

# Loading multiple PromethION Flow Cells

Version: PFC\_9097\_v1\_revM\_18Mar2020  
 Last update: 07/02/2024



Flow Cell Number: .....

DNA Samples: .....

Before start checklist		
Materials	Consumables	Equipment
<input type="checkbox"/> DNA libraries	<input type="checkbox"/> PromethION Flow Cell (FLO-PRO114M)	<input type="checkbox"/> PromethION 24/48 device
<input type="checkbox"/> Flow Cell Priming Kit V14 (EXP-FLP004)	<input type="checkbox"/> 1.5 ml Eppendorf DNA LoBind tubes	<input type="checkbox"/> Ice bucket with ice
<input type="checkbox"/> Sequencing Auxiliary Vials V14 (EXP-AUX003)	<input type="checkbox"/> 2 ml Eppendorf DNA LoBind tubes	<input type="checkbox"/> Pipettes and pipette tips P20, P200, P1000
<input type="checkbox"/> Flow Cell Wash Kit XL (EXP-WSH004-XL)		

  

INSTRUCTIONS	NOTES/OBSERVATIONS
<p><b>Priming and loading multiple flow cells on a PromethION</b></p> <p>Loading multiple PromethION Flow Cells</p> <p><b>IMPORTANT</b></p> <p><input type="checkbox"/> Scale up reagent volumes as needed</p> <p><b>IMPORTANT</b></p> <p><input type="checkbox"/> After taking flow cells out of the fridge, wait 20 minutes before inserting the flow cell into the PromethION for the flow cell to come to RT. Condensation can form on the flow cell in humid environments. Inspect the gold connector pins on the top and underside of the flow cell for condensation and wipe off with a lint-free wipe if any is observed. Ensure the heat pad (black pad) is present on the underside of the flow cell.</p> <p>Insert all flow cells into the docking ports within the PromethION:</p> <ul style="list-style-type: none"> <li><input type="checkbox"/> Line up the first flow cell in position 1A with the connector horizontally and vertically before smoothly inserting into position.</li> <li><input type="checkbox"/> Press down firmly onto the flow cell and ensure the latch engages and clicks into place.</li> <li><input type="checkbox"/> Repeat for the next flow cell until all flow cells are loaded.</li> </ul> <p><b>IMPORTANT</b></p> <p><input type="checkbox"/> Insertion of the flow cells at the wrong angle can cause damage to the pins on the PromethION and affect your sequencing results. If you find the pins on a PromethION position are damaged, please contact support@nanoporetech.com for assistance.</p> <p><input type="checkbox"/> If not already completed, perform a flow cell check on all flow cells.</p> <p><input type="checkbox"/> Thaw the Sequencing Buffer (SB), Library Beads (LIB) or Library Solution (LIS, if using), Flow Cell Tether (FCT) and Flow Cell Flush (FCF) at RT, before mixing by vortexing. Then spin down before storing on ice.</p> <p>Prepare the flow cell priming mix in a suitable vial for the number of flow cells to flush. Once combined, mix well by vortexing and keep at RT.</p> <ul style="list-style-type: none"> <li><input type="checkbox"/> Flow Cell Tether (FCT) 780 µl @ 30 µl (1,500 µl cycle)</li> <li><input type="checkbox"/> Flow Cell Flush (FCF) 30,420 µl @ 1,170 µl (58,500 µl cycles)</li> </ul> <p><input type="checkbox"/> Slide the inlet port cover clockwise to open the inlet port of the first flow cell inserted. Repeat for all flow cells.</p>	

# Loading multiple PromethION Flow Cells

Version: PFC\_9097\_v1\_revM\_18Mar2020  
 Last update: 07/02/2024



Flow Cell Number: .....

DNA Samples: .....

