

· Part No.: H-020-000881-00

· Version: 2.0

· Release date: April 2024 ©MGI All rights reserved.

**RT-PCR** multiplex amplification

RT-PCR product purification

Preparing Fast PCR-FREE library

Making OS DNBs

Sequencing and analysis

### Applicable sets/kits

ATOPlex HIV-1 Library Preparation Set (16 RXN)

Name	Component		Cat. No.
ATOPlex HIV-1	ATOPlex HIV-1 Drug Resistance Primer Pool		940-000722-00
Amplification Kit	RT-PCR Buffer		
	RT-PCR Enzyme Mix		
MGIEasy Fast PCR- FREE FS DNA Library Prep Set (16 RXN) (Cat. No.:	MGIEasy Fast PCR-FREE FS DNA Library Prep Kit	20* Elute Enhancer Fast FS Buffer Fast FS Enzyme Ligation Enhancer Fast Ligation Buffer Ad Ligase	940-000017-00
940-000019-00)	MGIEasy UDB PF Adapter Kit	UDB Adapters	940-000018-00
	MGIEasy DNA Clean Beads	TE Buffer DNA Clean Beads	1000005278

• ATOPlex HIV-1 Library Preparation Set (96 RXN)

Name	Component		Cat. No.
ATOPlex HIV-1	ATOPlex HIV-1 Drug Resistance Primer Pool		940-000722-00
Amplification Kit × 6	RT-PCR Buffer		
	RT-PCR Enzyme M	lix	
MGIEasy Fast PCR-FREE FS DNA Library Prep Set (96 RXN) (Cat. No.: 940-	MGIEasy Fast PCR-FREE FS DNA Library Prep Kit	20* Elute Enhancer Fast FS Buffer Fast FS Enzyme Ligation Enhancer Fast Ligation Buffer Ad Ligase TE Buffer	940-000020-00
000021-00)	MGIEasy UDB PF Adapter Kit A	UDB Adapters A	940-000023-00
	MGIEasy DNA Clean Beads × 2	TE Buffer DNA Clean Beads	1000005278
MGIEasy DNA Clean Beads	TE Buffer DNA Clean Beads		1000005278



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 DNBSEQ OneStep DNB Make Reagent Kit (OS-DB) (Cat. No.: 1000026466)

Name	Quantity
Low TE Buffer	300 μL × 1 tube
Make DNB Buffer (OS DB)	80 μL ×1 tube
Make DNB Enzyme Adapters Mix I (OS)	160 μL × 1 tube
Make DNB Enzyme Mix II (OS)	16 µL × 1 tube
Stop DNB Reaction Buffer	100 μL × 1 tube

Sequencing sets

Name	Cat. No.
DNBSEQ-G99RS High-throughput Sequencing Set (G99 SM FCL PE150)	940-000410-00
DNBSEQ-G99RS High-throughput Sequencing Set (G99 FCL PE150)	940-001269-00

#### **Recommended consumables**

Name	Cat. No.	Brand
0.5 mL SC micro tube	1000001558	
2 mL SC micro tube	1000001553	
250 µL automated filter tips	100000723	- MGI
0.2 mL 96 well full-skirt PCR plates	091-000165-00	IVIGI
1.3 mL U-bottom deep-well plates	1000004644	_
Break-away 8-strip PCR tubes and caps	100-000016-00	-

#### RT-PCR multiplex amplification (manually)

### **Preparing samples**

Prepare the total RNA samples, including plasma and serum. It is recommended to use high-quality genomic RNA samples with the CT value less than or equals to 32. The sample amount is recommended to be 20  $\mu$ L, but should not exceed 20.5  $\mu$ L.



Use a buffer that does not contain EDTA to dissolve RNA samples.

#### Performing RT-PCR multiplex amplification

1. Take out the following reagents from ATOPlex HIV-1 Amplification Kit. Prepare the RT-PCR multiplex amplification mixture on ice:

Reagent	Volume (μL)
RT-PCR Buffer	25
RT-PCR Enzyme Mix	2.5
ATOPlex HIV-1 Drug Resistance Primer Pool	2
Nuclease-Free Water	0.5
Total	30

- 2. Vortex the mixture, briefly centrifuge it, and transfer it into a new 0.2 mL PCR tube.
- 3. Add 20  $\mu L$  RNA sample to the PCR tube. The total volume reaches 50  $\mu L$ .
- 4. Pipette the mixture in the PCR tube for 10 times, briefly centrifuge it, and collect the mixture at the bottom of the tube.

