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Nuclei Isolation for 10X Multiome Sequencing

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ABSTRACT

This protocol is for isolating nuclei for 10X multiome sequencing.

GUIDELINES

Keep tissue/nuclei on ice as much as possible.

OPEN ACCESS

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

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Prepare Stock Solutions

- 1 Make 20 mL **10% BSA** by combining 2 mL of BSA with 18 mL of MilliQ water in a 50 mL falcon

tube. (4°C - 2 weeks)

- 2 Make 20 mL **10% Triton X-100** by combining 18 mL MilliQ water with 2 mL Triton X-100 in a 50 mL tube. Vortex and then incubate at room temperature for 20 minutes. Filter it through a 0.22 µm filter with a syringe into a clean 50 mL tube. (4°C - 1 month)
- 3 Make 250 mL **Nuclear Isolation Media** by filling a 250 mL bottle with 200 mL of MilliQ water and then adding 2.5 mL 1M Tris, 6.26 mL 1M KCl, 1.25 mL 1M MgCl₂, and 21.45 g Sucrose. Shake until sucrose is dissolved then fill to 250 mL with MilliQ water. (4°C - 2 weeks)

Prepare Fresh Solutions

- 4 Make 3 mL **Homogenization Buffer** per sample by adding 2.895 mL Nuclear Isolation Media (filtered via syringe) to a 5 mL eppendorf. Then add 3 µL 100 mM DTT and 30 µL 10% Triton X-100. Add 15 µL RNasin and invert to mix. Store on ice.
- 5 Make 200 µL **Blocking Buffer** per sample by dividing your total desired volume of blocking buffer by 10 to get the amount of 10% BSA in µL. Add this amount to a tube and then fill the remainder with 1X PBS.

Homogenization

- 6 Clean dounce, scalpel, and forceps using MilliQ water, ethanol, RNase Zap, then MilliQ again. The red-tape forceps are for unfixed tissue.
- 7 Get tissue sample from -80°C freezer and place on dry ice. Weigh it on a sterile, tared weigh boat.
- 8 Add tissue to dounce and push it to the bottom using 1 mL of Homogenization Buffer and the pestle. Homogenize the tissue without creating bubbles. Then add the remaining 2 mL of the Homogenization Buffer and continue to dounce until homogenized.

- 9 Pass all of the nuclei suspension through three FlowMi filters, 1000 μ L at a time into a new 5 mL eppendorf.
- 10 Centrifuge at 900 g/rcf for 10 minutes at 4°C.
- 11 Discard the supernatant and resuspend the pellet in 200 μ L Blocking Buffer. Incubate for 10 minutes on ice.

Cell Count

- 12 Add 9 μ L of sample to a PCR tube and then add 1 μ L of acridine orange.
If sample is very concentrated, instead add 2 μ L sample to 2 μ L of acridine orange and 16 μ L 1X PBS.
- 13 Pipette mix and then add 10 μ L to a three-chamber cell counting chip and make note of the channels used (A, B, and/or C).
- 14 On the cell counter, select Fluorescence Cell Counting -> Cell Lines & Primary Cells, Advanced-> Protocol -> and then choose "NUCLEI" from the list of protocols. Load the protocol.

If the sample is very concentrated and you are adding 2 μ L, select Fluorescence Cell Counting -> Cell Lines & Primary Cells-> Protocol -> and then choose "NUCLEI" from the protocol list. Load the protocol.

Then go to Settings and choose the appropriate number of channels.

Then hit "Count" and then "Start Count."
- 15 When the cell count is complete, you will get a reading in cells/mL. Convert this to cells/ μ L by dividing this number by 1,000.

- 16** Dilute this nuclei stock to a concentration of 7,000 nuclei/uL.

Example:

$$(2,000 \text{ n/uL})(500 \text{ uL}) = (7,000 \text{ n/uL})(x \text{ uL})$$

$$x = 2,286 \text{ uL}$$

2,286 uL - 498 uL remaining after cell count = 1,788 uL blocking buffer needed to add to sample

- 17** If there are channels left on the cell counter chip, mark the used channels on the back and place it back in the drawer for future use.

- 18** Proceed with step 1 of the 10X Multiome protocol.