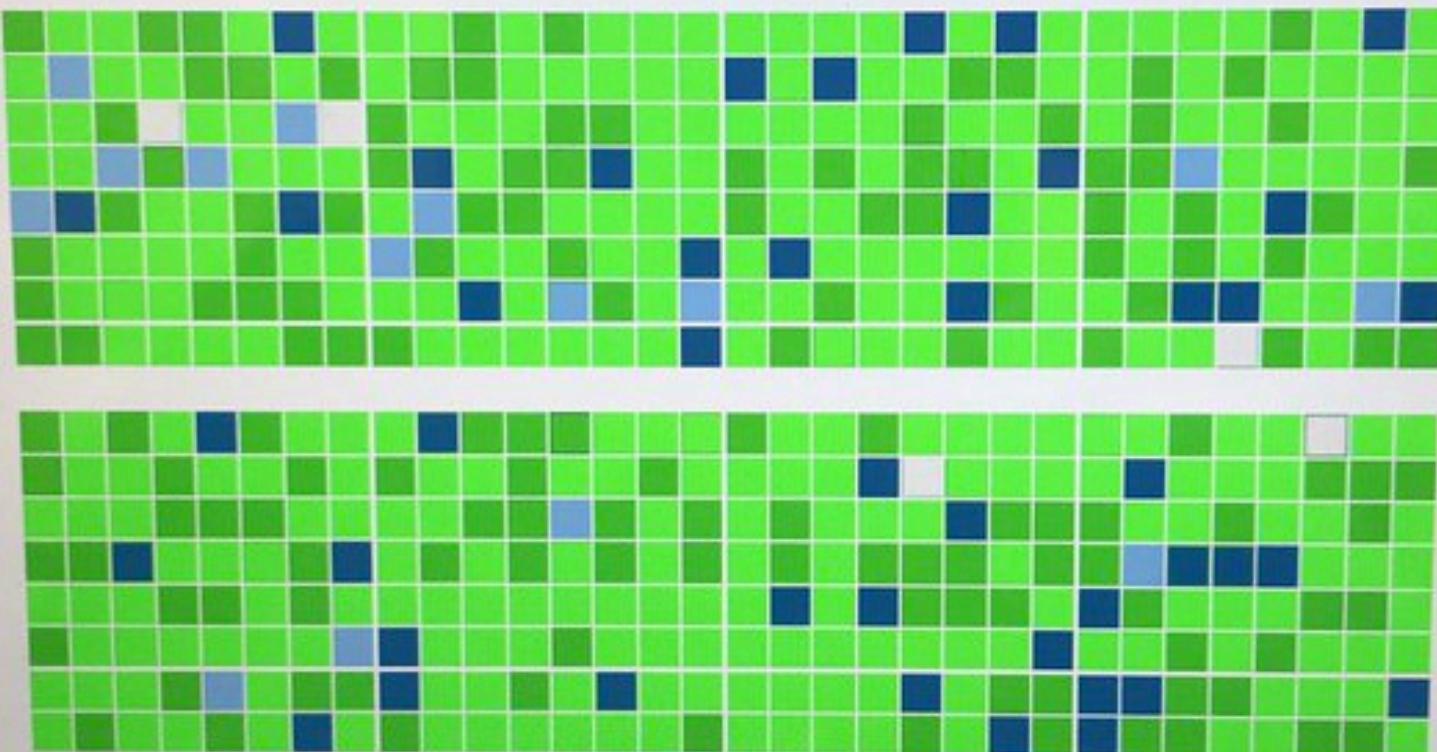


Channels Panel

Live status of each channel's state during sequencing



301
Sequencing



146
Pore



45
Recovering



15
Inactive

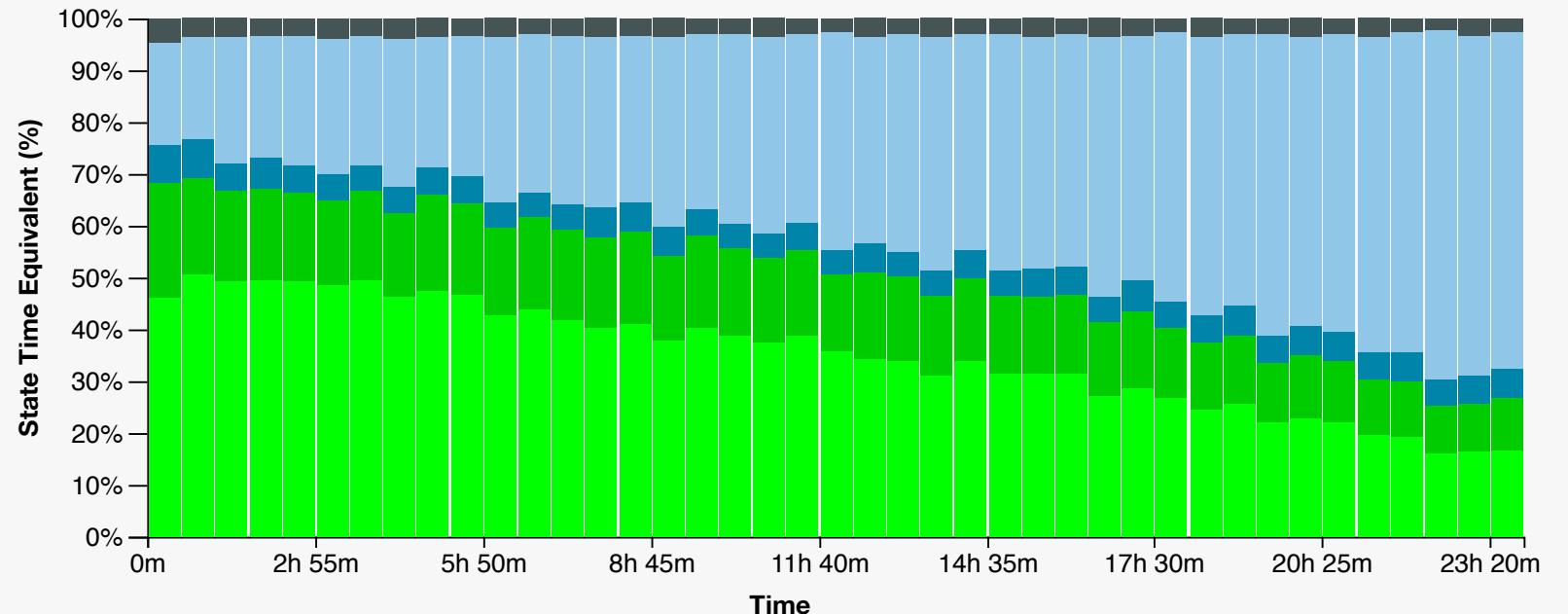