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Do not vortex or shake the tube or pipette the solution vigorously.

5. Place the PCR tube in the PCR device and start a reaction according to the conditions in the table below:

Temperature	Time	Cycle number
105 °C (Heated lid)	On	/
50 ℃	30 min	1
94 ℃	3 min	·
94 ℃	30 s	
58 ℃	45 s	42
72 °C	2 min	
72 °C	5 min	. 1
12 ℃	Hold	.

6. After the reaction is completed, take out the PCR tube and place it on ice until use.

### **RT-PCR** product purification

#### **Preparing samples**

Transfer 50 µL of the RT-PCR product to one set of 8-strip tubes and put them on ice until use.



### Tips

- Ensure that no bubble exists at the bottom of the tube and no liquid exists on the inner wall.
- For 8 samples, one set of 8-strip tubes is needed. For 16 samples, 2 sets of 8-strip tubes are needed.
- If the number of samples is less than 8 or 16, fill in the remaining tubes with water. Prepare the reagents and consumables according to 8 or 16 samples.

### Preparing reagents

1. Take out the following reagents from MGIEasy DNA Clean Beads. Mix them thoroughly by using a vortex mixer.

Doowant	Volum	e (µL)
Reagent	8 RXN	16 RXN
DNA Clean Beads	380	660
TE Buffer	500	900



Take out DNA Clean Beads 30 min in advance and put them at room temperature. Vortex them thoroughly before using them.

- 2. Take two new 2 mL SC micro tubes. Transfer the reagents of the volume listed in the foregoing table above into the tubes, cap the tubes, and mark them as "DNA Beads 1" and "TE".
- 3. Prepare 25 mL of 80% ethanol by using absolute ethanol and Milli-Q water.
- Tips

Use 80% ethanol immediately after preparation.



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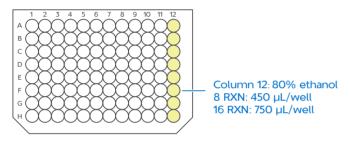
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4. Take out a 1.3 mL U-bottom deep-well plate. Add 80% ethanol to wells in column 12 of the plate.



### **Initializing MGISP-100**

- 1. Turn on the computer and the desktop is displayed. Double-click the icon on the control software to open the software.
- 2. Select Real mode and click Create.
- 3. Click **User Entry** to open the main interface.
- 4. Click **Initialize** on the top of the interface to start initializing. Once the initialization is completed, a prompt message will appear on the interface.

### Performing a pre-clean



Perform a pre-clean before conducting experiments every day.

- 1. Click the menu button on the left of the interface, and select Pre-post clean.
- 2. From the Clean list, select **Pre-clean**, and click **Start**.

- 3. Perform the required operations by following the on-screen instructions. Then, click Continue.
- 4. The UV lamp and air filter start working.

### Performing RT-PCR product purification

1. Set **Solution** and **Script** in the Run Wizard interface of MGISP-100 according to the table below:

	Solution	JB-A06-117 MGI HIV-1 Drug Resistance Sequencing Package_RV1.0_SV1.0	
Script 1.ATOPlex_RNA_HIVDR_PCR_Purification_step1			



Ensure that the automation script and PCR program have been properly installed according to MGISP-100 MGI HIV-1 Drug Resistance Sequencing Package User Manual before running the script.

- 2. Mix DNA Clean Beads and EN-Beads thoroughly by using a vortex mixer and centrifuge them briefly.
- **Tips**

Ensure that no bubble exists at the bottom of other tubes, no liquid exists on the inner wall, and all tubes are uncapped.



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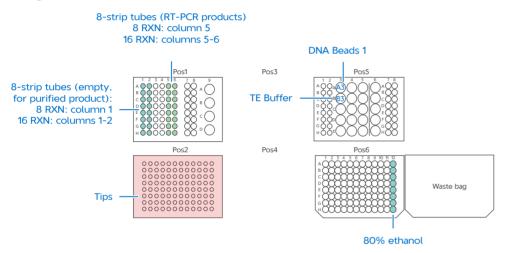
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3. Open the door of MGISP-100. Place the consumables according to the figures below:



- 4. Close the door of the device.
- 5. Click **Run**. The Sample\_Num window pops up. Then select the desired number of samples on the pop-up window and click **Continue**.
- 6. After the process is completed, take out the 8-stripe tubes with the RT-PCR products at Pos 1, and cap the tubes. The volume should be 30 µL for each tube.

