

# Histomorphometric analyses of human adipose tissues using intact, flash-frozen samples

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46 **Abstract**

47 Histomorphometric analyses of adipose tissue usually require formalin fixation of fresh  
48 samples. Our objective was to determine if intact, flash-frozen whole adipose tissue  
49 samples stored at  $-80^{\circ}\text{C}$  could be used for measurements developed for fresh-fixed  
50 adipose tissues. Portions of adipose tissue samples were either formalin-fixed  
51 immediately upon sampling or flash-frozen and stored at  $-80^{\circ}\text{C}$  and then formalin-fixed  
52 during the thawing process. Mean adipocyte diameter was measured.  
53 Immunohistochemistry was performed on additional samples to identify macrophage  
54 subtypes (M1, CD14+ and M2, CD206+) and total (CD68+) number. All slides were  
55 counterstained using haematoxylin & eosin (H&E). Visual inspection of H&E-stained  
56 adipose tissue slides performed in a blinded fashion showed little or no sign of cell  
57 breakage in 74% of frozen-fixed samples and in 68% of fresh-fixed samples ( $p>0.5$ ).  
58 There was no difference in the distribution frequencies of adipocyte sizes in fresh-fixed  
59 vs. frozen-fixed tissues in both depots ( $p>0.9$ ). Mean adipocyte size from frozen-fixed  
60 samples correlated significantly and positively with adipocyte size from fresh-fixed  
61 samples ( $r=0.74$ ,  $p<0.0001$ , for both depots). The quality of staining/immunostaining and  
62 appearance of tissue architecture were comparable in fresh-fixed vs. frozen-fixed  
63 samples. In conclusion, intact flash-frozen adipose tissue samples stored at  $-80^{\circ}\text{C}$  can be  
64 used to perform techniques conventionally applied to fresh-fixed samples. This approach  
65 will allow retrospective studies with frozen human adipose tissue samples.

66 **Introduction**

67 Adipose tissue expansion occurs through increases in the volume of adipocytes  
68 (adipocyte hypertrophy) and/or increases in the total number of adipocytes (Cristancho  
69 and Lazar 2011; Lowe et al. 2011). Hypertrophy of mature adipocytes is linked to altered  
70 adipose tissue function, a phenomenon that is independent of concomitant differences in  
71 total adiposity (Arner et al. 2010; Bjorntorp et al. 1971; Hoffstedt et al. 2010; Ledoux et  
72 al. 2010; Lundgren et al. 2007; Veilleux et al. 2011; Weyer et al. 2000). The  
73 accumulation of visceral fat is also a strong predictor of the metabolic complications  
74 associated with obesity (reviewed in (Tchernof and Després 2013)). Expansion of  
75 visceral adipose tissue is thought to be related to limited expandability of subcutaneous  
76 fat, which limits the storage of excess dietary fatty acids (Tchernof and Després 2013).

77 Adipocyte size represents a major marker of the metabolic profile of obese patients and  
78 varies according to the metabolic status of their various fat compartments (Laforest et al.  
79 2015; Laforest et al. 2017). We have previously shown that adipocyte size can be  
80 assessed through three techniques currently utilized in the literature and these methods  
81 generate similar associations with adiposity and metabolic variables (Laforest et al.  
82 2017). However, only histological analysis can be used to study tissue architecture  
83 (Laforest et al. 2017). Adipocyte hypertrophy is associated with increased macrophage  
84 infiltration in adipose tissue, thus contributing to the local and systemic inflammation  
85 observed in obesity and insulin resistance (Michaud et al. 2012; Michaud et al. 2013).  
86 Using histomorphometric analyses of adipose tissue, we have shown that pericellular  
87 fibrosis in adipose tissue, but not total fibrosis, is associated with a detrimental metabolic  
88 profile (Michaud et al. 2016), thus enhancing the importance of studying *in situ* markers

89 of adipose tissue dysfunction to better understand the physiopathology of obesity and  
90 concomitant diseases such as type 2 diabetes.

91

92 According to available literature, no attempt has been made to measure the size of  
93 adipocytes after flash freezing, regardless of the use and type of cryoprotective agents  
94 (Son et al. 2010; Wolter et al. 2005). Because adipocytes contain less water than most  
95 cell types, there is a reduced possibility of ice crystals formation, the main cause of cell  
96 membrane breakage during the thawing process. Our working hypothesis was that it is  
97 feasible to preserve the integrity of tissue architecture and consequently quantify cell size  
98 in tissue that was frozen in liquid nitrogen and stored at -80°C without cryoprotectant.  
99 We also postulated that immunohistochemistry (IHC) characterization of the infiltrating  
100 macrophages is also possible in these samples. We tested the hypothesis that cell size  
101 measurements and macrophage quantification in frozen and subsequently fixed adipose  
102 tissue samples is comparable to measurements obtained in portions of the same samples  
103 that were fixed in the fresh state. The usefulness of frozen adipose tissue in  
104 histomorphometric analysis is of the utmost importance in the field of obesity and would  
105 allow for retrospective studies in large tissue banks.

## 106 **Materials and Methods**

### 107 *Participant recruitment*

108 Women undergoing gynecological surgery were enrolled at the Gynecology Unit of CHU  
109 de Quebec-Laval University Medical Center. The study was approved by the Research  
110 Ethics Committees of Laval University Medical Center (C09-08-086). Samples were also

111 obtained from research volunteers participating in IRB-approved studies of obesity at  
112 Mayo Clinic. All procedures performed were in agreement with the ethical guidelines of  
113 the Declaration of Helsinki and its later amendments. All subjects provided written  
114 informed consent before their inclusion in the study.

115 ***Adipose tissue samples***

116 Subcutaneous (SC) (n=29) and omental (OM) (n=29) adipose tissue samples were  
117 collected during the surgical procedure at the site of incision (lower abdomen) and at the  
118 distal portion of the greater omentum, respectively at the CHU de Quebec-Laval  
119 University Medical Center. At Mayo Clinic, SC adipose tissue from abdomen (n=4) for  
120 direct comparison of freshly fixed vs. frozen-fixed tissue and an additional set of samples  
121 (n=6) and thigh (n=7) were collected for frozen-fixed analysis. All biopsies were  
122 performed under local anesthesia using sterile technique. Samples were  
123 immediately carried to the laboratory. To measure adipocyte size, a portion of the sample  
124 was fixed in 10% buffered formalin for 24 to 48 hours at room temperature and then  
125 processed for standard paraffin embedding. The remaining portion of the sample was  
126 immediately flash frozen with liquid nitrogen and subsequently kept at -80°C.

