

The nanopore processes the length of **DNA or RNA** presented to it. The user can control this through the library preparation protocol utilised. (e.g. >2 MB DNAS has been recorded.)

An **enzyme motor** controls the translocation of the DNA or RNA strand through the nanopore. Once the DNA or RNA has passed through, the motor protein detaches and the nanopore is ready to accept the next fragment.

The **nanopore signal**, captured by the ASIC in the device, is characteristic of the sequence of the DNA or RNA fragment.
Algorithms are used to convert the signal into basecalls.

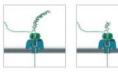
В



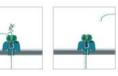
C

Translocation - 1D

The template and the complement strands are sequenced as individual strands



Template... Template...



.. (Exit)



Next molecule...

Translocation - 1D²

The 1D² library preparation deploys special adapters that increase the probability that the complement strand will immediately follow the template strand. This method of sequencing when used with 1D² analysis produces a higher accuracy read.







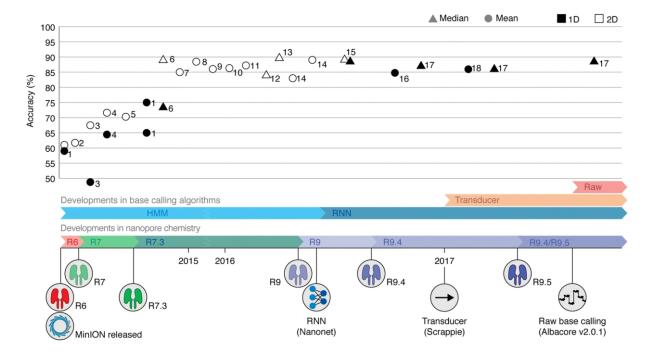


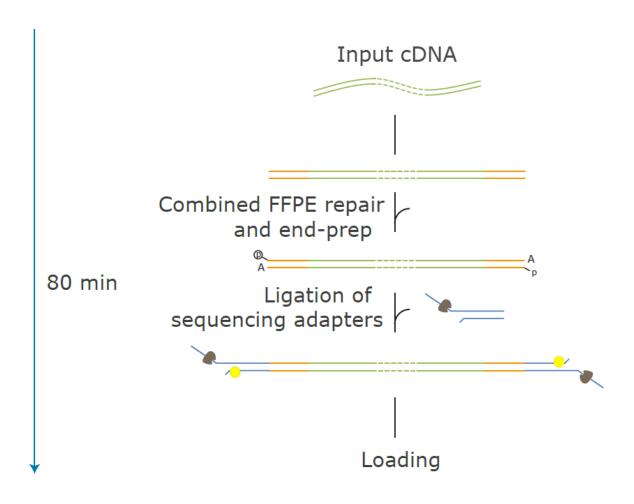
Template...

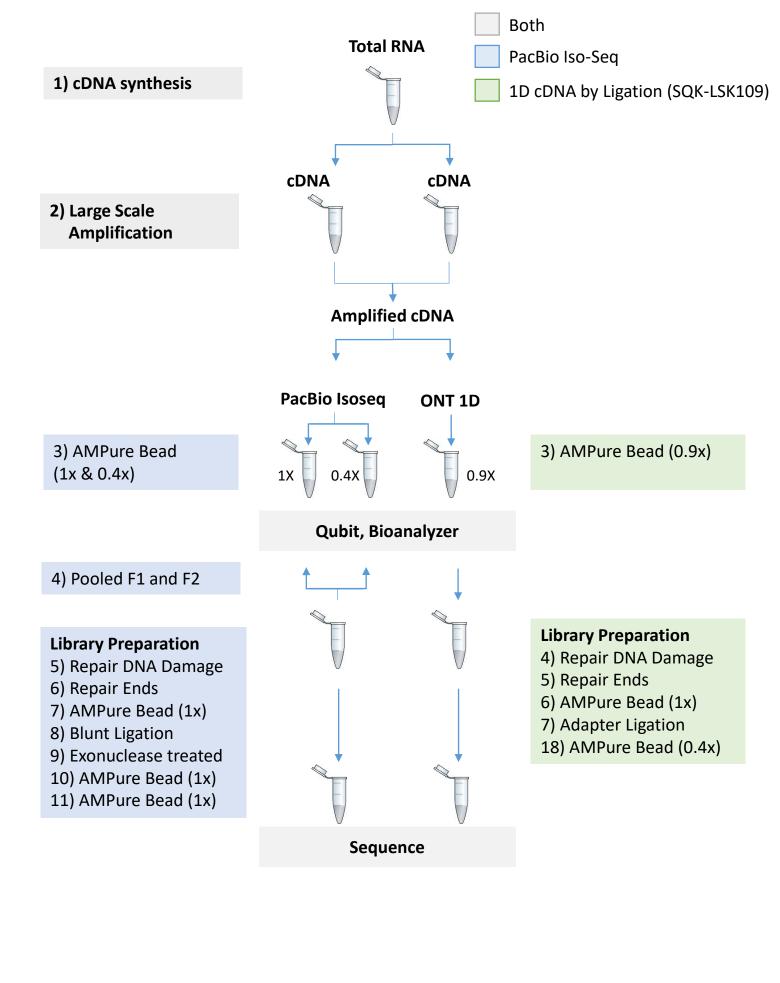
Template...

