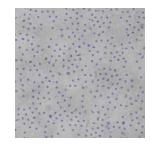


Nov 21, 2023

Nuclei extraction for 10x Genomics Single Cell Multiome ATAC + Gene Expression from frozen tissue using Singulator™ 100 or 200 (S2 Genomics) V2.0





DOI

dx.doi.org/10.17504/protocols.io.j8nlkkmq6l5r/v1

Ignas Masilionis¹, ojasvi chaudhary², Brigita Meskauskaite Urben¹, Ronan Chaligne¹

¹Single-Cell Analytics Innovation Lab, Memorial Sloan Kettering Cancer Center;

²Memorial Sloan Kettering Cancer Center



Ignas Masilionis

Memorial Sloan Kettering Cancer Center

Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account





DOI: https://dx.doi.org/10.17504/protocols.io.j8nlkkmq6l5r/v1

Protocol Citation: Ignas Masilionis, ojasvi chaudhary, Brigita Meskauskaite Urben, Ronan Chaligne 2023. Nuclei extraction for 10x Genomics Single Cell Multiome ATAC + Gene Expression from frozen tissue using Singulator™ 100 or 200 (S2 Genomics) V2.0. **protocols.io** <u>https://dx.doi.org/10.17504/protocols.io.j8nlkkmq6l5r/v1</u>



License: This is an open access protocol distributed under the terms of the **Creative Commons Attribution License**, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

Protocol status: Working

We use this protocol and it's working

Created: May 06, 2022

Last Modified: November 21, 2023

Protocol Integer ID: 62125

Keywords: Single Cell, 10x Genomics, Nuclei, Multiome, Singulator, S2 Genomics, snRNAseq, snATACseq, Single Nuclei, 10x genomics single cell multiome atac, suitable for single cell multiome atac, single cell multiome atac, extraction from snap frozen tissue, gene expression from frozen tissue, nuclei extraction, snap frozen tissue, nuclei fac, washing nuclei suspension, resulting nuclei suspension, nuclei suspension in buffer, cancer snap, frozen tissue, gene expression, s2 genomic, mouse tissue, single cell, core needle biopsy, gene

Abstract

This protocol describes nuclei extraction from snap frozen tissue using Singulator, including washing nuclei suspension in buffer supplemented with sucrose, and nuclei FACS sorting. The resulting nuclei suspension is suitable for Single Cell Multiome ATAC + Gene Expression using 10x Chromium platform. The protocol has been validated on various healthy and cancer snap frozen human and mouse tissues (lung, brain, prostate, core needle biopsies etc).

Guidelines

Make sure to use Swinging bucket centrifuges to spin nuclei.



Materials

MATERIALS

- DTT Merck MilliporeSigma (Sigma-Aldrich) Catalog #D0632
- \$\times 4,6-Diamidino-2-Phenylindole, Dihydrochloride (DAPI) Thermo Fisher Scientific Catalog #D1306
- Trizma hydrochloride solution Merck MilliporeSigma (Sigma-Aldrich) Catalog #T2194-100ML
- 🔯 Sodium Chloride Solution 5 M Merck MilliporeSigma (Sigma-Aldrich) Catalog #59222C
- Magnesium chloride solution for molecular biology (1.00 M) Merck MilliporeSigma (Sigma-Aldrich) Catalog #M1028
- Digitonin (5%) Thermo Fisher Catalog #BN2006
- Albumin, Bovine Serum, 10% Aqueous Solution, Nuclease-Free Merck MilliporeSigma (Sigma-Aldrich) Catalog #126615-25ML
- X Protector RNase Inhibitor Merck MilliporeSigma (Sigma-Aldrich) Catalog #3335402001

