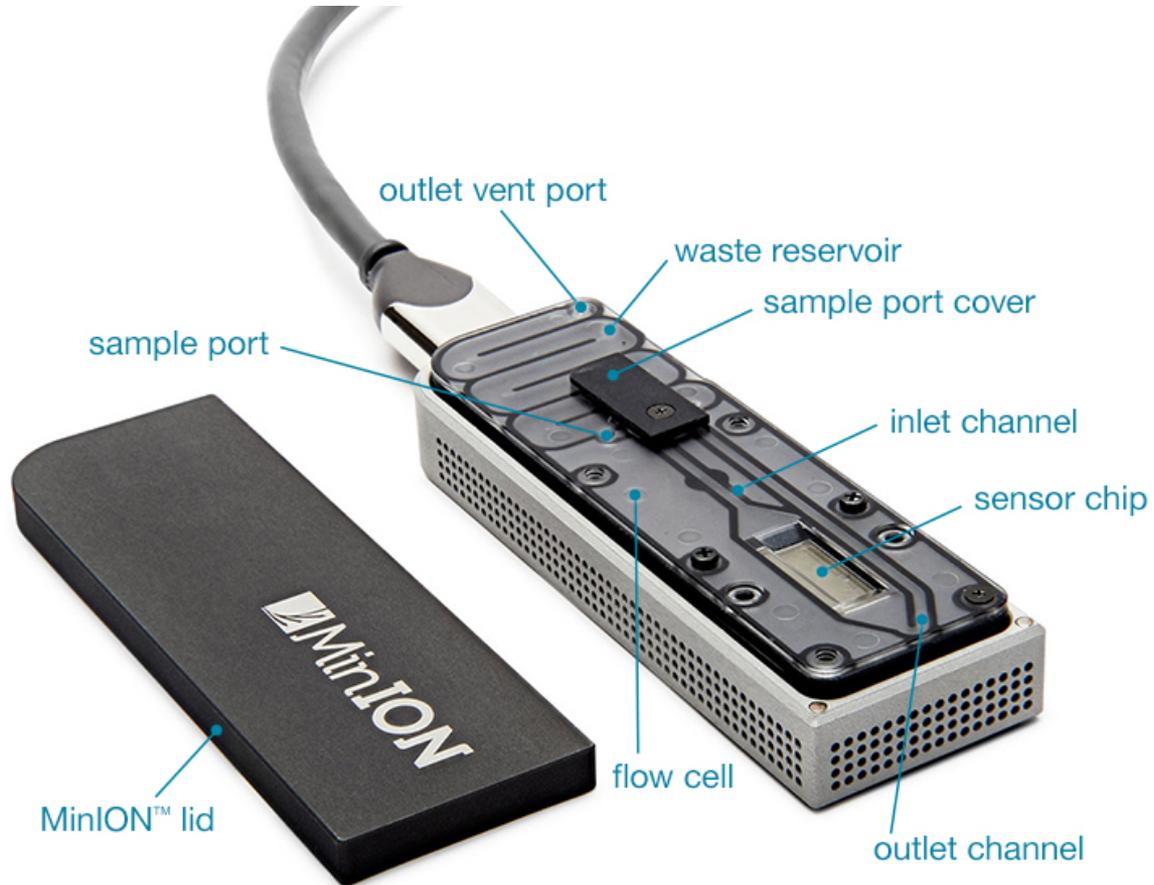
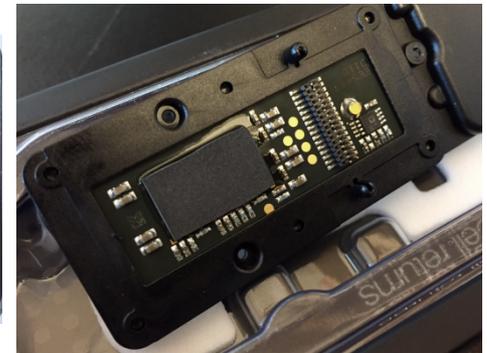


MinION Anatomy



Pre-Run Check List

- ① **Make sure required Software is installed:-**
 - MinKNOW** Control of MinION device & run parameters
 - Metricor** Cloud basecalling of event data
 - MinoTour** Live monitoring / control of run while sequencing
 - Chronolapse** Screen image grabber for record keeping
 - Poretools, poRe** Sequence extraction and data summaries
 - TeamViewer** Remote control of MinION computer
- ② **Confirm automatic software updates and sleep modes are disabled**
- ③ **Check computer SSD for available storage space >150Gb**
- ④ **Flowcell inspection** Remove bubbles in fluid lines and on surface of flowcell where possible if present, & confirm conductive heat pad is installed on bottom surface of ASIC chip.



