


NANOME
Report



Running information

Base-calling results

Methylation-calling results

NANOME pipeline report for Oxford Nanpore sequencing data EcoliDemo

Running information

Running summary

Title	Information
dsname	EcoliDemo
projectDir	/pod/2/li-lab/yang/workspace/nanome
workDir	/fastscratch/li-lab/nanome/work-ecoli
commandLine	nextflow run main.nf -resume -profile singularity,hpc -work-dir /fastscratch/li-lab/nanome/work-ecoli --outputDir /fastscratch/li-lab/nanome/outputs-ecoli -config conf/jax_hpc.config,conf/ecoli_demo.config --singularity_cache /fastscratch/li-lab/nanome/singularity-cache --cleanCache false
runName	intergalactic_bassi
start	2021-09-28T07:49:25.050-04:00
input	https://zenodo.org/record/5483859/files/ecoli_data_from_meteore.tar.gz
outputDir	/fastscratch/li-lab/nanome/outputs-ecoli

Base-calling results

Base-calling report

Title	Information
General summary	EcoliDemo
Active channels	46.0
Mean read length	13,320.6
Mean read quality	7.8
Median read length	13,178.5
Median read quality	7.2
Number of reads	50.0
Read length N50	13,605.0
STDEV read length	3,298.3
Total bases	666,028.0
Number, percentage and megabases of reads above quality cutoffs	
>Q5	50 (100.0%) 0.7Mb
>Q7	32 (64.0%) 0.4Mb
>Q10	6 (12.0%) 0.1Mb
>Q12	1 (2.0%) 0.0Mb
>Q15	0 (0.0%) 0.0Mb

Methylation-calling results

Methylation-calling report


Tool	CpGs
Nanopolish	1,598
Megalodon	2,482
DeepSignal	2,858
Guppy	1,344
Tombo	2,940
METEORE	2,302
DeepMod	3,259

Methylation-calling plots

Methylation % and coverage distribution for Nanopolish

NANOME

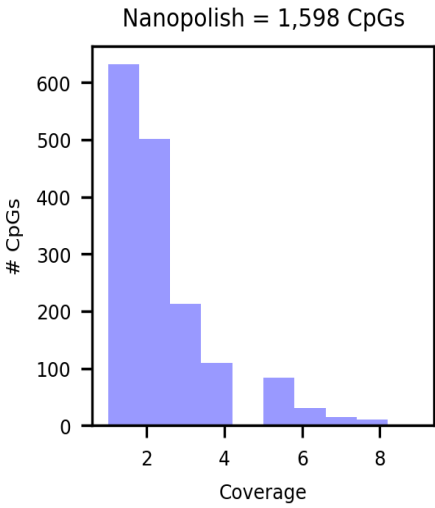
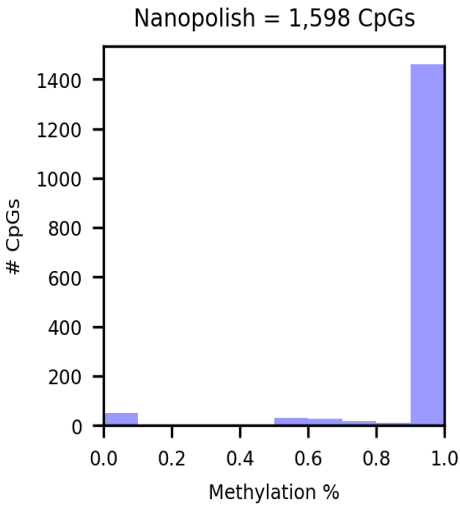
Report



