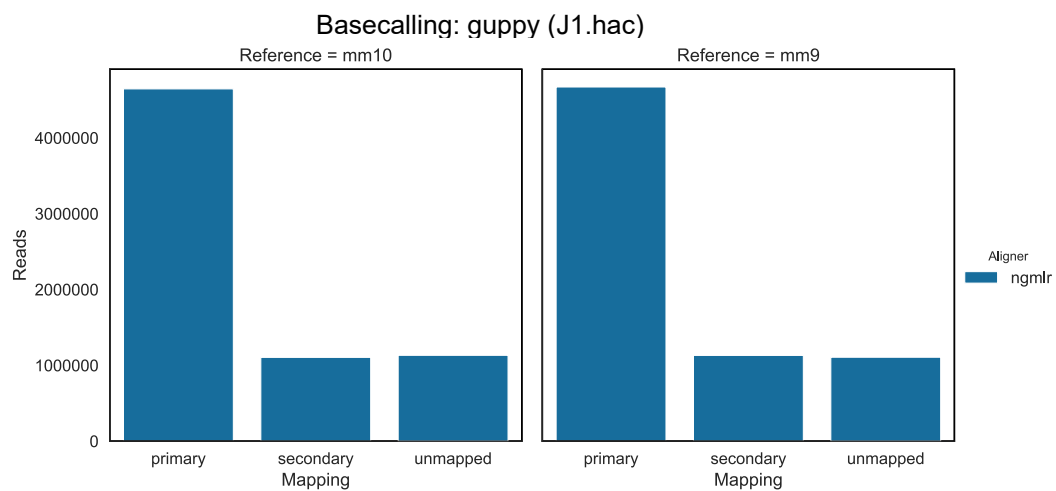

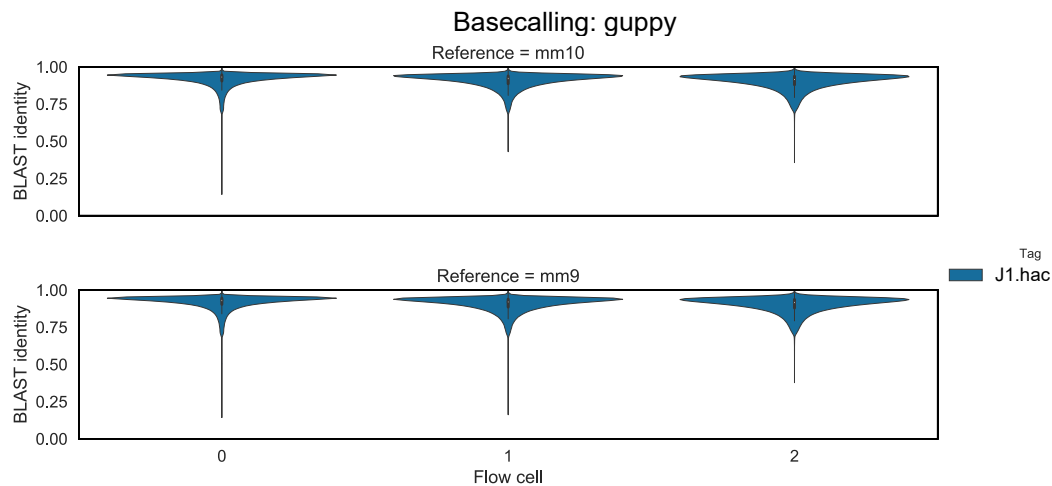
3 Alignments

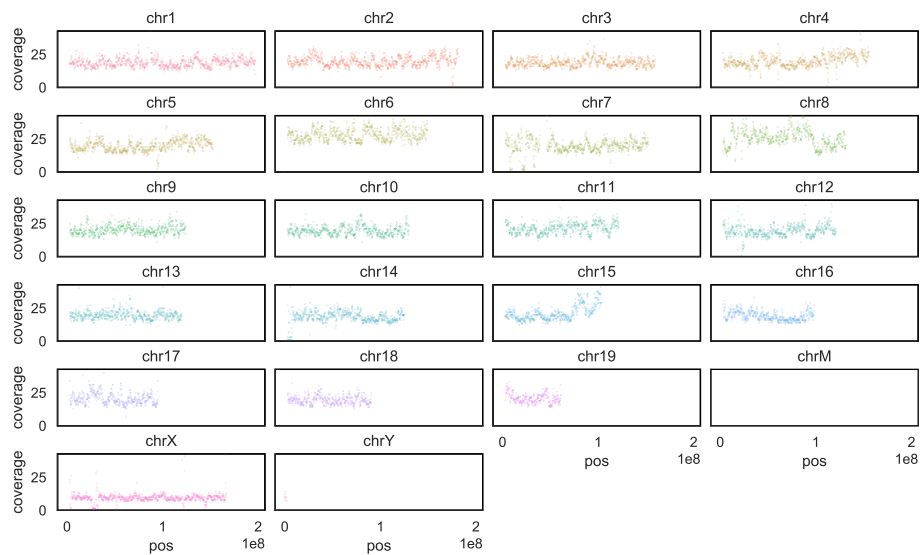## 3.1 Mapped bases (primary alignments)

Basecalling: guppy

3.2 Mapped reads

3.3 Read identity (primary alignments, all aligners)

3.4 Coverage

4 Methylation

4.1 CpG coverage

Basecalling: guppy, Alignment: ngmlr