Preparing & Loading MinION Device

- 1 Attach Flowcell to MinION device and plug into USB3 port



- 2 Start MinKNOW software and start device



- 3 Name Run and start platform-QC protocol



If flow-cell good proceed
(Single Good pores >650)

- 4 Prime flow-cell with 2x 500 μ l RB+Fuel for 10 mins each
(500 μ l 2xRB + 473.4 μ l H₂O + 26.6 μ l Fuel)



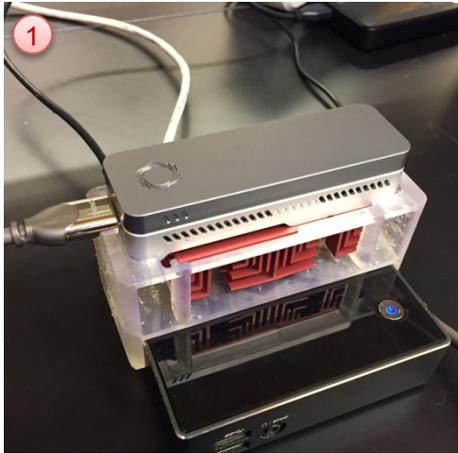
- 5 Load 150 μ l of Library in RB+Fuel
(75 μ l 2xRB + 65 μ l H₂O + 4 μ l Fuel + 6 μ l Pre-sequencing Mix)



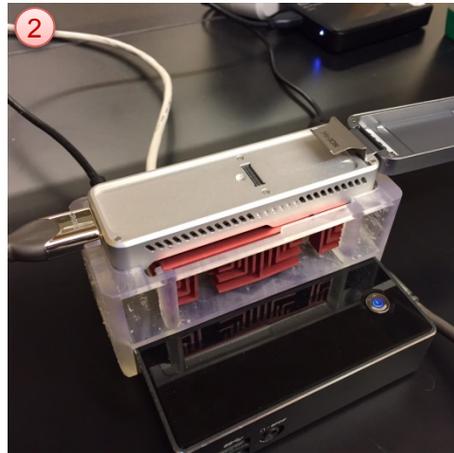
- 6 Name run and start Sequencing Protocol – Standard / Modified :o)
(Start Metrichor & required workflow plus screen capture software)



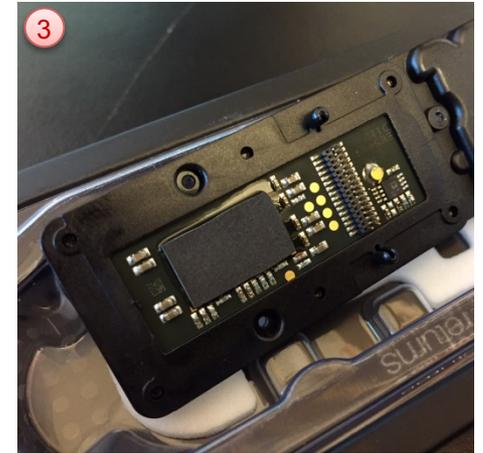
Inserting Flow-cell into MinION Device



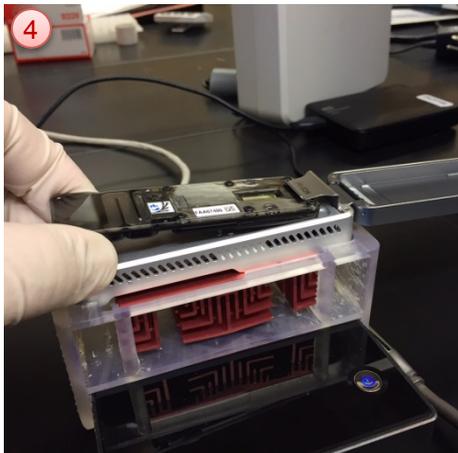
Ready MinION



Open lid



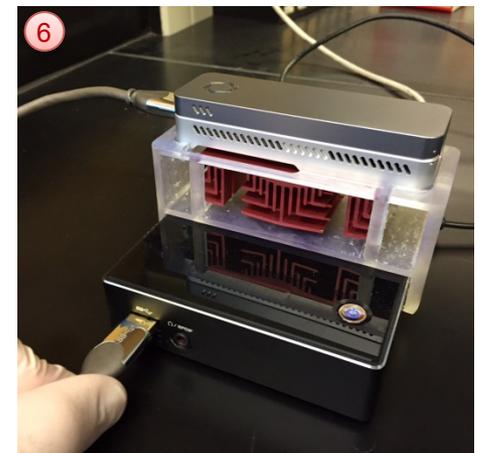
Unpack flow-cell, check heat pad intact & ASIC bubble free



Slide flow-cell into place



Make sure flow-cell is seated properly



Plug MinION into USB3 port

Flow-cell Quality Assessment

Platform-QC scans the 2048 channels as 4 Mux groups @ -180mV & reports back single good channel assignment to each of g1 to g4 groups with up to 512 wells each. These are best case numbers and you will have lower numbers as sequencing script start @ -140mV.

