

Video Article

Immunomagnetic Separation of Fat Depot-specific Sca1^{high} Adipose-derived Stem Cells (ASCs)

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Abstract

The isolation of adipose-derived stem cells (ASCs) is an important method in the field of adipose tissue biology, adipogenesis, and extracellular matrix (ECM) remodeling. *In vivo*, ECM-rich environment consisting of fibrillar collagens provides a structural support to adipose tissues during the progression and regression of obesity. Physiological ECM remodeling mediated by matrix metalloproteinases (MMPs) plays a major role in regulating adipose tissue size and function^{1,2}. The loss of physiological collagenolytic ECM remodeling may lead to excessive collagen accumulation (tissue fibrosis), macrophage infiltration, and ultimately, a loss of metabolic homeostasis including insulin resistance^{3,4}. When a phenotypic change of the adipose tissue is observed in gene-targeted mouse models, isolating primary ASCs from fat depots for *in vitro* studies is an effective approach to define the role of the specific gene in regulating the function of ASCs. In the following, we define an immunomagnetic separation of Sca1^{high} ASCs.

Video Link

The video component of this article can be found at <https://www.jove.com/video/53890/>

Introduction

Stem cell antigen 1 (Sca1, or Ly6A/E) was first identified as a cell surface marker expressed by hematopoietic and mesenchymal stem cells^{5,6}. The stromal vascular fraction (SVF) of adipose tissue obtained from mouse fat depots is a heterogeneous population of cells comprising of fibroblasts, macrophages, vascular endothelial cells, neuronal cells, and adipocyte progenitor cells⁷. Adipocyte progenitor cells, or adipose-derived stem cells (ASCs) are non-lipid-laden cells that reside in the collagen-rich perivascular extracellular matrix (ECM)⁸. Approximately 50% of the SVF consist of ASCs, which are characterized as lineage-negative (Lin⁻) and CD29⁺: CD34⁺: Sca1⁺⁹. Most of these cells are Sca1⁺: CD24⁻ adipocyte progenitors, which are capable of adipocyte differentiation *in vitro*; however, only a fraction of cells (0.08% of SVF) constitutes Sca1⁺: CD24⁺ cells that are fully capable of proliferating and differentiating into adipocytes in the *in vivo* conditions⁹. Despite the potential caveat of using Sca1⁺ SVF without discriminating CD24⁺ cells from CD24⁻ cells, isolating Sca1⁺ ASCs from fat depots using immunomagnetic cell separation is an efficient and practical approach to determine the cell-autonomous phenotype of primary adipocyte progenitor cells.

In the field of obesity and diabetes, tissue fibrosis and inflammation play a critical role in the development and maintenance of type-2 diabetes³. Recently, Tokunaga *et al.* showed that Sca1^{high} cells isolated from inguinal (or subcutaneous, SQ) and perigonadal (or visceral, VIS) C57BL6/J fat depots exhibit different gene signatures and ECM remodeling *in vitro*¹⁰. MMP14 (MT1-MMP), a prototypical member of the membrane-type matrix metalloproteinase (MMP) family mediates the development of white adipose tissue (WAT) through its collagenolytic activity¹.

Examples of experiments that may be conducted with the cells isolated and enriched through the following protocol include three-dimensional culture, differentiation studies, collagen degradation assays, and RNA sequencing^{10,11}. Degradation assays should be conducted with acid-extracted collagen to ensure the preservation of telopeptide^{11,12}. The following protocol will demonstrate the methods to isolate primary vascular stromal cells from different fat depots and enrich adipocyte progenitor cells using immunomagnetic cell separation. The validity of the cell sorting will be assessed with flow cytometry and through using Sca1-GFP mice that express GFP in Sca1⁺ cells, driven by a Sca1 promoter¹³.

Protocol

Ethics Statement: The University of Michigan Committee on Use and Care of Animals (UCUCA) has approved all methods and protocols in accordance with the Guide for the Care and Use of Laboratory Animals (Institute for Laboratory Animal Research, National Research Council). Mice are maintained in a University of Michigan vivarium and are given free access to food and water and kept on a 12 hr dark/light cycle.

