Visualization and Analysis of Whole Depot Adipose Tissue Neural Innervation

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Summary

Little is known about the diversity and function of adipose tissue nerves due, in part, to the

inability to effectively visualize the various nerve subtypes residing within these tissues. The

tools currently available for researchers to image and quantify adipose tissue innervation are

limited and dependent on optical clearing techniques and light sheet microscopy. Here we

present a method of tissue processing that uses mechanical force to compress tissue to

decrease tissue thickness in the z-axis by expanding it in the x and y-axes whilst leaving cells

intact. This has been combined with autofluorescence quenching techniques to permit intact

whole tissues to be mounted on slides and imaged on any microscope, with a complementary

means to perform whole tissue neurite density quantification. We have included examples of

how this technique can be used to further our current knowledge of adipose-nerve

communication by characterizing the nerves, nerve-subtypes, and neurovascular interactions

within subcutaneous white adipose tissue in mice.

Keywords:

White adipose tissue, innervation, neurovascular interaction, myelination, confocal microscopy,

widefield microscopy, whole mount, maximum projection, clearing, mouse

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Introduction:

Historically overlooked as a location of diverse peripheral nerve innervation [1], the adipose

organ was most prominently inspected for innervation in the mid 1960's when sympathetic

nerve fibers were visualized within brown adipose tissue (BAT) [2]. BAT nerves were later

comprehensively investigated by the Bartness research group [3]. Energy expending BAT was

the first adipose tissue to be identified as being highly innervated due to its important role in

thermogenesis [4], which requires significant sympathetic nervous system (SNS) input and

release of norepinephrine [2, 5, 6]. Sensory innervation has also been documented in BAT,

particularly around vasculature, and has been proposed to play a role in lipolysis [7]. More

recently white adipose tissue (WAT), associated more with energy storage, but which is a

dynamic and plastic tissue capable of undergoing transformation during a cold-induced

"browning" process, was demonstrated to be highly innervated by sympathetic [8, 9] and

sensory nerves [5, 10], but not parasympathetic nerves [11].

In order to visualize adipose innervation, it had been common practice to slice adipose

tissue into 7-10 μm thick sections and immunolabel for various neuronal markers [11-14], which

include the sympathetic nerve activation marker tyrosine hydroxylase (TH); the rate-limiting

enzyme for the synthesis of catecholamine neurotransmitters such as norepinephrine. Several

important findings emerged from this practice, but it was not without limitations. Thin sections of

tissue reduced peripheral nerves to puncta, leaving investigators unable to accurately determine

arborization or the ability to quantify innervation across an intact tissue. Importantly, our

laboratory has revealed a new map of adipose anatomy in the inguinal subcutaneous white

adipose tissue (scWAT) depot, and we and others have demonstrated that the pattern of

innervation in scWAT is heterogeneous across the tissue [15] similar to the heterogenous

browning potential also observed [16]. This warrants a more comprehensive look at adipose

innervation across an intact depot or more anatomically-specific regional changes that may

occur with physiological or pathophysiological stimuli.

Adipose tissue is notoriously difficult to image. Tissue resident lipids scatter light, and

lipofuscin (especially prominent in unhealthy adipose) autofluoresces. Adipose tissue is also

home to a dense vascular network which includes arteries, veins, capillaries, and lymphatics; all

of which are also highly autofluorescent. Adipose depots can also be quite large in comparison

to other organs, especially with the increased adiposity of obesity. Metabolically healthy and

lean mice can have inguinal scWAT depots that are several millimeters thick and this increases

drastically in metabolically unhealthy mice. Because of this, combating tissue autofluorescence

has been a necessity for researchers, especially in whole depot imaging.

Methods for optically clearing and delipidating whole organs for imaging have been

developed, particularly for brains, which are full of light scattering lipids similar to adipose.

These techniques are currently being applied to imaging adipose tissue and specifically imaging

and quantifying innervation within whole adipose depots [15, 17-22]. While useful for getting

crisp images of the large myelinated nerve bundles in adipose, this technique does have its

drawbacks. The current methods typically rely on quantifying only a few small so-called

'representative' three-dimensional tissue sections, which can miss the heterogeneity and

regional anatomy of the intact tissue. Until the regional variation in neurite density in scWAT [15,

20] is better understood, it cannot be accurately reduced to a few representative images or a

small tissue block. Furthermore, some of the most effective methods of tissue clearing use

caustic reagents which, for optimal results, require immersion of specialized objective lenses (ie:

"BABB-safe") in the clearing media for refractive index matching, and many of these shorten

fluorescence lifespan and can be detrimental to endogenous fluorophores. This can be costly

and prevents tissue from being mounted on slides for easy widefield epifluorescence imaging.

Some clearing media can also solidify at room temperature which causes yet another hurdle for

researchers. There is also the separate issue of obtaining homogenous staining through thick

cleared tissue. Traditionally, poor antibody penetration has had to be combatted with drastically

increased incubation times and more recently with stochastic electrotransport [23] and

barometric pressure [24].

For these reasons we have developed an alternative approach to adipose tissue

processing that uses mechanical pressure to homogenously compress entire adipose depots to

reduce tissue thickness. Z-axis compression has been used for other tissues as a means of

reducing tissue thickness to allow for faster image acquisition [25]. We aimed at developing a

method that will be accessible for a wide population of laboratories to use, which is a whole

mount imaging technique that does not require optical clearing or tissue sectioning and can be

used with either standard widefield or confocal microscopy without the need for specialized

BABB-safe objective lenses. This is accompanied by a simple option for quantification of

adipose neurites using methods that we have made publicly available (see Methods). Widefield

microscopy is sufficient for many qualitative analyses of entire adipose depots, but for precise

colocalization and quantitative analysis confocal imaging is required which can pose a

significant increase to image acquisition times (see Methods). However, the cost for light sheet

microscopes can create a barrier to entry that we wish to ameliorate, allowing more research

groups to investigate adipose innervation and the role of adipose nerves in physiology and

pathophysiology.

Like all of the other methods mentioned above, ours also has some drawbacks and

limitations, which is that the tissue depot must be compressed mechanically for it to be imaged

effectively on both widefield and confocal microscopes. Otherwise, our technique presented

here maintains an intact adipose depot and allows for detailed visualization of up to five

fluorescent channels at a time, either from endogenous fluorescent mouse reporter lines or from

fluorescent antibody immunostaining (both of which maintain fluorophore integrity, in this

method).

We have tested this technique using two separate scWAT depots (inguinal and axillary)

which have been suggested to demonstrate different responses to neurotrophic stimuli and

browning potential [15]. This whole mount technique has been optimized for the pan-neuronal

markers PGP9.5 [26, 27] and β3-tubulin [28, 29]; the sympathetic nerve marker, tyrosine

hydroxylase (TH) [14]; markers for sensory innervation that included advillin (AVIL) [30, 31] and

Na_v 1.8 [32]; the myelination marker myelin protein zero (MPZ) [33]; and others. This technique

has also been optimized for non-antibody based fluorescent labeling approaches such as

nuclear labeling with DAPI and vascular labeling with Isolectin IB_4 (IB4). This has allowed us to

further our understanding of scWAT in mice by characterizing the innervation that exists within

this tissue with greater scrutiny, demonstrating neurovascular interactions, parenchymal

innervation, and neuroimmune interactions.

Results and Discussion

<u>Development of the whole mount technique</u>

We have tested numerous clearing techniques with scWAT and BAT (Suppl. Table 1) which

included ScaleA2 [34], BABB [35, 36], CUBIC [37], CUBIC CB-perfusion [37], iDISCO [38],

uDISCO [39], UbasM [40], and a sucrose gradient method [41]. Each clearing method had its

own set of trade-offs, with some methods distorting tissue morphology, limiting fluorescence

lifespan, and requiring costly objective lenses to image (summarized in Suppl. Table 1). Not all

clearing techniques sufficiently clear adipose, as well. These pros and cons have been well

documented in various other tissues [42-45]. To circumvent these issues, we have developed a

method of whole tissue processing, outlined in figure 1a, that allowed for whole adipose depots

to be imaged on any epifluorescence widefield microscope or confocal microscope without the

need for immersion lenses or caustic clearing agents. Entire processing time for an intact

adipose depot with a single directly conjugated antibody takes as little as 6-days. This is

significantly shorter than the majority of adipose clearing methods (10-14days) and is

comparable to the remainder.