Protocol materials

- Digitonin (5%) Thermo Fisher Catalog #BN2006
- 🔯 Protector RNase Inhibitor Merck MilliporeSigma (Sigma-Aldrich) Catalog #3335402001
- DTT Merck MilliporeSigma (Sigma-Aldrich) Catalog #D0632
- X Trizma hydrochloride solution Merck MilliporeSigma (Sigma-Aldrich) Catalog #T2194-100ML
- 🔯 Sodium Chloride Solution 5 M Merck MilliporeSigma (Sigma-Aldrich) Catalog #59222C
- Magnesium chloride solution for molecular biology (1.00 M) Merck MilliporeSigma (Sigma-Aldrich) Catalog #M1028
- Albumin, Bovine Serum, 10% Aqueous Solution, Nuclease-Free Merck MilliporeSigma (Sigma-
- 🔯 7-AAD (7-Aminoactinomycin D) **Thermo Fisher Catalog** #A1310
- Protector RNase Inhibitor Merck MilliporeSigma (Sigma-Aldrich) Catalog #3335402001
- 🔯 7-AAD (7-Aminoactinomycin D) **Thermo Fisher Catalog #**A1310

Troubleshooting



Preparation

15m

1

10m

Note

IMPORTANT

AFTER NUCLEI ISOLATION, YOU STILL NEED TO PERMEABILIZE THEM (step. 14) BEFORE TRANSPOSITION

- Turn on and pre-cool Singulator 👏 00:10:00
- Make sure there is sufficient number of Nuclei cartridges (or NIC+ cartridges for samples <20mg) stored / pre-cooled at 4 °C
- Take the tissue out of LN2 or -80C freezer and place it on dry ice to keep it frozen Ideally, piece should be <100mg (if not cut it in smaller pieces: <100mg and >10mg).

10m

Cutting can be done with razor blade and within a petri dish on dry ice. Hold the tissue with cooled and RNAse free tweezers (be careful it might fly off). Work fast to not let the tissue thaw while cutting. Place tissue back in a tube on dry ice immediately after cutting.

3 Prepare 2mL of Nuclei Wash Buffer per sample:

5m

```
Tris-HCI (pH 7.4): [M] 10 millimolar (mM) final
```

NaCl: [M] 10 millimolar (mM) final

MgCl₂: [M] 3 millimolar (mM) final

Purified BSA: 1% final

DTT: [M] 1 millimolar (mM) | final

0.2 - 1.0 U/ul RNAse inhibitor (depending on tissue type, see RNase Activity in Mouse Tissue below)

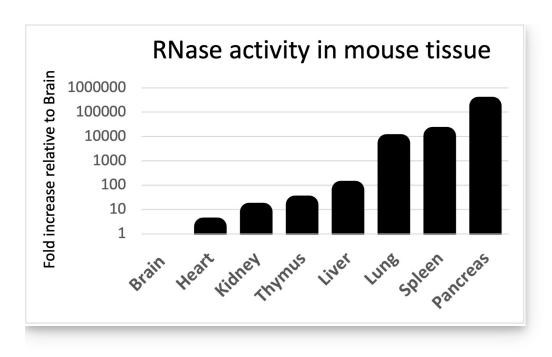
Tissue below) (

Protector RNase Inhibitor Merck MilliporeSigma (Sigma-Aldrich) Catalog #3335402001

)

Top-off to 2mL with Ambion Nuclease-free water

https://www.thermofisher.com/us/en/home/references/ambion-tech-support/nuclease-enzymes/tech-notes/rnase-activity-in-mouse-tissue.html



Singulator

10m

4 Add the following directly into Nuclei Isolation Cartridge before adding your tissue:

10m

RNAse inhibitor (0.2-1.0 U/uL final depending on tissue type - Final volume \sim 3.5 mL for standard nuclei isolation protocol)

DTT, 1mM final concentration (3.5uL of IM) 1 Molarity (M) DTT for standard nuclei isolation protocol).

Transfer tissue into the cooled Nuclei Isolation Cartridge and start standard nuclei isolation protocol.