127 ***Fixation of frozen adipose tissue***

128 A container was filled with dry ice to keep surgical instruments cold (tweezers, scalpel  
129 and scissors). Precautions were also taken so that frozen adipose tissue samples never  
130 thawed during weighing and preparation. Surgical instruments were used to collect a  
131 subsample of 50-100 mg from frozen tissue pieces kept on dry ice. The subsamples were  
132 then transferred to a 5 mL tube kept cold and immediately stored in a -80°C freezer until  
133 fixation. The frozen tissue pieces were then fixed in 10% buffered formalin kept at 4°C

134 for 24 to 48 hours. No thawing of the adipose tissue samples was performed prior to  
135 incubation in cold buffered formalin. Samples for IHC underwent the same processing,  
136 but paraffin incubation time was limited to 24 hours to limit cross-linking of proteins  
137 which may affect the quality of IHC.

138 *Cell sizing*

139 All adipose tissue slides (tissue slice thickness: 5  $\mu$ m) were stained with haematoxylin &  
140 eosin (H&E) for adipocyte diameter measurements. The histological H&E were  
141 photographed with the help of a QImaging Retiga 2000R FAST camera (QImaging,  
142 Surrey, C.B., Canada) coupled to a BX51 microscope (Olympus, Shinjuku, Tokyo,  
143 Japan). Cell areas were measured with ImageJ software (according to (Parlee et al. 2014))  
144 with slight modifications to the protocol (**Figure 1**). Briefly, analyses were performed in  
145 a blinded fashion, without knowledge of the original patient, the type of depot (OM or  
146 SC) and condition (frozen or fresh). Visual inspection of H&E-stained adipose tissue  
147 slides was performed prior to cell sizing to assess the quality and quantity of the tissue.  
148 For each folder of pictures, one image was randomly selected and was converted to 8-bit.  
149 The background was subtracted and background noise was eliminated. A threshold was  
150 set for the binary conversion, with values that were different from a group pictures to  
151 another depending on the specifics of the tissue and due to uneven H&E staining. This  
152 value was noted for programming of automated image analysis.

153 Pictures were treated with the steps described above in an automated manner according to  
154 their threshold value. They were converted into binary, contours of the cells were dilated  
155 and each cell was selected with the Wand tool, measured and labeled with the extension  
156 module 'Measure and Label' of ImageJ. To prevent bias, all image files were placed in a

157 computer-generated random order. Cells had to show complete membranes to be  
158 measured. The results were then converted in diameter assuming a circular surface of  
159 adipocytes measured according to the following formula:  $D = 2\sqrt{A/\pi}$ . The average of  
160 100 cells was used when possible.

### 161 *Automated high-performance histological analysis of serial adipose tissue sections*

162 Additional SC (n=1) and OM (n=1) adipose tissue samples were collected during the  
163 surgical procedure at the site of incision and greater omentum, respectively. Consent was  
164 obtained through the management framework of the Quebec Heart Lung Institute Obesity  
165 Tissue Bank. Samples were processed as described in *Adipose tissue samples* and the  
166 *Fixation of frozen adipose tissue* sections above. Thirty consecutive 5  $\mu\text{m}$ -thick sections  
167 for each paraffin block (OM and SC fresh- and frozen-fixed) were cut, covering 150  $\mu\text{m}$   
168 of thickness per block and stained with H&E.

169 Subsequently, slides were scanned using a Zeiss AxioScan.Z1 whole slide scanner  
170 equipped with a 20X plan apochromat objective lens (NA 0.8). Since tissues stained with  
171 H&E are typically autofluorescent in the red channel, the image acquisition (Zen 2.3  
172 acquisition software) was performed using an AlexaFluor 555 filter set to generate  
173 grayscale images. Mosaic of images were exported as BigTiff files and processed using a  
174 modified algorithm used for the morphological analysis of lung alveoli as published  
175 previously (Sallon et al. 2015). Briefly, Matlab software version 2014a (Mathworks) was  
176 used to perform a distributed computing analysis of the images, and results were exported  
177 as CSV files. Grayscale images contrast was adjusted, then images were segmented using  
178 a specific threshold to generate binarized images. Closed structures were identified and  
179 sorted based on their area ( $>1200 \mu\text{m}^2$ ) and solidity values ( $>0.9$ ). Thus, an adipocyte

180 with a solidity value close to 1 will be very round and smooth, while adipocytes with low  
181 solidity value will be distorted or broken, hence excluded from our analysis. The  
182 frequency distribution of adipocyte area was calculated using Prism 7 and distribution  
183 plots were generated.

#### 184 *Immunohistochemistry*

185 All samples were paraffin-embedded and 5  $\mu$ m sections were placed onto slides and  
186 stained at the Pathology Research Core at Mayo Clinic, Rochester, MN. Each sample was  
187 stained with a total macrophage and monocyte marker antibody against CD68; an M1  
188 macrophage or pro-inflammatory macrophage marker antibody against CD14 and an M2  
189 macrophage or anti-inflammatory macrophage marker antibody against CD206. The IHC  
190 staining procedure was performed on-line using the Leica Bond III Stainer (Leica,  
191 Buffalo, IL). The tissue slides were dewaxed using Bond Dewax (Leica, Buffalo, IL).  
192 Tissue slides for CD14 and CD68 stain were retrieved for 20 minutes using Epitope  
193 Retrieval 2 (Leica, Buffalo, IL) and slides for CD206 stain were retrieved for 20 minutes  
194 using Epitope Retrieval 1 (Leica, Buffalo, IL). All primary antibodies were diluted in  
195 Bond Antibody Diluent (Leica, Buffalo, IL). The primary antibody CD14 (Sigma-  
196 Aldrich, St. Louis, MO) was used at 1:300; CD68 (Clone PG-M1, Dako, Carpinteria,  
197 CA) was used at 1:200; and CD206 (Clone 685645, R&D Systems, Minneapolis, MN)  
198 was used at 1:200. The detection system used was the Polymer Refine Detection System  
199 (Leica, Buffalo, IL). This system includes the hydrogen peroxidase block, secondary  
200 antibody polymer, DAB and hematoxylin. Immunostaining visualization was achieved by  
201 incubating slides 10 minutes in DAB and DAB buffer (1:19 mixture) from the Bond  
202 Polymer Refine Detection System. To this point, slides were rinsed between steps with 1'



203 Bond Wash Buffer (Leica, Buffalo, IL). Before and after DAB incubation, slides were  
204 rinsed in distilled water. Slides were counterstained for 5 minutes using Schmidt  
205 hematoxylin and molecular biology grade water (1:1 mixture), followed by several rinses  
206 in 1' Bond wash buffer and distilled water; this is not the hematoxylin provided with the  
207 kit. Following completion of the IHC, processed slides were removed from the stainer  
208 and rinsed in tap water for 5 minutes. Slides were dehydrated in increasing concentrations  
209 of ethyl alcohol and cleared in 3 changes of xylene prior to placing a permanent coverslip  
210 in xylene-based medium.

