

Rapid Barcoding Sequencing (SQK-RBK004)

Version: RBK_9054_v2_revV_14Aug2019
 Last update: 07/10/2021



Flow Cell Number:

DNA Samples:

Before start checklist		
Materials	Consumables	Equipment
<input type="checkbox"/> ~400 ng high molecular weight genomic DNA	<input type="checkbox"/> 1.5 ml Eppendorf DNA LoBind tubes	<input type="checkbox"/> Ice bucket with ice
<input type="checkbox"/> Rapid Barcoding Sequencing Kit (SQK-RBK004)	<input type="checkbox"/> 0.2 ml thin-walled PCR tubes	<input type="checkbox"/> Microfuge
<input type="checkbox"/> Flow Cell Priming Kit (EXP-FLP002)	<input type="checkbox"/> Nuclease-free water (e.g. ThermoFisher, cat # AM9937)	<input type="checkbox"/> Timer
	<input type="checkbox"/> Agencourt AMPure XP beads (optional)	<input type="checkbox"/> Thermal cycler or heat block at 30°C and 80°C
	<input type="checkbox"/> Freshly-prepared 70% ethanol in nuclease-free water (optional)	<input type="checkbox"/> Pipettes and pipette tips P2, P20, P100, P200, P1000
	<input type="checkbox"/> 10 mM Tris-HCl pH 8.0 with 50 mM NaCl (optional)	
INSTRUCTIONS		NOTES/OBSERVATIONS
Library preparation		
<p>Thaw kit components at RT, spin down briefly using a microfuge and mix by pipetting as indicated by the table below:</p> <ul style="list-style-type: none"> <input type="checkbox"/> Fragmentation Mix RB01-12: not frozen, briefly spin down, mix well by pipetting <input type="checkbox"/> Rapid Adapter (RAP): not frozen, briefly spin down, mix well by pipetting <input type="checkbox"/> Sequencing Buffer (SQB): thaw at RT, briefly spin down, mix well by pipetting* <input type="checkbox"/> Loading Beads (LB): thaw at RT, briefly spin down, mix by pipetting or vortexing immediately before use <input type="checkbox"/> Flush Buffer (FLB) - 1 tube: thaw at RT, briefly spin down, mix well by pipetting* <input type="checkbox"/> Flush Tether (FLT): thaw at RT, briefly spin down, mix well by pipetting <p>Prepare the DNA in Nuclease-free water.</p> <ul style="list-style-type: none"> <input type="checkbox"/> Transfer ~400 ng genomic DNA into a DNA LoBind tube <input type="checkbox"/> Adjust the volume to 7.5 µl with Nuclease-free water <input type="checkbox"/> Mix by flicking the tube to avoid unwanted shearing <input type="checkbox"/> Spin down briefly in a microfuge <p>In a 0.2 ml thin-walled PCR tube, mix the following:</p> <ul style="list-style-type: none"> <input type="checkbox"/> 7.5 µl 400 ng template DNA <input type="checkbox"/> 2.5 µl Fragmentation Mix RB01-12 (one for each sample) <ul style="list-style-type: none"> <input type="checkbox"/> Mix gently by flicking the tube, and spin down. <input type="checkbox"/> Incubate the tube at 30°C for 1 minute and then at 80°C for 1 minute. Briefly put the tube on ice to cool it down. <input type="checkbox"/> Pool all barcoded samples in your desired ratio, noting the total volume. 		

