

PycoQC report

Generated on 02/03/22 with pycoQC 2.5.0.23

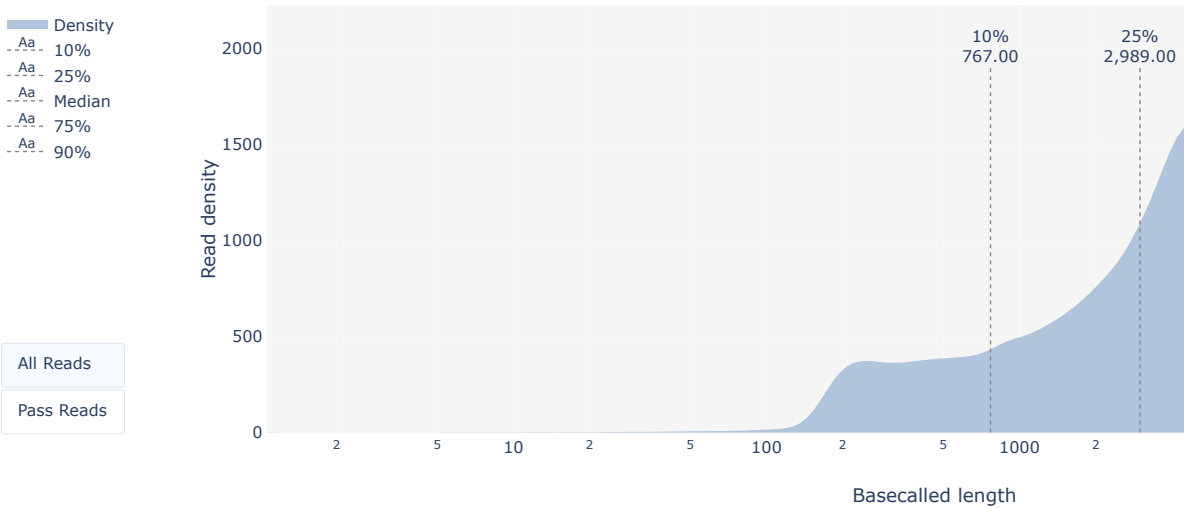
General run summary

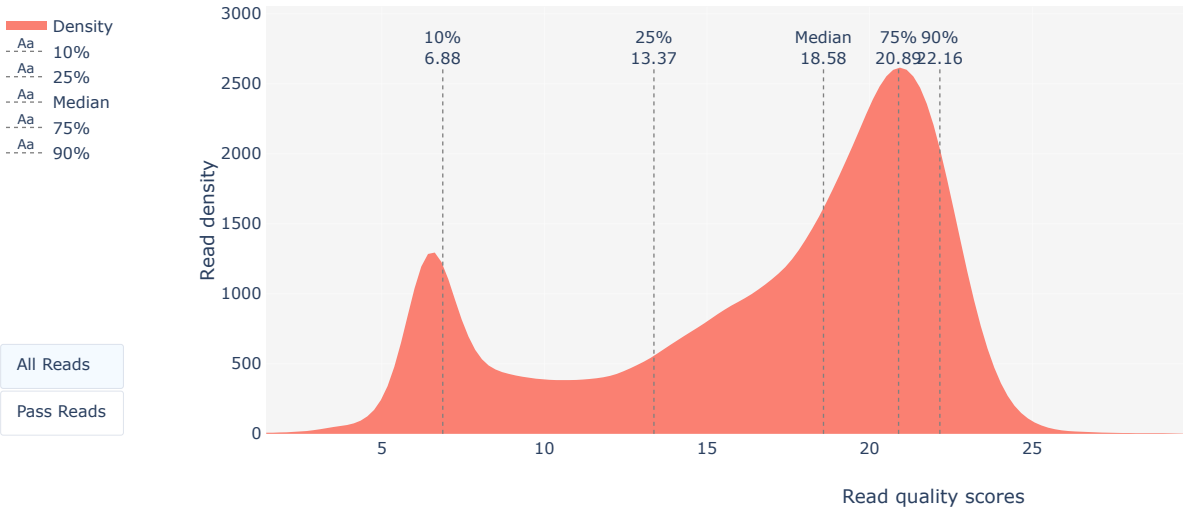
Status	Run Duration (h)	Active Channels	Number of
All Reads	287.97	2963	4
Pass Reads	287.97	2961	4