Washes and Prep for FACS



When run is over on Singulator, transfer nuclei suspension (~3.5mL for standard protocol) into 5mL Protein LoBind eppendorf Tubes (or split 3.5mL evenly into four 1.5mL Protein LoBind eppendorf Tubes, conical bottom of the 1.5mL and 5mL tubes increase nuclei recovery).

5m



6 Add 500uL [M] 2 Molarity (M) Sucrose solution to a tube ([M] 250 millimolar (mM)

final conc). Adjust volume if using 1.5mL tubes.

Invert the tube slowly at least 10 times to mix sucrose and nuclei suspension (Do not vortex!).

Make sure sucrose is well mixed, if sucrose cushion is visible after centrifugation, mix again and repeat centrifugation.

7

\$\infty\$ 500 x q, 4°C, 00:05:00 , Swinging bucket centrifuge

5m

Nuclei Washes and FACS



8 After centrifugation, a layer of debris might be visible at the top and a pellet of nuclei at the bottom (for lower yields it might not be noticeable).

2m

Aspirate supernatant with a P1000 pipette, be really careful when removing the debris layer on the meniscus without disrupting nuclei pellet at the bottom of the tube (some liquid can be left to prevent pellet aspiration ~50uL).

1m

9 Re-suspend Nuclei pellet in ~400uL Nuclei Wash Buffer. This can be adjusted based on tissue input and pellet size.

9.1 OPTIONAL:

A second wash with Nuclei Wash buffer can be performed to remove traces of sucrose (as it might impact FACS sorting).

A second wash with 250mM sucrose can be performed if high amount of debris or fat is still present in the sample.

10

2m

Filter nuclei suspension through a 35 µm FACS tube cap filter (blue) or 30 µm Preseparation filters (Miltenyi).

11 Count nuclei using DAPI on automated cell counter (Countess II for instance) Trypan can also be used to count in Bright field, but in our experience DAPI count is more accurate.

5m

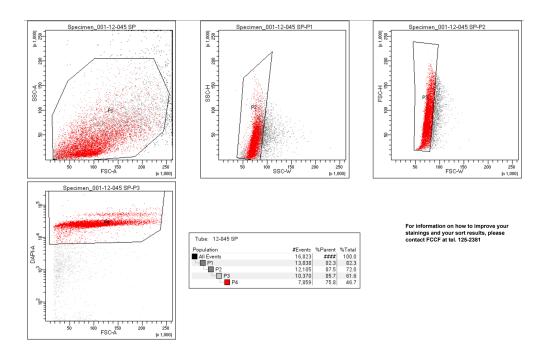
12 Nuclei should be stained with

30m

Positive nuclei population can be FACS into 500uL of Nuclei wash buffer using 1.5mL Lobind Eppendorf as collection tube (see attached PDF for gating setup example). Suggested nozzle size: 100um.

Ideally, aim to collect >200 000 nuclei.





10x Genomics

35m

13 Concentrate the nuclei suspension post-sorting:



€ 500 x g, 4°C, 00:05:00 , Swinging bucket centrifuge

Then, aiming for >=1000 nuclei/uL using FACS numbers, re-suspend sorted nuclei with nuclei Wash buffer .

Count nuclei using DAPI on Countess II cell counter.

Trypan can also be used to count in Bright field, but DAPI counts are more accurate.

14 Aliquot 200k nuclei in 0.2mL PCR tube.



₱ 500 x g, 4°C, 00:05:00 , Swinging bucket centrifuge

Remove supernatant leaving 5uL precisely (calculate: (Aliquot volume minus 5uL) and aspirate that amount, eyeballing it is highly discouraged).

Then continue to 10X Genomics Nuclei Isolation for Single Cell Multiome ATAC + Gene Expression Sequencing protocol <u>CG000365</u>. Following "Low Cell Input Nuclei Isolation" in Appendix starting at step e., omit NP40 from lysis buffer and perform lysis for 30s.