Good

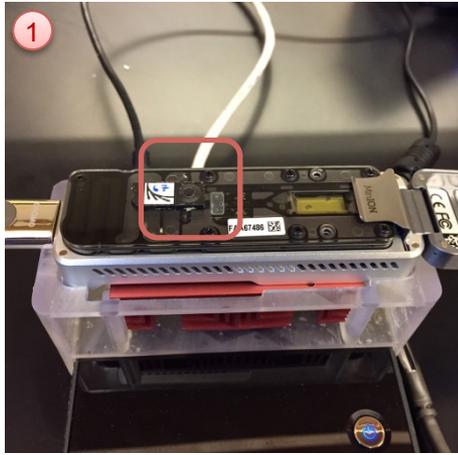


Bad

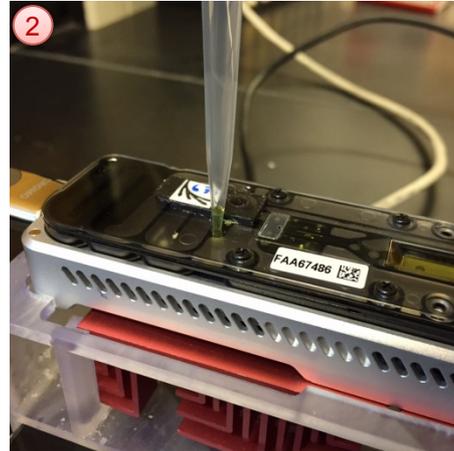


>650 Pore guarantee for
purchased flowcells

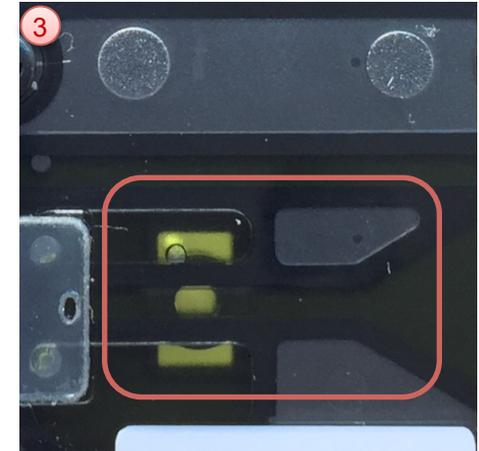
Priming and Loading Flow-cell



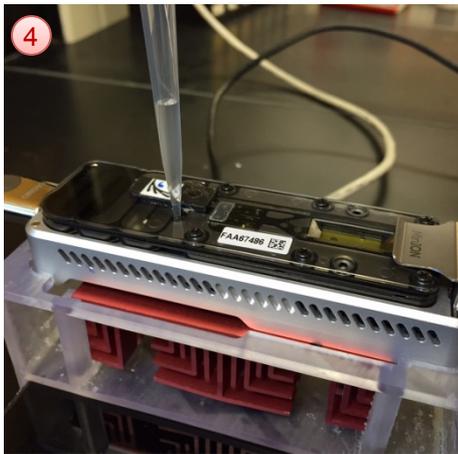
Open sample port



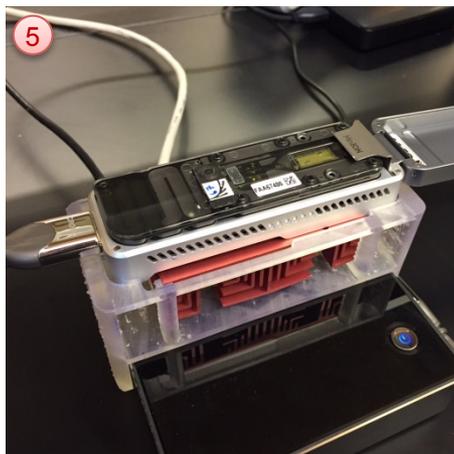
Slowly remove any air from sample port with pipette



Confirm fluidic channel is free of bubbles



2 x 10 min 500µl RB+Fuel followed by 150µl Library



Close sample port



Close lid and GO!

