# Fragment DNA

- $\Box 1$  Quantify gDNA.
- $\square$ 2 Normalize gDNA with RSB in a final volume of 15  $\mu$ l in a new plate:
  - ▶ For a 350 bp insert size—1.67 ng/µl of sample
  - ▶ For a 550 bp insert size—5 ng/µl of sample
- $\square$ 3 Transfer 15  $\mu$ l DNA to a microTUBE-15.
- $\Box 4$  Centrifuge at 3000 × g for 1 minute.
- $\Box$ 5 Fragment the DNA on a Covaris.

Table 1 Covaris S220 or E220 Settings

Setting	350 bp Insert	550 bp Insert
Duty factor	20%	
Peak Incident Power	18 W	
Cycles per burst	50	
Duration	45 seconds	22 seconds
Temperature	20°C	
Water Level—S220	15	
Water Level—E220	10	

Table 2 Covaris M220 Settings

Setting	350 bp Insert	550 bp Insert
Duty factor	20%	
Peak Incident Power	30 W	
Cycles per burst	50	
Duration	42 seconds	23 seconds
Temperature	20°C	

- $\Box$ 6 Centrifuge at 600 × g for 5 seconds.
- $\Box$ 7 Transfer 15 µl DNA to a new plate.

SAFE STOPPING POINT

If you are stopping, seal the plate and store at -25°C to -15°C for up to 7 days.

# Prepare Samples for Loading

- □1 Add 1 of the following to a new 1.5 ml microcentrifuge tube:
  - ▶ For a 350 bp insert size—900 µl SC350
  - ▶ For a 550 bp insert size—700 µl SC550
- $\square$ 2 Vortex DMB.
- $\square$ 3 Add 100 µl DMB. Vortex for 5 seconds.
- □4 Pour the DMB and SC mixture into a new reservoir.
- Add 35 μl DMB and SC mixture to the sample plate. Pipette to mix.
- $\Box$ 6 Shake or vortex at 1400 rpm for 12 minutes.

# Set Up Run and Load Library Card

- $\Box$ 1 Vortex the reagent plate for 3 seconds.
- Centrifuge at  $600 \times g$  for 5 seconds.
- ☐ 3 Select **Prepare Libraries**.
- $\Box 4$  Select a protocol or run. Select **Next**.
- □5 Configure the run. Select **Next**.
- ☐ Review the run and sample information. Select **Next**.
- $\Box$ 7 Enter tracking information. Select **Next**.
- $\square 8$  Place the library card on the stage.
- □9 Close the door. Select **Verify Library Card**.
- $\Box$ 10 Place the guide on the library card.
- $\Box$ 11 Load the oil.
- $\square$ 12 Transfer 45  $\mu$ l of samples 1–8 and 9–16.
- $\Box$ 13 Add 45 µl RSB to empty sample wells.
- □14 Transfer 125 µl of reagents i–iv and v–vii.
- $\Box$ 15 Vortex DMB.
- □16 Add 60 µl DMB to reagent well viii.
- $\Box$ 17 Transfer 15 µl of reagents 1–4 and 5–8.
- $\Box$ 18 Transfer 5 µl of reagents a–d and e–h.
- $\Box$ 19 Transfer 3 µl of adapters A–H and I–P.
- $\square$ 20 Remove the guide.
- $\Box$ 21 Close the door. Select **Start Run**.
- $\square$ 22 When the run is complete, select **Next**.

# TruSeq Nano DNA Library Prep for NeoPrep

For Research Use Only. Not for use in diagnostic procedures.

### **Unload Libraries**

$\Box 1$ Add 10 $\mu$ l 1	RSB to a new pla	ate
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- $\Box$ 2 Place the guide on the library card.
  - Transfer 20  $\mu$ l from 1L–8L and 9L–16L to the plate. Pipette to mix.
- $\Box$ 4 Centrifuge briefly.
- ☐5 Transfer from plate wells 1–8 and 9–16 to the library separation tubes.
- $\Box$ 6 Let stand for 10 seconds.
- $\square$ 7 Transfer from library separation tubes 1–8 and 9–16 to a new plate.
- $\square 8$  Remove the library card and guide.
- $\square$ 9 Select **Home**.

#### SAFE STOPPING POINT

If you are stopping, seal the plate and store at -25°C to -15°C for up to 2 months.

## **Pool Libraries**

- $\Box$ 1 Determine the number of samples to combine.
- $\square$ 2 Transfer 5  $\mu$ l to a single well of a new plate. Pipette to mix.
- $\square$ 3 Proceed to cluster generation.

#### SAFE STOPPING POINT

If you are stopping, seal the plate and store at -25°C to -15°C for up to 2 months.

# Acronyms

Acronym	Definition
DMB	Digital Microfluidics Beads
RSB	Resuspension Buffer
SC350	Sample Concentration Solution 350
SC550	Sample Concentration Solution 550