INSTRUCTIONS	NOTES/OBSERVATIONS
<p><b>IMPORTANT</b></p> <p><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>	
<p>After opening the inlet port, draw back a small volume to remove any air bubbles for all flow cells:</p> <ul style="list-style-type: none"> <li><input type="checkbox"/> Set a P1000 pipette tip to 200 µl.</li> <li><input type="checkbox"/> Insert the tip into the inlet port of the first flow cell.</li> <li><input type="checkbox"/> Turn the wheel until the dial shows 220-230 µl, or until you see a small volume of buffer entering the pipette tip.</li> <li><input type="checkbox"/> Repeat for all the flow cells using the same pipette tip.</li> </ul> <p>Complete the first flush as follows for all flow cells:</p> <ul style="list-style-type: none"> <li><input type="checkbox"/> Load 500 µl of the priming mix into the first flow cell via the inlet port, avoiding the introduction of air bubbles.</li> <li><input type="checkbox"/> Start a timer for 5 minutes. This is to ensure 5 minutes passes for the first flow cell before the second flush is started.</li> <li><input type="checkbox"/> Immediately repeat step 1 for the remaining flow cells using the same pipette tip.</li> </ul> <p>Complete a second flush once all the flow cells have been flushed once, as follows:</p> <ul style="list-style-type: none"> <li><input type="checkbox"/> Check that the timer has finished to ensure it has been at least 5 minutes since the first flow cell was flushed.</li> <li><input type="checkbox"/> Load 500 µl of the priming mix into the inlet port of the first flow cell, avoiding the introduction of air bubbles.</li> <li><input type="checkbox"/> Repeat the previous step for the remaining flow cells in the same order as the first flush, using the same pipette tip.</li> </ul> <p><input type="checkbox"/> Freshly prepare the DNA libraries for loading as outlined in the relevant protocol.</p> <p>Load your samples:</p> <ul style="list-style-type: none"> <li><input type="checkbox"/> Gently mix the first library and load the full volume through the inlet port of the first flow cell.</li> <li><input type="checkbox"/> Using a new tip between each flow cell, repeat for the remaining flow cells, ensuring to gently mix each library before loading.</li> </ul> <p><input type="checkbox"/> Close the valve to seal the inlet port.</p>	
<p><b>IMPORTANT</b></p> <p><input type="checkbox"/> Install the light shield on your flow cell as soon as library has been loaded for optimal sequencing output.</p>	
<p>If the light shield has been removed from the flow cell, install the light shield as follows:</p> <ul style="list-style-type: none"> <li><input type="checkbox"/> Align the inlet port cut out of the light shield with the inlet port cover on the flow cell. The leading edge of the light shield should sit above the flow cell ID.</li> <li><input type="checkbox"/> Firmly press the light shield around the inlet port cover. The inlet port clip will click into place underneath the inlet port cover.</li> </ul>	
<p>Close the PromethION lid when ready to start a sequencing run on MinKNOW.</p>	
<p><input type="checkbox"/> For multiple flow cell washing, use the same experiment name and identifying sample IDs for all runs to enable all flow cells to be paused simultaneously.</p>	

# Loading multiple PromethION Flow Cells

Version: PFC\_9097\_v1\_revM\_18Mar2020  
 Last update: 07/02/2024



Flow Cell Number: .....

DNA Samples: .....