5
Unclassified



Duty Time

Summary of channel states over time



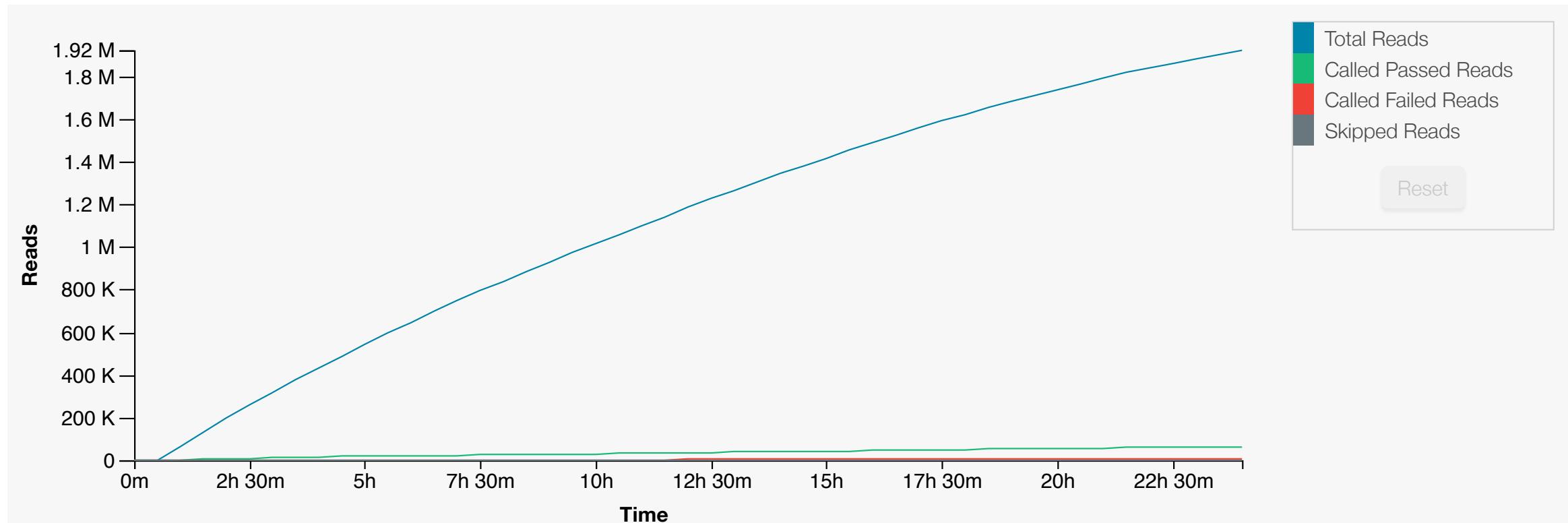
- Bucket size (minutes)
- 35
- Auto scale bucket size
- Display channels proportionately