1. Preparations

1. Prepare primary culture media with DMEM, 10% FBS, 1x P/S/G, and 1x antibiotics/antifungals. This will be used to keep tissues in before digestion. Place stock and aliquot in 37 °C water bath.
2. Prepare five milliliters of Type III collagenase solution at 5mg/ml dissolved in HBSS (+Ca, +Mg) for each type of fat pad being isolated, up to five mice. For example, when isolating SQ and VIS from a single genotype, make 10 ml, if wild-type and mutant SQ and VIS, make 20 ml. Adjust pH to 7.4 and sterile filter. Aliquot into 50 ml tubes. Set aside away from light.
3. Use 70% ethanol to disinfect surgical field. Wipe down and cover with a blue pad. Spray some ethanol on the blue pad.
4. Use the 22 G needles to pin down an absorbent pad to the styrofoam board and trim with scissors. Spray the pad with ethanol then set the board on the blue pad and cover with another blue pad until beginning dissection.
5. Fill two 50 ml tubes with ethanol and place in a stand adjacent to the surgical field. Place the scissors and forceps in the ethanol.
6. Fill another 50 ml tube with PBS plus 1x anti-anti for rinsing ethanol off surgical tools.
7. In the hood, label 60 mm dishes for each tissue type and fill with culture media warmed to 37 °C. Place the plates adjacent to the surgical area.

2. Isolation of Subcutaneous (SQ) Fat Pads

1. Euthanize mouse with an overdose of isoflurane and pneumothorax.
2. Gently spray mouse with 70% ethanol and lay supine. Pin the paws to the foam board with 22 G needles.
3. Make a small cut in the skin of the lower abdomen. Holding the top of the cut with the forceps, take the scissors and separate the skin from the peritoneum.
4. Once the skin is separated from the peritoneum from the abdomen to the thorax, reflect the skin away from the peritoneum towards the head by making lateral cuts in the skin along the side of the body.
5. Reflect the remaining skin holding the inguinal fat away from the body and pin down to the board with 22 G needles.
6. Switch to fine scissors and forceps. Grab the inguinal fat pad with the forceps at the origin near the peritoneum and snip away between the skin and inguinal fat progressing towards the groin avoiding contaminating the SQ with skin.
7. Place the isolated inguinal fat pad in the labeled 60 mm dish.

3. Isolation of Visceral (VIS) Fat Pads

1. In a similar fashion to step 2.5, cut the peritoneum open to expose the visceral fat pads and gut.
2. Move the gut away toward the thorax.
3. Grab the VIS fat pad at the distal end and gently pull up. Dissect out the VIS fat pad while being careful to exclude epididymal (or uterine, if using female mice) tissue.
4. Place in labeled dish and repeat step 3.3 for the remaining VIS pad.

4. Collagenase Digestion of Fat Pads

1. Move 60 mm dishes to the tissue culture hood and aspirate off media.
2. Mince the fat pads with curved scissors then add to collagenase solution. Be sure to rinse the scissors and forceps in between tissue types with ethanol and PBS.
3. Shake at 300 rpm and 37 °C for 10-20 min until tissues are digested. It is acceptable if some pieces of SQ are still visible. This will be addressed in a later step.
4. Add 25 ml of culture media to the collagenase solution to stop the collagenase treatment, and pipette up and down 10 times with a 10 ml serological pipette to disperse the digested tissues.
5. Strain cells using a 100 µm cell strainer. Centrifuge for 10 min at 300 x g.
6. Carefully decant off media not to disturb the pellet and add 5 ml of sterile water to the pellet and gently pipette to suspend cells. Wait 2 min to lyse erythrocytes.
7. Add 25 ml of media with 10% FBS and strain cells through a new 100 µm cell strainer.
8. Centrifuge for 10 min at 300 x g. Decant media and resuspend pellet in 1 ml of culture media.
9. Count cells with a hemocytometer using 1:1 trypan blue.
10. Plate 1×10^6 cells in one well on a 6-well plate.

