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Protocol status: Working
 We use this protocol and it's working

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🌐 S.O.F protocol for nuclei isolation from fresh and frozen tissues using OptiPrep® discontinuous gradient V.3

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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 (S.O.F means Son 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® discontinuous gradient to remove debris. It is the alternative protocol when FACS is not available.

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46581

Keywords: snRNAseq, 10x,
nuclei, isolation, FACS

ATTACHMENTS

[___Frankenstein___proto
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ue_FINAL.pdf](#)

GUIDELINES

Use RNA techniques.

The use of RNase Inhibitor is HIGHLY recommended.

MATERIALS

Required Buffers and Reagents

1. Nuclei EZ lysis buffer **Sigma Catalog #EZ PREP NUC-101** (keep 4°C)
2. Nuclei wash and resuspension buffer (prepare fresh, keep 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

G1: OptiPrep®: 60% (w/v) solution of iodixanol in water.

G2: 150 mM KCl, 30 mM MgCl₂, 120 mM, Tricine-KOH pH 7.8

G3: solution containing 50% (w/v) of iodixanol --> mix 5 volumes of G1 with 1 volume of solution G2 (final: 50% iodixanol, 25 mM KCl, 5 mM MgCl₂, 20 mM Tricine-KOH pH 7.8).

G4: 0.25 M Sucrose, 25 mM KCl, 5 mM MgCl₂, 20 mM Tricine-KOH pH 7.8.

G5: solution containing 25% (w/v) of iodixanol --> mix equal volumes of G3 and G4 (final: 25% iodixanol, 25 mM KCl, 5 mM MgCl₂, 20 mM Tricine-KOH pH 7.8).

G6: solutions of 30% --> mix 6 volumes of G3 with 4 volumes of G4 (final: 30% iodixanol, 25 mM KCl, 5 mM MgCl₂, 20 mM Tricine-KOH pH 7.8).

G7: solutions of 35% --> mix 7 volumes of G3 with 3 volumes of G4 (final: 35% iodixanol, 25 mM KCl, 5 mM MgCl₂, 20 mM Tricine-KOH pH 7.8).

4. Stock solutions and G2/G4 Set-Up

Keep the following Stock Solutions at 4°C:

-500 mM Tricine: 8.96 g in 100 ml water

-1 M KCl: 7.45 g in 100 ml water

-1 M $\text{MgCl}_2 \cdot 6\text{H}_2\text{O}$: 20.3 g in 100 ml water

G2: To 50 ml water add 24 ml, 15 ml and 3 ml respectively of the Tricine, KCl and $\text{MgCl}_2 \cdot 6\text{H}_2\text{O}$ stock solutions (above); adjust to pH 7.8 with 1 M KOH and make up to 100 ml.

G4: 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 $\text{MgCl}_2 \cdot 6\text{H}_2\text{O}$ stock solutions (above); adjust to pH 7.8 with 1 M KOH and make up to 100 ml.

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 to ice.

- 2 Add 300 μL of chilled Nuclei EZ Lysis Buffer (supplemented with RNase Inhibitor 0.2-0.5 U/ μL) to the tissue in 1.5 mL tube.
- 3 **Gently** homogenize the sample using a douncer by stroking 10-20 times. Keep tube on ice at all times.

Nuclei Isolation

10m

- 4 Add an extra 900 μL of chilled Nuclei EZ Lysis Buffer (supplemented with RNase Inhibitor 0.2-0.5 U/ μL), mix gently by pipetting using **wide-bore tips** and incubate on ice for 00:05:00 . Repeat mixing 2-3 times during the incubation. RNase inhibitor will now be in the range of 0.2-0.5 U/ μL .
- 5 Filter homogenate using a 70 μm -strainer mesh to fit a 15 ml Falcon tube (e.g. pluriStrainer Mini 70 μm Cell Strainer).

Equipment

pluriStrainer Mini 70 μm

NAME

Cell Strainer

TYPE

pluriSelect

BRAND

43-10070-40

SKU

<https://www.pluriselect.com/us/pluristrainer-mini-70-m-cell-strainer.html#size=27>

LINK

- 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 $^{\circ}\text{C}$ and remove supernatant leaving behind ~ 50 μL .

- 7 **Optional:** add 1 mL of EZ Lysis buffer (supplemented with RNase Inhibitor 0.2-0.5 U/ μL),

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gently resuspend pellet and incubate for 00:05:00 on ice. Then centrifuge the nuclei at 500 x g for 00:05:00 at 4 °C .

- 8 Remove supernatant as much as possible without disturbing pellet (if pellet looks loose leave ~ 50 µL behind) and add 1 mL of Nuclei Wash and Resuspension Buffer (supplemented with RNase Inhibitor 0.2-0.5 U/uL) **without resuspending**.
- 9 Centrifuge the nuclei at 500 x g for 00:05:00 at 4 °C .
- 10 Optional: Remove supernatant as much as possible without disturbing pellet (if pellet looks loose leave ~ 50 µL behind) and add 500 µL of Nuclei Wash and Resuspension Buffer (supplemented with RNase Inhibitor 0.2-0.5 U/uL) **without resuspending**. Centrifuge the nuclei at 500 x g for 00:05:00 at 4 °C .
- 11 Remove supernatant as much as possible without disturbing pellet (if pellet looks loose leave ~ 50 µL behind) and resuspend the pellet in 1 mL of G5 buffer and transfer onto a 10 mL LoBind centrifuge tube. Then add another mL of G5 (final volume 2 mL).

Debris Removal

- 12 Carefully, underlayer the sample with 2 mL of the 30% iodixanol and then 2 mL of the 35% iodixanol, as shown in the picture below.

Application Sheet S10."/>

Layering the gradient. Image adapted from *OptiPrep™ Application Sheet S10*.

- 13** Centrifuge the nuclei at 5000 x g for 00:20:00 at 4 °C using swinging bucket centrifuge with break set to OFF. (Note: using 10000 x g is also possible).

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Application Sheet S10."/>

Position on nuclei pellet. Image adapted from *OptiPrep™ Application Sheet S10*.

- 14** After centrifugation, a white-ish band of nuclei between the 30%-35% iodixanol interface should be visible. Aspirate the top layers down until the white nuclei band at the interphase of 29%-35%. Using 200 uL tip, collect the nuclei band and transfer to a fresh 1.5 mL LoBind tube.
- 15** Then top up to ~1.3 mL with Wash and Resuspension Buffer (supplemented with RNase Inhibitor 0.2-0.5 U/uL), mix well but gently, and centrifuge the nuclei at 500 x g for 00:05:00 at 4 °C.
- 16** Remove supernatant and repeat once more for a total of 2 washes.
- 17** 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).

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18 Proceed to your amazing snRNA-Seq experiment!