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We use this protocol and it's working

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Daughter of Frankenstein protocol for nuclei isolation from fresh and frozen tissues using OptiPrep® *continuous* gradient V.2

In 1 collection

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ABSTRACT

This protocol is the result of the combination of various nuclei isolation protocols for single cell RNA-seq experiments using droplet-based methods, and is an extension of the Frankenstein (D.O.F means Daughter Of Frankenstein).

Developed to prepare nuclei isolates from fresh and frozen material of small-to-large sizes. The good thing is that it does not use FACS but OptiPrep® *continuous* gradient to remove debris. It is the alternative protocol when FACS is not available.

This protocol works better with smaller samples. After centrifugation the nuclei will form a pellet and contaminating membranes will float at the top. This continuous gradient can be applied for as many times as needed to the same prep to continue cleaning up your nuclei suspensions.

GUIDELINES

Use RNA techniques.

The use of RNase Inhibitor is HIGHLY recommended.

MATERIALS

Required Buffers and Reagents

1. Nuclei EZ lysis buffer **Merck MilliporeSigma (Sigma-Aldrich) Catalog #EZ PREP NUC-101** (keep at 4°C)
2. Nuclei Wash and Resuspension Buffer (prepare fresh, keep at 4°C)

1x PBS

1.0% BSA (MACS® BSA Stock Solution, Miltenyi)

0.2-0.5 U/uL RNase Inhibitor (Protector RNA Inhibitor, Millipore Sigma)

3. Gradient Solutions

G60: OptiPrep®: 60% (w/v) solution of iodixanol in water.

GD: 150 mM KCl, 30 mM MgCl₂, 120 mM, Tricine-KOH pH 7.8.

GW: 50% of iodixanol --> mix 5 volumes of G60 with 1 volume of GD (final: 50% iodixanol, 25 mM KCl, 5 mM MgCl₂, 20 mM Tricine-KOH pH 7.8).

GH: 0.25 M Sucrose, 25 mM KCl, 5 mM MgCl₂, 20 mM Tricine-KOH pH 7.8. (keep at 4°C).

G30: solutions of 30% --> mix 6 volumes of GW with 4 volumes of GH (final: 30% iodixanol, 25 mM KCl, 5 mM MgCl₂, 20 mM Tricine-KOH pH 7.8). **Prepare fresh.**

4. Stock solutions and GD/GH Set-Up

-500 mM Tricine: 8.96 g in 100 ml water

-1 M KCl: 7.45 g in 100 ml water

-1 M MgCl₂·6H₂O: 20.3 g in 100 ml water

GD: To 50 ml water add 24 ml, 15 ml and 3 ml respectively of the Tricine, KCl and MgCl₂·6H₂O stock solutions (above); adjust to pH 7.8 with 1 M KOH and make up to 100 ml. Keep at RT.

GH: Dissolve 8.5 g of sucrose in 50 ml of water; add 4 ml, 2.5 ml and 0.5 ml respectively of the Tricine, KCl and MgCl₂·6H₂O stock solutions (above); adjust to pH 7.8 with 1 M KOH and make up to 100 ml. Keep at 4°C.

SAFETY WARNINGS

See SDS (Safety Data Sheet) for safety warnings and hazards.

BEFORE START INSTRUCTIONS

All samples and reagents are kept on ice or at 4 °C (wet ice).

Prepare all buffers and reagents as described in the "Materials" section.

Tissue Homogenization

- 1 Mince/chop tissue with a razor blade to small pieces. The tissue may be as small as a grain of rice.

Note

For mincing the tissue, you may take the tube out of ice, however, be quick and return it to ice.

- 2 Add $300\ \mu\text{L}$ of chilled Nuclei EZ Lysis Buffer (supplemented with RNase Inhibitor 0.2-0.5 U/uL) to the tissue in 1.5 mL tube.
- 3 **Gently** homogenize the sample using a douncer by stroking 10-20 times. Keep nuclei suspension on ice at all times.
- 4 Add an extra $700\ \mu\text{L}$ of chilled Nuclei EZ Lysis Buffer (supplemented with RNase Inhibitor 0.2-0.5 U/uL), mix gently by pipetting using **wide-bore tips** and incubate on ice for 00:05:00. Repeat mixing 2-3 times during the incubation.

Nuclei Isolation

10m

- 5 Filter homogenate using a 70 μm -strainer mesh to fit a 15 ml Falcon tube (e.g. pluriStrainer Mini 70 μm Cell Strainer).
- 6 Transfer flow through into a 1.5 mL LoBind tube and centrifuge the nuclei at 500 x g for 00:05:00 at 4 °C and remove supernatant leaving behind ~ $50\ \mu\text{L}$.

Homogenate after filtering.

Nuclei pellet.

- 7 **Optional:** add 1 mL of EZ Lysis buffer (supplemented with RNase Inhibitor 0.2-0.5 U/uL), gently resuspend pellet (optional: incubate for $00:05:00$ on ice for additional lysis). Then centrifuge the nuclei at $500 \times g$ for $00:05:00$ at 4°C .

10m

Homogenate after re-pelleting.

- 8 Remove supernatant as much as possible without disturbing pellet (if pellet looks loose leave $\sim 50 \mu\text{L}$ behind) and add $300 \mu\text{L}$ of G30 buffer. Mix well but gently to resuspend the pellet.
- 9 Carefully, **underlay** the sample with 1 mL of the G30. A clear separation between homogenate and added G30 will form. Do not disturb.

Homogenate after resuspending and underlying G30. Arrow indicates the separation of both layers.

- 10 Centrifuge the nuclei at $8000 \times g$ for 00:20:00 at 4°C . After centrifugation the nuclei will form a pellet and contaminating membranes will float at the top. This continuous gradient can be applied for as many times as needed to the same prep to continue cleaning up your nuclei suspensions (Optional Step 11).

20m

Clean pelleted nuclei.

- 11 Optional: repeat steps 8-10) for additional cleaning.

Nuclei Wash and Resuspension

10m

12

5m

After centrifugation, remove supernatant and add 1 mL of ice-cold Nuclei Wash and Resuspension Buffer supplemented (supplemented with RNase Inhibitor 0.2-0.5 U/uL) **without resuspending the nuclei**. Then centrifuge the nuclei at $500 \times g$ for $00:05:00$ at 4°C .

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5m

Remove supernatant as much as possible without disturbing pellet (if pellet looks loose leave $\sim 50 \mu\text{L}$ behind) and resuspend the pellet in 1 mL of ice-cold Nuclei Wash and Resuspension Buffer supplemented (supplemented with RNase Inhibitor 0.2-0.5 U/uL). Then centrifuge the nuclei at $500 \times g$, $00:05:00$ for $00:05:00$ at 4°C .

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Remove supernatant and resuspend with $300 \mu\text{L}$ of ice-cold Nuclei Wash and Resuspension Buffer (supplemented with RNase Inhibitor 0.2-0.5 U/uL) and filter using 40-um Flowmi® Cell Strainer and transfer to a new LoBind 1.5 mL Eppendorf tube.

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Check integrity and purity under microscope and count manually or using an automatic counter. For automatic counter I recommend LUNA-FL™ Dual Fluorescence Cell Counter and Acridine Orange/Propidium Iodide (AO/PI) Cell Viability Kit (F23001).

DAPI-stained nuclei post-clean up.

BioA traces of total RNA from HEK293 cell line and RNA extracted from a representative nuclei suspension; in this case liver as an example. This sample was successfully run on the 10x Genomics Chromium using the 3' v3.1 Dual Index kit.

16 Proceed to your amazing snRNA-Seq experiment!