



#### Eab 21 2020

## HMW gDNA purification and ONT ultra-long-read data generation

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#### **ABSTRACT**

This protocol describes the purification of high-molecular-weight genomic DNA from mammalian cells and the generation of ultra-long (N50 >100 kbp) Oxford Nanopore data. It is based on the Sambrook and Russell protocol and Josh Quick's protocol with additional modifications. This protocol improves upon previous protocols developed for ultra-long read sequencing, as it gives longer reads with greater yield. In our hands, we obtain about 1-2 Gb of ultra-long-read ONT data with an N50  $\sim$ 150 kbp on one MinION flow cell.

#### **MATERIALS**

NAME V	CATALOG #	VENDOR V
UltraPure™ Phenol:Chloroform:Isoamyl Alcohol (25:24:1, v/v)	15593031	Thermo Fisher Scientific
Tris-EDTA, pH 8.0	AM9849	Ambion
Proteinase K	19131	Qiagen
RNAse A	19101	Qiagen
200 Proof Ethanol pure	E7023	Sigma Aldrich
Disposable Inoculating Loops and Needles, Flexible Loop; Volume:10µ; Color: Yellow; Individual wrapped	22363600	Thermo Fisher
Ammonium Acetate (5 M), RNase-free	AM9070G	Thermo Fisher
Lysis buffer [10 mM Tris-Cl (pH 8.0) 0.1 M EDTA (pH 8.0) 0.5% w/v SDS]		
Phase Lock Gel Light	2302820	Quantabio
Rapid Sequencing Kit	SQK-RAD004	Oxford Nanopore Technologies
FRA Buffer		Oxford Nanopore Technologies
RAP buffer		Oxford Nanopore Technologies

# Cell collection and lysis

- 1 Freeze down 2-7 x 10^7 cells as a cell pellet, and store at -80C.
- 2 When you are ready to purify the DNA, thaw the cell pellet on ice (usually takes ~30 mins).
- While the cells are thawing, add RNAse A to the lysis buffer at a final concentration of 20 ug/mL. This must be done fresh each time. Keep the lysis buffer + RNase A solution at RT until ready to use.
- 4 Resuspend thawed cells in ice-cold TE (pH 8.0) at a concentration of 5 x 10^7 cells/mL on ice.

- 5 Transfer the cell suspension to a glass Erlenmeyer flask.
  - For 1 mL of cell suspension, use a 50-mL flask; for 2 mL of cell suspension, use a 125-mL flask.
  - Make sure that the cells are well-dispersed over the inner surface of the Erlenmeyer flask. This dispersal minimizes the formation of intractable clumps of cells.
- 6 Quickly add 10 mL of lysis buffer + RNase A for each mL of cell suspension, drop-wise in a circular motion.
- 7 Incubate the cell suspension for 1 hr at 37°C.
- 8 Add proteinase K to a final concentration of 200 ug/mL in a drop-wise manner.
  - For 10 mLs of cell suspension, add 100 uL Proteinase K.
- 9 Swirl the flask to mix the enzyme gently into the viscous cell lysate.
- 10 Incubate the lysate in a water bath for 2 hours at 50°C. Swirl the viscous solution once per hour.
- 11 Cool the solution to RT.

#### Phenol-chloroform extraction

- 12 Add light phase-lock gel to 4 15-mL polypropylene tubes.
  - Phase-lock gels typically come in 2-mL tubes. Transfer the gel by cutting the lid off 3 2-mL phase-lock tubes, placing them upside down into the 15 mL tube, and spinning it at 4000 rpm for 1 min each.
- 13 Split the viscous lysate into the 2 15-mL Falcon tubes prepared with phase-lock gel.
  - This is easiest using a 10 mL serological pipette at low speed.
- 14 Add an equal volume of ultra-pure phenol:chloroform:isoamyl alcohol (~5-6 mLs) to each Falcon tube containing lysate.
- Gently mix the two phases by slowly turning the tube end-over-end for 20 mins on a tube mixer. If the two phases have not formed an emulsion at this stage, place the tube on a roller apparatus for 1 hr.
- 16 Spin in a centrifuge at 4500 rpm for 10 mins.
- 17 Pour the aqueous phases into two new 15-mL Falcon tubes containing phase-lock gel.
- $18 \qquad \text{Add an equal volume of ultra-pure phenol:} chloroform: isoamyl alcohol ($\sim 5-6$ mLs) to each Falcon tube containing lysate.$
- 19 Gently mix the two phases by slowly turning the tube end-over-end for 20 mins on a tube mixer. If the two phases have not formed an emulsion at this stage, place the tube on a roller apparatus for 1 hr.
- 20 Spin in a centrifuge at 4500 rpm for 10 mins.
- 21 Combine the aqueous phases from the two tubes by slowly pouring them into a new 50-mL Falcon tube.

