## illumina

## NovaSeq Xp Workflow Checklist

This document is not a replacement for the *NovaSeq 6000 System Guide (document # 100000019358).* Use the most recent versions of the system guide and NovaSeq Control Software.

## **Prepare Reagents**

□ 1 Thaw SBS and cluster (CPE) cartridges in room temperature water as follows.

| Cartridge                        | Duration of Thaw |
|----------------------------------|------------------|
| SP, S1, and S2 SBS cartridge     | 4 hours          |
| SP, S1, and S2 cluster cartridge | Up to 2 hours    |
| S4 SBS cartridge                 | 4 hours          |
| S4 cluster cartridge             | Up to 4 hours    |
|                                  |                  |

- 2 Thoroughly dry the cartridge bases.
- $\square$  3 Blot the foil seals dry if needed.
- $\Box$  4 Invert each cartridge 10 times.
- 5 Gently tap the bottom of each cartridge on the bench.
- ☐ 6 [Optional] Store thawed reagents at 2°C to 8°C for up to 24 hours.

## Prepare Instrument

- $\Box$  1 Complete the post-run wash.
- □ 2 Empty used reagent bottles and return them to the buffer drawer.

## Pool and Dilute Library

- □ 1 Bring a flow cell package to room temperature for at least 10 minutes.
- 2 Thaw DPX1, DPX2, and DPX3, and then place on ice.
- □ 3 Pool libraries to the desired plexity.

| Mode  | Total Volume of Pool Per Lane (µI) |
|-------|------------------------------------|
| SP/S1 | 18                                 |
| S2    | 22                                 |
| S4    | 30                                 |

4 Dilute library to final loading concentration as follows.

| Library<br>Type                       | Final Loading<br>Concentration<br>(pM) | Pooled Loading<br>Concentration<br>(nM) |
|---------------------------------------|--|---|
| DNA PCR-<br>free library<br>pool      | 115–235                                | 0.575–1.175                             |
| DNA PCR-<br>amplified<br>library pool | 200–400                                | 1.0–2.0                                 |
| Single Cell                           | 175–275                                | .875–1.375                              |

## Denature Library

- □ 1 Prepare 0.2 N NaOH by diluting stock NaOH with laboratory-grade water.
- 2 [Optional] Spike-in 1% nondenatured PhiX as follows.
  - a Dilute 10 nM PhiX to 0.25 nM using 10 mM Tris-HCl, pH 8.5.
  - b For each lane, add the appropriate volume of PhiX to the tube of nondenatured library pool.

| Mode  | Nondenatured<br>0.25 nM PhiX<br>(µl) | Nondenatured<br>Library Pool<br>(µl) |
|-------|--------------------------------------|--------------------------------------|
| SP/S1 | 0.7                                  | 18                                   |
| S2    | 0.8                                  | 22                                   |
| S4    | 1.1                                  | 30                                   |

□ 3 For each lane, add 0.2 N NaOH to the nondenatured library as follows.

| Mode  | 0.2<br>Ν ΝaΟΗ<br>(μl) | Nondenatured Library<br>Pool (µl) |
|-------|-----------------------|-----------------------------------|
| SP/S1 | 4.0                   | 18.0                              |
| S2    | 5.0                   | 22.0                              |
| S4    | 7.0                   | 30.0                              |

- $\Box$  4 Cap and then vortex briefly.
- □ 5 Incubate at room temperature for 8 minutes.
- 6 For each lane, add 400 mM Tris-HCl, pH 8.0 to neutralize as follows.

| Mode  | 400 mM Tris-HCl,<br>pH 8.0 (μl) | Resulting<br>Volume              |
|-------|---------------------------------|----------------------------------|
| SP/S1 | 5.0                             | 27.0 μl, or 27.7 μl<br>with PhiX |
| S2    | 6.0                             | 33.0 µl, or 33.8 µl<br>with PhiX |
| S4    | 8.0                             | 45.0 μl, or 46.1 μl<br>with PhiX |

 $\Box$  7 Cap and then vortex briefly.

■ 8 Place on ice until use.

#### SAFE STOPPING POINT

If you cannot immediately proceed, cap the tube and store at -25°C to -15°C for up to three weeks.

