

Targeted Single-Cell DNA Sequencing using the Tapestri® Platform V2

For more information, consult the Tapestri Single-Cell User Guide (PN 3354)

- NOTE**
- **Vortex all reagents.**
 - **Thaw -20 °C reagents on ice.**
 - **Avoid sources of static and pipette slowly and carefully when handling emulsions.**

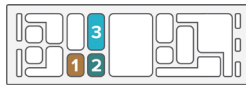
Prepare Single Cell Suspension

- IMPORTANT** Provide debris-free cell suspension with > 80% viability. Final cell concentration is 3,000 - 4,000 cells/μl.

- Using **Cell Buffer** dilute cells to ~ 3,500 cells/μl in a total volume of at least 50 μl.

Encapsulate Cells

- Mount the Base Plate onto the Tapestri Instrument.
- Place 0.2 mL Axygen MAXYmum Recovery PCR tube into the middle of the slot at the left side of the Base Plate.
- Place the DNA cartridge onto the base plate.
- In a new tube, prepare **Lysis Mix** by adding **7.3 μl of Reverse Primer Pool** (●) into **92.7 μl of Lysis Buffer** (●).
- Pipette **100 μl of Lysis Mix** into **reservoir 1**.
- Pipette **35 μl of Cell Suspension (3,500 cells/μl)** into **reservoir 2**.
- Pipette **200 μl of Encapsulation Oil** into **reservoir 3**.
- Apply DNA Gasket** on top of the cartridge and close instrument lid.
- Run the Cell Encapsulation program by pressing **Step 1: Encapsulation** on the Tapestri instrument touch screen.
- Once program is completed, press **DONE** and remove cartridge and collection tube from base plate.
- Use a **gel loading tip** to carefully **remove up to 100 μl of the oil layer** at the bottom of the sample leaving no more than 5 μl of oil.



Lysis + Protease Digest

- Place sample tube into thermal cycler and run the Lysis/Digest protocol:

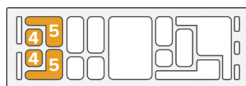
Step	Temperature	Time
1	50 °C	60 min
2	80 °C	10 min
3	4 °C	HOLD

- Once complete, store the lysed and digested sample at 4 °C until required in the next step.

Barcode Cells

Prime the DNA Cartridge

- Label eight 0.2 mL Axygen MAXYmum Recovery PCR tubes with the sample number and load them into the eight bottom slots of the Base Plate.
- Mount the DNA Cartridge again onto the Base Plate.
- Pipette **200 μl of Electrode Solution** into **reservoirs 4** and **500 μl of Electrode Solution** into **reservoirs 5**.
- Apply DNA Gasket** on top of the cartridge and close instrument lid.
- Run the Priming program by pressing **Step 2: Barcoding** on the Tapestri Instrument touch screen.
- Once program is completed, press **DONE** and proceed to step 9.
- In the meantime retrieve **Barcoding Beads V2** (●) and **leave at room temperature (protect from light)** and prepare **300 μl Barcode Mix** as shown in the table on the right top side of this page.
- Briefly **vortex the Barcode Mix** and centrifuge to collect the contents and **store on ice**. *Continued on second column of this page.*



Reagent	Volume (μl)
Barcoding MM V2	295
Forward Primer Pool (○)	5
Total Volume	300

Load the DNA Cartridge

- Retrieve lysed and digested sample from thermal cycler.
- Pipette the entire **cell encapsulation sample** (~80 μl), including any oil at the bottom, into **reservoir 6**.
- Vortex **Barcoding Beads V2** (●) for 1 minute at high speed and carefully pipette **200 μl of Barcoding Beads V2** (●) into **reservoir 7**.
- Pipette **250 μl of Barcode Mix** into **reservoir 8**.
- Pipette **1.25 mL of Barcoding Oil** into **reservoir 9**.
- Apply DNA Gasket** on top of the cartridge and close instrument lid.
- Run the Cell Barcoding program by pressing **NEXT** on the Tapestri Instrument touch screen.
- Once the program is completed, press **DONE** and remove the base plate together with the cartridge and the eight collection tubes
- Remove the DNA Cartridge from the Base Plate.
- Use a **gel loading tip** to carefully **remove up to 120 μl of oil** from the bottom layer of each sample tube.