This approach has allowed us to image entire tissue depots and construct 2D visualizations

from 3D data acquisition by tiling z-max projection images taken at low magnifications using a

laser scanning confocal microscope (Fig. 1b) and used for quantifying total nerve density (as

described in Methods section) (Fig. 1c). Our quantification method is based on pixel intensity.

By capturing data as a z-max projection we retain the highest pixel intensities from each z-plane

to greatly decrease computation time for the quantitative analyses while keeping potentially lost

data to a minimum. We have developed our quantification method using open access software

and Matlab and all code has been made publicly available (Fig. 1d). To acquire a micrograph of

an entire intact depot on a point scanning confocal microscope times are subject to the desired

resolution, number of fluorescent markers, and size of the tissue. Acquisition time has averaged

13-16hrs when a single fluorophore is imaged at 10X objective magnification per tile at

resolutions of 720x720 and a z-step sampling size of 12μm. We also experimented with

confocal resonant scanning and subsequent denoising with Denoise.ai to reduce image

acquisition time at the cost of resolution (Suppl. Fig 1). A whole depot was imaged in only 2hrs

35min (Suppl. Fig. S1a). Unfortunately, we found that by reducing resolution to image faster, we

lost most of the small neurites when compared to the slower point scanning approach (Suppl.

Fig S1b). Because of this we have determined that higher resolutions and longer imaging times

are required to capture an accurate representation of parenchymal innervation.

Although the use of light sheet microscopy can decrease image acquisition time from hours

to minutes, in our experience (using multiple platforms) imaging an intact whole adipose depot

from various physiological states was extremely challenging with commercially available

models. This led us to develop an adipose tissue processing technique that made imaging

entire intact adipose depots (no need to bifurcate the depot) on a confocal microscope possible.

Whole mount processing of inguinal and axillary scWAT was achieved by first excising intact

depots from the mouse and fixing in 2% PFA overnight. Tissue thickness was reduced to

shorten incubation times, enhance antibody penetration, and reduce light scatter by flattening

the fixed tissue between 2 large glass slides with binder clips to apply gentle pressure (Fig. 1a)

in a process we have termed 'Z-depth reduction'.

Tissues are then quenched for autofluorescence and mounted on glass slides with a

glycerol based mountant. The glycerol based mountant acts, in part, as a clearing agent, though

its use was dictated by the increased coverslip adherence the high viscosity provided. Mounting

in a non-clearing aqueous based mountant offers the same quality images but we found that the

coverslip adherence is poor, resulting in tissue oxidation and subsequent loss of signal, as well

as increased autofluorescence.

To investigate how z-depth reduction altered tissue morphology we first measured the

change in tissue area that occurred by z-depth reduction and subsequent mounting for axillary

and inguinal scWAT depots (Fig. 2a). Z-depth reduction and mounting resulted in a significant

increase to tissue area for both axillary (p=0.0006) and inguinal (p=0.0002) depots (Fig. 2b). In

both cases XY area was increased approximately by a factor of three with tissue area

increasing after z-depth reduction and again when mounted (Fig. 2b). Tissue thickness in the Z-

dimension was measured at the thickest point of each tissue before and after Z-depth reduction.

Tissue thickness was reduced significantly (axillary p=0.0230, inguinal p=0.0192) from

approximately 2-3 mm before Z-depth reduction to less than 200 μm after mounting (Fig. 2c). It

is important to specify that the tissues used here are from metabolically healthy young adult

mice. Tissues from obese or lipodystrophic mice will likely follow the same trend, but exact

changes may be different.

Next, we were interested in how the adipocyte size was affected by the increased tissue

area, and whether cells remained intact. Because tissue area and thickness were altered

similarly between both inguinal and axillary depots we narrowed our analysis to inguinal depots

moving forward. Unstained whole inguinal scWAT depots were either Z-depth reduced and

mounted or remained in their physiological state post-dissection and left unmounted.

Transmitted light was used to visualize unstained cell structure (Fig. 2d). Tissue thickness was

reduced by Z-depth reduction which increased light transmission and softened the cell outlines

visible in the control tissues (Fig 2d). Cell size was averaged across four representative images

of each tissue to quantify possible changes in cell size (30-120 cells counted per image). Only

cells that had clear and in-focus boundaries (given this was not 3D imaging) were included in

the quantification. The average cell area was increased in every instance (p=0.0265) (Fig 2d).

Under closer inspection of individual cells, we found that intracellular gaps were reduced to

make room for the expanding cell area and that cells, due to being pressed against one another,

adopted a polygonal shape (Fig 2e). Hematoxylin staining of 7 μm thick inguinal scWAT

sections showed that cells remained intact and structurally similar following z-depth reduction

(Fig 2f). A caveat of this analysis is that we were only able to investigate cell integrity following

Z-depth reduction and not tissue mounting, which was demonstrated to cause additional strain

to scWAT depots.

Adipose depots are malleable by nature, especially the subcutaneous adipose that melds to

changes in skin pressure, and thus this method of tissue processing takes advantage of that by

increasing tissue area in the X and Y directions to reduce tissue thickness in the Z. This

structural manipulation introduces slight changes to individual adipocytes but leaves cells intact,

not dissimilar to natural changes caused by animal movements such as lying down or

stretching. We believe the benefits of tissue penetration and imaging ease outweigh these

drawbacks, however slight morphological changes to adipose using this approach cannot be

ruled out.

Tissue autofluorescence has always been a significant problem when imaging adipose due

to the inherent autofluorescecent nature of lipids and lipofuscin that reside within it [46]. Sudan

Black B (henceforth referred to by its historic name Typogen Black) staining has been used as a

treatment to quench tissue autofluorescence for decades in 7-10 μm thick sections [47]. We

applied Typogen Black and two other autofluorescence quenching techniques, TrueBlack® and

hydrogen peroxide (used in many optical clearing protocols [42]), and compared quenching

abilities against fluorescence filter cubes spanning the visible spectra: DAPI (blue), GFP/FITC

(green), Cy3/TRITC (orange), and Cy5 (far-red; purple) (Fig. 3a). All methods were able to

reduce autofluorescence in the blue, green, and orange wavelengths similarly, but Typogen

Black was the least effective. The far-red Cy5 channel posed interesting barriers. Though far-

red excitation results in the least autofluorescence of unstained tissue, autofluorescence is

intensified strongly by Typogen Black staining and was left unquenched by TrueBlack®.

Hydrogen peroxide was the only effective means of quenching autofluorescence in all 4-

channels (Fig 3a) but adds significant time to the overall protocol and could be utilized only if all

channels are needed. We next wanted to see how 647nm fluorophores were affected by the

various autofluorescence quenching techniques (Fig 3b). All methods appeared to reduce

fluorophore brightness when compared to the control tissue. Hydrogen peroxide was the least

masking, followed by Typogen Black, and then TrueBlack® which resulted in a very diminished

signal respectively (Fig. 3b).

Although Typogen Black had the least autofluorescence quenching potential it was found to

be sufficient for our uses and was preferred over TrueBlack®, which tended to mask true signal,

and hydrogen peroxide, which increased processing times significantly. For these reasons

Typogen Black was used when not staining with a 647nm fluorophore and TrueBlack® was

used only when a 647nm fluorophore was used concurrently with other fluorophores.

Quenching dilutions were optimized in conjunction with pan-neuronal PGP9.5 staining to

find the ideal balance between reduced tissue autofluorescence and undiminished true

fluorescent signal (Suppl. Fig. S2a). Optimal concentrations, as determined by autofluorescence

reduction and signal intensity, were found to be 0.1% Typogen Black (in 70% EtOH) and 1X or

0.1% TrueBlack® (2% TrueBlack® dye in 98% dimethyl formamide, diluted from 20X to 1X with

70% EtOH.)

An additional means of reducing tissue autofluorescence was performed by washing with 1X

PBS with 10U/mL Heparin, which removed highly autofluorescent red blood cells (Suppl. Fig.

S2b). Alternatively, perfusion can be performed to remove blood cells prior to tissue collection.

Though the use of a glycerol-based mountant was decided upon due to its increased viscosity, it

is also a known clearing agent [48], and we found that adding only a couple of drops to the

tissue was enough to provide some level of optical clarity (Fig. 2a). This clearing was made

more evident when tissues remained at room temperature for incubation and wash steps, prior

to mounting, which resulted in reduced background autofluorescence (Suppl. Fig. S2c). It

should be noted that autofluorescence when imaging with a widefield epifluorescent microscope

is still quite present even after application of the aforementioned quenching techniques.

Because of this, it is strongly recommended that confocal microscopy be used whenever

possible to further reduce this background and non-specific signal.

