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# Economical Nanopore LSK Sequencing: Adapted Protocols with Homebrew Reagents V.2

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**Protocol status:** Working

**We use this protocol and it's working**

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## Abstract

This protocol introduces a cost-effective alternative for end repair and dA-tailing in DNA library preparation, particularly tailored for PCR samples with an N50 of 3 kb. By employing a homebrew end repair reagent solution, this method replaces the use of the NEBNext® Ultra™ II End Repair/dA-Tailing Module recommended by Nanopore. Optimization of reagent quantities and a bead-free purification method are combined to achieve efficient adapter ligation while minimizing costs. Notably, an extended incubation time during adapter ligation enhances efficiency. This resource provides a strategic approach for researchers aiming to customize their sequencing workflows while achieving optimal results and substantial savings.

## Guidelines

- 1. Optimized Homebrew Approach:** This protocol features a unique homebrew end repair reagent solution as a cost-effective alternative to the Nanopore-recommended NEBNext® Ultra™ II End Repair/dA-Tailing Module. Adhere to the specified volumes meticulously to achieve successful results with your customized reagent.
- 2. Bead-Free Purification:** The protocol employs a bead-free PEG/NaCl precipitation method for efficient purification. Ensure precise centrifugation to maintain DNA pellet integrity and maximize recovery rates.
- 3. Strategic Ligation Enhancement:** An extended incubation period during adapter ligation enhances efficiency. Dedicate adequate time to this step to optimize adapter ligation results.

## Materials

### 1. 4X Quick Ligase Buffer

|   | A | B                                     | C                             |
|---|---|---------------------------------------|-------------------------------|
|   |   | A                                     | B                             |
| 1 |   | H <sub>2</sub> O                      | 70 ml                         |
| 2 |   | Tris-Base                             | 2.42 g<br>(200 mM)            |
| 3 |   | MgCl <sub>2</sub> · 6H <sub>2</sub> O | 0.8 g<br>(40 mM)              |
| 4 |   | DTT                                   | 0.6 g<br>(40 mM)              |
| 5 |   | PEG 6000                              | 40 g<br>(40% w/w)             |
| 6 |   | HCl                                   | Adjust pH to 7.6              |
| 7 |   | H <sub>2</sub> O                      | bring the volume up to 100 mL |

Add the materials in the following table sequentially

### 2. 33.5% (w/v) PEG 8000

### 3. NP (5M NaCl + 6.7% PEG8000)

|   | A | B        | C                |
|---|---|----------|------------------|
| 1 |   | PEG 8000 | 3.35 g<br>(6.7%) |
| 2 |   | NaCl     | 14.61 g<br>(5M)  |

### 4. 80% (v/v) ethanol

### 5. 99.5% ethanol

### 6. Elution buffer (EB, 1X TE, pH=8.0)

### 7. LFB wash buffer

|   | A | B                    | C     |
|---|---|----------------------|-------|
| 1 |   | 33.5% (w/v) PEG 8000 | 80 uL |

|  | A | B   | C      |
|--|---|-----|--------|
|  | 2 | NP  | 40 uL  |
|  | 3 | H2O | 320 uL |

## Safety warnings

- ❗ **1. Ethanol Flammability:** Ethanol is highly flammable. Use caution when handling and storing ethanol solutions. Work in well-ventilated areas away from open flames, sparks, or heat sources.

## Homebrew End Repair/dA-Tailing

50m

- 1 Add the materials in the following table sequentially.

| Materials              | Quantity                                  |
|------------------------|-------------------------------------------|
| DNA                    | 1600 ng<br>(for samples with N50 of 3 kb) |
| H2O                    | Bring up to a volume of 68 ul             |
| 4X Quick Ligase Buffer | 25 ul                                     |
| ATP (25 mM)            | 1 ul                                      |
| dNTP (10 mM)           | 5 ul                                      |
| Taq polymerase         | 0.5 ul (2.5 U)                            |
| T4 PNK                 | 0.5 ul (5 U)                              |

### Note

1. Due to the high concentration of PEG in the solution, it may be slightly viscous. Gently invert several times to ensure thorough and uniform mixing.

- 2 Incubate at  37 °C for  00:30:00

30m

### Note

1. T4 PNK will add phosphate to the 5' ends of both strands of DNA

- 3 Incubate at  65 °C for  00:20:00

20m

### Note

T4 PNK will be inactivated, and Taq polymerase will begin repairing DNA ends and adding A-tails.



## Purification

30m

- 4 For the subsequent purification, commercially available **spin-column** or **magnetic bead-based methods** can be used.

Here, to save costs, we will use the **PEG/NaCl precipitation method**.

- 5 Add  15  $\mu\text{L}$  of **NP**

### Note

1. Due to the high concentration of PEG in the solution, it may be slightly viscous. Gently invert several times to ensure thorough and uniform mixing.