## Prepare ExAmp Master Mix

- □ 1 Place flow cell onto the flow cell dock and place the manifold over the flow cell.
- $\Box$  2 Close the clamp.
- $\Box$  3 Invert or vortex briefly to mix DPX1 and DPX2.
- $\Box$  4 Briefly vortex DPX3 to mix.
- $\Box$  5 Briefly centrifuge DPX1, DPX2, and DPX3.
- 6 For each flow cell, combine the following volumes in a suitable microcentrifuge tube in the order specified.

| Addition<br>Order | Reagent* | Volume for<br>Two-Lane<br>Flow Cell<br>(SP/S1/S2)<br>(µl) | Volume<br>for<br>Four-<br>Lane<br>Flow<br>Cell<br>(S4) (µl) |
|-------------------|----------|---|---|
| 1                 | DPX1     | 126   | 315   |
| 2                 | DPX2     | 18  | 45  |
| 3                 | DPX3     | 66  | 165   |
|                   |          |   |   |

\*DPX reagent tube caps may be color coded (red, yellow, and blue for DPX1, DPX2, and DPX3, respectively). Make sure that color coding is preserved when replacing tube caps.

- These volumes result in 210 µl ExAmp master mix for SP, S1, or S2 mode, or 525 µl Master Mix for S4 mode.
- $\square$  8 Pipette and dispense slowly to avoid bubbles.
- $\Box$  9 Vortex for 20–30 seconds.
- $\Box$  10 Centrifuge at up to 280 × g for up to 1 minute.
- 11 For the best sequencing performance, immediately proceed to the next step. If necessary, ideal storage of the master mix is up to 1 hour on ice. Use within 30 minutes if storing at room temperature.

## Load Library Onto the Flow Cell

□ 1 For each lane, add ExAmp Master Mix to each denatured library pool as follows.

| Mode  | Denatured<br>Library Pool<br>(µl) | ExAmp<br>Master<br>Mix (µl) | Resulting<br>Volume<br>(µl) |
|-------|-----------------------------------|-----------------------------|-----------------------------|
| SP/S1 | 27                                | 63                          | 90                          |
| S2    | 33                                | 77                          | 110                         |
| S4    | 45                                | 105                         | 150                         |

- 2 If using tube strips, pipette to mix until homogenous.
- $\Box$  3 Centrifuge at up to 280 × g for up to 1 minute.
- 4 Add the appropriate volume of library/ExAmp mixture to each manifold well.

| Mode  | Library/ExAmp Mixture per Well<br>(µl) |
|-------|--|
| SP/S1 | 80                                     |
| S2    | 95                                     |
| S4    | 130                                    |

- ☐ 5 After adding the ExAmp/library mixture to all manifold wells, wait approximately 2 minutes for the mixture to reach the opposite end of each lane.
- 6 Start the sequencing run within 30 minutes of loading libraries onto the flow cell.

## Load Flow Cell Onto the Instrument

- □ 1 From the Home screen, select **Sequence**, and then select a single or dual flow cell run:
  - ► A+B-Set up a dual flow cell run.
  - ► A-Set up a single flow cell run on side A.
  - ▶ **B**-Set up a single flow cell run on side B.
- $\square$  2 Remove flow cell from the flow cell dock:

- a Open the clamp that secures the flow cell and manifold.
- b Carefully remove and discard the manifold.
- ☐ c Grasp the sides of the flow cell and remove it from the dock.
- □ 3 Load flow cell onto the instrument:
  - $\Box$  a Invert the flow cell around the long axis.
  - b Place the flow cell on the flow cell stage, and then select Close Flow Cell Door.

# Load Cartridges Onto the Instrument

- □ 1 Place an empty library tube into position #8 of the cluster cartridge.
- 2 Remove the used SBS and cluster cartridges.
- □ 3 Load the SBS and cluster cartridges into the reagent chiller drawer.
- $\Box$ 4 Remove the used buffer cartridge.
- □ 5 Place a new buffer cartridge into the buffer drawer.

## Start Sequencing

- □ 1 [Optional] Sign in to BaseSpace Sequence Hub.
- $\Box$  2 Select **Run Setup**, and then **NovaSeq Xp**.
- $\Box$  3 Enter the required parameters and settings.
- 4 Select Start Run.