To summarize: Typogen Black should be avoided when using far red excitation. TrueBlack®

can be an acceptable replacement but it can mask fluorescent signal so avoid using with dim

fluorophores. Hydrogen peroxide yields the best results all around but adds significant time to

an already lengthy procedure. If only using one fluorophore it is best to use far-red excitation

and do all washes and incubations at room temperature to avoid the use of autofluorescence

quenching reagents entirely.

Secondary antibody non-specificity is likely to increase when immunostaining an entire

tissue, and when primary-conjugated antibodies are not available or specific enough. We tested

numerous secondary antibodies to confirm that we were only using those antibodies least likely

to bind non-specifically (Suppl. Fig. S2d). Goat anti-rabbit IgG secondary antibodies, regardless

of being highly cross adsorbed or not, did not result in significant levels of non-specific binding.

Because goat anti-mouse secondary antibodies had such high levels of non-specific binding, we

decided to avoid the use of all mouse primary antibodies for whole mount immunostaining

moving forward.

Uniform staining of large whole mount tissues has always been problematic due to the

increased incubation times required for antibodies to completely diffuse throughout the tissue

[23, 49]. Incubation times were reduced by 1-2 days by reducing tissue thickness with z-depth

reduction. Depth coding of immunostained scWAT following z-depth reduction indicated

homogenous penetration of antibodies throughout the whole tissue (140µm total depth) (Fig.

3c). This was confirmed for various fluorescent staining approaches, including directly labeled

β3-tubulin, indirectly labeled TH, and IB4 staining. Depth information can even be somewhat

retained following z-maximum projecting by creating a height color coded projection, though no

accompanying color-coded depth scale is provided at this time (Suppl. Fig. S3). Although we

have not tried this ourselves, the addition of sucrose to 4% paraformaldehyde fixation can

improve axonal staining by decreasing the appearance of "beading" along the axon [50].

<u>Characterization of peripheral nerves in scWAT</u>

An inguinal scWAT depot was co-stained with two different pan-neuronal markers (PGP9.5

and β3-tubulin) and counterstained with DAPI to emphasize tissue cellular structure, and tiled

images were z-max projected.

To demonstrate the versatility and capability of this technique we imaged tissues in two

ways. Firstly, the whole adipose tissue was imaged at low magnification (5X) with limited

resolution to observe overall morphology and fluorescence of the larger nerve bundles (Fig. 4a).

Secondly, to visualize detailed innervation, areas of interest of the same tissue were imaged at

63X with much higher resolutions (Fig. 4b-c).

Whole tissue imaging exhibited near overlap of staining of the largest nerve bundles by both

pan-neuronal markers (Fig. 4a). Nerves are unevenly distributed throughout the tissue in the X

and Y-axis, and in the anterior-posterior anatomical directions. Because the tissues are

compressed and subsequently z-max projected we were unable to accurately report nerve

distribution within the z-axis. The highest concentration of large nerve bundles were present in

the centermost third of the tissue (in the x-y direction), surrounding the subiliac lymph node

(SiLN), as reported previously [15]. At greater magnification, z-max projection images showed

an extensive network of neurites varying in diameter, running throughout the tissue and

surrounding adipocytes as well as vasculature (Fig. 4b), also as previously noted[15]. Again,

there appears to be almost complete fluorescent overlap of both pan-neuronal markers, with

PGP9.5 being more readily visible on the smaller nerve fibers, which fits with the use of this

antibody to assess small fiber neuropathy in the skin. Variation in immunofluorescence staining

between the two pan-neuronal markers can be attributed in part to β3-tubulin being directly

labeled and PGP9.5 being indirectly labeled (and thus, some off-target binding of the secondary

antibody). Having a reporter mouse line instead of relying on antibodies would improve this

technique. The innate autofluorescent quality of adipose tissue, though greatly reduced by

Typogen Black or TrueBlack® staining, is still visible in most images and can help provide

anatomical outlines. In fact, in most adipose clearing techniques in the published literature,

autofluorescence of the tissue is exploited to reveal cellular architecture [20].

Digital cross section imaging of large nerve bundles demonstrated complete staining of the

bundle but with gradation resulting in a weaker signal at the bundle's center (Fig. 4c). Paraffin

embedded cross sections do not show the same gradation (Suppl. Fig. S4) , so we have

attributed this observation to incomplete antibody penetration into the bundle. DAPI stained

nuclei can be seen both surrounding the nerve bundle and residing between the numerous

axons (Fig. 4d) (Suppl. Fig. S4). Since the nuclei of the neurons are located in the ganglia of the

spine, the presence of nuclei within the nerve bundle suggests the presence of supporting cells

(e.g. Schwann cells), immune cells, or potentially perineural adipocytes residing within the

bundles.

Transmission electron microscopy (TEM) of scWAT sections revealed that putative nerves

not only traverse through the tissue (Suppl. Fig. S5a) but may also come in direct contact with

the adipocytes (Suppl. Fig. S5b-c). However, we have not yet confirmed synapses or tissue

junctions directly onto adipocytes. Potential synapses have been observed in the stromal

vascular fraction (SVF) of perigonadal adipose tissue (Suppl. Fig. S6a) as well as on blood

vessels (Suppl. Fig. S6b), and on myeloid lineage SVF cells in inguinal scWAT (Suppl. Fig. 6c)

using the post-synaptic marker PSD95. In general, the understanding of adipose tissue

innervation has not extended to characterizing cellular interactions and whether they are

synaptic (or a junction similar to the neuromuscular junction), or simply result from diffusion of

neuropeptides and neurotransmitters from nearby free nerve endings.

To further characterize the nerves within scWAT, numerous immunostaining experiments

were conducted by co-staining β3-tubulin with markers for either sympathetic nerves, sensory

nerves, or myelination using the markers tyrosine hydroxylase (TH), advillin (AVIL), and myelin

protein zero (MPZ) respectively. Whole mount imaging demonstrated fluorescence overlap of

β3-tubulin and TH in the largest nerve bundles with extensive TH+ axons that spanned

throughout the tissue (Fig. 5a). Nerve bundle digital cross sectioning showed that only some of

the axons in each nerve bundle were TH+ (Fig. 5b). The small individual axons branching

through the tissue, however, were nearly all TH+ (Fig. 5c), consistent with previous reports [11,

22].

Digital cross sectioning of AVIL+ sensory nerve bundles revealed similar findings to cross

sectioning TH+ bundles, with only some of the axons within bundles presenting as AVIL+ (Fig.

5d). This further suggested the presence of mixed nerves in adipose. It is worth noting that the

AVIL antibody used here was observed to mark structural proteins in some large blood vessels

as well, which accounted for the presence of AVIL+ regions that were β3-tubulin- when viewed

as a tiled image. At greater magnification these large blood vessels did not interfere with

imaging, and the lack of AVIL+ nerve fibers became apparent. Due to this, AVIL would appear

to be a less-than-ideal marker for quantifying sensory innervation in conditions that may also

alter vascularity. It is thought at this time that an AVIL reporter mouse or a Na_v 1.8 reporter

mouse would be a far superior method of labeling sensory axons.

Therefore, *Na_v 1.8-Cre x tdTomato* reporter mice were also used to investigate the presence

of sensory innervation in scWAT. Na_v 1.8 marks sodium channels specific to sensory nerves

[32] and fluorescence imaging showed a number of Na$_v$ 1.8+ sensory nerves throughout the

tissue (Suppl. Fig. S7a). These were both large bundles and smaller parenchymal fibers, which

complements our own results with AVIL imaging and current literature suggesting sensory

innervation plays a significant role in WAT metabolic function [14, 51]. As observed with AVIL

immunostaining, imaging of inguinal scWAT from *Na$_v$ 1.8-Cre x tdTomato* reporter mice also

suggest the presence of mixed nerve bundles in this adipose depot (Suppl. Fig. S7a).

Whole mount staining with β3-tubulin and MPZ revealed that nerve bundles 25 µm and

greater in diameter in scWAT are myelinated (Fig. 5e). In concordance with immunostaining,

Luxol fast blue (myelin stain) staining of whole inguinal scWAT depots was performed which

showed that the most highly myelinated nerves traverse through or nearby the SiLN (Suppl. Fig.