INSTRUCTIONS	NOTES/OBSERVATIONS
<p><b>Washing multiple flow cells on a PromethION</b></p>	
<p><b>IMPORTANT</b></p> <p><input type="checkbox"/> Scale up reagent volumes as needed</p>	
<p><b>IMPORTANT</b></p> <p><input type="checkbox"/> A P1000 pipette must be used for all flushing steps to create a seal with the flow cell ports.</p> <p><input type="checkbox"/> Place the tube(s) of Wash Mix (WMX) on ice. Do not vortex the tube.</p> <p><input type="checkbox"/> Thaw the tube(s) of Wash Diluent (DIL) at RT.</p> <p><input type="checkbox"/> Mix the contents of Wash Diluent (DIL) thoroughly by vortexing, then spin down briefly and place on ice.</p> <p>Prepare the Wash Mix in a suitable vial for the number of flow cells to wash. Mix well by pipetting, and place on ice. Do not vortex the tube.</p> <p><input type="checkbox"/> Wash Mix (WMX) 52 µl @ 2 µl (100 µl cycles)</p> <p><input type="checkbox"/> Wash Diluent (DIL) 10,348 µl @ 398 µl (19,900 µl cycles)</p> <p><input type="checkbox"/> Stop or pause the sequencing run in MinKNOW for all flow cells to be washed. Leave the flow cells in the device.</p>	
<p><b>IMPORTANT</b></p> <p><input type="checkbox"/> It is vital that the inlet port is closed before removing waste to prevent air from being drawn across the sensor array area, which would lead to a significant loss of sequencing channels.</p>	
<p>Remove the waste buffer, as follows:</p> <p><input type="checkbox"/> Ensure the inlet port is closed.</p> <p><input type="checkbox"/> Insert a P1000 pipette into a waste port and remove the waste buffer</p> <p><input type="checkbox"/> Slide the inlet port cover clockwise to open the inlet port of the first flow cell inserted. Repeat for all flow cells.</p>	
<p><b>IMPORTANT</b></p> <p><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>	
<p>After opening the inlet port, draw back a small volume to remove any air bubbles:</p> <p><input type="checkbox"/> Set a P1000 pipette to 200 µl.</p> <p><input type="checkbox"/> For the first flow cell, insert the tip into the inlet port.</p> <p><input type="checkbox"/> Turn the wheel until the dial shows 220-230 µl, or until you can see a small volume of buffer/liquid entering the pipette tip.</p> <p><input type="checkbox"/> Repeat for the remaining flow cells using a new pipette tip.</p> <p>Slowly load 200 µl of the prepared flow cell wash mix into the inlet port, as follows:</p> <p><input type="checkbox"/> Using a P1000 pipette, take 200 µl of the flow cell wash mix</p> <p><input type="checkbox"/> Insert the pipette tip into the inlet port, ensuring there are no bubbles in the tip</p> <p><input type="checkbox"/> Slowly twist the pipette wheel down to load the flow cell (if possible with your pipette) or push down the plunger very slowly, leaving a small volume of buffer in the pipette tip.</p> <p><input type="checkbox"/> Set a timer for a 5 minute incubation.</p>	

# Loading multiple PromethION Flow Cells

Version: PFC\_9097\_v1\_revM\_18Mar2020  
 Last update: 07/02/2024



Flow Cell Number: .....

DNA Samples: .....