211 The stained tissue sections were then visualized by light microscopy using an Olympus  
212 BX43 microscope. Ten images per slide were obtained by randomly selecting fields at  
213 40x magnification. Two independent observers counted positively stained macrophages  
214 and total adipocytes for each field of view. Data are expressed as mean (standard  
215 deviation) number of positive cells per 100 adipocytes.

216 All slides were marked with a code rather than the source of the sample to ascertain that  
217 the independent observers were blinded to the other reader's data, to the participant, the  
218 research protocol, and biopsy site.

### 219 *Statistical analyses*

220 Differences between positive cells/mean cell diameter in fresh- vs. frozen-fixed were  
221 assessed by paired t-test. Cell size frequency distributions were assessed by the  
222 Kolmogorov-Smirnov test. Pearson correlations were computed with normally distributed  
223 variables according to the Shapiro-Wilk test. Distribution correction was performed  
224 according to Lenz et al. (2016). This correction considers the cut-off parameter of the  
225 imaging software, the use of cross-sectional images instead of 3D samples and the tissue

226 thickness. The normal distribution algorithm was used but the gamma distribution  
227 algorithm generated similar results (Lenz et al. 2016). A chi-squared test was used to  
228 assess differences in the proportion of damaged and/or unusable samples between the  
229 fresh- and frozen-fixed methods. A  $p$ -value  $\leq 0.05$  was considered significant. All  
230 statistical analyses were performed with JMP and/or SAS software (SAS Institute, Cary,  
231 NC, USA).

## 232 **Results**

### 233 *Visual inspection*

234 Histological examination of fresh-fixed and frozen-fixed tissues was performed in all  
235 samples to determine whether intact, flash-frozen adipose tissue showed signs of cell  
236 breakage and alteration of the architecture of adipose tissue after thawing and fixation.  
237 **Figure 2** shows representative pictures of frozen- and fresh-fixed adipose tissue. Slightly  
238 more apparent tissue shrinkage, damage and other alterations were observed in the fresh  
239 samples. However, the proportion of unusable samples was not different in the frozen-  
240 fixed vs. fresh-fixed methods ( $p=0.54$ ). Samples with large amounts of stromal tissue  
241 (capillaries, fibroblasts, etc.) appeared to show more damage.

242 As the morphology of the tissue was retained in most samples, both fresh- and frozen-  
243 fixed, we performed IHC to label macrophages. Macrophage detection is one of the most  
244 used procedure performed on fresh-fixed fat samples to further characterize the tissue.  
245 Preliminary tests showed that the antibodies still retain the ability to bind corresponding  
246 cell surface markers in frozen adipose tissue (**Figure 3**).

247 *Immunohistochemistry*

248 We compared 4 slides that had been frozen and then fixed with slides from the same  
249 freshly fixed tissue for CD68, CD14 and CD206 stained macrophages. Freshly fixed  
250 tissue had 27(16), 8(6), and 22(10), CD68, CD14 and CD206 macrophages/100  
251 adipocytes, respectively. For the frozen-fixed tissue these values were 18(15), 7(4), and  
252 20(4), CD68, CD14 and CD206 macrophages per 100 adipocytes, respectively (standard  
253 deviation values in parentheses). Representative images are shown in **Figure 3, 4 and 5**,  
254 for CD68, CD206 and CD14, specifically. Paired t-tests showed no difference between  
255 fresh- vs. frozen-fixed for CD68-, CD206- or CD14- positive cells (**Figure 6**).

256 We used frozen tissue from an additional 7 volunteers (13 slides, 7 thigh and 6 abdomen)  
257 to further test the frozen-fixation protocol. For these 13 slides, we found that there were  
258 14(5), 7(2), and 13(3), CD68, CD14 and CD206 macrophages per 100 adipocytes,  
259 respectively (data not shown).

260 *Adipocyte size*

261 Mean cell size distribution curves from OM and SC fresh- and frozen-fixed tissue are  
262 shown in **Figure 7 and 8**. There was no difference in cell size distributions ( $p>0.90$ ) and  
263 in mean adipocyte sizes (**Figure 7 and 8**). Furthermore, in both OM and SC samples,  
264 adipocyte mean diameters from fresh- and frozen-fixed samples were strongly  
265 intercorrelated ( $r=0.74$ ,  $p<0.0001$ , for both) (**Figure 7 and 8**). We also presented the same  
266 analysis with the corrected cell size distributions for both depots and both conditions  
267 (**Figure 7 and 8**) as proposed by Lenz and collaborators to reduce measurement bias  
268 (Lenz et al. 2016). We performed automated, high-performance histological analysis of  
269 30 consecutive cross-sectional adipose tissue sections covering 150  $\mu\text{m}$  for both depots

270 and both conditions. Distribution of adipocyte area was not significantly different across  
271 conditions for each depot (**Figure 9**). In **Figure 10**, we present cell size distributions and  
272 main characteristics from two different patients to illustrate individual heterogeneity.

## 273 **Discussion**

274 We investigated if it was possible to evaluate adipocyte size, tissue architecture and  
275 macrophage infiltration in intact, frozen whole adipose tissue stored at -80°C. We  
276 demonstrated here for the first time that flash-frozen adipose tissue, without use of  
277 cryoprotective agents, can be utilized to assess fat cell size mean diameter and  
278 distribution. Furthermore, antigen labeling can be performed in a similar fashion, as  
279 shown by macrophage detection by IHC. These results are consistent with those reported  
280 in fresh-fixed adipose tissue in our study. We acknowledge the inherent limitations  
281 related to histological assessment of adipocyte cell size (Laforest et al. 2017). Our  
282 attempts to apply stereological principles to cell sizing analyses in the present study were  
283 unsuccessful. Our assessment is that the very large size of human adipocytes, especially  
284 in obese individuals, makes the use of stereological analysis extremely difficult. It  
285 requires registering pairs of very large images, and overlaying unbiased stereological  
286 frames over precisely matching regions in both the reference and look-up sections is not  
287 possible, or introduces bias in the quantification.

288 The main strength of our study is that we were able to perform this experiment and  
289 validate its use in two different research centers. Moreover, all independent observers  
290 were blinded to other reader's data, to the biopsy site, to the state of tissue (fresh-fixed or  
291 frozen-fixed), to the research protocol and to the participant. We also added corrected

292 adipocyte diameter, as proposed by Lenz et al. (2016) and found comparable results. In  
293 additional samples, we also performed an automated analysis of multiple slices covering  
294 150  $\mu\text{m}$  in both depots and conditions and found no difference in cell size distribution or  
295 mean cell size.