Number of samples	Position
8 RXN	Column 1
16 RXN	Columns 1-2

7. Click **Continue** to end the process.

- 8. Quantify the RT-PCR products by using Qubit® dsDNA HS Assay Kit or Quant-iT PicoGreen dsDNA Assay Kit. The product concentration should be no less than 2 ng/µL.
- Stoppoint

The RT-PCR products can be stored in a -20 °C refrigerator and should be used in two weeks.

- 9. Dispose of the discarded sample tubes, reagent tubes, deep-well plates, and waste bag to the designated waste area.
- 10. (Optional) If no experiment will be conducted on the day, clean the device with pure water and 75% ethanol. Then, perform a post-clean.

#### **Preparing Fast PCR-FREE library**

### **Preparing samples**

- 1. The starting amount of each sample is recommended to be 215 ng for Fast PCR-FRE library preparation. Add TE Buffer to the sample until the total volume reaches 48 µL. If the starting amount is less than 215 ng, put all of the sample for library preparation and add TE Buffer to the sample until the total volume reaches 48 µL.
- 2. Transfer 48  $\mu$ L of each sample to a set of break-away 8-strip PCR tubes, and place the tubes on ice until use.



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- Ensure that no bubble exists at the bottom of the tubes and no liquid exists on the inner wall.
- For 8 samples, one set of 8-strip tubes is needed. For 16 samples, 2 sets of 8-strip tubes are needed.
- For the number of samples less than 8 or 16, fill in the remaining tubes with water. Prepare the reagents and consumables according to 8 or 16 samples.

### **Preparing reagents**

- 1. Prepare 1\* Elute Enhancer, En-TE, and En-Beads.
- **Tips**

1\* Elute Enhancer should be stored at room temperature, and En-TE and En-Beads should be placed at a 4 °C refrigerator. Use all solutions within 7 days.

■ 1× Elute Enhancer

Component	Volume (µL)
20× Elute Enhancer	1.5
Nuclease-Free Water	28.5
Total volume	30

■ En-TE

Component	Volume (µL)
1× Elute Enhancer	5.4
TE Buffer	2694.6
Total volume	2700

#### ■ EN-Beads

Component	Volume (µL)
1× Elute Enhancer	20
DNA Clean Beads	1980
Total volume	2000

2. Mix En-Beads thoroughly. Transfer them into a new 2 mL SC micro tube according to the table below. Mark the tube as "DNA Clean Beads 2", and cap the tubes.

Mark Consumable	Volume (μL)		
Iviai K	Consumable	8 RXN	16 RXN
DNA Clean Beads 2	2 mL SC micro tube	950	1800

3. Take out the reagents according to the table below and thaw them. Mix each reagent thoroughly, centrifuge them, and place them on ice until use. Take out 0.5 mL SC micro tubes to aliquot the reagents.

### Tips

- If you prepare libraries for 8 RXN by using 16 RXN set, take out five 0.5 mL SC micro tubes to aliquot the reagents.
- If you prepare libraries for 16 RXN by using 16 RXN set, you do not need to aliquot the reagents, but directly use them for library preparation. However, you need to prepare two 0.5 mL SC micro tubes.
- If you prepare libraries for 16 RXN by using 96 RXN set, take out five 0.5 mL SC micro tubes to aliquot the reagents. You need to prepare two 0.5 mL SC micro tubes.

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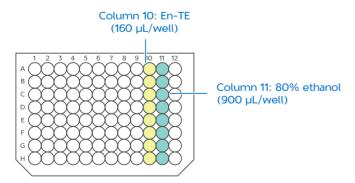
Sequencing and analysis

Reagent	Volume (μL)	
	8 RXN	16 RXN
Fast FS Buffer	102	200
Fast FS Enzyme	47.5	110
Ligation Enhancer	25	48
Fast Ligation Buffer	220	430
Ad Ligase	45	105

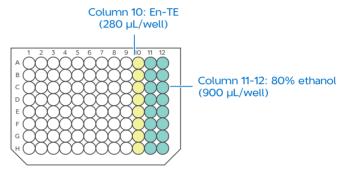
- 4. Prepare 25 mL of 80% ethanol by using absolute ethanol and Milli-Q water.
- **Tips**

Use 80% ethanol immediately after preparation.