S7b). It is not yet clear if these are mainly sensory axons that are myelinated, or a mix of

sensory and sympathetic. Z-max projection imaging of parenchymal nerve fibers revealed that

the majority of the smaller nerve endings are unmyelinated (Fig. 5f). Digital cross sectioning of

MPZ+ nerve bundles showed that all of the axons within the bundle appeared myelinated, with

the greatest fluorescence intensity being on the exterior of the bundle (Fig. 5g). At this time, it is

thought that the gradation of the MPZ staining is caused by incomplete antibody penetrance but

it cannot be ruled out that the exterior axons of these bundles are more myelinated than the

interior axons. This potent MPZ+ staining on the bundle periphery also acts to mask some of the

β3-tubulin+ nerves that reside within. TEM imaging was used to show both myelinated and

unmyelinated putative neurites in contact with adipocytes (Suppl. Fig. S5b-c).

Our whole mount imaging technique has validated the presence of mixed bundle nerves

residing in WAT as shown previously [17] and allowed for further investigation by creating digital

cross sections of these mixed bundles. High magnification z-max projections revealed that the

majority of neurites within the scWAT parenchyma are TH+, however, this was shown using TH-

antibody which is known to be somewhat non-specific. Studies should be redone using a TH-

reporter mouse to be certain. A relative minority of axons in the tissue are myelinated, and those

that are reside mostly in large nerve bundles or run in close proximity, however, the contribution

of myelinated versus unmyelinated axons to tissue function is currently unknown. Quantified

colocalization of the various neuronal markers outlined above (PGP9.5, β3-tubulin, TH, MPZ,

AVIL) at the whole tissue level and on nerve bundle cross sections can serve as another route

for continued investigation.

<u>Neurovascular interaction in scWAT</u>

The autonomic nervous system, comprised of sympathetic, parasympathetic, and

sensory nerves, is required for regulating vascular tone [52] throughout the body, and WAT is

no different. Generally, parasympathetic fibers are responsible for vasodilation and sympathetic

nerve fibers are responsible for vasoconstriction, though this response is receptor specific and

sympathetic nerves can mediate vasodilation as well [52]. It has been suggested that WAT

lacks parasympathetic innervation [11] indicating that precise control of vasodilation is either not

required in WAT or regulated entirely by sympathetic and sensory innervation. Neurovascular

staining of whole mount scWAT was performed by co-staining with pan-neuronal markers and

IB4, a marker for vasculature, as it binds to erythrocytes and endothelial cells [53-57] and

effectively marks vessels smaller than 50 μm in diameter. Wholemount axillary scWAT tiled z-

max projections exposed a dense vascular and lymphatic network residing in the scWAT depot

(Fig. 6a). In general, the highest concentration of nerve bundles and blood vessels that are

greater than 25 μm in diameter are located within close proximity of one another. These

observations are consistent with what we have observed in inguinal scWAT [15]. Close

inspection of the neurovascular interactions within scWAT revealed three specific and

reoccurring types of interactions (Fig. 6b): i) Large nerve bundles and blood vessels that run

next to each other within the tissue but do not seem to interact or intersect, ii) Large nerve

bundles that have a vascular supply, likely providing nutrients; or a vasa nervorum, iii) Blood

vessels that are highly innervated by smaller neurites which may regulate vasoconstriction; like

a perivascular sympathetic plexus. In support of this, TEM imaging revealed a small putative

axon in the stromal vascular fraction in between adipocytes in close proximity to a capillary

(Suppl. Fig. S5a). Sympathetic innervation of scWAT blood vessels was also analyzed by co-

staining β3-tubulin with TH and IB4. Small TH+ nerve fibers were found lining many of the

vessels (Fig. 6c). Although small capillaries had significantly less innervation compared to larger

arterioles, some TH+ nerves were found running along them. This supports the current literature

that capillaries are relatively lacking in sympathetic innervation, whereas larger vessels have

more sympathetic innervation [58-60]. Larger blood vessels likely also have sensory innervation

as well [61].

Not all nerves within the tissue are lining blood vessels; many can be found branching

the gaps from one blood vessel to another or disassociated from the blood vessels entirely. The

vasa nervorum, or the blood vessels supplying nutrients to the nerve bundle, were digitally cross

sectioned to show a blood vessel branching around a nerve bundle which contained 2

sympathetic axons (Fig. 6d). Extensive innervation of arterioles could be found surrounding both

large and small diameter vessels (Fig. 76e). Of the nerves running throughout scWAT, the

majority of MPZ+ nerves were found running along blood vessels and also had nuclei directly

contacting them, as indicated by co-staining with DAPI, β3-tubulin, MPZ, S100β (a Schwann cell

marker), and IB4 (Fig. 6f). Only one or two of the largest nerves lining each vessel tended to be

myelinated. The smallest sympathetic projections that often engulfed many of the arterioles,

when present, were found to be unmyelinated.

It is important to note several caveats with IB4 staining of vasculature. Mouse blood

vessel diameter ranges from 250μm to 4μm, depending on type of vessel. Veins encompass the

upper limit, while microvasculature such as capillaries encompass the lower limit [62]. Large

diameter blood vessels (>100um) often lose their erythrocytes during tissue preparation.

However, erythrocytes tend to remain in the microvasculature. Because of this, the

microvasculature is doubly stained (erythrocytes and endothelium) whereas arteries and veins

present only with stained endothelium. This reduction in fluorescence, most prominent in the

largest vasculature, can even be seen starting as small as 50 µm in diameter though these

vessels are still readily visible (Fig. 6). Also, large sensory nerve bundles, immunostained for

AVIL, an actin binding protein expressed specifically in somatosensory neurons [30, 31], tended

to be co-stained with IB4. IB4 binding non-specifically to sensory neurons has been reported in

the literature [63, 64]. However, only large sensory nerve bundles were observed to be labeled

by IB4, eliminating the risk of misidentifying capillaries and, due to tissue morphology, positively

stained nerve bundles can be easily distinguished from the vasculature.

<u>Whole mount imaging allows for quantification of total depot innervation and vascularity</u>

477   Axillary scWAT depots taken from male C57BL/6J mice, housed at room temperature or

cold exposed at 5ºC for 7-days, were processed for whole mount imaging using the pan-

neuronal marker PGP9.5 (Fig. 7a). The neurite densities per tile were then averaged for the

entire area of the tissue for comparison among treatments. Quantification of images allowed for

statistical analysis between experimental groups which showed that three of the four RT mice

had less neurite density than all of the cold exposed mice but that overall this difference was not

statistically significant (p=0.414) (Fig. 7b). This is similar to our previous studies using inguinal

scWAT depots which used an alternative, but complementary, approach to whole tissue neurite

density quantification [15]. Whole depot quantification might blunt regional differences in

innervation, that may occur, for example, around the lymph node. Skeletonizing and quantifying

neurite density from z-maximum projections, by definition, results in under sampling data. To

verify the extent of this potential loss of spatial information, we calculated the neurite density on

the individual z-slices that make up the 3D confocal data cube and then synthesized these

quantities in two different ways: by taking the average neurite density score across the confocal

stack (Suppl. Figure S8a), or by taking the maximum neurite density score across the confocal

stack (Suppl. Figure S8b). In both cases, there is a very high correlation between the

synthesized neurite density score (either obtained by averaging or by taking the maximum) and

the neurite density score obtained from the z-max projection image ($p<0.0001$.) Of course, this

does not allow us to conclude that no information is being lost by restricting the analysis to 2D z-

max projections. However, it does allow us to conclude that the potential gain that one might

make by considering all of the individual 2D slices instead of only the z-max projection would

have to be explored beyond simple statistical measures such as taking the average or the

maximum. And additionally, this potential gain in information would have to be weighed against

the additional computational analysis time and effort. Neurite density per z-max tile was

portrayed as a heatmap to better visualize region specific differences in neurite densities (Fig.

7c). This quantification and analysis approach can also be applied to vasculature (Fig. 7d).

<u>Second harmonic generation imaging</u>

We also visualized the changes to scWAT innervation of morbidly obese mice which we

have demonstrated to have adipose neuropathy (BTBR$^{ob/ob}$); [15]. This proved technically

difficult to accomplish using epifluorescent or standard confocal microscopy, largely due to the

fibrotic nature of obese adipose tissue that led to severe autofluorescence that could not be

mitigated with quenching. However, with second generation harmonic label-free imaging using

2-photon microscopy, it was possible to visualize nerves and collagen within scWAT of

BTBR$^{ob/ob}$ animals (Suppl. Fig. S9). While collagen is abundant in both BTBR$^{+/+}$ and BTBR$^{ob/ob}$

animals, a greater degree of colocalization and apparent alignment between collagen fibers and

nerves is visible in BTBR$^{ob/ob}$ animals (Suppl. Fig. S9), likely contributing to our inability to image

the adipose nerves successfully in this mouse model.