296 Previous studies used fresh-fixed tissue for obvious and practical reasons. However, the  
297 utility of frozen-fixed tissue sections of adipose tissue may be of great importance for  
298 researchers in the field of obesity (Ashwell et al. 1975). Some large cohorts may have  
299 only flash-frozen adipose tissue due to various rationales (cell size assessed by  
300 collagenase or by osmium instead of histological analysis, tissues kept only for RNA or  
301 DNA analyses, etc.) and are possibly lacking important *in situ* characterization. This  
302 novel method offers a unique opportunity to study markers of adipose tissue function or  
303 dysfunction in samples that could not be studied before. Investigators have used frozen  
304 adipose tissue in previous publications. However, it was always fixed prior to the  
305 freezing (Berry et al. 2014; Sjostrom et al. 1971) or placed in OCT compounds (Xue et  
306 al. 2010). Yuan and collaborators have published an interesting hypothesis in which they  
307 suggested that freezing adipose tissue at  $-20^{\circ}\text{C}$  without cryoprotectant and thawing it at  
308 room temperature (3 days later) affected cell viability and caused necrosis of the tissue  
309 without damaging whole tissue architecture (Yuan et al. 2015). Our results suggest that it  
310 is in fact possible to observe and characterize *in situ* markers of adipose tissue in flash-  
311 frozen samples stored for at least three years at  $-80^{\circ}\text{C}$ .

312 We anticipate that this approach will allow retrospective studies with flash-frozen human  
313 adipose tissue samples and will shed some light on the importance of adipose tissue in the  
314 pathogenesis of obesity, especially excess abdominal adiposity.

315 **Conflict of interest**

316 AT is the recipient of research grant support from Johnson & Johnson Medical  
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390 **Fig. 1** Image processing for adipocyte size measurements

391 **Fig. 2** Representative images of frozen-fixed and fresh-fixed adipose tissue. (a) Frozen-  
 392 fixed and (b) Fresh-fixed sample from the same woman showing little or no sign of cell  
 393 breakage (representative of 74% and 68% of samples, respectively). (c) Frozen-fixed and  
 394 (d) Fresh-fixed sample from the same woman showing (arrows) signs of cell breakage  
 395 (representative of 26% and 32% of samples, respectively). Scale bars from complete scan  
 396 represented 500  $\mu\text{m}$  and from zoom image, Scale bars 50  $\mu\text{m}$

397 **Fig. 3** Representative images of fresh-fixed (a, b) and frozen-fixed (c, d) adipose tissue  
 398 showing CD68+ cell labeling (IHC) Scale bar: 50  $\mu\text{m}$

399 **Fig. 4** Representative images of frozen-fixed adipose tissue showing CD206+ cell  
 400 labeling (IHC) Scale bar: 50  $\mu\text{m}$

401 **Fig. 5** Representative images of frozen-fixed adipose tissue showing CD14+ cell labeling  
 402 (IHC) Scale bar: 50  $\mu\text{m}$

403 **Fig. 6** Number of CD68-, CD206- or CD14-positive cells per 100 adipocytes in fresh- vs.  
 404 frozen-fixed adipose tissue as measured by IHC

405 **Fig. 7** Adipocyte size frequency distribution from (a) OM fresh-fixed and frozen-fixed  
 406 adipose tissue samples (n=17) and (d) corrected OM, showing similar profiles ( $p>0.05$ ).  
 407 Correlation between adipocyte mean diameter in fresh- vs. frozen-fixed (b) OM ( $r=0.72$ ;  
 408  $p<0.01$ ) and (e) corrected OM ( $r=0.73$ ;  $p<0.001$ ). No significant difference between  
 409 adipocyte mean diameter in (c) OM fresh- vs. frozen-fixed (Student paired t-test;  $p>0.05$ )  
 410 and (f) corrected OM (Student paired t-test;  $p>0.05$ ). These analyses were performed in a  
 411 subset of patients with at least 20 adipocytes per depot measured

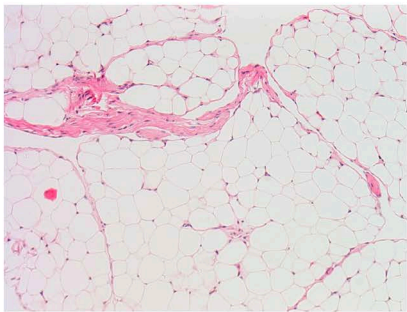
412 **Fig. 8** Adipocyte size frequency distribution from (a) SC fresh-fixed and frozen-fixed  
 413 adipose tissue samples (n=13) and (d) corrected SC, showing similar profiles ( $p>0.05$ ).  
 414 Correlation between adipocyte mean diameter in fresh- vs. frozen-fixed (b) SC ( $r=0.81$ ;  
 415  $p<0.001$ ) and (e) corrected SC ( $r=0.72$ ;  $p<0.01$ ). No significant difference between  
 416 adipocyte mean diameter in (c) SC fresh- vs. frozen-fixed (Student paired t-test;  $p>0.05$ )  
 417 and (f) corrected SC (Student paired t-test;  $p>0.05$ ). These analyses were performed in a  
 418 subset of patients with at least 20 adipocytes per depot measured

419 **Fig. 9** Adipocyte area frequency distribution derived from automated, high-performance  
 420 histological analysis of serial adipose tissue sections covering 150  $\mu\text{m}$  of both (a) OM  
 421 and SC fresh- and frozen-fixed tissues. (b) No significant difference in adipocyte cell size  
 422 frequencies in OM or SC fresh- vs. frozen-fixed was observed (Student t-test;  $p>0.05$ )

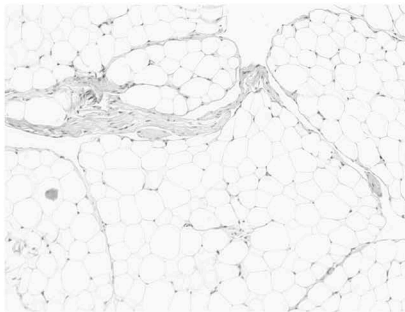
423 **Fig. 10** Adipocyte size frequency distribution from Patient A (a) OM; (b) SC; and from  
 424 Patient B (d) OM; (e) SC; fresh-fixed and frozen-fixed adipose tissue samples. Main  
 425 characteristics are shown for both patients (c, f)

426

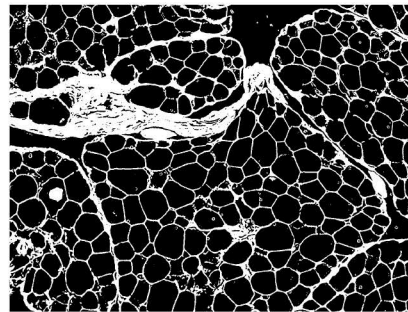




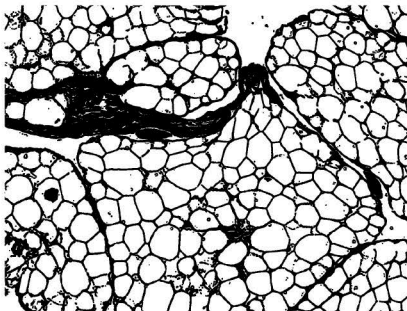
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(RGB-color)