- NOTE** Volumes may vary. The final volume of oil must be ~35 μl per tube & the total volume in each tube must not exceed 100 μl.

UV Treatment

- Place entire Base Plate with tubes (closed) containing emulsions on ice.
- Place a UV light on top of the ice bucket and expose the samples to non-irradiating **UV light for 8 minutes**.

- IMPORTANT** Make sure to use the **Analytik Jena Blak-Ray XX-15L UV light source**.

Targeted PCR Amplification

- After UV exposure, remove the Base Plate from the ice, transfer the sample tubes to the thermal cycler, and run the Targeted PCR protocol:

		Panel Size	20 - 100	100 - 200	200 - 300	> 300	
Step	Ramp	Temp.	Time	Time	Time	Time	Cycle
1	4 °C/s	98 °C	6 min	6 min	6 min	6 min	
2	1 °C/s	95 °C	30 sec	30 sec	30 sec	30 sec	10
3		72 °C	10 sec	10 sec	10 sec	10 sec	
4		61 °C	3 min	4.5 min	6 min	9 min	
5	72 °C	20 sec	20 sec	20 sec	20 sec		
6	1 °C/s	95 °C	30 sec	30 sec	30 sec	30 sec	
7		72 °C	10 sec	10 sec	10 sec	10 sec	
8		48 °C	3 min	4.5 min	6 min	9 min	
9	72 °C	20 sec	20 sec	20 sec	20 sec		
10	4 °C/s	72 °C	2 min	2 min	2 min	2 min	
11		4 °C	HOLD	HOLD	HOLD	HOLD	

- IMPORTANT** Make sure to set up two separate 10 cycle programs each with a ramp rate of 1 °C/s.

Break Emulsions

- Pool the entire contents of tubes 1 - 4 and tubes 5 - 8 into two new 1.5 mL Eppendorf tubes.
- Add **40 µl of Extraction Agent** (●) to each tube. Vortex and quick spin.
- Incubate at room temperature for **3 minutes**.
- Add **180 µl of nuclease-free water** to each tube.
- Briefly vortex and spin down for 10 seconds in a mini centrifuge to separate the aqueous and oil layers.
- Pipette **168 µl of the aqueous top layer** in each tube into two new 1.5 mL Eppendorf tubes. **Do not transfer any oil.**

Digest PCR Product

- For each of the two 168 µl samples prepare a **200 µl digestion mix** by adding **20 µl DNA Clean up Buffer** (○) and **12 µl Clean up Enzyme** (●).
- Mix by pipetting up and down and quick spin the tubes.
- Transfer both tubes to a heat block and **run the Digest protocol**:

Step	Temperature	Time
1	37 °C	60 min
2	4 °C	HOLD

Clean Up PCR Product

NOTE Equilibrate Ampure XP reagent to room temperature. Prepare 5 mL fresh 80% ethanol using nuclease-free water.