5. Magnetic Cell Separation

1. After about 4 to 6 hr of cell adhesion, rinse cells twice with HBSS (-Ca, -Mg). Dissociate adherent cells using 0.05% trypsin. Dilute trypsin cell suspension in 1 ml separation buffer and spin for 5 min at 300 x g.
2. Aspirate supernatant. Resuspend cells in 500 µl buffer. Remove a 10 µl aliquot and count cells using a hemocytometer.
3. Spin cells for 5 min at 300 x g.
4. Aspirate supernatant completely. Resuspend cells in 90 µl buffer. Add 10 µl anti-Sca1-FITC primary antibody. Mix well and incubate for 10 min at 4 °C in the dark.
5. Wash cells with 1 ml buffer and centrifuge cells for 5 min at 300 x g. Remove supernatant completely.
6. Resuspend cells in 80 µl buffer. Add 20 µl anti-FITC microbeads. Mix well and incubate for 15 min at 4 °C in the dark.
7. Wash cells with 1 ml buffer and centrifuge cells for 5 min at 300 x g.
8. Remove supernatant and resuspend pellet in 500 µl buffer.
9. Place column in the magnetic holder and rinse column with 500 µl buffer.

10. Add cell suspension to column. Avoid bubbles while pipetting.
11. Collect the unlabeled Sca1⁺ cells and wash column 3 times with 500 μ l buffer. Only add new buffer when the reservoir is empty. Avoid bubbles while pipetting.
12. Remove column from the magnetic holder and place into a 15 ml conical tube.
13. Add 1 ml buffer to column and elute Sca1⁺ by pushing supplied plunger through the column reservoir.
14. Spin down cell fractions for 5 min at 300 x g. Resuspend cells in 1 ml culture media.
15. Remove a 10 μ l aliquot from the labeled and unlabeled fractions and count cells using a hemocytometer.
16. Plate each cell fraction in 1 well of a 6-well plate.

6. Verification of Immunomagnetic Separation of Sca1^{high} ACSs with Flow Cytometry

1. Rinse cells twice with HBSS (-Ca, -Mg) and dissociate cells using 0.05% trypsin.
2. Centrifuge cells at 300 x g for 5 min. Resuspend cells in 1 ml culture media and count using a hemocytometer.
3. Obtain $>10^6$ cells in 1 ml culture media and centrifuge at 300 x g for 5 min.
4. Remove supernatant and repeat step 6.3 two more times.
5. Resuspend cells with 1 ml of 2% goat serum + 2% BSA and block for 30 min at RT.
6. Centrifuge cells at 300 x g for 5 min.
7. Remove blocking solution.
8. Add rat IgG2a Alexa Fluor 647 (0.25 μ g, 1:400) or anti-Sca1 Alexa Fluor 647 (0.25 μ g, 1:400), in 100 μ l of PBS with 2% goat serum and 2% BSA + PBS at 4 °C for 30 min in the dark.
9. Add 1 ml of cold PBS to cells. Centrifuge 300 x g for 5 min at 4 °C.
10. Remove supernatant and repeat step 6.9 two more times.
11. Resuspend cells in 1 ml PBS. Pass cell suspensions through a 100 μ m cell strainer in preparation for flow cytometric analysis.

Representative Results

Enrichment of Sca1^{high} ASCs from Different Fat Pads.

The vascular stromal cells isolated from SQ fat display fibroblast-like, stretched cell shape regardless of Sca1 expression level (**Figure 1A**). On the other hand, VIS (eWAT-derived) Sca1^{high} and Sca1^{low} cells demonstrate distinct difference in their cell shape. Like SQ (iWAT-derived) Sca1^{high} cells, VIS (eWAT-derived) Sca1^{high} cells display stretched, fibroblast-like cell shape, whereas VIS Sca1^{low} cells demonstrate epithelioid shape. Sca1^{high} cells isolated from Sca1-GFP mice are easily identified as GFP-positive cells in tissue culture (**Figure 1B**). When these cells were assessed with flow cytometry, most of the GFP positive cells were confirmed to express Sca1 proteins on the cell surface as detected with Anti-Sca1 antibody (**Figure 1C**). Sca1^{high} cells derived from inguinal fat pad maintain increased capacity of adipocyte differentiation whereas eWAT-derived Sca1^{high} cells are more difficult to differentiate into adipocyte with conventional adipogenic mix¹⁰.