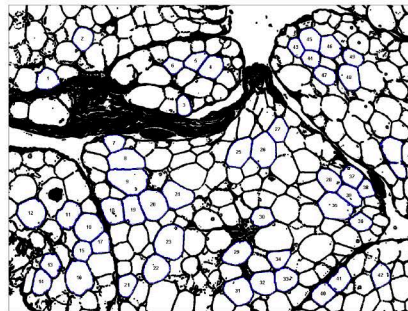
Step 2:  
8-bit, Subtract Background,  
Despeckle



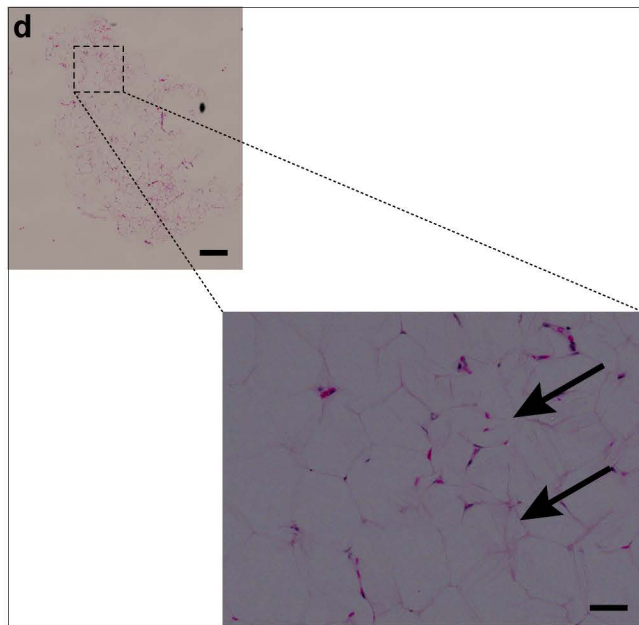
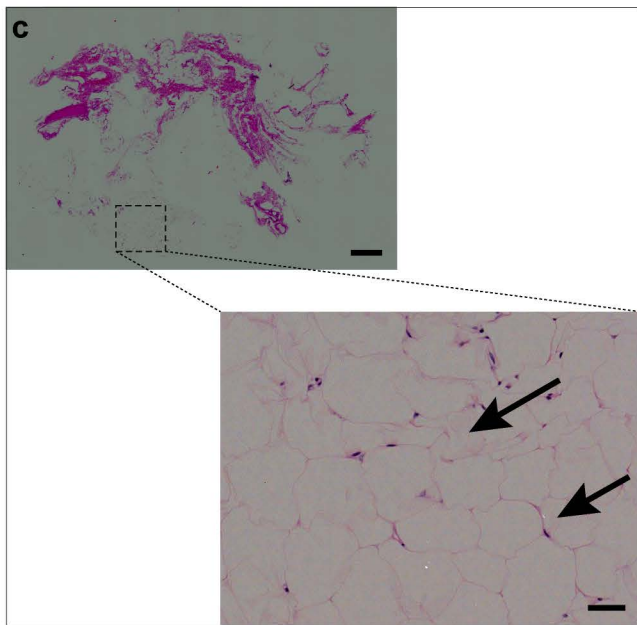
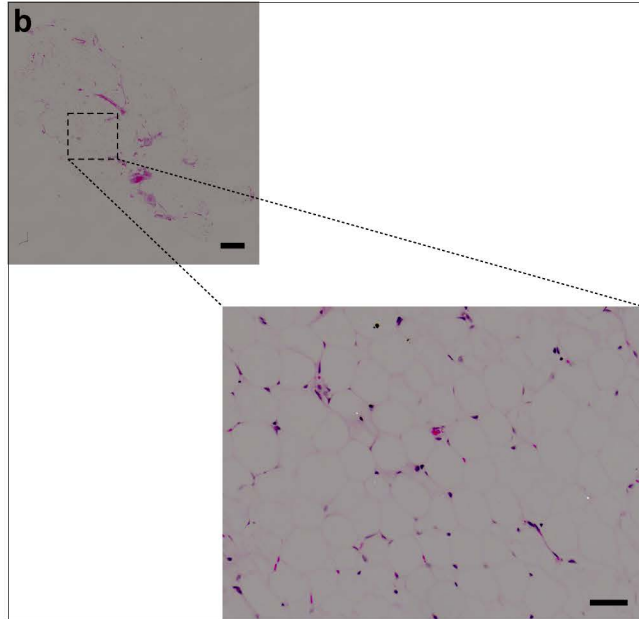
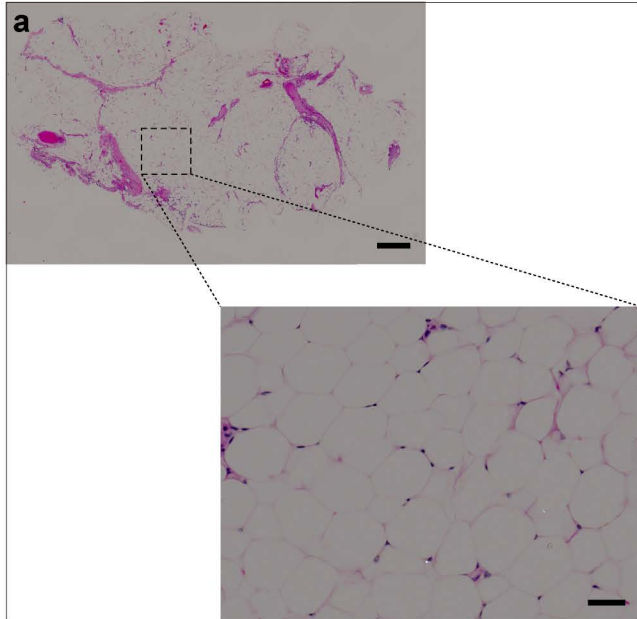