Start Run

① Start Sequencing Recipe Script

Name run and select required Recipe Script

② Start Chronolapse screen capture

Image grab every 30 seconds and as required at start

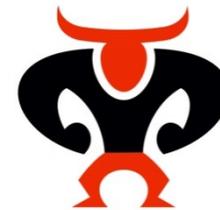
③ Start Metrichor workflow

Select desired workflow:- 2D basecall or WIMP

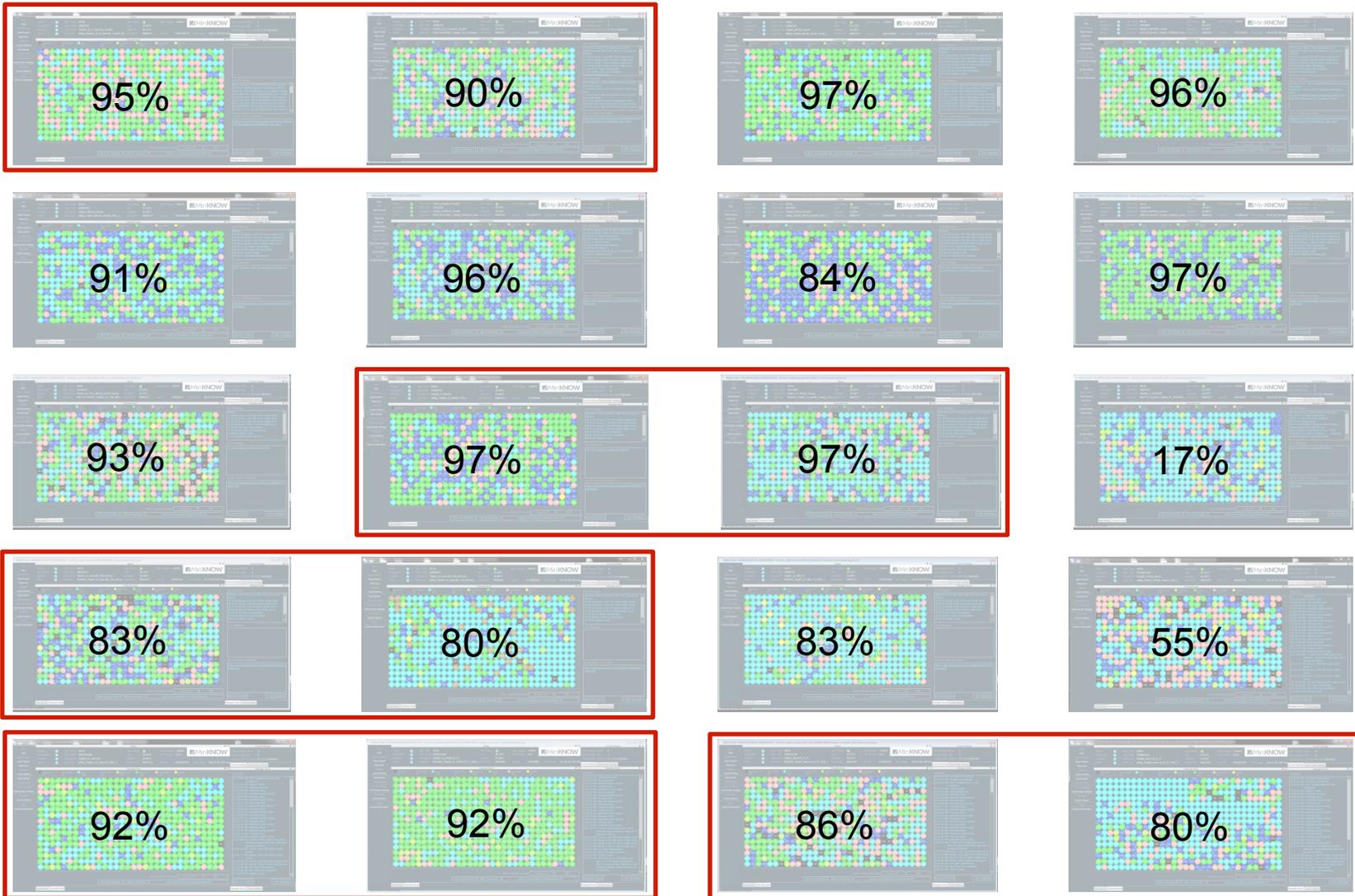
④ Start MinUp & Open MinoTour Browser Window

Issue command line parameters for read mapping and device control as below or use MinUP-GUI:

```
minUP.exe -dbh minotour.nottingham.ac.uk -dbu USER -pw PWD -w d:\data -f c:\reference\REF.txt -u USER -c -bwa -d -ip xxx.xxx.xxx.xxx -pin XXX -s minion
```



Good or Bad Library/Flowcell ?



% of Single Good pores in Strand 1hr post addition

Red boxes indicate same library addition

Single Good Pore % in Strand as Measure of Library Quality

Use the relative quantity of in Strand reads as a measure of library quality independent of overall pore number.

You would like >90% ~30mins after addition



$$\frac{292}{10 + 292} = 97\%$$

Native Client - MinKNOW 0.49.3.7 b201504271121 - Running /python/recipes/MAP_4x8hrs_Incrementing_Sequencing_Run.py

Status: Exp. Time 3611s, Raw Yield 10850752, Sample ID Chip54_GAERS10_Snutch, Data Set SNUTCH-GB-BX17_Chip54_GAERS10_Snut...

Asic Status: 24.46°C, Heatsink 34.88°C, MinION ID MN02521, Asic ID 3976896244, Disk Usage 95 of 232 GB Free

Channels Panel: saturated (7), -inf to 20pA (0), zero (36), SpA to +inf (119), Single Pore (10), strand (292), unavailable (46), multiple (2)

Grid of pore channels with numbers 1-388 and colors (green, blue, red, yellow).

Buttons: Show All Channels, Hide All Channels, Select Surrounding 8 Channels Mode OFF, Show Channels by State, Freeze OFF, OFF

Message Viewer: User Messages, Error Messages, Experiment Report, Clear Messages

2015-04-30 09:34:57.10

Script Tinkering

① Bias-Voltage Setting & Remux

Bias voltage directly controls induced current flow across the membrane, and current flow is used to assign pores to different categories. With time greater bias-voltage is required to produce current that is in the “single good” pore range. Bias-voltage is the master control. Selection or “remux” of pores at a particular targeted voltage must be carried out for maximum efficiency.