Fat Depot-dependent Gene Expression of Sca1^{high} ASCs (**Figure 2**).

Genome-wide transcriptome analyses with RNA sequencing demonstrated the enrichment of genes related to extracellular matrix proteins and modifiers (GO:0031012, GO:0005578) in those Sca1^{high} ASCs¹⁰. Coupled with real-time PCR analyses, we were able to demonstrate the differential expression of the collagenolytic MMPs (MMP2, MMP8, MMP13, MMP14) between iWAT- and eWAT-derived Sca1^{high} ASCs. When fluorescein-labeled type I collagen gels were used to assess pericellular degradation activity, we observed the markedly increased collagen remodeling activity mediated by VIS Sca1^{high} ASCs¹⁰.

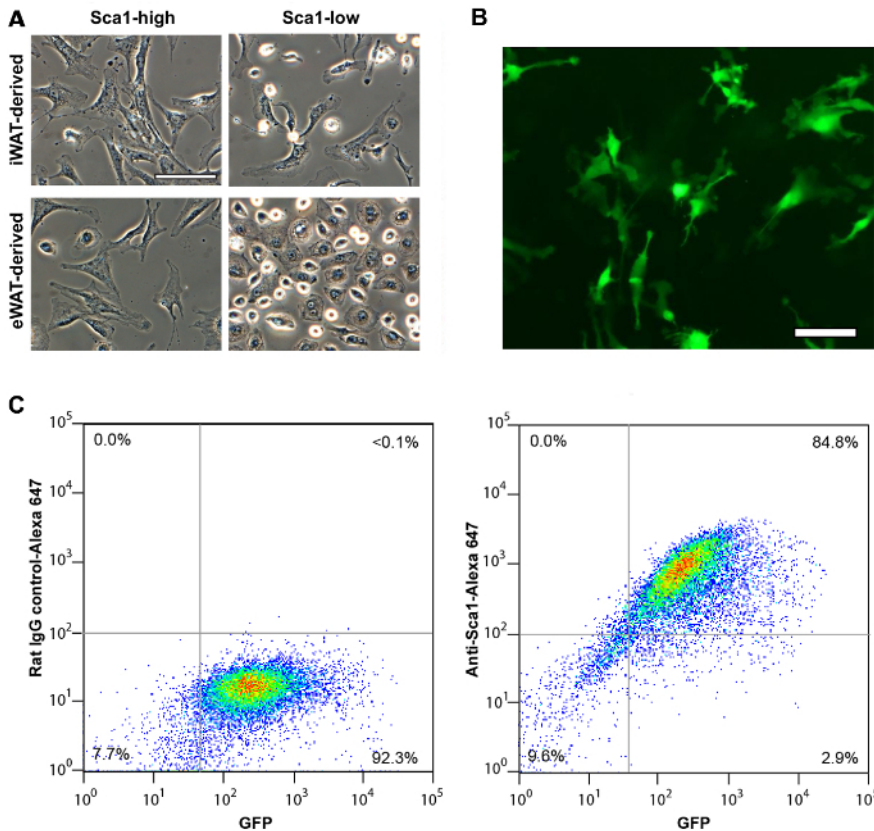


Figure 1: Immunomagnetic Separation of Murine Sca1^{high} ASCs from Different Fat Depots. (A) Sca1^{high} and Sca1^{low} ASCs isolated from SQ (iWAT) and VIS (eWAT). Scale = 100 μ m. (B) Sca1-GFP cells isolated from iWAT of Sca-GFP mice. Scale = 100 μ m. (C) Cell surface expression of Sca1 in Sca1-GFP-positive cells assessed with flow cytometry. (left) control rat IgG (right) anti-Sca1 antibody. X-axis, GFP intensity. Y-axis, Alexa-Fluor 647. Panel A shown previously in Tokunaga, M. *et al.*, (2014). [Please click here to view a larger version of this figure.](#)

