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③ Preparation of Nuclei Suspension from Human Musculoskeletal Tissues

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Protocol status: Working

We use this protocol and it's working

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Abstract

An adapted version of the 'Frankenstein' protocol (Luciano G Martelotto 2020. 'Frankenstein' protocol for nuclei isolation from fresh and frozen tissue for snRNAseg. protocols.io https://dx.doi.org/10.17504/protocols.io.bgxymxpw) to prepare single nuclei suspensions from freshly frozen human musculoskeletal tissues before nuclei sorting by flow cytometry.

Protocol described for one sample.

This protocol has been used in the following available studies:

PREPRINT: Distinct myofibre domains of the human myotendinous junction revealed by single nucleus RNA**seq.** Anders Karlsen et al. https://doi.org/10.1101/2022.12.16.519020

Guidelines

- Avoid freeze-thawing of tissues
- Keep materials and reagents chilled
- Keep work space RNAse free
- All pipetting steps are performed on ice



Materials

Reagents

- PBS, sterile
- Penicillin-Streptomycin (5,000 U/mL) (15070063, Thermo Fisher Scientific)
- Nuclei Isolation Kit: Nuclei EZ Prep (NUC101-1KT, Sigma-Aldrich)
- PBS without magnesium
- Phosphate Buffered Saline with 10% Bovine Albumin (SRE0036-250ML, Sigma-Aldrich)
- 10x PBS (AM9625, ThermoFisher Scientific)
- 1 M MgCl₂ (M1028-10X1ML, Sigma)
- Protector RNase Inhibitor (03335399001, Roche)
- Trypan Blue Solution, 0.4% (15250061, Thermo Fisher Scientific)

Materials

- Scalpel
- 50 ml tube
- 15 ml tube
- Cryotube
- 2 ml screw cap tubes
- 2 ml microcentrifuge tubes
- 1.5 ml microcentrifuge tube
- 2.3 mm diameter stainless steel (11079123ss, BioSpec Products)
- 1.0 mm silicon carbide beads (11079110sc, BioSpec Products)
- pluriStrainer Mini 70 μm (Cell Strainer) (43-10070-40, PluriSelect)
- pluriStrainer Mini 40 μm (Cell Strainer) (43-10040-40, PluriSelect)

Equipment

- Liquid nitrogen
- Freezer cabinet or dry ice
- MP Biomedicals FastPrep-24
- Microcentrifuge with a cooling function
- Centrifuge that can hold 15 ml tubes and has a cooling function
- Neubauer Improved haemocytometer

Protocol materials

X Nuclei EZ lysis buffer	Merck MilliporeSigma (Sigma-Aldrich) Catalog #EZ PREP NUC-101
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Troubleshooting

Before start

- Pre-cool centrifuges
- Obtain ice
- Label and pre-chill tubes on ice



Tissue Preparation

- 2 Using a scalpel, dissect the tissue as desired \(\bigsim \) On ice.

Note

E.g. For hamstring tendons, use a scalpel to scrape off any muscle fibres. For muscle tissues, cut away tendon tissue.

Cut up tissue into small pieces of around 100 μm^3 , remove excess PBS from tissue pieces and snap freeze in cryotubes in liquid nitrogen. Store samples at \$\$\$ -80 $^{\circ}$ C.

Note

Each tissue sample may be divided into multiple small pieces, each frozen separately.

In a freezer cabinet or on dry ice, cut each piece of frozen tissue into smaller pieces using a scalpel. Transfer ~ 50 mg tissue into a 2 ml screw cap tube containing homogenising beads.

Note

E.g. For tendon tissue use five stainless steel balls of 2.3 mm. For muscle tissue use five stainless steel balls of 2.3 mm and an additional 1.0 mm silicon carbide bead.

Nuclei Wash and Suspension Buffer

Freshly prepare Nuclei Wash and Suspension Buffer (2% bovine serum albumin in PBS without magnesium, 2 mM MgCl₂, 0.2 U/I RNA inhibitor). Keep On ice



Note

For each sample prepare 5 ml Nuclei Wash and Suspension Buffer.

Tissue Homogenisation

- 7 Pipette 4 1 mL chilled
 - Nuclei EZ lysis buffer Merck MilliporeSigma (Sigma-Aldrich) Catalog #EZ PREP NUC-101

to each 2 ml screw cap tube containing ~50 mg cut up frozen tissue and homogenising beads.

- Homogenise the tissues at 4.0 M/S for 00:00:20 in a FastPrep 24. Immediately after, transfer the tubes and incubate On ice for 5 min.
- Repeat homogenisation step. Immediately after, transfer the tubes and incubate

 On ice for 5 to 15 min for the bubbles to settle.

Note

The tissue may not be completely homogenised. Extra homogenisation steps could lead to the sample overheating. This step will need to be optimised for other tissue types.

- Transfer the homogenate from each tube to a pre-chilled 2 ml microcentrifuge tube and add Δ 500 μ L chilled
 - Nuclei EZ lysis buffer Merck MilliporeSigma (Sigma-Aldrich) Catalog #EZ PREP NUC-101

to each tube. Mix gently using a wide bore 1000 µl pipette. Incubate the tubes

On ice and gently mix two more times using a wide bore 1000 µl pipette to help release more nuclei from the remaining tissue.

- - Nuclei EZ lysis buffer Merck MilliporeSigma (Sigma-Aldrich) Catalog #EZ PREP NUC-101
 - . Centrifuge at \$\infty\$ 500 x g, 4°C, 00:05:00 \cdot\$.

5m

20s



12 Remove as much supernatant as possible without disturbing the pellet. Pipette

5m

5m

□ 1.5 mL chilled

Nuclei EZ lysis buffer Merck MilliporeSigma (Sigma-Aldrich) Catalog #EZ PREP NUC-101

to the tube. Using a wide bore 1000 μ l pipette gently resuspend the pellet by pipetting up and down 10 times. Transfer the suspension to a pre-chilled 2 ml microcentrifuge tube. Centrifuge at $\$500 \times 9$, 4° C, 00:05:00.

- Remove as much supernatant as possible without disturbing the pellet. Pipette 500 μ l chilled Nuclei Wash and Suspension Buffer **without disturbing the pellet**. Incubate On ice for 5 min
- Add Δ 1 mL Nuclei Wash and Suspension Buffer. Resuspend the pellet using an uncut 1000 μl pipette tip, gently pipetting up and down 20 times.
- Place a 40 μm cell strainer over a pre-chilled 2 ml microcentrifuge tube. Filter the nuclei suspension. After, wash the cell strainer with Δ 0.5 mL chilled Nuclei Wash and Suspension Buffer. Centrifuge 500 x g, 4°C, 00:05:00 and resuspend the pellet in the desired volume of Nuclei Wash and Suspension Buffer. Keep samples on ice.

Nuclei Visualisation and Counting

Mix a small volume nuclei suspension with Trypan blue. E.g. 10 μl of each. Visualise on a Neubauer Improved Haemocytometer.

Note

Nuclei will be stained blue. They can be distinguished from debris by their clear outlines and bean shape.

Nuclei purification

To purify the nuclei suspension from tissue debris, e.g., collagen extracellular matrix in tendon samples, samples may be further purified by fluorescence-activated nuclei sorting or by ultracentrifugation using an iodixanol gradient (DOI: 10.1126/sciadv.abn836).