Step 3:  
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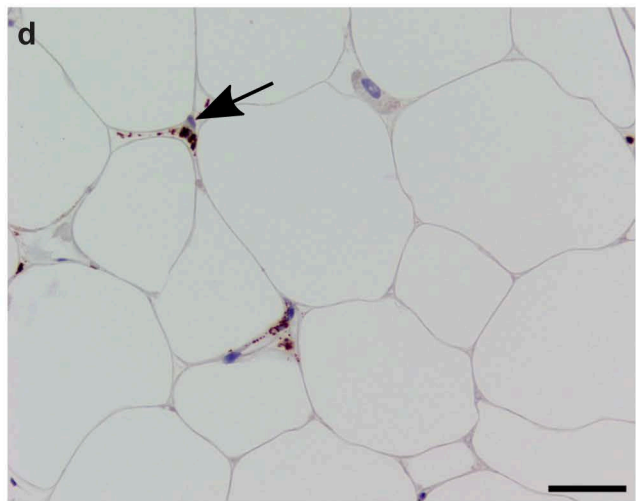
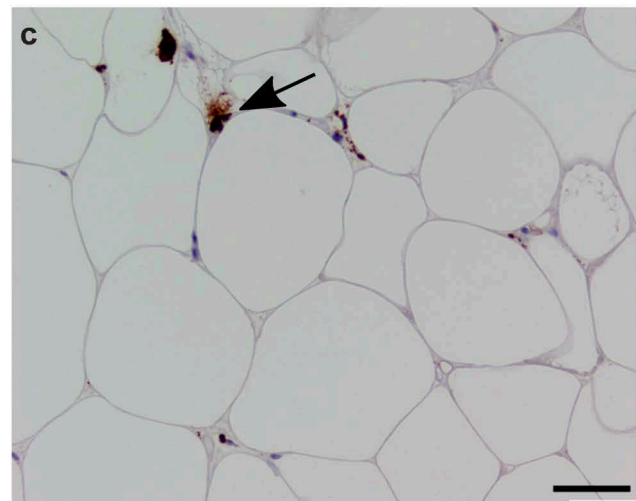
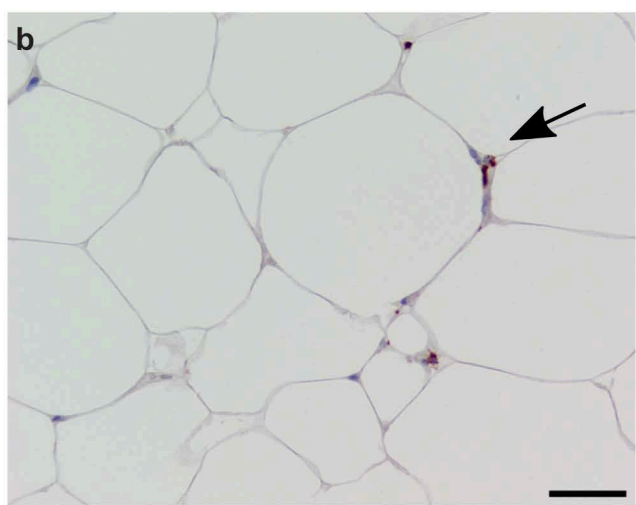
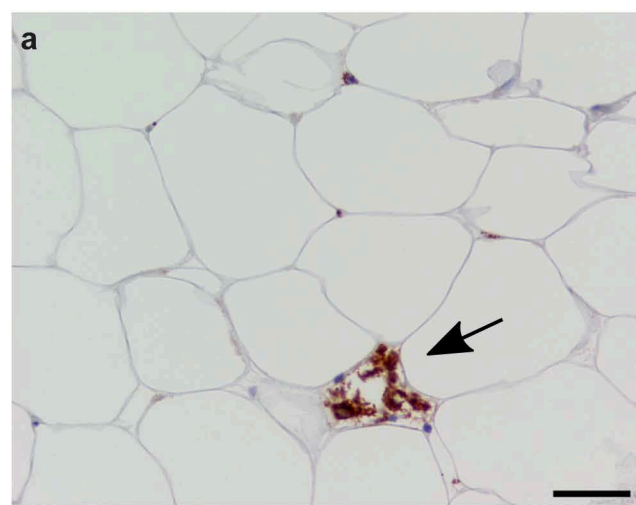