INSTRUCTIONS	NOTES/OBSERVATIONS
<p>Once the 5 minute incubation time is complete, carefully load the remaining 200 µl of the prepared flow cell wash mix into the inlet port, as follows:</p> <ul style="list-style-type: none"> <li><input type="checkbox"/> Using a P1000 pipette, take 200 µl of the flow cell wash mix</li> <li><input type="checkbox"/> Insert the pipette tip into the inlet port, ensuring there are no bubbles in the tip</li> <li><input type="checkbox"/> Slowly twist the pipette wheel down to load the flow cell (if possible with your pipette) or push down the plunger very slowly, leaving a small volume of buffer in the pipette tip.</li> </ul> <p><input type="checkbox"/> Close the inlet port and start a 60 minute timer once the first flow cell has been loaded.</p> <p><input type="checkbox"/> Repeat the two-step addition of the wash mix with the 5 minute break in between and the hour incubation for all of the flow cells.</p> <p><input type="checkbox"/> After each row (8 flow cells), thoroughly mix the Wash Mix.</p>	
<p><b>IMPORTANT</b></p> <p><input type="checkbox"/> It is vital that the inlet port is closed before removing waste to prevent air from being drawn across the sensor array area, which would lead to a significant loss of sequencing channels.</p>	
<p>For each flow cell, remove the waste buffer as follows:</p> <ul style="list-style-type: none"> <li><input type="checkbox"/> Ensure the inlet port is closed.</li> <li><input type="checkbox"/> Insert a P1000 pipette into a waste port and remove the waste buffer</li> </ul>	
<p>Follow one of the two options described in the next steps of the protocol.</p>	
<p><b>To run a second library on multiple PromethION Flow Cells straight away</b></p>	
<p><b>IMPORTANT</b></p> <p><input type="checkbox"/> Install the light shield on your flow cell as soon as library has been loaded for optimal sequencing output.</p> <p>To run a second library straight away, follow the instructions in the 'Priming and loading multiple flow cells on a PromethION' of this protocol, with the recommendations below.</p> <ul style="list-style-type: none"> <li><input type="checkbox"/> Pipette very slowly when loading priming mix into the flow cell.</li> <li><input type="checkbox"/> Wait five minutes between priming mix flushes.</li> <li><input type="checkbox"/> After the five minute pause, close the priming port, ensure the SpotON port is closed and remove the waste from waste port 1. This prevents the nuclease from diffusing through the flow cell.</li> </ul>	
<p><b>IMPORTANT</b></p> <p><input type="checkbox"/> When priming a flow cell after a nuclease wash with the Flow Cell Wash Kit, it is vital to wait five minutes between the priming mix flushes and to remove the waste for effective removal of the nuclease.</p>	
<p><b>To store multiple PromethION Flow Cells for later use</b></p>	
<p>Storage Buffer (S) can be used to flush flow cells for storage for later use or to check number of available nanopores before loading another library.</p>	
<p><b>IMPORTANT</b></p> <p><input type="checkbox"/> Scale up reagent volumes as needed</p>	
<p><input type="checkbox"/> Thaw Storage Buffer (S) at RT.</p>	

# Loading multiple PromethION Flow Cells

Version: PFC\_9097\_v1\_revM\_18Mar2020  
 Last update: 07/02/2024



Flow Cell Number: .....

DNA Samples: .....

INSTRUCTIONS	NOTES/OBSERVATIONS
<ul style="list-style-type: none"> <li><input type="checkbox"/> Mix contents thoroughly by pipetting and spin down briefly.</li> <li><input type="checkbox"/> Slide the inlet port cover clockwise to open the inlet port of the first flow cell inserted. Repeat for all flow cells.</li> </ul>	
<p><b>IMPORTANT</b></p> <ul style="list-style-type: none"> <li><input type="checkbox"/> Take care when drawing back buffer from the flow cell. Do not remove more than 20-30 <math>\mu</math>l, 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.</li> </ul>	
<p>After opening the inlet port, check for a small air bubble under the cover. Draw back a small volume to remove any air bubbles:</p> <ul style="list-style-type: none"> <li><input type="checkbox"/> Set a P1000 pipette to 200 <math>\mu</math>l.</li> <li><input type="checkbox"/> For the first flow cell, insert the tip into the inlet port.</li> <li><input type="checkbox"/> Turn the wheel until the dial shows 220-230 <math>\mu</math>l, or until you can see a small volume of buffer entering the pipette tip.</li> <li><input type="checkbox"/> Repeat for the remaining flow cells.</li> <li><input type="checkbox"/> Slowly add 500 <math>\mu</math>l of Storage Buffer through the inlet port of the first flow cell and repeat for the remaining flow cells.</li> <li><input type="checkbox"/> Once all the flow cells are loaded, close the inlet port covers, and remove any buffer from the waste port.</li> <li><input type="checkbox"/> Remove the flow cells from the device by pressing down on the latch to release the flow cells before sliding out smoothly.</li> <li><input type="checkbox"/> Store the flow cells at 4-8° C.</li> </ul>	
<p>When you wish to reuse the flow cell, remove the flow cell from storage, and allow it to warm to RT for ~5 minutes.</p>	
<p><b>IMPORTANT</b></p> <ul style="list-style-type: none"> <li><input type="checkbox"/> After performing a flow cell wash or storing your flow cell, we recommend using running a 'Flow cell check' to check number of available nanopores.</li> </ul>	