- Add **200 µl of nuclease-free water** to each tube (total volume = 400 µl).
- Thoroughly vortex Ampure XP reagent at high speed immediately prior to usage.
- Add **288 µl of Ampure XP reagent** to each of the two tubes. Vortex for **5 seconds** and quick-spin to collect contents.
- Incubate tubes at room temperature for **5 minutes**.
- Place on magnet and wait 5 minutes for the beads to separate from solution.
- Without removing the tubes from the magnet, remove the clear liquid from each tube and discard.
- To each tube carefully add **800 µl of the freshly prepared 80% ethanol**, wait 30 seconds, and **remove 800 µl of ethanol** without disturbing the Ampure beads.
- Repeat step 7 once, for a total of two wash cycles.
- Keeping the tubes on the magnet, remove all residual ethanol from each tube without disturbing the beads.
- Dry Ampure bead pellets in the tubes on the magnet by incubating at room temperature for **5 minutes**. **Avoid overdrying beads.**
- Remove the tubes from the magnet. Add **65 µl of nuclease-free water** into each tube. Vortex and quick-spin to collect the contents.
- Incubate tubes at room temperature for **2 minutes**.
- Place the tubes onto the magnet and wait for at least **2 minutes** or until solutions are clear.
- Transfer and combine **50 µl of purified PCR product** from each tube to a single new 0.2 mL PCR tube. **Avoid transfer of beads.**
- Store the tube with 100 µl purified PCR product on ice.
- Quantify **1 µl of purified PCR product** from each sample, using the High Sensitivity Qubit Kit (or equivalent assay) according to the manufacturer's instructions.

NOTE The DNA quantity in each sample tube may vary between **0.2 ng/µl to 4.0 ng/µl**. If yields are outside this range, contact support@missionbio.com for additional support.

- Store purified PCR product sample at -20 °C until proceeding to the next step.

Library PCR

- In a Pre-PCR area in a new 0.2 mL PCR tube normalize the sample to 2 ng in 15 µl, using the **Tapestri Sample Quantification Tool (PN 40676E1)**.
- Add Library MM V2 and V2 Index Primer (●) as follows:

Reagent	Volume
Library MM V2	25
V2 Index Primer (●)	10
Normalized PCR product (2 ng total)	15
Total Volume	50

- Vortex and quick-spin tube to collect contents.
- Transfer sample tube to a thermal cycler and run the Library PCR protocol:

Step	Temperature	Time	Cycle
1	95 °C	3 min	
2	98 °C	20 sec	9
3	62 °C	20 sec	
4	72 °C	45 sec	
5	72 °C	2 min	
6	4 °C	HOLD	

Clean Up Library PCR Product

NOTE Equilibrate Ampure XP reagent to room temperature. Prepare 5 mL fresh 80% ethanol using nuclease-free water.

- Add **34.5 µl of Ampure XP reagent** to the 50ul sample tube. Vortex for **5 seconds** and quick-spin to collect contents.
- Follow Steps 4 - 15 of Section **Clean Up PCR Product** with the following modifications: **Wash with 200 µl ethanol, elude DNA in 12 µl of nuclease-free water and transfer 1 x 10 µl of purified product** to a new 0.2 mL PCR tube.

Quantify and Pool Library

- Quantify the library using a Agilent Bioanalyzer. DNA 1000 chips may be used with 1 µl of undiluted samples or DNA HS chips may be used with 10x diluted samples.
- Use the **Tapestri Sample Quantification Tool (PN 40676)** to pool 5 nM of each of the sample libraries.
- Verify the concentration of the pooled library using a Qubit Fluorometer or equivalent instrument.

IMPORTANT Refer to the **Tapestri Single-Cell DNA Sequencing User Guide (PN 3354)** for additional information on how to accurately quantify Tapestri libraries that include large-size off-target products.

Sequence Tapestri Single-Cell DNA Library

IMPORTANT The final library consists of target-specific amplicons ranging from **350 - 550 bp**.

- Illumina MiSeq, HiSeq 2500, HiSeq 4000, NextSeq 550, and NovaSeq 6000 are currently supported.
 - Paired-end Illumina sequencing chemistry is required (2x150 bp).
 - Each sample is split into two dual-indexed libraries.
 - When multiplexing more than 8 samples use Illumina's Nextera dual indices.
 - Please refer to the following Illumina User Guides.
- Sequence the library following manufacturer's instructions.

For technical support visit www.missionbio.com/support or email support@missionbio.com.