Conclusion

The whole mount processing and imaging technique presented here is meant to broaden

the imaging capabilities currently available to researchers. We do not present a method meant

to fully replace the need for optically clearing adipose depots or the use of light sheet

microscopy but wish to provide an alternative method when such imaging techniques are not

required or available. We have presented data to support that these more complex techniques

are in fact not always required to get high quality images of adipose innervation. We have

demonstrated how our method can be applied to imaging on affordable widefield

epifluorescence microscopes as well as laser scanning confocal microscopes, to illustrate that

this method of tissue processing can be applicable to researchers regardless of the tools they

have available. The ease and accessibility of being able to take an entire adipose tissue and

mount it on a slide to be imaged on any microscope cannot be overlooked.

Though the time to image an entire depot at resolutions sufficient for analyzing the

smallest parenchymal neurites is substantial, this is only necessary when whole tissue

quantifications are desired. For all other instances capturing a traditional single tile micrograph

will take only seconds. Due to the broad adaptability of this technique, it is even possible to

analyze gross tissue morphology and aberrant changes in the entire tissue in only a couple of

hours, if desired, by lowering magnifications, decreasing resolutions, or by using a resonant

scanning microscope.

The data collected herein reveal a rich and diverse neural innervation in mouse inguinal

scWAT, likely indicating physiological roles that are yet to be uncovered. For example,

differences in myelinated vs unmyelinated axons, heterogeneity of TH+ nerves, and the

contribution of sensory nerves in adipose are yet to be clarified. In addition, the localization of

true synapses or tissue junctions, versus release of nerve products from free nerve endings,

and whether these exist on adipocytes or cells in the stromal vascular fraction (SVF) of the

tissue, will also be important to uncover. The availability of new techniques allowing nerve

visualization under various physiological and pathophysiological conditions will help the field

advance an understanding of adipose innervation and brain-adipose communication, including

neurovascular and neuroimmune contributions, and plasticity of neurites in adipose.

We have recently demonstrated diminished innervation of adipose tissue under

circumstances of obesity, diabetes, and aging, a condition we have called 'adipose neuropathy'

[15]. Neuropathy of adipose tissue (ie: a reduction in neurite density that may represent

pathological nerve die-back) could have effects beyond the removal of sympathetic nerves

releasing norepinephrine. Given that other nerve products, including from adipose tissue

sensory nerves (ex: Substance P, VIP, CGRP, etc.) are found in adipose depots, the loss of

proper innervation could also impact the physiological contributions of these neuromodulatory

substances in adipose. Whether or not adipose sensory nerves directly communicate fuel status

to the brain, perhaps in complement to endocrine factors such as leptin, is still an open

question. Incoming sympathetic nerves may release norepinephrine in order to affect

vasoconstriction, and also to impact local adipocytes and immune cells via synaptic and non-

synaptic connections. We hope that by providing our method as a resource for investigators that

these questions can be answered.

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Author Contributions

JWW wrote the manuscript, created and optimized protocols, designed experiments, and conducted the majority of the microscopy studies and technical troubleshooting/optimization. MB wrote the manuscript, designed experiments, and optimized protocols. EG conducted TEM experiments. AL and ALD conducted clearing experiments. SB and AK developed neurite density quantification methods. WPB and KBT conducted 2-photon imaging. KLT wrote the manuscript, designed experiments, analyzed data, and oversaw the project.

Declarations of Interests

The authors declare no competing interests.

Figure Legends:

Figure 1: Adipose tissue processing, imaging, and quantification method. Diagram depicting steps of wholemount tissue processing technique (a). See methods for detailed description of each step. Tiled z-stacks were imaged for the entire tissue with either a 5X or 10X objective and a 2D maximum projection image was rendered (b). Post processing of single tile z-max projections for neurite density quantification (c). Each z-max projection tile was further processed by subtracting the background away and adding light smoothing. Next a thresholded mask was applied and the image was skeletonized. Total neurite length was calculated for each tile and averaged for the entire viewing area (c.) Link to whole mount protocols and script for quantification (d.)

Figure 2: Z-depth reduction and tissue morphology. Axillary and inguinal scWAT depots

excised from adipose were Z-depth reduced and mounted on slides (a). Tissue area was

measured before Z-depth reduction, after z-depth reduction, and after mounting (b). N=3, paired

two-tailed Student's t-test, alpha level 0.05, error bars are SEMs (b). Tissue thickness in the Z-

axis was measured at the thickest point before and after Z-depth reduction and mounting (c).

Mounted tissues were measured by 3D projecting tissue autofluorescence captured on a Nikon

A1R with a 20X objective (c). N=3, paired two-tailed Student's t-test, alpha level 0.05, error bars

are SEMs (c). Inguinal scWAT was either whole mount processed (Z-depth reduced) or left

unaltered (control) and placed on a slide and imaged with transmitted light (d). Four

representative 10X micrographs were captured for each tissue and the area for 30-120 cells

was measured per micrograph and analyzed using a two-tailed Student's t-test, Alpha level

0.05, error bars are SEMs (d). 3.5X Digital zoom of 10X transmitted light micrograph illustrating

difference in cell shape (c). Hematoxylin staining of 7 µm inguinal scWAT sections that either

received z-depth reduction before paraffin embedding or did not (control)(f). Transmitted light

micrographs were captured on a Nikon E400 upright microscope (d-f). Scale bars are 186 µm

and 174 µm (c), 200 µm (d,f) and 50 µm (e).

Figure 3: Whole mount adipose autofluorescence quenching and fluorescence staining.

scWAT depots excised from *C57BL/6J* mice were whole mount processed and had

autofluorescecne quenched with either Typogen Black, TrueBlack®, or 5% hydrogen peroxide.

Autofluorescence was evaluated for each blocking method for 4 different fluorescent filter cubes

spanning the visible spectra: DAPI, GFP/FITC, Cy3/TRITC, and Cy5 (a). Control tissue

brightness was normalized in-post across all four filters and these settings were then applied to

each image for a given filter: DAPI (used as control brightness), GFP/FITC (+10% brightness),

Cy3 (+20% brightness), Cy5 (+40% brightness). Quenching methods were further compared for

the Cy5 filter by staining the tissues with the vascular marker isolectin IB_4 (IB4) conjugated to a

647nm fluorophore (b). Inguinal scWAT depot was whole mount processed and stained with β3-

Tubulin, Tyrosine Hydroxylase (TH), and IB4 and depth coded (c). Whole depot was imaged as

a tiled z-maximum projection with a 10X objective, (0-140um), a single tile captured at 10X (0-

45um), and a high magnified section captured at 63X (0-70um) (c). Captured on Leica TCS SP8

DLS microscope (c). Scale bars are 200 µm (a), 200 µm and 50 µm (b). X and Y axes are 35

mm (c, whole mount), 1.1 mm (c, 10X) and 180 µm (c, 63X).

Figure 4: Peripheral innervation of scWAT. Inguinal subcutaneous depots from *C57BL/6J*

mice were whole mount processed and stained with DAPI (blue) to show tissue morphology and

the pan-neuronal markers β3-Tubulin (green) and PGP9.5 (red) captured on Leica TCS SP8

DLS microscope (a-c). Whole inguinal scWAT depot tiled z-max projections captured using 5X

objective (a). White boxes show single tile expanded for visualization

with 3.57X digital zoom-ins (i-ii). Representative images of small fiber innervation imaged as z-

max projections with 63X objective (b). Nerve bundle imaged with 63X objective and digital

cross sectioning performed with a 2.54X digital zoom (c). Nuclei on surface of nerve bundle and

interlaced between nerve fibers captured in widefield on Nikon E400 with 60X objective (d).

White box was digitally zoomed 3.9X (d). Scale bars are 8 mm (a) and 25 µm (b-d).

Figure 5: Characterization of whole mount scWAT nerves. scWAT depots from *C57BL/6J*

mice were wholemount processed and stained with β3-Tubulin (green) and either TH (red) (a-c),

AVIL (red) (d), or MPZ (red) (e-g). Whole mount tiled z-max projections of inguinal scWAT (a,d)

and axillary scWAT (g) imaged with 10X objective. White boxes show a single tile expanded for

visualization 5.75X (a,e). Nerve bundle cross sections imaged with 63X objective with a 2.54X

digital zoom applied to the digital cross section rendering (b,e). Representative images of small

fiber innervation imaged as z-max projections with 63X objective (c,f). All images captured on

Leica TCS SP8 DLS confocal microscope (a-g). Scale bars are 2 mm (a), 25 μm and 10 μm (b),

25 μm and 50 μm (c), 50 μm and 10 μm (d), 5 mm (e), 25 μm (f), an 25 μm and 10 μm (g).