Basecall summary

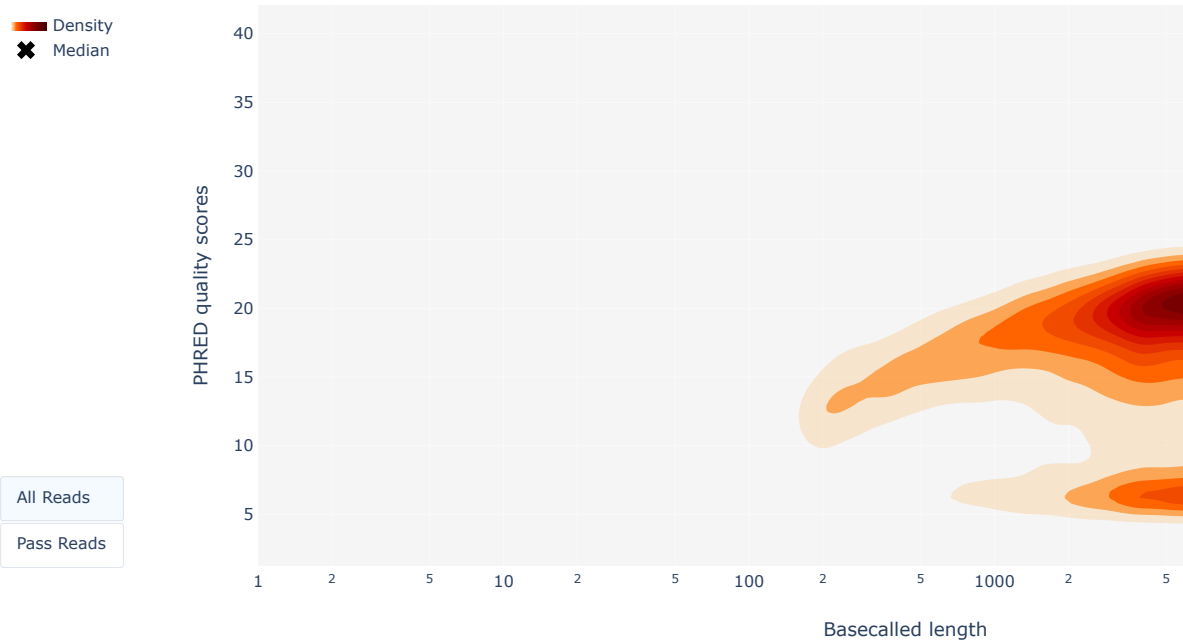
Status	Reads	Bases	N50	Media
All Reads	2.214248e+7	2.677626e+11	22300	
Pass Reads	1.976850e+7	2.420634e+11	22400	

Basecalled reads length





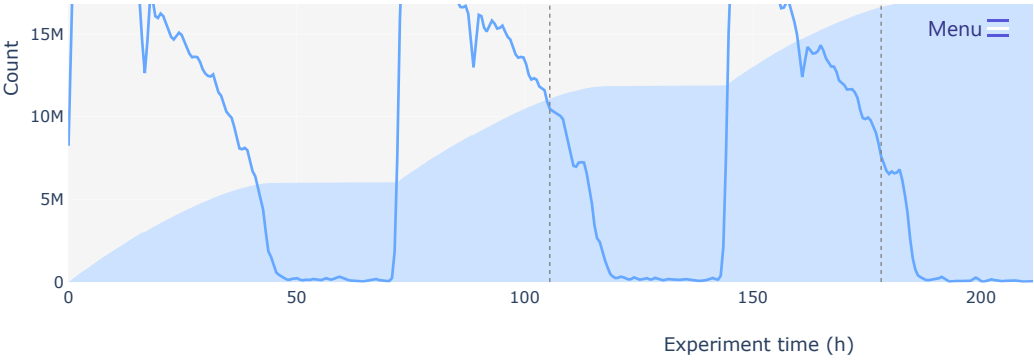
Basecalled reads length vs reads PHRED quality