② Yield Monitoring

As pore numbers fall and/or reagent is depleted you will see a drop in the event yield over time. Gaining access to still functional, but unselected, pores once event accumulation rate drops below a set value makes much better use of a flow-cell. This lessens unproductive electrochemical gradient deterioration observed with standard recipe scripts.

③ Pore Shepherding

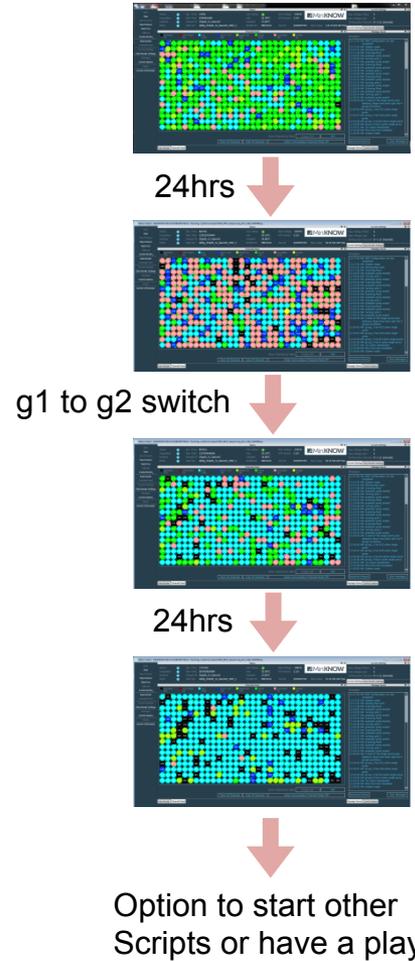
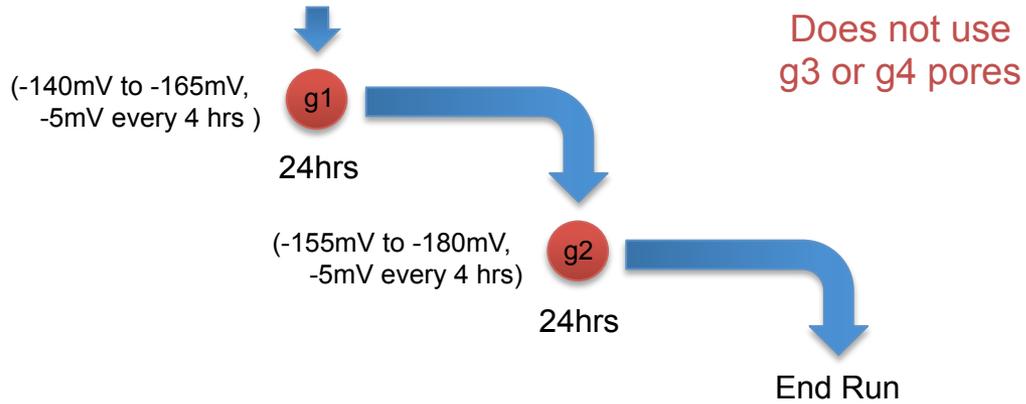


Wells actively sequencing experience an electrochemical gradient deterioration that is twice that of inactive pores, but even inactive pores show a deterioration that requires bias-voltage increases for optimal functioning. Because of the 4-1 (wells - electrical channel) nature of current flow-cells this results in a spreading of the optimal bias-voltage required for a particular well depending on its on/off history. To mitigate this, sub-peak bias-voltage selection can “shepherd” well population more tightly by attempting to deplete least used wells.

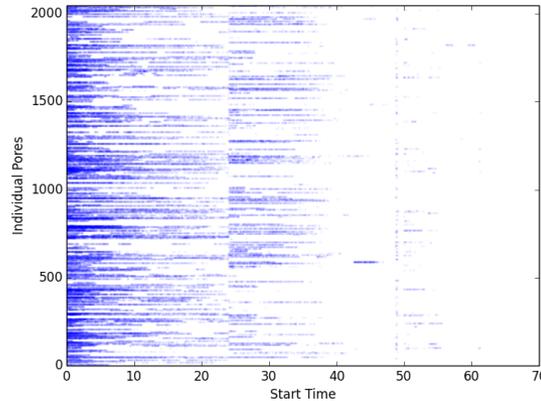
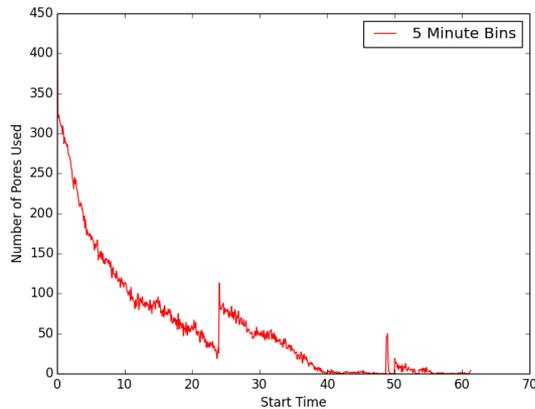
Standard 48hr Sequencing Recipe Script