Sequencing Pore Recovering Inactive Unclassified

More

Cumulative Output

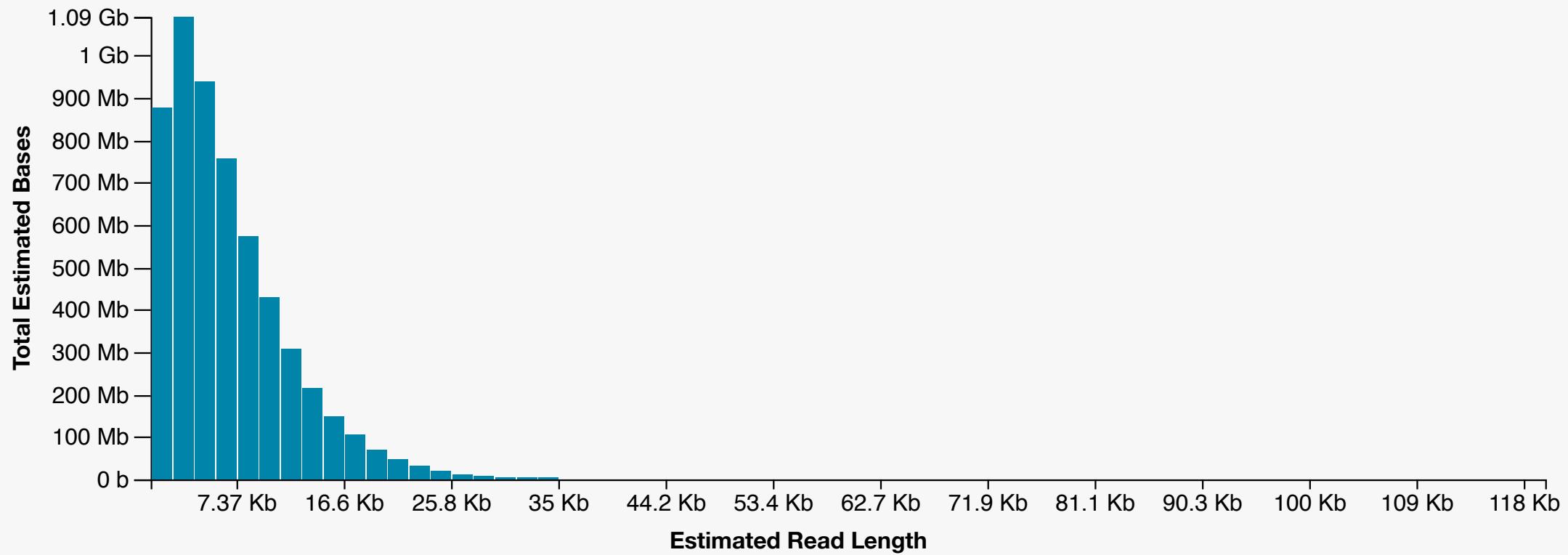
Output of run over time



Read Length Histogram

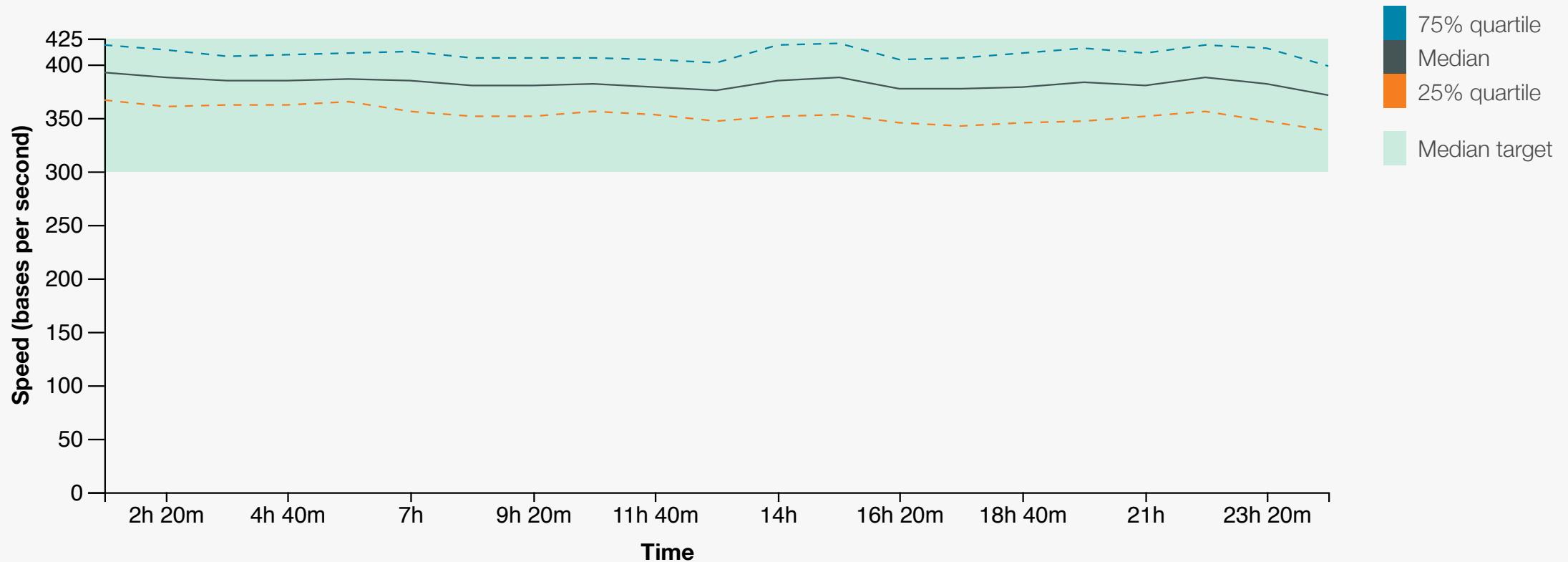
Summary read length distribution