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<p>IMPORTANT</p> <p><input type="checkbox"/> If barcoding four or more samples, increased throughput can be achieved through cleaning up and concentrating the pooled material using AMPure XP beads as outlined in Steps 6-15. Otherwise, for a more rapid sample preparation, transfer 10 µl of pooled sample from Step 5 into a clean 1.5 ml Eppendorf DNA LoBind tube, and proceed directly to Step 16.</p>	
<p><input type="checkbox"/> Resuspend the AMPure XP beads by vortexing.</p> <p><input type="checkbox"/> To the entire pooled barcoded sample from Step 6, add an equal volume of resuspended AMPure XP beads, and mix by flicking the tube.</p> <p><input type="checkbox"/> Incubate on a Hula mixer (rotator mixer) for 5 minutes at RT.</p> <p><input type="checkbox"/> Prepare 500 µl of fresh 70% ethanol in Nuclease-free water.</p> <p><input type="checkbox"/> Spin down the sample and pellet on a magnet. Keep the tube on the magnet, and pipette off the supernatant.</p> <p><input type="checkbox"/> Keep the tube on the magnet and wash the beads with 200 µl of freshly prepared 70% ethanol without disturbing the pellet. Remove the ethanol using a pipette and discard.</p> <p><input type="checkbox"/> Repeat the previous step.</p> <p><input type="checkbox"/> Spin down and place the tube back on the magnet. Pipette off any residual 70% ethanol. Briefly allow to dry.</p> <p><input type="checkbox"/> Remove the tube from the magnetic rack and resuspend pellet in 10 µl of 10 mM Tris-HCl pH 7.5-8.0 with 50 mM NaCl. Incubate for 2 minutes at RT.</p> <p><input type="checkbox"/> Pellet the beads on a magnet until the eluate is clear and colourless, for at least 1 minute.</p> <p>Remove and retain 10 µl of eluate into a clean 1.5 ml Eppendorf DNA LoBind tube.</p> <p><input type="checkbox"/> Remove and retain the eluate which contains the DNA in a clean 1.5 ml Eppendorf DNA LoBind tube</p> <p><input type="checkbox"/> Dispose of the pelleted beads</p> <p>End of optional steps.</p> <p><input type="checkbox"/> Add 1 µl of RAP to 10 µl of barcoded DNA.</p> <p><input type="checkbox"/> Mix gently by flicking the tube, and spin down.</p> <p><input type="checkbox"/> Incubate the reaction for 5 minutes at RT.</p>	
<p>The prepared library is used for loading into the MinION flow cell. Store the library on ice until ready to load.</p>	
<p>Priming and loading the SpotON Flow Cell</p>	
<p>IMPORTANT</p> <p><input type="checkbox"/> Please note that the Sequencing Tether (SQT) tube will NOT be used in this protocol. It is provided in the kit for potential future product compatibility.</p>	
<p><input type="checkbox"/> Thaw the Sequencing Buffer (SQB), Loading Beads (LB), Flush Tether (FLT) and one tube of Flush Buffer (FB) at RT.</p>	

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<ul style="list-style-type: none"> <input type="checkbox"/> Mix the Sequencing Buffer (SQB), Flush Tether (FLT) and Flush Buffer (FB) tubes by vortexing and spin down at RT. <input type="checkbox"/> Open the MinION Mk1B lid and slide the flow cell under the clip. <input type="checkbox"/> Slide the priming port cover clockwise to open the priming port. 	
<p>IMPORTANT</p> <ul style="list-style-type: none"> <input type="checkbox"/> Take care when drawing back buffer from the flow cell. Do not remove more than 20-30 μl, and make sure that the array of pores are covered by buffer at all times. Introducing air bubbles into the array can irreversibly damage pores. 	
<p>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 μl):</p> <ul style="list-style-type: none"> <input type="checkbox"/> Set a P1000 pipette to 200 μl <input type="checkbox"/> Insert the tip into the priming port <input type="checkbox"/> Turn the wheel until the dial shows 220-230 μl, or until you can see a small volume of buffer entering the pipette tip <ul style="list-style-type: none"> <input type="checkbox"/> To 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 at RT. <input type="checkbox"/> 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. <input type="checkbox"/> Thoroughly mix the contents of the Loading Beads (LB) tubes by vortexing. <p>In a new tube, prepare the library for loading as follows:</p> <ul style="list-style-type: none"> <input type="checkbox"/> 34 μl Sequencing Buffer (SQB) <input type="checkbox"/> 25.5 μl Loading Beads (LB), mixed immediately before use <input type="checkbox"/> 4.5 μl Nuclease-free water <input type="checkbox"/> 11 μl DNA library 	
<p>IMPORTANT</p> <ul style="list-style-type: none"> <input type="checkbox"/> The Loading Beads (LB) tube contains a suspension of beads. These beads settle very quickly. It is vital that they are mixed immediately before use. 	
<p>Complete the flow cell priming:</p> <ul style="list-style-type: none"> <input type="checkbox"/> Gently lift the SpotON sample port cover to make the SpotON sample port accessible. <input type="checkbox"/> 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. <input type="checkbox"/> Mix the prepared library gently by pipetting up and down just prior to loading. <input type="checkbox"/> 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. <input type="checkbox"/> Gently replace the SpotON sample port cover, making sure the bung enters the SpotON port, close the priming port and replace the MinION Mk1B lid. 	

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INSTRUCTIONS	NOTES/OBSERVATIONS
Ending the experiment	
<input type="checkbox"/> 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 <input type="checkbox"/> Follow the returns procedure by washing out the flow cell ready to send back to Oxford Nanopore.	
IMPORTANT <input type="checkbox"/> 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.	