Flow Cell Number: ..... DNA Samples: .....

Before start checklist

Materials

- [ ] 500 ng polyA-tailed RNA in 9 µl
- [ ] Direct RNA Sequencing Kit (SQK-RNA002)
- [ ] Flow Cell Priming Kit (EXP-FLP002)

Consumables

- [ ] 1.5 ml Eppendorf DNA LoBind tubes
- [ ] 0.2 ml thin-walled PCR tubes
- [ ] Nuclease-free water (e.g. ThermoFisher, cat # AM9937)
- [ ] Freshly prepared 70% ethanol in nuclease-free water
- [ ] SuperScript III Reverse Transcriptase (Thermo Fisher Scientific, 18080044)
- [ ] 10 mM dNTP solution (e.g. NEB N0447)
- [ ] NEBNext Quick Ligation Module (E6056)
- [ ] Agencourt RNAClean XP beads
- [ ] Qubit RNA HS Assay Kit (ThermoFisher Q32852)
- [ ] Qubit dsDNA HS Assay Kit (ThermoFisher Q32851)

Equipment

- [ ] Hula mixer (gentle rotator mixer)
- [ ] Magnetic separator, suitable for 1.5 ml Eppendorf tubes
- [ ] Microfuge
- [ ] Vortex mixer
- [ ] Ice bucket with ice
- [ ] Timer
- [ ] Thermal cycler
- [ ] Qubit fluorometer (or equivalent for QC check)
- [ ] Pipettes and pipette tips P2, P10, P20, P100, P200, P1000

INSTRUCTIONS	NOTES/OBSERVATIONS
Library preparation	

Prepare the RNA in Nuclease-free water.

- [ ] Transfer 500 ng RNA into a 1.5 ml Eppendorf DNA LoBind tube
- [ ] Adjust the volume to 9 µl with Nuclease-free water
- [ ] Mix thoroughly by flicking the tube to avoid unwanted shearing
- [ ] Spin down briefly in a microfuge

In a 0.2 ml thin-walled PCR tube, mix the reagents in the following order:

- [ ] 3.0 µl NEBNext Quick Ligation Reaction Buffer
- [ ] 9.0 µl RNA
- [ ] 0.5 µl RNA CS (RCS), 110 nM
- [ ] 1.0 µl RT Adapter (RTA)
- [ ] 1.5 µl T4 DNA Ligase

- [ ] Mix by pipetting and spin down.
- [ ] Incubate the reaction for 10 minutes at RT.

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Mix the following reagents together to make the reverse transcription master mix:

- ☐ 9.0 µl Nuclease-free water
- ☐ 2.0 µl 10 mM dNTPs
- ☐ 8.0 µl 5x first-strand buffer
- ☐ 4.0 µl 0.1 M DTT

☐ Add the master mix to the 0.2 ml PCR tube containing the RT adapter-ligated RNA from the "RT Adapter ligation" step. Mix by pipetting.

☐ Add 2 µl of SuperScript III reverse transcriptase to the reaction and mix by pipetting.

☐ Place the tube in a thermal cycler and incubate at 50° C for 50 min, then 70° C for 10 min, and bring the sample to 4° C before proceeding to the next step.

☐ Transfer the sample to a clean 1.5 ml Eppendorf DNA LoBind tube.

☐ Resuspend the stock of Agencourt RNAClean XP beads by vortexing.

☐ Add 72 µl of resuspended RNAClean XP beads to the reverse transcription reaction and mix by pipetting.

☐ Incubate on a Hula mixer (rotator mixer) for 5 minutes at RT.

☐ Prepare 200 µl of fresh 70% ethanol in Nuclease-free water.

☐ Spin down the sample and pellet on a magnet. Keep the tube on the magnet, and pipette off the supernatant.

Keep the tube on magnet, and wash the beads with 150 µl of freshly prepared 70% ethanol without disturbing the pellet as described below.

- ☐ Keeping the magnetic rack on the benchtop, rotate the bead-containing tube by 180°. Wait for the beads to migrate towards the magnet and form a pellet.
- ☐ Rotate the tube 180° again (back to the starting position), and wait for the beads to pellet.

☐ Remove the 70% ethanol using a pipette, and discard.

☐ Spin down and place the tube back on the magnet. Pipette off any residual 70% ethanol.

☐ Remove the tube from the magnetic rack and resuspend pellet in 20 µl Nuclease-free water. Incubate for 5 minutes at RT.

☐ Pellet the beads on a magnet until the eluate is clear and colourless.

☐ Pipette 20 µl of eluate into a clean 1.5 ml Eppendorf DNA LoBind tube.