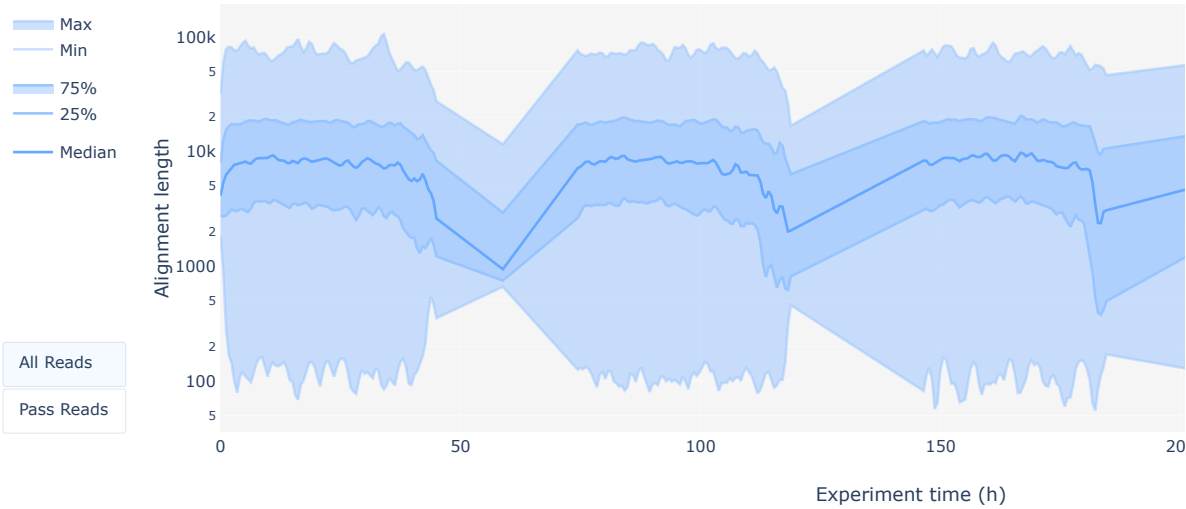
Output over experiment time



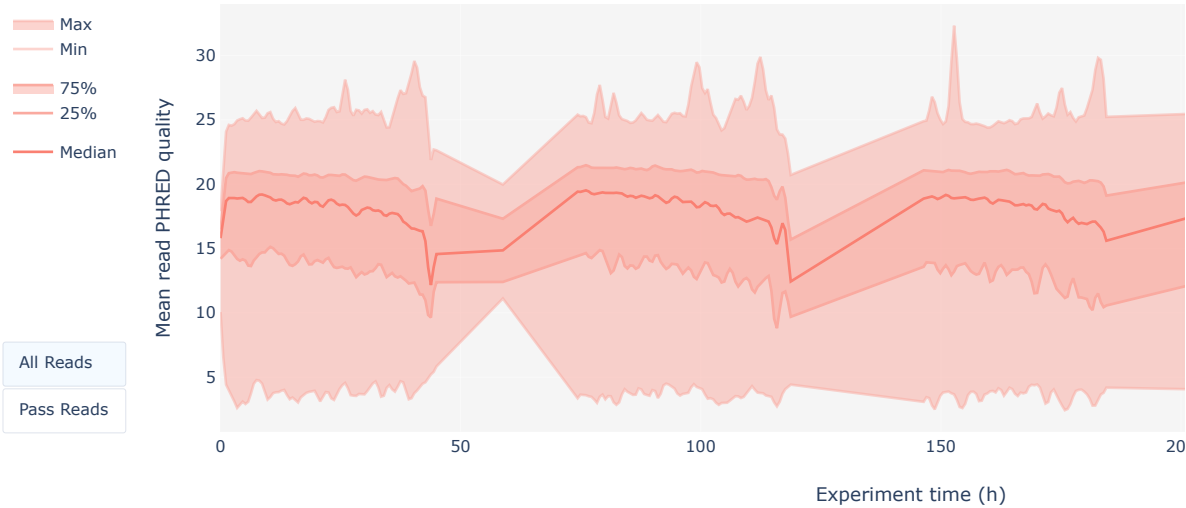
- All Reads
- Pass Reads
- All Bases
- Pass Bases