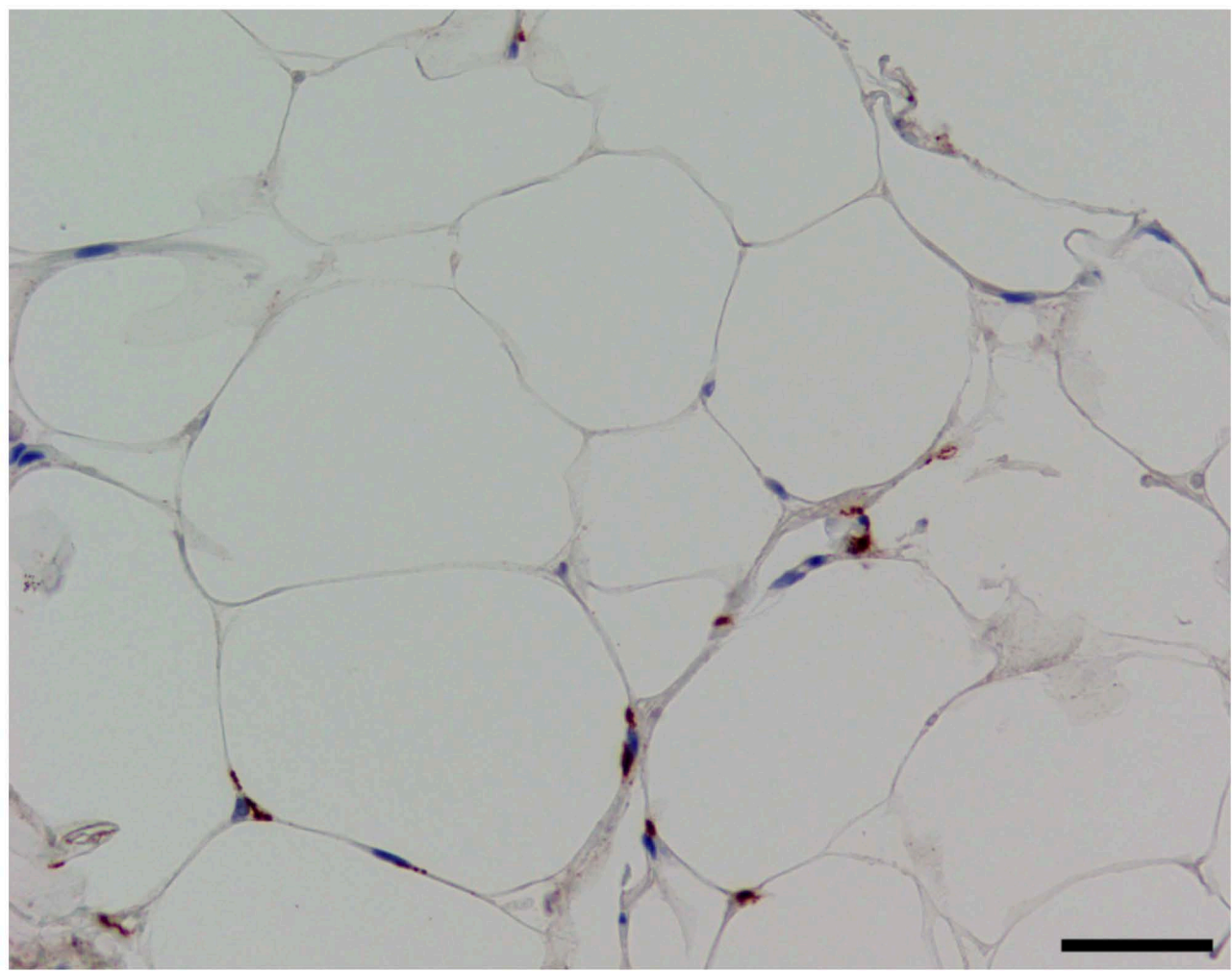
Step 4:  
Make Binary, Dilate

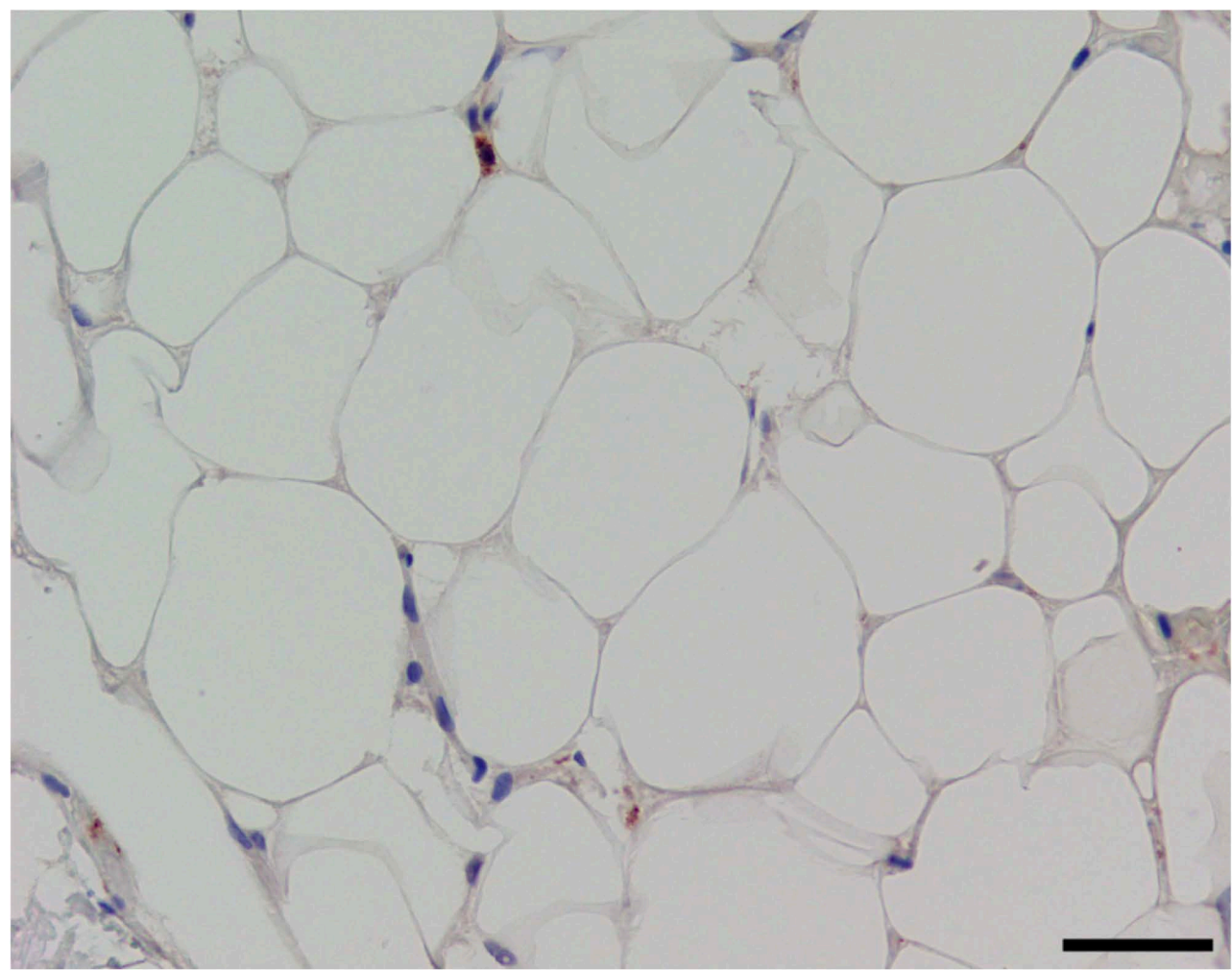


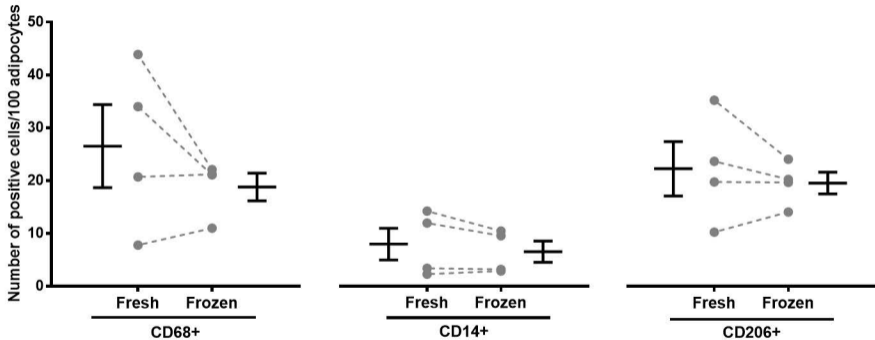
Step 5:  
Quantification



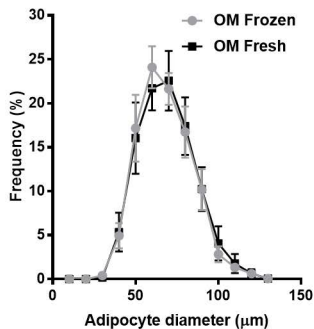
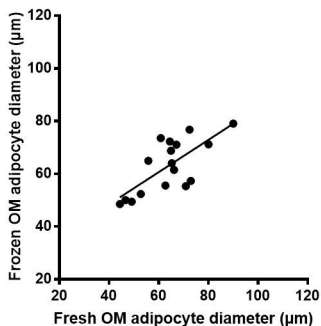
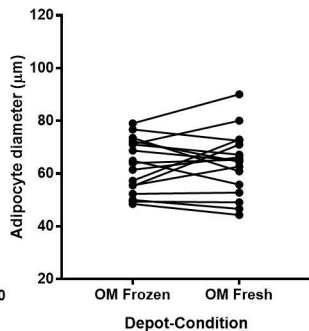
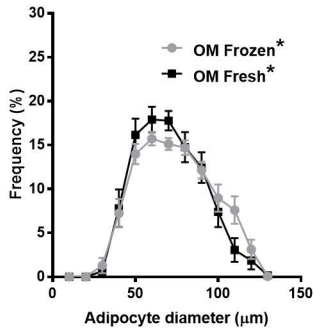
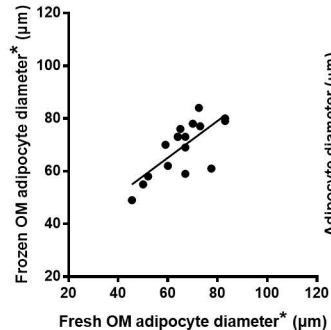
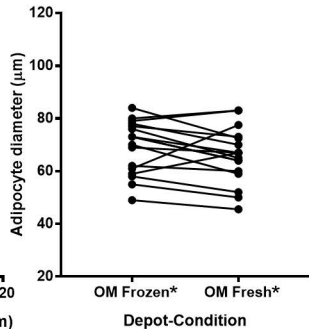


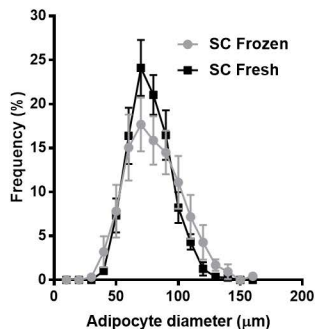
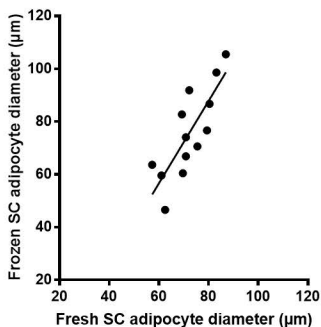
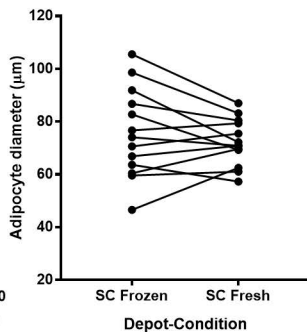
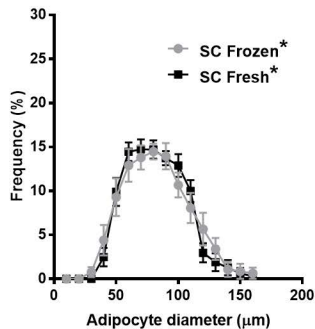
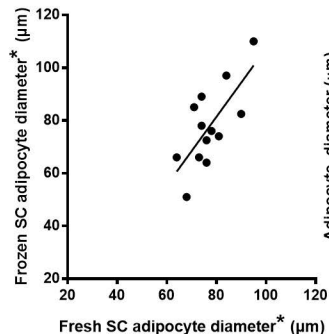
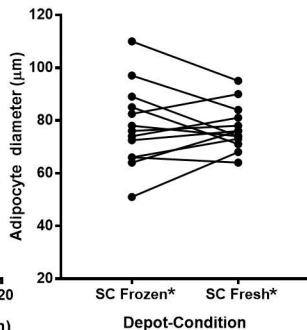