- 6 Centrifuge at maximum speed (at least 13000 rpm) for  00:15:00 .

15m

- 7 Carefully remove the supernatant using a pipette.

### Note

Most of the time, the pellet is not visible to the naked eye, so pay attention to the orientation during centrifugation

- 8 Add  200  $\mu\text{L}$  of 80% ethanol.

### Note

Sometimes the pellet becomes more visible after adding ethanol.

- 9 Centrifuge at maximum speed (at least 8000 rpm) for  00:02:00

2m

- 10 Carefully remove the supernatant using a pipette.



11 Add  200  $\mu\text{L}$  of 80% ethanol.

12 Centrifuge at maximum speed (at least 8000 rpm) for 2 minutes.

2m

13 Carefully remove the supernatant using a pipette.

14 Wait for approximately  00:10:00 to ensure complete ethanol evaporation.

10m

#### Note

Prevent excessive drying of the DNA pellet. Prolonged periods can result in DNA breakage.

15 Add  25  $\mu\text{L}$  of **elution buffer**.

#### Note

Optional: Measure the DNA concentration.

Normally, the DNA concentration **should be >30 ng/ $\mu\text{L}$** .

## Adapter ligation

30m

16 **When using LNB (with ATP) provided by Nanopore**

|  | Reagent               | Volume                                      |
|--|-----------------------|---------------------------------------------|
|  | DNA                   | 400 ng (for samples with N50 of 3 kb)       |
|  | H <sub>2</sub> O      | Bring up to a volume of 15.25 $\mu\text{l}$ |
|  | Ligation Buffer (LNB) | 6.25 $\mu\text{l}$                          |
|  | T4 Ligase             | 2.5 $\mu\text{l}$                           |
|  | Ligation Adapter      | 1 $\mu\text{l}$                             |

### When using homemade 4X Quick Ligase Buffer (without ATP)

| A                      | B                                     |
|------------------------|---------------------------------------|
| Reagent                | Volume                                |
| DNA                    | 400 ng (for samples with N50 of 3 kb) |
| H2O                    | Bring up to a volume of 14.25 µl      |
| 4X Quick Ligase Buffer | 6.25 µl                               |
| ATP (25 mM)            | 1 ul                                  |
| T4 Ligase              | 2.5 µl                                |
| Ligation Adapter       | 1 µl                                  |

#### Note

Note that the reagent amounts used in this step are about **1/4** of the manufacturer's recommended volume. This means that the ligation sequencing kit, which was originally designed for 6 uses, **can now be utilized for 24 uses**. This adjustment is primarily due to our utilization of a **bead-free purification** method later on, which significantly enhances the recovery rate.

- 17 Incubate at room temperature for  01:00:00 .

1h

#### Note

Despite the official recommendation of a 5-minute reaction, based on our own experience, **a 60-minute reaction** significantly improves adapter ligation efficiency.

- 18 Add  3 µL of **NP**



**Note**

1. Due to the high concentration of PEG in the solution, it may be slightly viscous. Gently invert several times to ensure thorough and uniform mixing.  
2. Normally, PEG and NaCl buffer is required for DNA precipitation. However, it's evident that LNB buffer already contains a high concentration of PEG, so adding NP alone is sufficient for DNA precipitation. For more information, refer to:  
<https://dx.doi.org/10.17504/protocols.io.7erhjd6>.

- 19 Centrifuge at maximum speed (at least 13000 rpm) for at least  00:15:00 (If a higher recovery rate is needed, centrifuging for up to one hour is also possible).

15m

**Note**

To prevent overheating, it is recommended to use a refrigerated centrifuge. If a refrigerated centrifuge is unavailable, it is advised to perform centrifugation in two steps of 10 minutes each, with a 10-minute interval in between.

- 20 Carefully remove the supernatant using a pipette.

**Note**

Most of the time, the pellet is not visible to the naked eye, so pay attention to the orientation during centrifugation

- 21 Add 200  $\mu$ L of **LFB wash buffer**

**Note**

Note that in this step, use the **LFB wash buffer** instead of alcohol. This is because the DNA now has motor proteins attached, and using alcohol could disrupt the motor proteins.

- 22 Centrifuge at maximum speed (at least 8000 rpm) for  00:02:00

2m

- 23 Carefully remove the supernatant using a pipette.



24 Repeat the above washing step twice.

25 Add  20  $\mu\text{L}$  of **EB buffer** (provided by the Ligation Sequencing Kit).

#### Note

Optional: Measure the DNA concentration.  
Normally, the DNA concentration **should be**  $>15 \text{ ng}/\mu\text{L}$ .

## Protocol references

<https://www.protocols.io/view/bead-free-long-fragment-lsk109-library-preparation-rm7vz8mj5vx1/v1>

<https://www.protocols.io/view/size-selective-precipitation-of-dna-using-peg-amp-j8nlkeb8dl5r/v1>