- 5. Take out a 1.3 mL U-bottom deep-well plate and add the reagents according to the figures below:
  - For 8 RXN



■ For 16 RXN



6. Take out Adapters, mix them thoroughly and centrifuge them, and add 10 µL to 8-strip tubes.

### **Preparing libraries on MGISP-100**

1. Set **Solution** and **Script** in the Run Wizard interface of MGISP-100 according to the table below:

Solution	JB-A06-117 MGI HIV-1 Drug Resistance Sequencing Package_RV1.0_SV1.0
Script	2.ATOPlex_Fast_PCR_FREE_DNA_8RXN_step2
Script 2.ATOPle	2.ATOPlex_Fast_PCR_FREE_DNA_16RXN_step2

2. Mix DNA Clean Beads and EN-Beads thoroughly by using a vortex mixer and centrifuge them briefly.



Ensure that no bubble exists at the bottom of other tubes, no liquid exists on the inner wall, and all tubes are uncapped.

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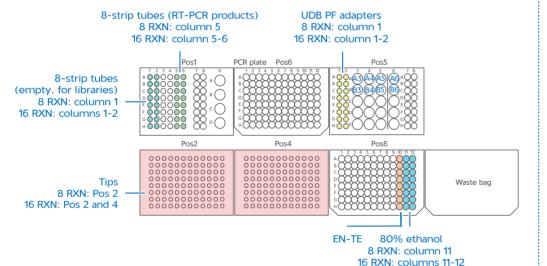
**Preparing Fast PCR-FREE library** 

Making OS DNB

Sequencing and analysis

- 3. Open the door of MGISP-100. Place the consumables according to the figures below:
- 🕜 Tips

Prepare the following consumables for both 8 RXN and 16 RXN samples unless it is specified.



Add the reagents to wells at Pos 5 according to the table below:

Well	Reagent
A3	DNB Beads 2
A4	Fast FS Buffer
A5&B5	Empty 0.5 mL SC micro tubes (16 RXN)
A6	Fast FS Enzyme
В3	Ligation Enhancer
B4	Fast Ligation Buffer
B6	Ad Ligase

- 4. Close the door of the device.
- 5. Click Run.
- 6. After the process is completed, take out the Fast PCR-FREE product from Pos 1. The volume should be 25  $\mu$ L for each tube. Then cap the 8-strip PCR tubes.

Number of samples	Position
8 RXN	Column 1
16 RXN	Columns 1-2

- 7. Click **Continue** to end the process.
- 8. Quantify the Fast PCR-FREE product by using Qubit dsDNA HS Assay Kit or Quant-iT PicoGreen dsDNA Assay Kit. The product concentration should be no less than 0.8 ng/µL.
- Stoppoint

The Fast PCR-FREE product can be stored in a -20 °C refrigerator.

- 9. Dispose of the discarded sample tubes, reagent tubes, deep-well plates, and waste bag in the designated waste area.
- 10. (Optional) If no experiment will be conducted on the day, clean the device with pure water and 75% ethanol. Then, perform a post-clean.



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### **Making OS DNBs**

### **Preparing samples**

Tips

If the samples are loaded on one flow cell, ensure that barcodes corresponding to the libraries are base-balanced.

- 1. Prepare all libraries in equal quantities and transfer them into a 0.2 mL PCR tube. Mix the libraries thoroughly by using a vortex mixer. Take 15 ng of the mixed library to make DNBs. The volume should be equal to or more than or 10  $\mu$ L.
- Tips

If the total volume is less than 10  $\mu L$ , add TE Buffer to the sample until the total volume reaches 10  $\mu L$ .

2. Take out the following reagents from DNBSEQ OneStep DNB Make Reagent Kit (OS-DB), and prepare the reaction solution:

Component	Volume (µL)
Mixed library	V
TE Buffer	10-V
Make DNB Buffer (OS-DB)	10
Total volume	20

3. Vortex and briefly centrifuge the PCR tube to collect the reaction solution at the bottom of the tube.

4. Place the PCR tube in the PCR device and start a reaction according to the conditions in the table below:

Temperature	Time
105 °C (Heated lid)	On
95 ℃	3 min
40 ℃	3 min
4 ℃	Hold

5. After the reaction is completed, take out the PCR tube, place it on ice, and add the following reagents:

Component	Volume (µL)
Make DNB Enzyme Mix I	20
Make DNB Enzyme Mix II (OS)	2
Total volume	22

- 6. Vortex and briefly centrifuge the PCR tube to collect the reaction solution at the bottom of the tube.
- 7. Place the PCR tube in the PCR device and start a reaction according to the conditions in the table below:

Temperature	Time
35 °C (Heated lid)	On
30 ℃	20 min
4 ℃	Hold

Tips

It is recommended to set the temperature of the heated lid to 35  $^{\circ}\text{C}$  or as close as possible to 35  $^{\circ}\text{C}$  .