In a clean 1.5 ml Eppendorf DNA LoBind tube, mix the reagents in the following order:

- ☐ 20.0 µl Reverse-transcribed RNA from the "Reverse Transcription" step
- ☐ 8.0 µl NEBNext Quick Ligation Reaction Buffer
- ☐ 6.0 µl RNA Adapter (RMX)
- ☐ 3.0 µl Nuclease-free water
- ☐ 3.0 µl T4 DNA Ligase

☐ Mix by pipetting.

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☐ Incubate the reaction for 10 minutes at RT. ☐ Resuspend the stock of Agencourt RNAClean XP beads by vortexing. ☐ Add 40 µl of resuspended RNAClean XP beads to the adapter ligation reaction and mix by pipetting. ☐ Incubate on a Hula mixer (rotator mixer) for 5 minutes at RT. ☐ Spin down the sample and pellet on a magnet. Keep the tube on the magnet, and pipette off the supernatant. ☐ Add 150 µl of the Wash Buffer (WSB) to the beads. Close the tube lid, and resuspend the beads by flicking the tube. Return the tube to the magnetic rack, allow beads to pellet and pipette off the supernatant. ☐ Repeat the previous step. ☐ Remove the tube from the magnetic rack and resuspend pellet in 21 µl Elution Buffer by the gently flicking the tube. Incubate for 10 minutes at RT. ☐ Pellet the beads on a magnet until the eluate is clear and colourless. ☐ Remove and retain 21 µl of eluate into a clean 1.5 ml Eppendorf DNA LoBind tube.	
Quantify 1 µl of reverse-transcribed and adapted RNA using the Qubit fluorometer DNA HS assay - recovery aim ~200 ng.	
The reverse-transcribed and adapted RNA is now ready for loading into the flow cell.	
Priming and loading the SpotON flow cell	
☐ Thaw the RNA Running Buffer (RRB), Flush Tether (FLT) and one tube of Flush Buffer (FB) at RT before placing the tubes on ice as soon as thawing is complete. ☐ Mix the RNA Running Buffer (RRB), Flush Buffer (FB) and Flush Tether (FLT) tubes thoroughly by vortexing, spin down and return to ice. ☐ Open the MinION Mk 1B lid and slide the flow cell under the clip. ☐ Slide the priming port cover clockwise to open the priming port. ☐ How to prime and load the SpotON Flow Cell	
IMPORTANT ☐ Take care when drawing back buffer from the flow cell. Do not remove more than 20-30 µls, and make sure that the array of pores is covered by buffer at all times. Introducing air bubbles into the array can irreversibly damage pores.	
After opening the priming port, check for a small air bubble under the cover. Draw back a small volume to remove any bubbles (a few µls): ☐ Set a P1000 pipette to 200 µl ☐ Insert the tip into the priming port ☐ Turn the wheel until the dial shows 220-230 µl, or until you can see a small volume of buffer entering the pipette tip	

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☐ Prepare the flow cell priming mix: add 30 µl of thawed and mixed Flush Tether (FLT) directly to the tube of thawed and mixed Flush Buffer (FB), and mix by vortexing. ☐ Load 800 µl of the priming mix into the flow cell via the priming port, avoiding the introduction of air bubbles. Wait for 5 minutes. During this time, prepare the library for loading by following the steps below.	
IMPORTANT ☐ Thoroughly mix the contents of the RRB tube by vortexing or pipetting, and spin down briefly.	
☐ Take 20 µl of the prepared RNA library and mix it with 17.5 µl of Nuclease-free water. In a new tube, prepare the library for loading as follows: ☐ 37.5 µl RRB ☐ 37.5 µl RNA library in Nuclease-free water Complete the flow cell priming: ☐ Gently lift the SpotON sample port cover to make the SpotON sample port accessible. ☐ Load 200 µl of the priming mix into the flow cell via the priming port (not the SpotON sample port), avoiding the introduction of air bubbles. ☐ Mix the prepared library gently by pipetting up and down just prior to loading. ☐ Add 75 µl of sample to the Flow Cell via the SpotON sample port in a dropwise fashion. Ensure each drop flows into the port before adding the next. ☐ Gently replace the SpotON sample port cover, making sure the bung enters the SpotON port, close the priming port and replace the MinION Mk 1B lid.	
Ending the experiment	
☐ After your sequencing experiment is complete, if you would like to reuse the flow cell, please follow the Wash Kit instructions and store the washed flow cell at 2-8 °C, OR ☐ Follow the returns procedure by washing out the flow cell ready to send back to Oxford Nanopore.	
IMPORTANT ☐ If you encounter issues or have questions about your sequencing experiment, please refer to the Troubleshooting Guide that can be found in the online version of this protocol.	