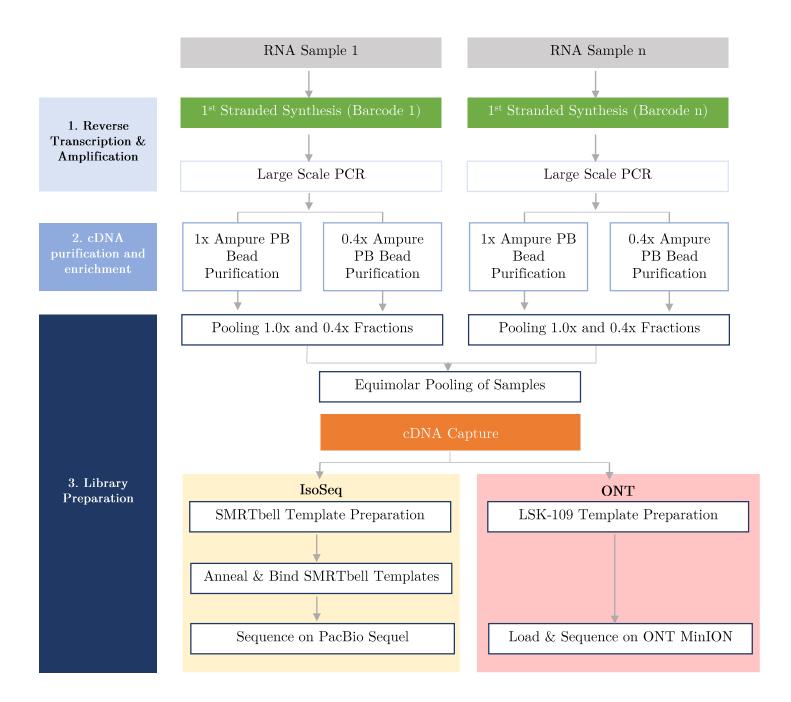
(Exit)

...Complement



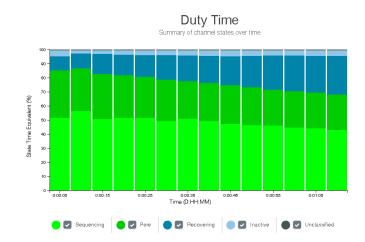




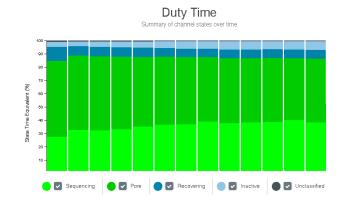


A Good Quality Library

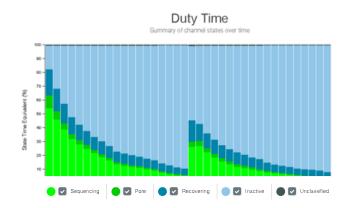
B Channel Blocking



C Low Pore Occupancy



D Flow Cell Failure



- 1) cDNA synthesis with barcode
- 2) Large Scale **Amplification**

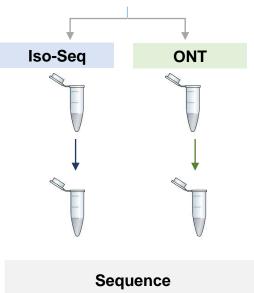
3) Pooling - AMPure Bead (1x & 0.4x)

RNA Sample 1 cDNA cDNA cDNA Amplified cDNA Qubit, Bioanalyzer Equimolar Pooling of Samples Captured DNA Iso-Seq ONT

4) Target Capture

SMRTbell Template **Library Preparation**

- 5) Repair DNA Damage
- 6) Repair Ends
- 7) AMPure Bead (1x)
- 8) Blunt Ligation
- 9) Exonuclease treated
- 10) AMPure Bead (1x)
- 11) AMPure Bead (1x)
- 12) Anneal and Bind SMRTbell templates



SQK LSK-109 Template Library Preparation

RNA Sample n

Amplified cDNA

cDNA

- 5) Repair DNA Damage
- 6) Repair Ends
- 7) AMPure Bead (1x)
- 8) Adapter Ligation
- 9) AMPure Bead (0.4x)