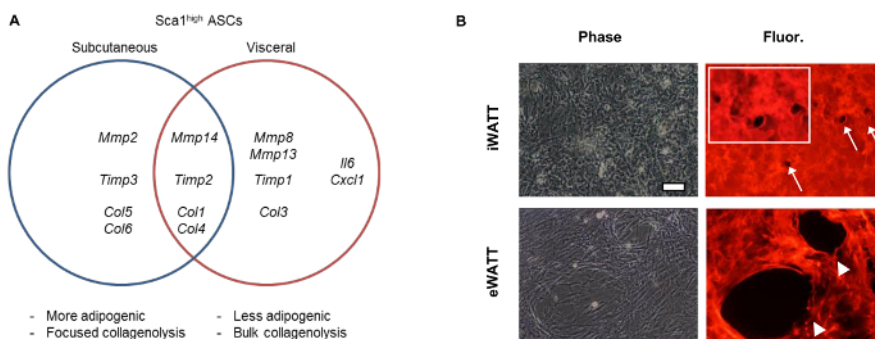


Figure 2: Fat Depot-Dependent Expression of Collagenolytic MMPs, TIMPs, and Collagens. (A) Differential gene expression of ECMs and ECM modifiers in Sca1^{high} ASCs isolated from different fat depots. (B) Increased collagen degradation activity of VIS (eWAT-derived) Sca1^{high} ASCs. Degradation of collagen is shown as the disappearance of fluorescent signals (arrows and arrowheads). Inset is an enlarged image of focused collagen degradation mediated by individual cell of SQ ASCs. Cells were cultured for 72 hr. Data shown previously in Tokunaga, M. *et al.*, (2014). [Please click here to view a larger version of this figure.](#)

Discussion

Herein we demonstrate the isolation and immunomagnetic cell separation of murine ASCs from different fat pads and their use for *in vitro* experiments. The presented method is effective for the quick isolation of large number of Sca1-positive ASCs, which is advantageous over the technically complex and expensive FACS-mediated isolation of ASCs^{9,14}. Unlike FACS, immunomagnetic cell separation does not allow the use of multiple antigen for the identification of a target cell population. Nonetheless, if the surface antigen is well-characterized, the use of immunomagnetic separation increases the number of cells to be analyzed without relying on the use of FACS equipment¹⁵, which is still not readily accessible to many biological researchers at small institutes without core facilities.

There are some critical aspects of the procedure that must be observed to ensure a successful outcome. The procurement of adipose-derived stem cells requires the surgical isolation of mouse adipose tissue depots. Therefore, the speed and accuracy of tissue retrieval is imperative to yield a high number of viable cells. Additionally, the maintenance of clean surgical fields and instruments are vital to the outcome of the procedure by preventing microbial contamination of cell and tissue samples. Dissected adipose tissues must be enzymatically dissociated in a type II collagenase solution. ECM composition differs between SQ and VIS fat pads^{10,16}, where VIS contain less collagens than SQ. Therefore, the duration of VIS fat pad digestion requires approximately half the amount of time as SQ. It is acceptable to stop tissue digestion after the stated incubation period even if small particles of tissue are still present in the collagenase solution. These pieces may be mechanically dissociated with pipetting once culture media has been added to inhibit collagenase activity. While it is possible to proceed directly to immunomagnetic sorting following SVF isolation, our group seeds 1×10^6 unsorted cells on plastic plates for about 4 to 6 hr before continuing with cell sorting. Despite multiple filtration steps during SVF isolation, debris will still be present within the cell suspension. Plating the cells before proceeding to immunomagnetic cell separation provides an opportunity to wash debris and unattached cells away, thus allowing magnetic sorting to only be applied to adherent cells that contain adipogenic progenitor cells.

While Sca1 is not found in human genome, the identification and validation of alternative cell surface antigens expressed on fat depot-dependent human adipose stem cells¹⁷, when coupled with this cell separation technique, may help us define the biology of human adipose stem cells.

Disclosures

The authors have nothing to disclose.

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