48hr Genomic DNA Sequencing Script

Mux @ -140mV to assign g1- g4 pores



Example Pore Usage



Modified Scripts to Maximise Flowcell Yields

Yield-Monitoring_Voltage-Tuning Scripts

(If new flowcell use -140mV to start)

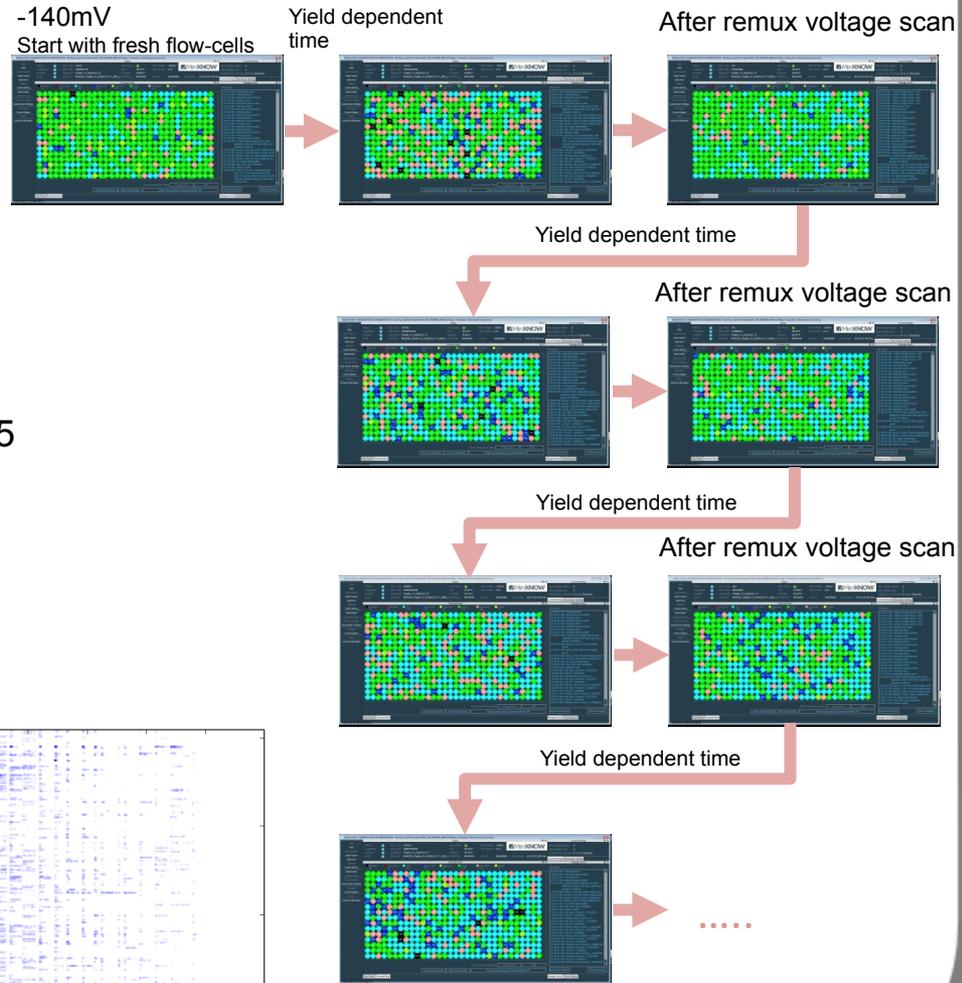
Multi-Voltage Re-mux scan selecting one with greatest "single good pore" count

30 min
 Mux / Re-Mux Scan
 (Voltage tuned across -140mV to -190mV range)

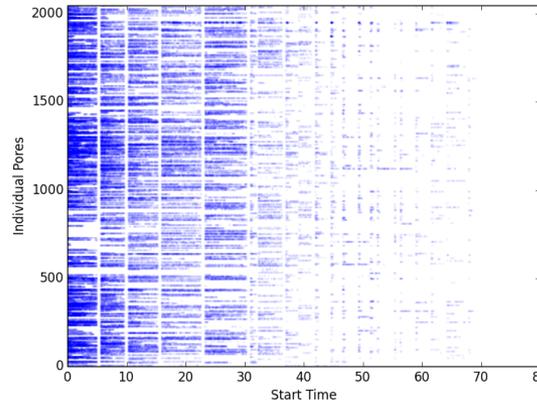
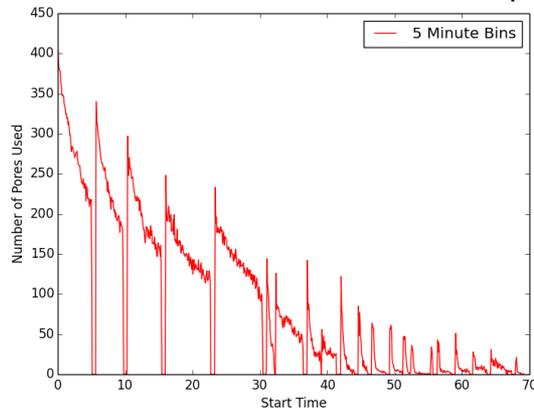
Switch when hourly event yield drops below 60% of first hour of this segment