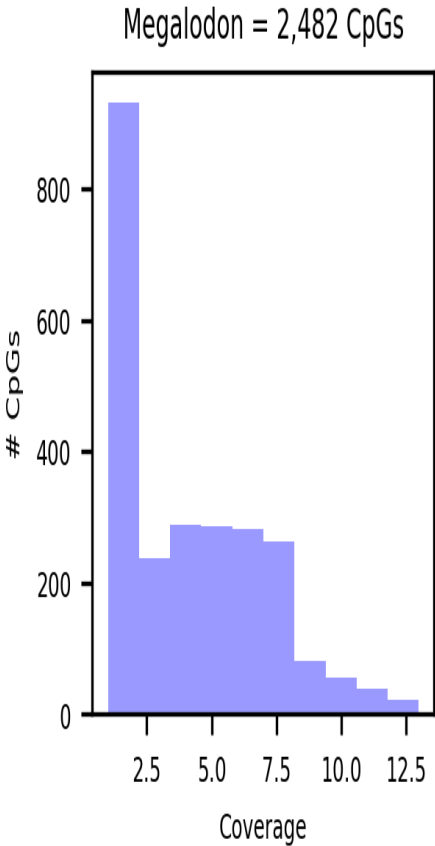
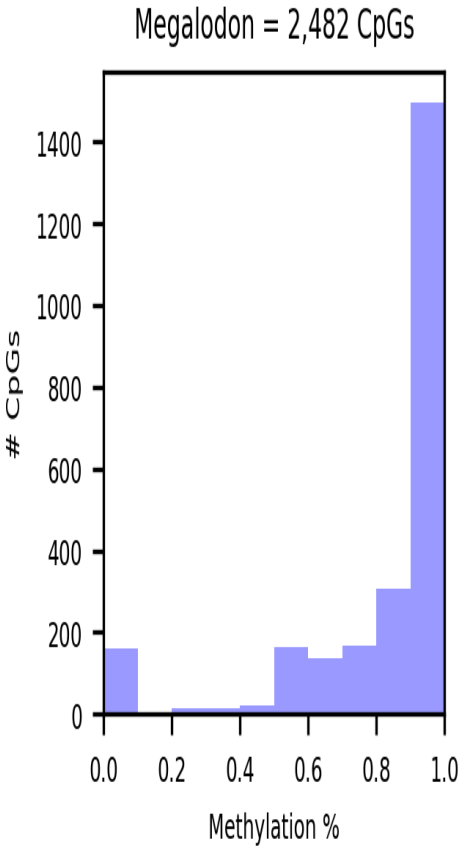
Running information

Base-calling results

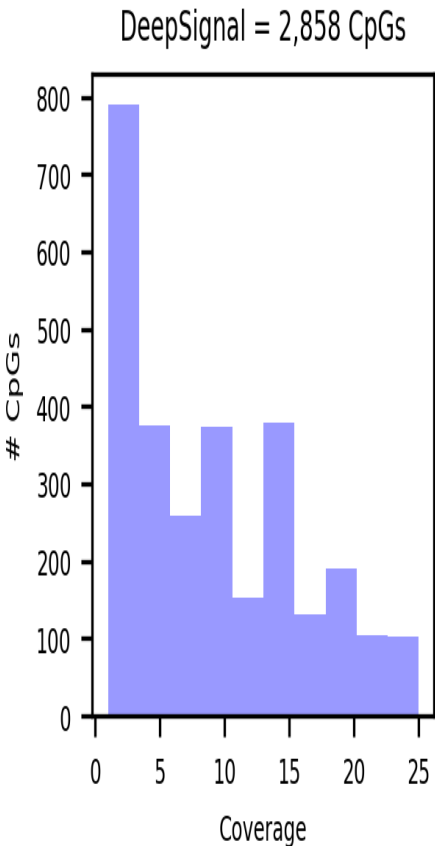
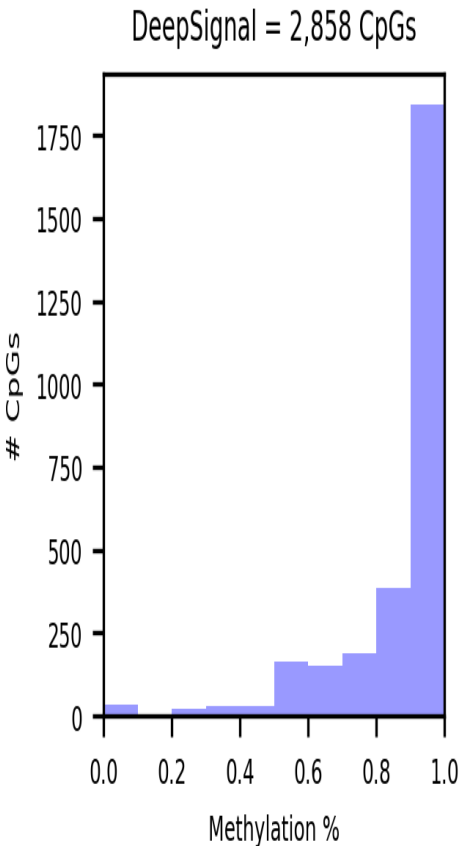
Methylation-calling results