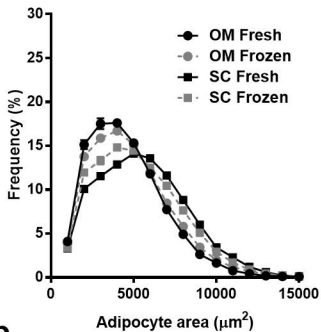
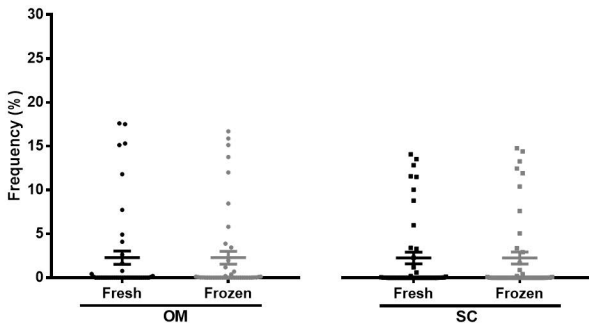


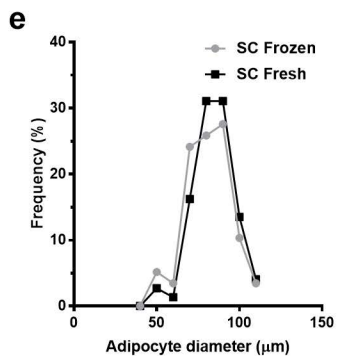
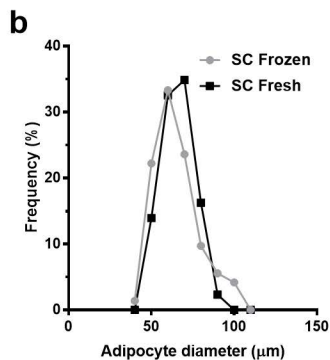
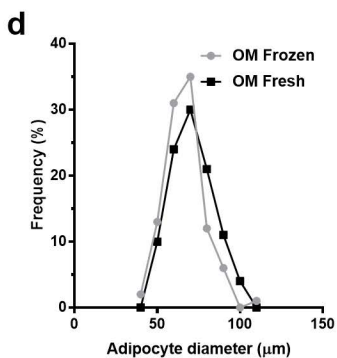
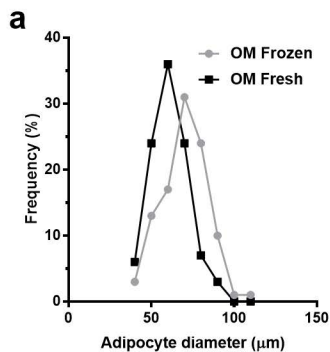


**a****b****c****d****e****f**

**a****b****c****d****e****f**



**a****b**



**c**

Patient A	
BMI ( $\text{kg}/\text{m}^2$ )	23
Age (y)	52
OM mean adipocyte diameter ( $\mu\text{m}$ )	56
SC mean adipocyte diameter ( $\mu\text{m}$ )	61

**f**

Patient B	
BMI ( $\text{kg}/\text{m}^2$ )	29
Age (y)	52
OM mean adipocyte diameter ( $\mu\text{m}$ )	66
SC mean adipocyte diameter ( $\mu\text{m}$ )	79