#### Ethanol precipitation

- 22 Add 0.4 volume of 5 M ammonium acetate to the purified DNA, and gently swirl to mix (usually takes several minutes).
  - It is very important to mix gently here. The gentler you mix, the longer your DNA will be.
- 23 Add 2 volumes of ethanol at RT and gently swirl to mix (usually takes  $\sim$ 30 mins).
  - Swirl gently, sometimes rocking back and forth slowly to get the DNA in solution with the salt and ethanol.
- 24 Store the solution O/N at 4°C.
  - We have found that storing the DNA overnight results in purer DNA that sequences better and gives more yield.

- The DNA forms a precipitate. Remove the precipitate in one piece from the ethanolic solution with a disposable inoculating needle shaped into a U.
  - If the DNA precipitate becomes fragmented, abandon the needle and collect the precipitate by centrifugation 4500 rpm for 5 mins at RT.
- 26 Wash the DNA precipitate 2x with 70% ethanol, and collect the DNA by centrifuging at max speed (~15000 rpm) for 15 secs.
- 27 Remove as much of the 70% ethanol as possible. Store the pellet of DNA in an open tube at RT until the last visible traces of ethanol have evaporated (~30 mins to 1 hr).
  - Do not allow the pellet of DNA to dry completely. Desiccated DNA is very difficult to dissolve.
- Add 125-250 uL EB + 0.02% Triton-X100 and incubate without mixing at 4°C for 2 days to allow the pellet to fully resuspend into a translucent viscous gel.

## ONT library preparation

- 29 Equilibrate an ONT MinION/GridION flow cell to room temperature.
- 30 Pre-set a thermocycler to the following program:
  - a. 30°C for 1.5 min
  - b. 80°C for 1 min
  - c. 4°C forever
- 31 Pre-warm the thermocycler by starting the program and then immediately pressing "Pause".
- 32 Add the following reagents to a PCR tube on ice (in order).

Note: Use a wide-bore pipette tip for the gDNA to avoid shearing.  $\label{eq:control} % \begin{center} \begin{$ 

10 uL HMW gDNA (2-3 ug) 5 uL dH $_2$ O 3 uL FRA Buffer

Total = 18 uL

- Pipette the gDNA mixture up and down slowly >20 times on ice with a wide-bore pipette tip. Do not worry about shearing the DNA. Try to avoid creating bubbles.
- In a lo-bind tube on ice, dilute FRA 1:12 in FRA buffer on ice
  - FRA buffer composition is proprietary, but ONT will send you free FRA buffer if you ask them.
  - This dilution can be modified depending on the needs of the sequencing experiment. In our hands, increased FRA concentration gives shorter reads but greater throughput.
- 35~ Add 1.5  $\mu l$  FRA (diluted) to the gDNA mixture on ice.
- Pipette the gDNA mixture up and down slowly >20 times with a wide-bore pipette tip while keeping the tube on ice. Try to avoid creating bubbles.
- 37 Place the tube in the thermocycler and let the program continue.
- 38 After the thermocycler is finished, move the tube to ice, and bury it in the ice to cool it down from all sides.
- 39 In a lo-bind tube on ice, dilute RAP 1:12 in RAP buffer.
  - RAP buffer composition is proprietary, but ONT will send you free RAP buffer if you ask them.
- 40 Add 0.5 μl RAP (diluted) to the gDNA mixture on ice.

- 41 Pipette the gDNA mixture up and down slowly >20 times with a wide-bore pipette tip while keeping the tube on ice. Try to avoid creating bubbles.
- 42 Incubate the gDNA mixture at room temperature for 2 hours.
  - This incubation time can be reduced to 30 min, but we obtain greater yield with an extended incubation.

# Flow cell priming and loading

- Place the flow cell on a MinION/GridION and run platform QC.
- Remove the flow cell from the GridION and place in its holder to let it cool down for at least 10 mins.
- 45 Add 30 µl FLT to the tube of FB. Invert to mix and spin down.
- 46 Using a P1000 set to ~800 μl, place the pipette tip in the inlet port of the flow cell and turn the wheel to the right to remove the air and a small amount of storage buffer. This will make the storage buffer flush with the inlet port opening.
- 47 Load 800 μl FLB + FLT via the inlet port slowly. Wait 5 minutes.
- 48 Meanwhile, thoroughly mix SQB by inverting.
- 49 Add 20 µl dH<sub>2</sub>O and 34 µl SQB to the gDNA mixture.
- 50 Pipette the gDNA mixture up and down slowly >20 times. Try to avoid creating bubbles.
- 51 Lift the cover off the SpotON port. Load 200 µl FLB + FLT via the inlet port slowly using a P1000 pipette. Try to dispense at a speed where a bead of liquid becomes visible over the SpotON port which then gets siphoned back in.
- 52 Using a P200 set to 75  $\mu$ I with a wide-bore tip, pipette the diluted library onto the SpotON port as it gets siphoned in.
- 53 Close the SpotON sample port cover, and close the priming port.
- Let incubate for 10 min before placing on the MinION/GridION and starting the run.

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