Methylation % and coverage distribution for Megalodon



Methylation % and coverage distribution for DeepSignal



Methylation % and coverage distribution for Guppy

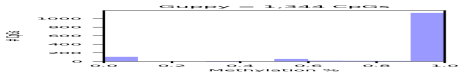
NANOME Report



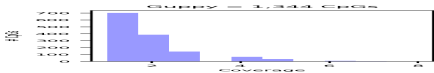
Running information

Base-calling results

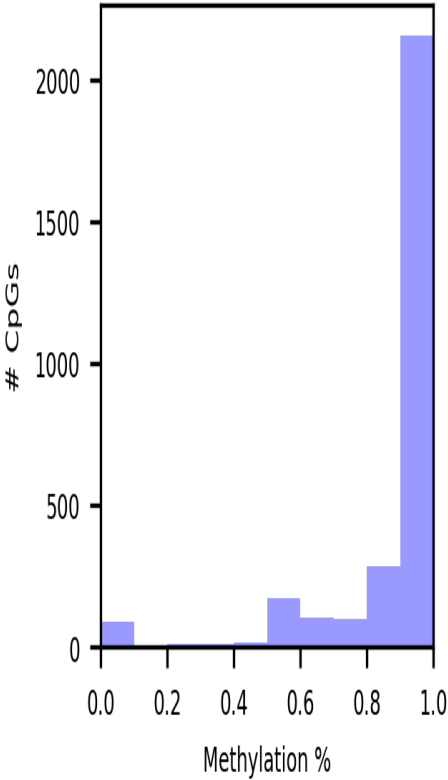
Methylation-calling results



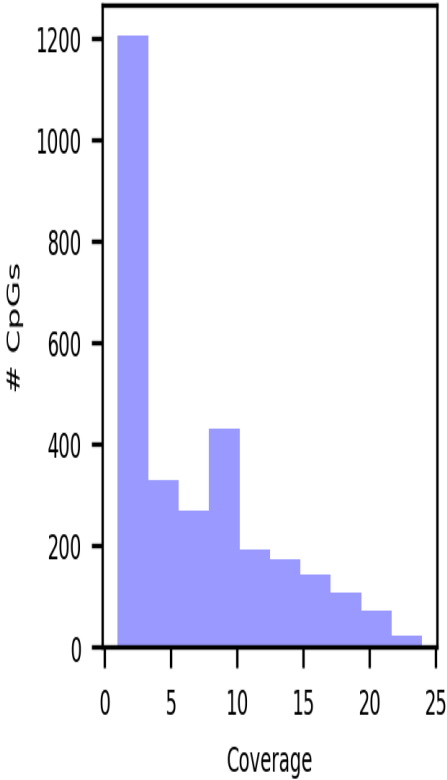
Methylation % and coverage distribution for Tombo