Figure 6: Neurovascular interaction within whole mount scWAT. Inguinal scWAT from

C57BL/6J mice were wholemount processed. Whole axillary scWAT depot stained with PGP9.5

(green) and IB4 (white) was captured as tiled z-max projections imaged with 10X objective on

Leica TCS SP8 DLS microscope (a). White box shows a single tile expanded for visualization

7.33X (a). PGP9.5 (green) and IB4 (red) used to display the 3 categories of neurovascular

interaction within scWAT, imaged on Nikon E400 with 4X, 10X, and 40X objectives respectively

(b). Representative images of the innervation of blood vessels imaged as z-max projections with

63X objective. β3-Tubulin (green), TH (red), IB4 (white) (b-d). Sympathetic innervation of small

blood vessels (c). Vasa nervorum of inguinal scWAT nerve bundle imaged as z-max projection

digitally cross sectioned with a 2.54X digital zoom (d). Large and small diameter blood vessel

digital cross sectioning (e). 2.54X digital zoom applied to large diameter vessel cross section,

6.96X digital zoom applied to small diameter vessel cross section (e). Representative images of

myelinated nerve fibers around vasculature imaged as z-max projections with 63X objective.

Nuclei in direct contact of myelinated nerve bundles present in both images and white boxes

expanded for visualization 3.67X (f). DAPI (blue), β3-Tubulin (green), MPZ (yellow), S100β

(red), IB4 (white) (f). Scale bars are 2 mm (a), 500 µm (b-i), 200 µm (b-ii), 50 µm (b-iii), 25 µm

(c), 25 µm and 10 µm (d), 25 µm, 10 µm, and 5 µm (e), and 25 µm (f).

Figure 7: Whole depot neurite density quantification and heat map generation. Female

C57BL/6J mice, aged 14-16wks, were either cold exposed (5⁰C) (n=6) or at room temperature

(n=4) for 7 days. Axillary scWAT depots were wholemount processed and stained with PGP9.5

and imaged with 10X objective. To demonstrate the difference between true signal and tissue

autofluorescence pixel intensity less than 55 was considered to be autofluorescence (grey) and

pixel intensity 55 or greater was considered true staining (green) (a). Images were z-max

projected, tiled, and thresholded and neurite density quantification was performed, compared

across the cohort, and analyzed using a two-tailed Student's t-test (b). Alpha level 0.05, error

bars are SEMs (b). Neurite density per z-max tile was calculated and used to generate a

heatmap (c). Inguinal scWAT depot was stained with IB4 to mark vasculature, imaged at 10X,

and vascular density per z-max tile was calculated and used to generate heat map (d) Each tile

is comprised of a 0.12cm x 0.12cm area (c-d). All imaging was performed with a Leica TCS SP8

DLS laser scanning confocal microscope. Scale bars are 2 mm (a).

STAR Methods:

Animals

The following mouse strains were obtained from The Jackson Laboratory: C57BL/6J

(Stock # 000664); BTBR$^{+/+}$ (WT) and BTBR$^{ob/ob}$ (MUT) (BTBR.Cg-Lep^{ob}/WiscJ, Stock # 004824);

LysM$^{cre/-}$ (B6.129P2-Lyz2/J, Stock # 004781); BDNF$^{fl/fl}$ (Bdnf^{tm3Jae}/J, Stock # 004339); R26R-

EYFP (B6.129X1-Gt(ROSA)26Sortm1(EYFP)Cos/J, Stock # 006148); TH$^{cre/-}$ (B6.Cg-

7630403G23RikTg(Th-cre)1Tmd/J, Stock # 008601. Myeloid lineage reporter mice were

generated by crossing LysM^{cre}x BDNF^{flfl} x R26R-EYFP. Nav1.8 reporter mice *(Na$_v$ 1.8^{cre} x*

tdTomato) were a kind gift from Dr. Ian Meng, (University of New England, Biddeford, ME,

USA). Animals were housed 3-5 to a cage (with the exception of BTBR WT and MUT animals

which were housed in pairs), in a monitored temperature and humidity-controlled environment

with 12/12hr light/dark cycle. Cages were cleaned weekly, ad libitum access to standard chow

diet and water was maintained. For all studies animals were sacrificed using CO_2 followed by

cervical dislocation.

Adult male and female C57BL/6J were used for majority of immunostaining studies.

Adult male BTBR WT and MUT were used in second harmonic generation imaging. Adult

female TH-Rosa-eYFP reporter mice were used in optical clearing, and transmission electron

microscopy studies. Adult male and female myeloid lineage reporter mice were used in high

magnification imaging of adipose SVF. Adult female Nav1.8 reporter mice were used for whole

mount immunostaining with PGP9.5 (GFP).

Cold exposure experiments

All cold exposure was carried out in a diurnal incubator (Caron, Marietta, OH, USA) at

5°C. Adult male C67BL/6J mice were housed two to a cage and either maintained at room

temperature or continuously cold exposed for 7 days. Inguinal and/or axillary scWAT was

collected for wholemount tissue processing.

<u>Mouse adipose tissue collection and processing for immunofluorescence</u>

Mice were euthanized using CO_2 followed by cervical dislocation. Whole scWAT depots were carefully removed to remain fully intact and immediately fixed in 2% PFA at 4°C for 4hr-16hrs depending on thickness of tissue. The tissues were then rinsed for 30 minutes with 1X PBS w/ 10U/mL heparin, twice at 4°C. Tissues next underwent z-depth reduction in which tissues were flattened by being placed between two large glass slides bound tightly together with large binder clips, for 1.5hrs at 4°C. Tissues were incubated in blocking buffer (1XPBS/2.5% BSA/1% Triton X-100) at 4°C overnight but can be incubated up to 7-days if desired with blocking buffer replaced every 24hrs. After blocking period tissues underwent one of three methods of autofluorescence quenching: 1) Incubation in 0.1% Typogen Black for 20 minutes at room temperature on a rotator, 2) incubation in 1X (0.1%) TrueBlack® Lipofuscin Autofluorescence Quencher for 10 minutes at room temperature on a rotator, or 3) stepwise dehydration and subsequent bleaching with 5% hydrogen peroxide in 20% DMSO/MeOH as described by *Renier et al. 2014* [38]. Following quenching the tissues were washed with 1X PBS w/ 10U/mL heparin on rotating platform at 4°C replacing PBS every 1hr for a total of 4-6hrs, or until all unbound stain was removed. Tissues were incubated with primary antibody for 48hrs at 4°C. The following day tissues were washed with 1XPBS on a rotating platform at 4°C, replacing PBS every 1hr for a total of 4-6hrs followed by incubation with secondary fluorescent antibodies overnight. Tissues were then again washed with 1XPBS on a rotating platform at 4°C, replacing PBS every 1hr for a total of 4-6hrs. Following the immunostaining steps, when applicable, tissues were incubated with 1 µg/mL isolectin IB_4 (IB4) (ThermoFischer, cat#:I32450 or cat#:I21413) diluted in HEPES buffer (pH 7.4), overnight at room temperature on a rotator. Tissues were then washed with 1XPBS on a rotating platform at room temperature for one hour twice. At this time, if DAPI co-staining was desired, the tissues were incubated in 100ng/mL

DAPI for 10 minutes at room temperature on a rotator. Tissues then received four 1hr washes in

1XPBS at room temperature. At the conclusion of washing steps tissues were placed on large

glass slides medial side facing up. A few drops of glycerol based mounting media was added to

the tissue and coverslip was placed on top. Glycerol based mounting media was used because

aqueous based mounting fluid lacks the required viscosity to fully adhere the coverslip when

dealing with whole mount adipose. Slides were weighted down for 2 days and then sealed.

Protocols for tissue dissection, processing for immunofluorescence, and confocal imaging can

be found at https://dx.doi.org/10.17504/protocols.io.brs2m6ge.

<u>Widefield microscopy</u>

All widefield micrographs were captured on a Nikon E400 with either a DS-fi2 color

camera (Nikon, Japan) or Hamamatsu ORCA-Flash4.0 V2 Digital CMOS monochrome camera

(Hamamatsu Photonics K.K., C11440-22CU) in conjunction with filter cubes for DAPI,

GFP/FITC, Cy3/TRITC, and Cy5. Images captured on with Hamamatsu ORCA-Flash4.0 V2

were pseudo colored with NIS-Elements software (Nikon, Japan.) The following objectives were

used: Nikon CFI Plan Apochromat Lambda 4X (NA 0.20, WD 20.0mm, dry), Nikon CFI Plan

Apochromat Lambda 10X (NA 0.45, WD 4.00mm, dry), Nikon CFI Plan Fluor 20X (NA 0.50, WD

2.10mm, dry) and Nikon CFI Plan Fluor 40X (NA 0.75, WD 0.66mm, dry.)