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- 8. After the reaction is completed, add 10  $\mu$ L of Stop DNB Reaction Buffer to the PCR tube. Pipette up and down 5 to 8 times by using a wide-bore pipette tip to mix the solution thoroughly.
- ▼ Tips

Do not vortex or shake the tube or pipette the solution vigorously.

Stoppoint

The DNBs can be stored at a 4 °C refrigerator for up to 48 hours.

- 9. Take out 2  $\mu$ L of the DNBs and quantify it by using Qubit ssDNA Assay Kit. The concentration of the DNBs should be no less than 8 ng/ $\mu$ L.
- Tips

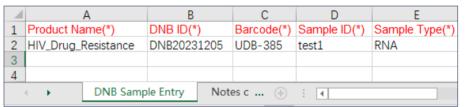
If the concentration of the DNBs exceeds 40 ng/ $\mu$ L, dilute the DNBs to a concentration of 20 ng/ $\mu$ L with DNB Load Buffer I.

### Sequencing and analysis

### Importing the sample information

- 1. Launch the Chrome browser, type the following IP address in the address bar, and press the **Enter** key: 192.168.1.5.
- 2. Enter the username (lite) and password (lite123456), and tap Login to open the Home page.
- 3. Tap **Sequencing + Analysis** on the Home page to open the New Sequence + Analysis page.
- 4. Select **HIV\_Drug\_Resistance** or set analysis product to **HIV\_Drug\_Resistance**, select **Import the Sample ID** (using this method as an example) to import DNB sample information, and tap **New**.

- 5. Tap **Excel Template** or **CSV Template** in the Sequencing + Analysis import window to download the sample template in .xlsx or .csv format.
- 6. Open the template, fill in the worksheet, and save it to the designated directory.



- Tips
  - A field with an asterisk (\*) is required, and others are optional.
  - Cells in the Excel template cannot be merged, and no spaces are allowed before or after the contents in each cell.
  - Some fields include drop-down options, and other fields require manual input. The inputting method varies among the fields.
  - **DNB ID**: The DNB ID should include both letters and numbers. It cannot be the same as the DNB ID that already exists in ZLIMS.
  - Barcode: When one sample ID corresponds to multiple barcodes, the multiple barcodes are separated by half-width commas (,). For multiple consecutive barcodes without letters, a tilde (~) can be used.
  - **Sample ID**: It should be a combination of letters and numbers and be unique for identifying the sample.
  - **Sample Type**: Select **RNA** as the sample type.
- 7. Return to the Sequencing + Analysis import window, tap **Choose File**. Select the completed worksheet and tap **Upload**. The New Sequence + Analysis page returns to view after importing the DNB sample information.



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8. After reviewing the sample information, tap **Save**, and then tap **OK** in the pop-up window.

### Preparing the flow cell and sequencing reagent cartridge

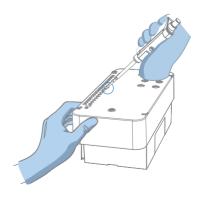
- 1. Take out the flow cell and place it at room temperature for at least 30 minutes for DNB loading.
- Tips
  - The flow cell should not be left at room temperature for more than 24 hours.
  - Do not open the outer plastic package of the flow cell at this time.
- 2. Take out the sequencing cartridge and thaw it in a water bath at room temperature for 3 to 4 hours. After thawing, place the cartridge in the  $2 \, ^{\circ}\text{C} 8 \, ^{\circ}\text{C}$  refrigerator until use.
- 3. Remove the package of the cartridge, open the cartridge cover and wipe the water condensation with a dust-free paper.
- 4. Press the cartridge wells M1, M2, M3, and M4 with a pressing tool.



5. Hold the A and B sides of the cartridge with both hands by following the marks A and B on the cartridge, and invert the cartridge for 10 to 20 times to mix the reagents thoroughly. Pierce the MDA well with a 1 mL pipette tip. Then, gently tap the cartridge against the desktop to remove bubbles.