Yield Dependent Time
 (1 - 24 hours)

x5

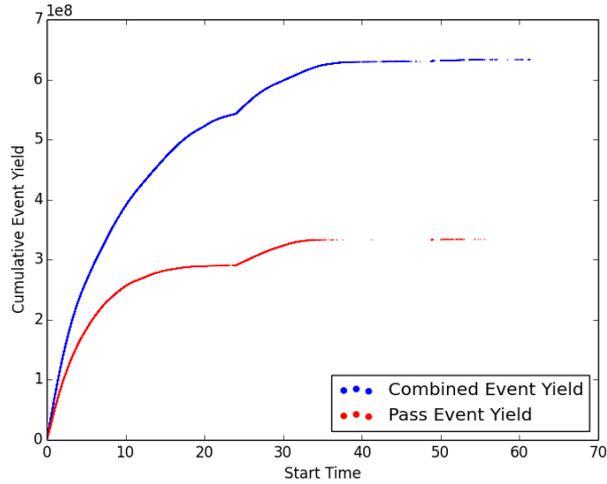


Example Pore Usage

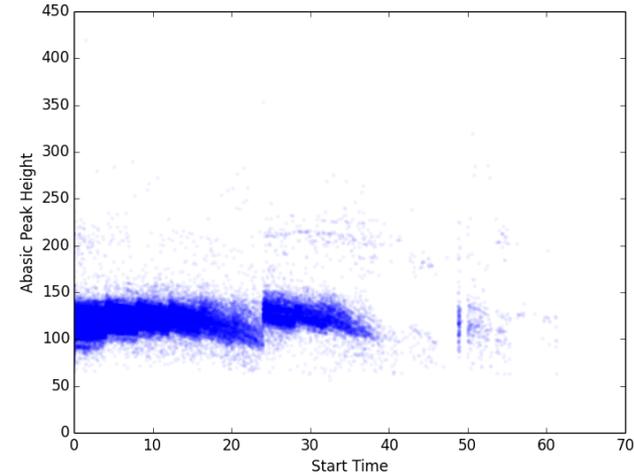


Run Dynamics:- SQK006 Standard Recipe Script

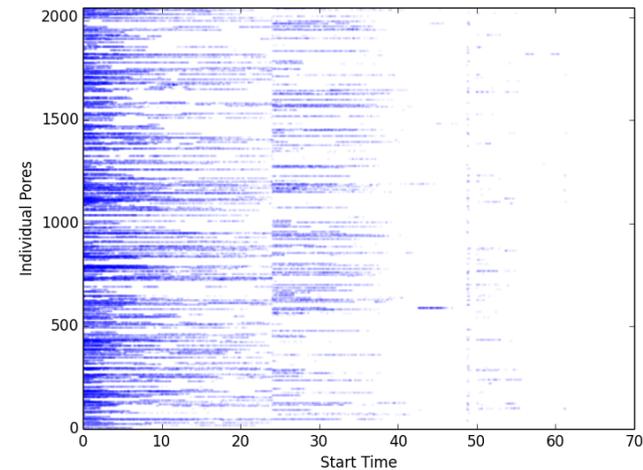
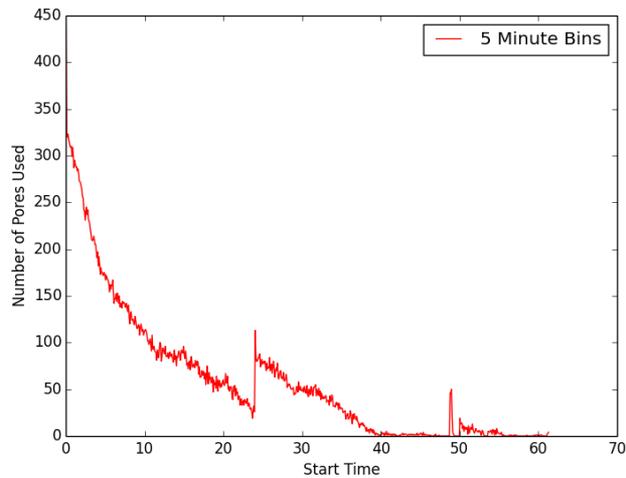
Event Accumulation