<u>Confocal microscopy</u>

All confocal micrographs were captured on a Leica TCS SP8 DLS (Leica Microsystems,

Wetzler, Germany) laser scanning confocal microscope. Fluorescent labels were excited with

either a diode 405nm laser (DAPI) or a white light laser (Alexa488, GFP, eYFP, Alexa532,

Alexa555, tdTomato, Alexa 568, Alexa647) with excitation and emission spectra tuned

specifically for each fluorophore or groups of fluorophores to reduce and eliminate spectral

overlap when possible. All imaging was performed sequentially by frame. Photons were

captured with both PMT and HyD detectors. The following objectives were used: HC PL

FLUOTAR 5X (NA 0.15, WD 13.70mm, dry), HC PL APO CS2 10X (NA 0.40, WD 2.74mm, dry),

HC PL APO CS2 20X (NA 0.75, WD 0.62mm, dry) and HC PL APO CS2 63X (NA 1.40, WD

0.14mm, oil.) Intact scWAT depots were imaged by tiling z-stacks across the full depth and area

of the tissue. Whole depot micrographs in this manuscript were scanned bidirectionally at either

400Hz or 600Hz at 720x720 resolutions. Line averages ranged between 3-8 and z-step size

ranged between 5-16 µm. Tissue thickness, though variable, was easily captured within the z-

step range of the software (±250 µm of the focal point.) Identical image acquisition settings were

applied for all tissues within cohorts that received neurite density quantifications. Anywhere

between 4,000-65,000 individual images were captured per tissue which ranged from 100-900

tiles. These tiles were then merged together and processed into a 2D maximum intensity

projection imaged either in LASX (Leica imaging software) or in Fiji [65] if being used for

quantification. Whole depot image insets consisted of randomly chosen 10X z-max tiles of the

merged image expanded for improved visualization. Representative images were captured at

1024x1024 resolution using a 63X objective and were z-max projected. Digital nerve bundle

cross sectioning was performed by using the XZY scanning mode in the LASX software which

applied a 2.54X digital zoom automatically to each image. All images were pseudo colored in

LASX software.

Neurite density quantification and heat map generation

Maximum intensity projections through the z-dimension (z-max projection) were

generated for each tile individually using Fiji [65]. All single tile z-max projections were further

processed using MATLAB x64 software (version 2018b, MathWorks). To remove low-frequency

background noise, a 2D Gaussian smoothing kernel was used to convolve each z-max

projection with a very large Gaussian blur using the *imgaussfilt* MATLAB command, with large

standard deviation (~150), which was then subtracted from its corresponding original z-max

projection. After background subtraction, a small gaussian blur (*imgaussfilt* with small standard

deviation, ~2-3) was used to broaden out the neurite signal slightly before thresholding the

image. Next, a binary thresholded mask was generated from the processed image. In the mask,

only regions having an area larger than 40 pixels were kept for further analysis (using the

bwareaopen MATLAB command). Then the *bwskel* MATLAB command was used to perform

the skeletonizing procedure, with the added feature of removing any branches less than 4 pixels

long. Except for an experimental analysis of 2D z-max projections vs. the individual 2D slices

making up the 3D confocal data (Suppl. Fig S8), total nerve length was calculated using values

measured for each 2D z-max projection. Total nerve arborization density was calculated as the

ratio of total nerve length divided by the total viewing area, resulting in nerve length per square

meter of tissue.

To generate heat maps a csv file was generated with the topological positions and the

associated arborization quantity of each tile using the HeatMapChart function. Heatmap

generation has multiple variables controlled by the user that could be explored further, such as

the size of the subregion size and whether or not (and to what extent) the subregions may

overlap. Code and instruction can be found at https://github.com/ktownsendlab/willows_et_al-

2020.

It is also worth noting that the Python and MATLAB scripts developed for this study,

provided above, allow the user to adjust the described parameters to account for batch

differences due to variation in fluorescence intensity and autofluorescence. These adjustments

can be made *a priori* or while looking at the skeletonized product of a single tile to quickly

visualize how each adjustment changes the skeletonization.

Antibodies and Fluorescing Stains

Primary antibodies included: PGP9.5 (1:1000, Abcam, Cambridge, U.K. Cat. #108986 and

#10404); PSD95 (1:1000, Abcam, Cambridge, U.K. Cat. #ab18258); MPZ (1:250, Abcam,

Cambridge, U.K. Cat. # ab31851); Advillin (1:500, Abcam, Cambridge, U.K. Cat. # ab72210);

Perilipin-1 (1:100, Abcam, Cambridge, U.K. Cat. #ab61682); TH (1:250, Millipore, Burlington, MA

USA; Cat. # AB152); neurofilament-M (2H3, 1:500) and synaptic vesicles (SV2, 1:250) from

Developmental Studies Hybridoma Bank, (University of Iowa, USA); β3-Tubulin conj. Alexa Fluor

488 (1:250, Abcam, Cambridge, U.K. Cat. # ab195879); S100β conj. Alexa Fluor 647 (1:250,

Abcam, Cambridge, U.K. Cat. # ab196175).

Secondary antibodies included: Goat anti-Rabbit IgG (H+L) Alexa Fluor 488 (1:500,

ThermoFisher Scientific, Waltham, MA, USA; Cat. # A-11008); Goat anti-Mouse IgG (H+L) Alexa

Fluor 488 (1:500, ThermoFisher Scientific, Waltham, MA, USA; Cat. # A-11001); Donkey anti-

Goat IgG (H+L) Alexa Fluor 488 (1:500, ThermoFischer Scientific, Waltham, MA, USA; Cat. #A-

11055). Goat anti-Rabbit IgG (H+L) Alexa Fluor 532 (1:500, ThermoFisher Scientific, Waltham,

MA, USA; Cat. #A-11009); Goat anti-Rabbit IgG (H+L) Alexa Fluor 555 (1:500 or 1:1000,

ThermoFisher Scientific, Waltham, MA, USA; Cat. #A-21428); Goat anti-Rabbit IgG (H+L) Alexa

Fluor 594 (1:1000, ThermoFisher Scientific, Waltham, MA, USA; Cat. # A-11005); Goat anti-

Rabbit IgG (H+L) Alexa Fluor 647 (1:1000, ThermoFisher Scientific, Waltham, MA, USA; Cat. #

A-11012).

Fluorescent stains include: For vascular staining, tissues were incubated in isolectin IB_4

stain conjugated to either Alexa Fluor 594 (1ug/ml, ThermoFisher Scientific, Waltham, MA, USA;

Cat. # I21413) or 647 (1ug/ml, ThermoFisher Scientific, Waltham, MA, USA; Cat. #I32450). For

nuclear staining, tissues were incubated in DAPI (100ng/ml, Sigma-Aldrich, St. Louis, MO, USA;

Cat. # D9564).

Luxol Fast Blue Myelin Staining of Whole Adipose

Mice were euthanized, whole subcutaneous white adipose tissue (scWAT) depots were

carefully removed to remain intact and fixed in 2% PFA at 4°C for 4hr-12hrs depending on

thickness of tissue. The tissues were then rinsed for 10 minutes with 1X PBS, twice at 4°C.

Tissues were stained with Luxol Blue Myelin stain kit (Abcam, Cambridge, U.K. Cat. #ab150675).

Tissues were incubated in Luxol Fast Blue Solution overnight at room temperature, briefly rinsed

with distilled water and differentiated in Lithium Carbonate Solution followed by 70% Alcohol

Reagent until the solution ran clear and adipocytes were colorless while nerves remained blue.

Tissue was then rinsed in distilled water, mounted, and imaged on a dissecting microscope.