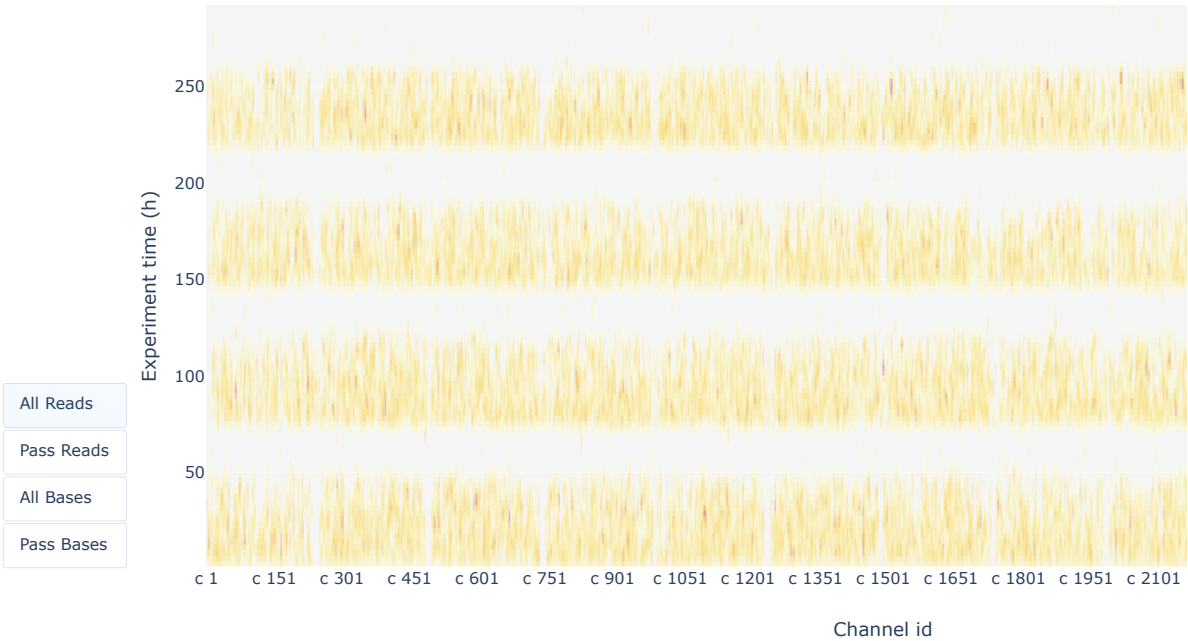


Read length over experiment time



Read quality over experiment time





Source summary files

- /binf-
isilon/sgarglab/people/rasmusab/data/whole_genome/QC/txts/all/guppy/sequencing_summary_5C.txt
- /binf-
isilon/sgarglab/people/rasmusab/data/whole_genome/QC/txts/all/guppy/sequencing_summary_5D.txt
- /binf-
isilon/sgarglab/people/rasmusab/data/whole_genome/QC/txts/all/guppy/sequencing_summary_5G.txt
- /binf-
isilon/sgarglab/people/rasmusab/data/whole_genome/QC/txts/all/guppy/sequencing_summary_5H.txt