Estimated N50: 3.85 Kb



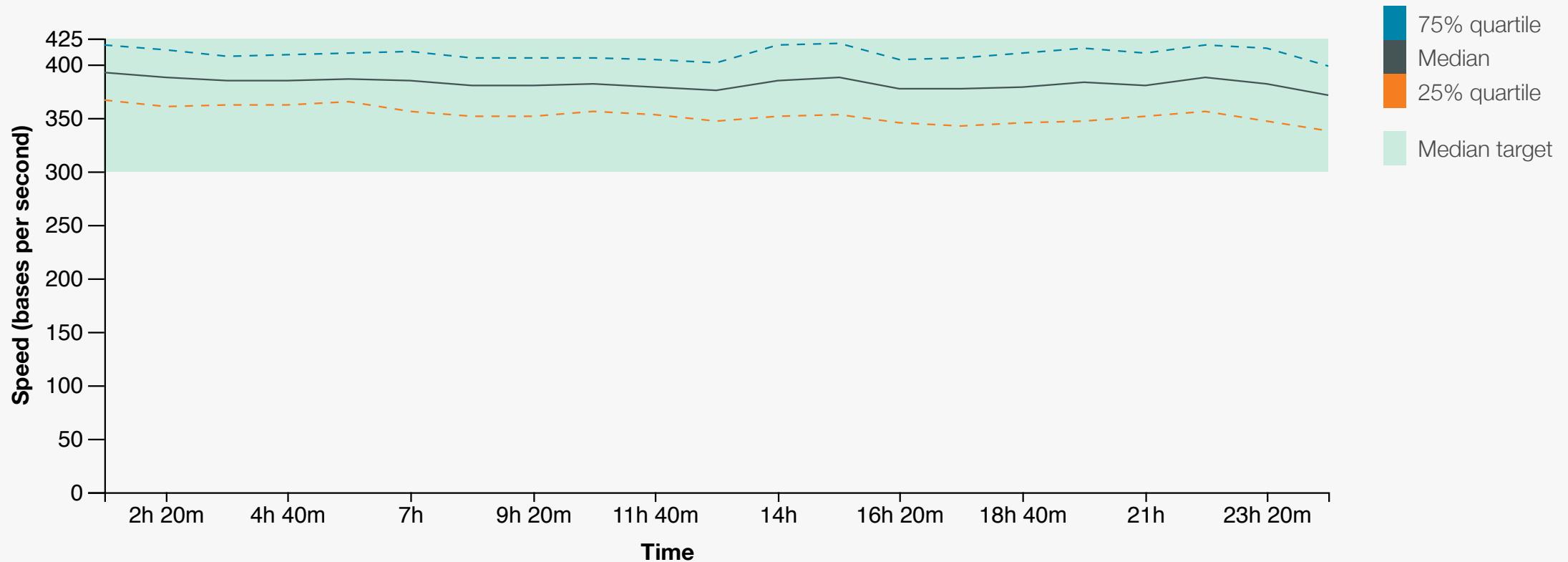
Translocation Speed and Qscore

Line charts displaying quartile values for translocation speed and qscore

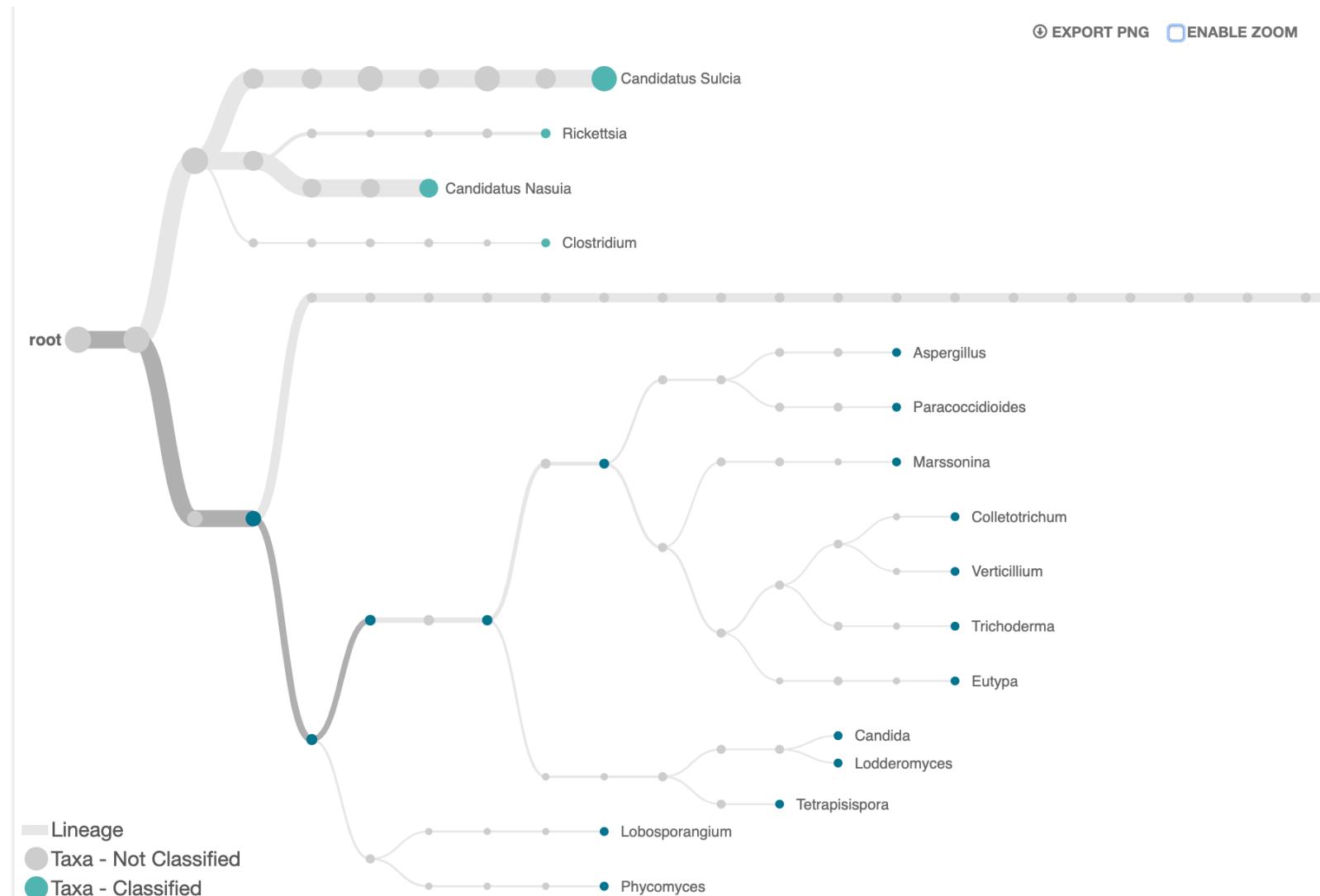
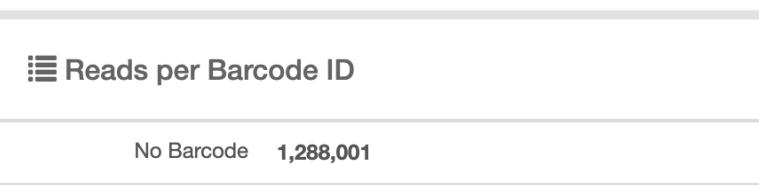


Translocation Speed and Qscore

Line charts displaying quartile values for translocation speed and qscore



Candidatus Sulcia	332,521
Candidatus Nasuia	202,121
Homo	67,795
Rickettsia	17,895
Phycomyces	4,412
Lobosporangium	2,205
Aspergillus	1,678
Paracoccidioides	1,492
Candida	1,386
Lodderomyces	1,294
Eutypa	1,254
Marssonina	1,200
« 1 2 3 4 5 ... 76 »	



https://epi2me.nanoporetech.com/workflow_instance/217061?token=5C5334B4-EC76-11E9-B6CA-A1E6BA8D1717