Tombo = 2,940 CpGs

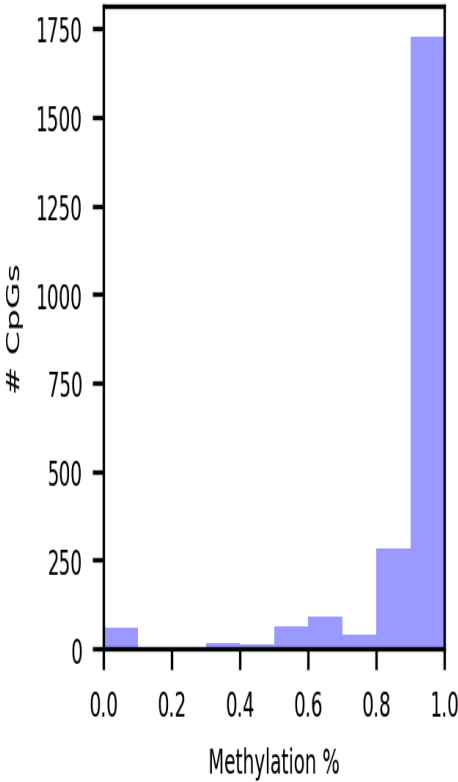


Tombo = 2,940 CpGs

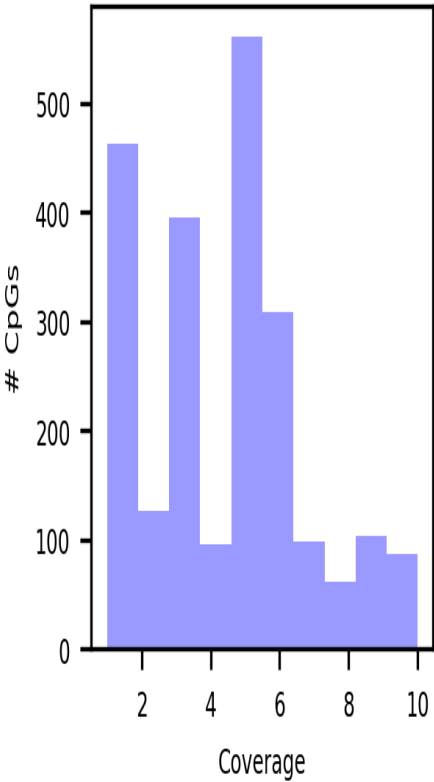


Methylation % and coverage distribution for METEORE

METEORE = 2,302 CpGs




METEORE = 2,302 CpGs



Methylation % and coverage distribution for DeepMod

NANOME

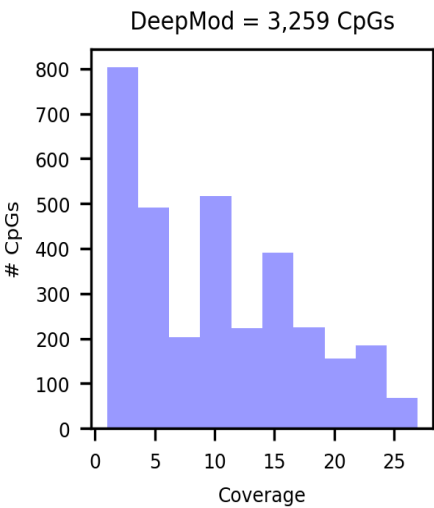
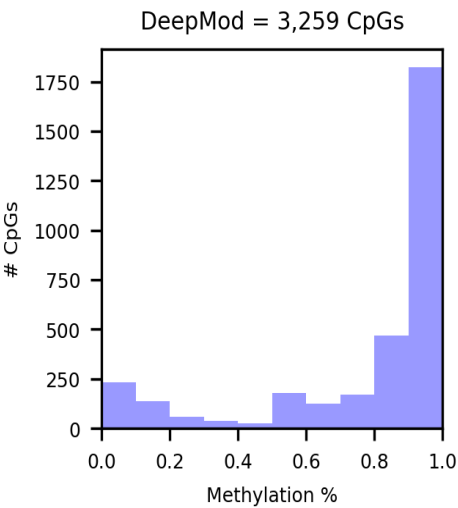
Report



Running information

Base-calling results

Methylation-calling results



-
1. Please use Chrome, Firefox or Safari for better browsing experience.
 2. Contact and report any bugs to <https://nanome.jax.org>.