Abasic Current

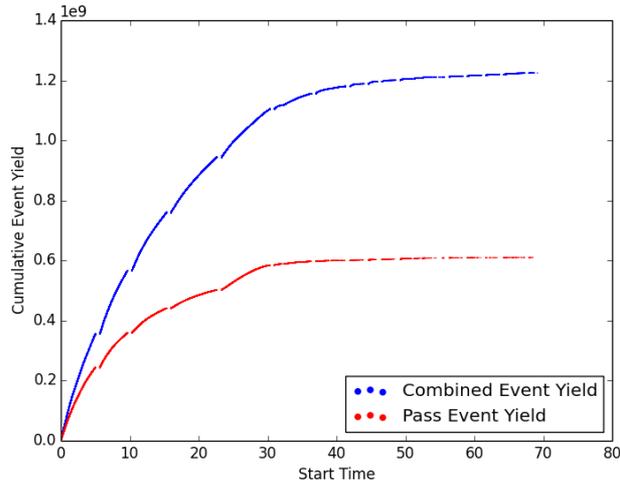


Pore Usage

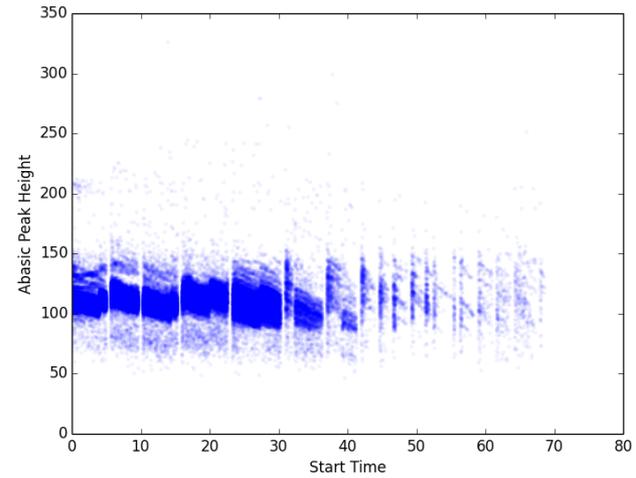


Run Dynamics:- SQK006 Tuning Recipe Script

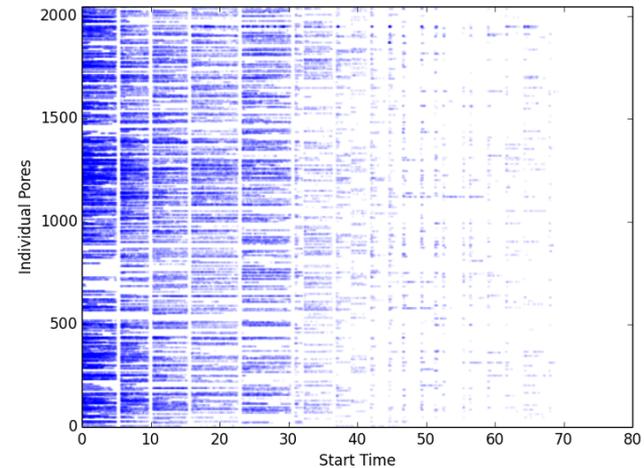
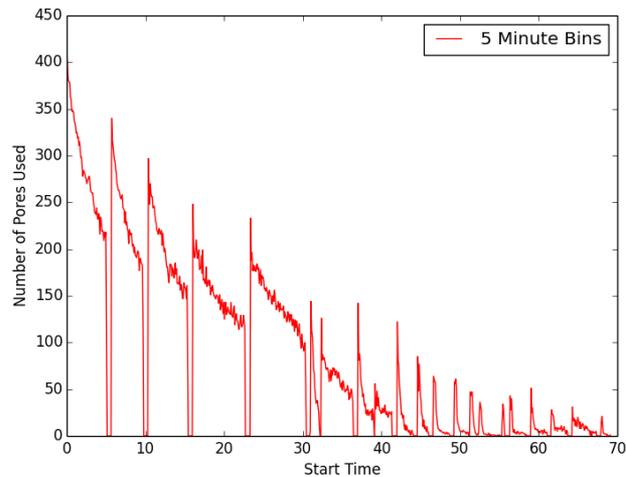
Event Accumulation



Abasic Current

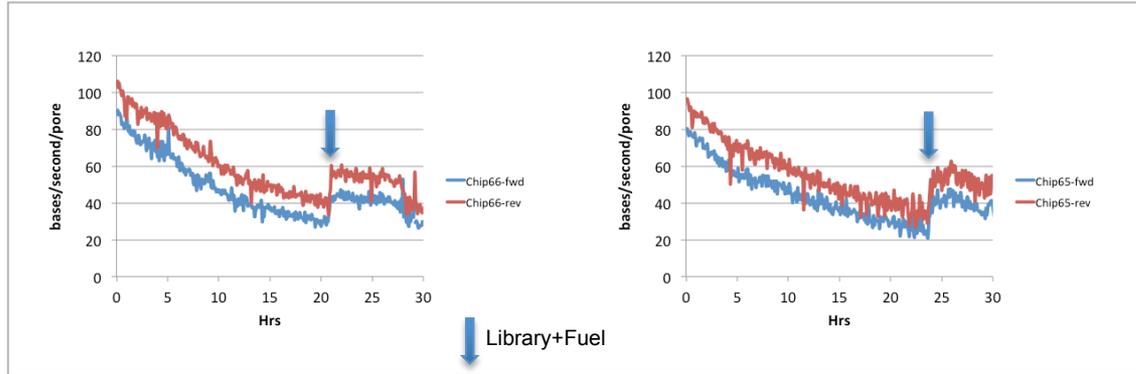


Pore Usage



Pore Speed Comparison

SQK6 Pore Speed



SQK5 Pore Speed