- 6. Take out MDA reagent and MDA Enzyme Mix from the reagent kit. Transfer 125 µL of MDA Enzyme Mix into MDA reagent tube by using a 200 µL pipette for PE sequencing.
- 7. Invert the cartridge 6 times to mix the solution thoroughly.
- 8. Add all solution into the MDA well.
- Tips
  - When adding MDA Enzyme Mix, ensure that the pipette tip is tilted against one side of the MDA well to avoid bubbles.
  - Add MDA Enzyme Mix slowly to prevent it from overflowing into other wells.



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### Sequencing and analysis

- 1. Tap in the upper-right corner of the main interface of the control software.
- 2. Enter the username (user) and password (123). Tap Login.
- 3. Tap **Sequence** in the flow cell A or B operation area. If you need simultaneously perform sequencing on the flow cell stages A and B, tap **Sequence A&B**.
- 4. After tapping **Sequence**, the system will automatically open the waste container compartment door. Check if the waste container is in place by following the on-screen prompts. After checking, close the waste container compartment door. The device automatically starts to check.
- Tips

Empty the waste container in the waste container compartment before starting sequencing.

5. After the check is completed, tap **Next** to open the setting parameters interface.

Select **Sequencing & Transmission** and input the DNB ID in the **DNB ID** box.

Tips

Ensure that DNB ID is consistent with that input in ZLIMS.

- 6. Select **PE150+10+10** in the **Recipe** box. Tap the dropdown box and select **UDB** (1-480).
- Tips

For instructions on importing the customized barcode **UDB** (1-480), refer to the relevant user manual.

- 7. Select **Yes** for both **Split Barcode** and **Auto Wash**. Tap **Next**. The auto-sliding screen moves up automatically.
- 8. Push the sequencing cartridge into the reagent compartment. The system will automatically scan the cartridge ID and display it in the **Reagent cartridge** box.
- Tips

If automatic scanning fails, you can manually input the ID.

- 9. Tap **Prime** and then **Yes** in the pop-up window. The system automatically closes the auto-sliding screen and starts priming.
- 10. Load the DNBs into the flow cell:
  - 1) Take out DNB Load Buffer II in advance and place it on ice for approximately 30 minutes. Mix it thoroughly by using a vortex mixer for 5 seconds, briefly centrifuge it, and place it on ice until use.
  - 2 Take out a new 0.2 mL tube and prepare the DNB loading mixture according to the table below.

Component	Volume (µL)
DNB Load Buffer II	7
Make DNB Enzyme Mix II (LC)	1
DNB	21
Total volume	29

- 3 Mix the DNB loading mixture slowly by using a wide-bore pipette tip for 5 to 8 times. Then, place it at a 4 °C refrigerator until use.
- Tips
  - Use the DNB loading mixture immediately after preparation.
  - Do not centrifuge, vortex, or pipette the DNB loading mixture vigorously.



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- 11. Take out the flow cell and check it. Pipette 10  $\mu$ L of DNB loading mixture by using a 100  $\mu$ L pipette. Gentle transfer the mixture into the inlet of the flow cell and press the release button on the pipette to discharge it. The mixture will automatically flow into the flow cell.
- Tips
  - Do not press the control button on the pipette.
  - During the DNB loading process, do not rotate the pipette tip or move the flow cell to avoid bubbles.
- 12. After the priming is completed, tap **Next**. The auto-sliding screen moves up automatically. Insert the flow cell into the flow cell compartment, and the system will automatically scan the ID.
- **Tips**

If automatic scanning fails, you can manually input the ID.

- 13. Tap **Next** to review the information.
- 14. After reviewing all information, tap **Sequence** and select **Yes** to start sequencing. The auto-sliding screen moves down and the Sequencing interface is displayed.
  - The interface shows the real-time sequencing progress, and you can view the status after refreshing the system.
- 15. After the sequencing is completed, tap **Finish**. The auto-sliding screen moves up. Take out the reagent cartridge and flow cell, and tap **Return home**.
- 16. After the sequencing is completed, the analysis software automatically starts the analysis.

### Viewing and downloading the report

- 1. Click the number under **Report Today** on the Home page. The Analysis Report page is displayed.
- 2. Set the query term in the **Query** area and click  $\mathbb{Q}$  to locate the sample.
- 3. Click the batch code under the **Batch Code** column to view project details.
- 4. Click in the **Result Path** column to open the result path.
- 5. Click **Result** to view and download the analysis reports.

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