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© Isolation of Stromal Vascular Fraction (SVF) from mouse brown adipose tissue (BAT) for single cell RNA-seq

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1 Works for me

dx.doi.org/10.17504/protocols.io.bpurmnv6

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ABSTRACT

This protocol outlines the procedure for the isolation of the Stromal Vascular Fraction (SVF) from mouse brown adipose tissue (BAT) for single cell RNA-seq. This protocol uses a combination of Collagenase I and Dispase II to digest freshly isolated BAT. Compared to using Collagenase I alone, this combination results in a more efficient dissociation of the adipose vasculature.

ATTACHMENTS

Isolation_of_Stromal_Vasc ular_Fraction_(SVF)_from_ mouse_brown_adipose_tis sue_(BAT)_for_single_cell_ RNA-seq.pdf

DO

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PROTOCOL CITATION

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KEYWORDS

stromal vascular fraction, mouse brown adipose tissue, brown adipose tissue, single cell RNA-seq, RNA-seq

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Biotec Catalog #130-090-101

S Corning® 40μm Cell

Strainer Corning Catalog #431750

⊠ MS Columns **Miltenyi**

Biotec Catalog #130-042-201

X ACK Lysing Buffer

(1X) Lonza Catalog #10-548E

Scientific Catalog #AM9780

⊠ Falcon® 100 μm Cell

Strainer Corning Catalog #352360

MACS Separator Miltenyi Biotec

Digestion Media:

⊠ Collagenase Type 1 Worthington Biochemical

Corporation Catalog #LS004196

Technologies Catalog # 07913

Bovine Serum Albumin (BSA): Gemini Bio Products BSA V FATTY ACID FREE 100G Fisher

Scientific Catalog #50-753-3073

₩ HBSS: Corning® Hanks Balanced Salt Solution 1X with calcium and magnesium

Corning Catalog #21-020-CM

Growth Media:

Fisher Catalog #11965118

SAFETY WARNINGS

For hazard information and safety warnings, please refer to the SDS (Safety Data Sheet).

ABSTRACT

This protocol outlines the procedure for the isolation of the Stromal Vascular Fraction (SVF) from mouse brown adipose tissue (BAT) for single cell RNA-seq. This protocol uses a combination of Collagenase I and Dispase II to digest freshly isolated BAT. Compared to using Collagenase I alone, this combination results in a more efficient dissociation of the adipose vasculature.

BEFORE STARTING

Prepare the digestion media containing 1.5 mg/ml Collagenase I, 2.5 U/ml Dispase, and %2 BSA in HBSS buffer.

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Warm to § 37 °C.

Prepare growth media by adding FBS (%10) to DMEM. Warm to § 37 °C.

- Sacrifice the mouse.
- 2 Spray the animal extensively with 70 % EtOH and RNaseZap™.
- 3 Dissect interscapular brown adipose tissue (BAT). If tissues from multiple animals are being dissected, store them in HBSS until all of them are dissected.
- 4

Mince the tissue to very fine pieces in a 50 ml Falcon tube. Add □10 mL digestion media for each BAT.

5

Place the tubes in a water bath or incubator with a shaker/rotator at § 37 °C for © 00:45:00 .

- 6 Remove the tissue from the incubator and vortex for **© 00:00:10**.
- 7

Centrifuge at 300 x g, 4°C, 00:10:00 in a swinging bucket centrifuge.

- Aspirate the supernatant carefully not to disturb the pellet of SVF cells.
- 9 Resuspend the pellets in **10 mL growth media**.
- 10

Filter through a 100 μ m cell strainer into a fresh 50 ml tube. Wash the tube with an additional \blacksquare 10 mL and filter through the cell strainer.



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Centrifuge at **300 x g, 00:07:00**.

- 12 Completely remove supernatant and re-suspend the pellet in **2 mL sterile ACK lysis buffer**; place § On ice for © 00:05:00.
- 13

Filter through a 40 μ m cell strainer into a fresh 50 ml tube. Wash the tube with \Box 20 mL growth media and filter through the cell strainer.

14

Centrifuge at \$300 x g, 00:07:00.

- 15 Resuspend the pellet in 11 mL %1.5 BSA in PBS.
- 16 Use $\square 10 \, \mu I$ of the cell suspension for cell counting and viability assessment.
- 17

Centrifuge the cell suspension 300 x g, 00:05:00.

18

Resuspend the cells in $\[\Box 100 \ \mu I$ dead cell removal bead solution . Incubate the samples for $\[\odot 00:15:00 \]$ at $\[\& Room \ temperature \]$.

- 19 Prepare the binding solution by diluting the 20X solution in sterile ddH20.
- Place the MS columns on the MACS separator. Prepare each column by rinsing it with **0.5 mL 1X binding solution**. Let the solution pass through the column.
- 21 Add **3900 μl 1X binding solution** to each sample and apply cell suspension onto the column.

- 22 Collect effluent in a 2 ml low bind tube as live cell fraction.
- 23

Rinse the column with an additional $\square 1$ mL 1X binding solution.

- 24 Use $\Box 10 \mu l$ sample for cell counting and viability assessment.
- 25

Centrifuge the cell suspension 300 x g, 00:05:00.

- 26 Resuspend the cells in \Box 50 μ I \Box 100 μ I %1.5 BSA in PBS .
- 27 Keep the cell suspension & On ice and proceed to 10x Genomics Single Cell Protocol. Minimize the time between cell preparation and chip loading.