Optical Clearing

Optical clearing methods were tested in inguinal scWAT and/or BAT from either _TH-Cre x_

ROSA YFP (Jackson Laboratory, Bar Harbor, ME, USA; Stock # 008601 and 006148) reporter

mice or _C57BL/6J_ mice (The Jackson Laboratory, Stock # 000664). ScaleA2 method described

by Hama, H. et al. [34]; sucrose gradient method as described by Tsai, P. et al. [66]; UbasM

method as described by Chen, L. et al. [40]. BABB method as described by Li, L. et al. [36]; CUBIC

and CUBIC CB-perfusion as described by Susaki, E. et al. [37]; uDISCO as described by Pan, C.

et al. [39]; iDISCO as described by Reinier, N. et al. [38]

Cell Size Quantification

Whole adipose depots were excised from male C57BL/6J mice and fixed in 2% PFA

overnight. Whole tissues were either z-depth reduced and mounted on a glass slide (z-depth

reduced) or remained unmanipulated and placed on a glass slide (control). Representative

images were taken of each tissue and cells with clear in-focus boundaries were outlined. Cell

area was measured in FIJI [65].

Hematoxylin Staining

Whole inguinal adipose depots were excised from male C57BL/6J, fixed in 10% buffered

formalin overnight and paraffin embedded. Paraffin embedded tissues were sectioned at 7μm and

stained with hematoxylin to increase cell membrane contrast. Tissues were then cover slipped,

sealed, and imaged.

Tissue Sectioning and IF for lysm cre Tissues

Both axillary and inguinal scWAT were collected from adult 7-day cold (5°C) exposed

LysM-Cre-Rosa eYFP reporter mice. Whole depots were cut into 100 µm-thick slices

longitudinally (from anterior to posterior). Slices from each depot were fixed in 2% PFA for 1hr,

then rinsed for 10 minutes with 1X PBS w/ 10U/mL heparin, twice at 4°C. Tissues were incubated

in blocking buffer (1XPBS/2.5% BSA/0.5-1% Triton) at 4°C overnight. Tissues were rinsed in rinse

buffer and stained with TrueBlack®. Tissue slices were stained with the primary antibody, PSD95

(ab18258 at 1:1000) for 48 hours. Then they were stained with goat anti-rabbit secondary

antibody bound to AlexaFluor 647 (1:1000) overnight. Finally, they were stained with the primary

antibody conjugated to GFP (ab6662 1:1000) for 72 hours. Sections underwent z-depth reduction

as previously describes in Methods Section prior to mounting. Tissues were either imaged on a

confocal microscope at 63X magnification. All tissues imaged with a confocal microscope were

stained with DAPI in addition to PSD95, AlexaFluor 647 and GFP.

Transmission Electron Microscopy

Contralateral inguinal and axillary tissue samples were collected as 1 mm^3 blocks. Tissues

were fixed in 2% glutaraldehyde-2% paraformaldehyde for 2 hours and then washed overnight in

t0.1 M PB (pH 7.4) buffer at 4°C. Samples were then post-fixed in 1% OsO^4 in PB, dehydrated in

an acetone series (50%, 70%, 95% and 100%) and embedded and polymerized at 60 °C for 48

hrs in Epon/Araldite resin. Once the tissue samples were in resin, an ultramicrotome (Leica UC6)

was used to cut the tissues into sections that did not exceed 60-90 nm in thickness.

Those sections were then collected directly on a 300-mesh nickel grid(s) and stained with uranyl

acetate and lead citrate. The samples were imaged using a CM10 transmission electron

microscope (Philips). Images were taken of both adipocytes and cells within the SVF.

Second Harmonic Generation Imaging

All 2-photon microscopy studies used a modified Olympus FV300 system with an upright

BX50WI microscopy stand (Olympus, Center Valley, Pennsylvania) and a mode-locked

Ti:Sapphire laser (Chameleon, Coherent, Santa Clara, California). Laser power was modulated

via an electro-optic modulator (ConOptics, Danbury, Connecticut). The fluorescence and SHG

signals were collected in a non-descanned geometry using a single PMT (H7422 GaAsP,

Hamamatsu, Hamamastu City, Japan). Emission wavelengths were separated from excitation

wavelengths using a 665 nm dichoric beamsplitter followed by 582/64 nm and 448/20 nm

bandpass filters for Alexa 488 and SHG signals respectively (Semrock, Rochester, New York).

Images were acquired using circular polarization with excitation power ranging from 1- 50 mW

and a 40X (0.8 NA) water immersion objective with 3X optical zoom with scanning speeds of

2.71s/frame. All images were 515 x 512 pixels with a field of view of 85 μm.

Image Processing and Figure Formatting

Images were compiled and formatted for figures in PowerPoint (Microsoft.) In some

cases, brightness was increased to enhance small structure visibility. Figure brightness was

adjusted identically when direct comparisons were made. Figures were each exported as a Tiff

(1650x2200) and then reduced to 300x600 DPI and 8-bit depth for publication.

Resource Availability

Lead Contact

Further information and requests for resources and reagents should be directed to and

will be fulfilled by the Lead Contact, Kristy Townsend (kristy.townsend@osumc.edu).

<u>Materials Availability</u>

This study did not generate new unique reagents.

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<u>Data and Code Availability</u>

Neurite density quantification and heat map generating code developed for this publication can be found on GitHub at https://github.com/ktownsendlab/willows_et_al-2020. Protocols used for this publication can be found on Protocols.io at https://dx.doi.org/10.17504/protocols.io.brs2m6ge. All other data is included in this manuscript.

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Supplemental Information

Supplemental Table S1: Comparison of methods for clearing adipose tissue

Supplemental Figure S1: Resonant scanning as an alternative imaging approach.

Supplemental Figure S2: Additional whole mount staining optimization.

Supplemental Figure S3: Height color coded z-maximum projection.

Supplemental Figure S4: 7µm cross section of peripheral nerve bundles in inguinal scWAT.

Supplemental Figure S5: Transmission Electron Microscopy (TEM) of scWAT Innervation.

Supplemental Figure S6: Synapsing within white adipose tissue.

Supplemental Figure S7: Additional nerve labeling in scWAT.

Supplemental Figure S8: Measuring neurite density from z-maximum projections is equivalent to

measuring from individual z-slices.

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Figure 1: Adipose tissue processing, imaging, and quantification method

a.   **Wholemount Tissue Processing for Immunostaining**

b.   **Tiled Z-Stack Imaging of Wholemount Tissue**

5 mm

c.   **Image Post-Processing for Quantification**

Original Image | Background Subtraction | Light Smoothing | Thresholded Mask | Skeletonization | Quantification

$$\frac{\text{Total Nerve Length (m)}}{\text{Viewing Area (m}^2\text{)}}$$

d.

Tissue Processing and Imaging:
https://dx.doi.org/10.17504/protocols.io.brs2m6ge
Quantification:
https://github.com/ktownsendlab/willows_et_al-2020

Figure 2: Z-depth reduction and tissue morphology

a. Z-depth Reduction and Mounting

b. Z-depth Reduction Increases Tissue Area

c. Z-depth Reduction Decreases Tissue Thickness

d. Z-Depth Reduction Increases Adipocyte Area

e. Whole Mount Z-Depth Reduction: Adipocyte Shape

f. 7µm Sections of Z-Depth Reduced Adipose Depot

Figure 3: Whole mount adipose autofluorescence quenching and fluorescence staining

Figure 4: Peripheral innervation of scWAT

a. Whole Depot Pan-neuronal Labeling of Tiled Z-stack scWAT

b. High Magnification of Pan-neuronal Labeling in scWAT

c. Nerve Bundle Cross Section

d. Nuclei Embedded in Nerve Bundle

Figure 5: Characterization of whole depot scWAT nerves

a. Whole Depot Sympathetic Innervation

b. TH⁺ Nerve Bundle Cross Section

c. Sympathetic Nerve Fibers

d. AVIL⁺ Nerve Bundle Cross Section

e. Whole Depot Myelination

f. MPZ⁺ Nerve Fibers

g. MPZ⁺ Nerve Bundle Cross Section

Figure 6: Neurovascular interaction within whole mount scWAT

a. Whole Depot Neurovascular Stain

b. Categories of Neurovascular Interactions in scWAT

c. SNS Innervation of scWAT Vasculature

d. scWAT Vasa Nervorum

e. scWAT Perivascular SNS

f. Myelinated Nerve Fibers Around scWAT Vasculature

Figure 7: Whole depot neurite quantification and heat map generation

a. Cold Exposed Whole Depot Axillary scWAT

RT — PGP9.5 | PGP9.5 | Autofluorescence

7 day 5°C — PGP9.5 | PGP9.5 | Autofluorescence

b. Neurite Density

p = 0.414

1/m — RT, 7-day Cold

c. Neurite Density Heatmap

PGP9.5 | Neurite Density (1/m)

d. Vascular Density Heatmap

IB4 | Vascular Density (1/m)