



The sexually dimorphic role of adipose and adipocyte estrogen receptors in modulating adipose tissue expansion, inflammation, and fibrosis

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ABSTRACT

Our data demonstrate that estrogens, estrogen receptor- α (ER α), and estrogen receptor- β (ER β) regulate adipose tissue distribution, inflammation, fibrosis, and glucose homeostasis, by determining that α ERKO mice have increased adipose tissue inflammation and fibrosis prior to obesity onset. Selective deletion of adipose tissue ER α in adult mice using a novel viral vector technology recapitulated the findings in the total body ER α null mice. Generation of a novel mouse model, lacking ER α specifically from adipocytes (AdipoER α), demonstrated increased markers of fibrosis and inflammation, *especially in the males*. Additionally, we found that the beneficial effects of estrogens on adipose tissue require adipocyte ER α . Lastly, we determined the role of ER β in regulating inflammation and fibrosis, by breeding the AdipoER α into the β ERKO background and found that in the absence of adipocyte ER α , ER β has a protective role. These data suggest that adipose tissue and adipocyte ER α protects against adiposity, inflammation, and fibrosis in both males and females.

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Keywords Estrogen receptor alpha ($ER\alpha$); White adipose tissue (WAT); Fibrosis; Inflammation

1. INTRODUCTION

Intra-abdominal white adipose tissue has been strongly correlated with insulin resistance, inflammation, cardiovascular disease, and the Metabolic Syndrome in animal models and humans [1-3]. Men, on average, have less total body fat but more intra-abdominal adipose tissue than women, whereas women, on average, have more total fat and subcutaneous adipose tissue [4-8]. Male and female adipose tissues are sexually dimorphic; female adipose tissue is more insulin sensitive, less susceptible to inflammation, and has higher expression of estrogen receptors (ERs) [9–11].

Estrogens are produced in the ovary and testes, as well as in adipocytes (by the action of aromatase on androgens), and are increased in proportion to total body adiposity [12,13]. Reduced circulating estrogens, as seen in post-menopausal females, result in the development of increased intra-abdominal adiposity and increased susceptibility to diseases associated with the Metabolic Syndrome. Importantly, women who receive estrogen replacement therapy are less likely to deposit adipose tissue in the intra-abdominal depot [14–16], and are relatively

more protected from the Metabolic Syndrome. To further demonstrate that ovarian hormones are responsible for the effects on adipose tissue distribution, we and others have data demonstrating removal of ovaries (ovariectomy (0VX)) results in body weight gain primarily as adipose tissue [17–19], which is deposited primarily in the intra-abdominal adipose depot and results in impaired glucose homeostasis, whereas, administration of estrogens reduces body weight, body adiposity, changes body fat distribution, and improves glucose homeostasis [18,20,21]. Increased adiposity and adipose tissue inflammation are now recognized as underlying mechanisms in the pathogenesis of dysregulated glucose homeostasis and the Metabolic Syndrome. As previously mentioned, females are protected from the Metabolic Syndrome, which may likely be due to the potent anti-inflammatory effects of estrogens [22,23].

Estrogens bind to two 'classical' estrogen receptor subtypes, estrogen receptor alpha (ER α) and estrogen receptor beta (ER β), with similar affinity; however these two receptors are believed to differ in their translational properties. Importantly, the relative direct importance of ER α and/or ER β in modulating adipose tissue inflammation is not known. Previous studies have described ER α protein [24,25], as well as

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Abbreviations: $ER\alpha$, estrogen receptor alpha; $ER\beta$, estrogen receptor beta; WAT, white adipose tissue; BAT, brown adipose tissue

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specific estrogen binding and ER α mRNA to be present in human subcutaneous adipose tissue [25]. However, others have not been able to detect estrogen receptors in human adipose tissue [26,27]. More recently, ER β mRNA has been detected in human subcutaneous adipose tissue, suggesting that direct effects of estrogen may involve both receptor subtypes [20].

The effects of estrogens on energy homeostasis are primarily mediated by ER α , as indicated by findings that women or female mice with mutations in the ER α gene display hyper-adiposity [28,29]. We recently reported that site-specific knockout of $ER\alpha$ in certain brain regions results in increased adiposity, changes in body fat distribution, reductions in energy expenditure, and alterations in fertility. Furthermore, global deletion of the ER α gene (α ERKO) results in increased adiposity in both males and females, with a near doubling of the intra-abdominal adipose tissues when compared to age-matched wild type (WT) mice [29,30]. However, mice with a deletion in ERB (BERKO) do not have increased adiposity or metabolic derangements. These data suggest an important role for $ER\alpha$ in regulating adipose tissue distribution and potentially inflammation. Therefore, we sought to determine the role of ER α in regulating adipose tissue distribution and 'function' as measured by levels of inflammation, and fibrosis. To do this, we first compared weight-matched wild type (WT) and α ERKO mice with respect to adipose tissue distribution, fibrosis, and inflammation. Our data suggest that α ERKO males and females have increased adipose tissue that has elevated levels of markers of inflammation and fibrosis. Since the α ERKO is a total body deletion of ER α and we were specifically interested in selective effects of ER α in adipose tissue we developed a novel approach to knock down ER α expression in an adipose tissue depot-specific manner using viral technology. We found that knockdown of $ER\alpha$ selectively in intra-abdominal adipose tissue results in increased adipose tissue mass, increased adipocyte size, and is associated with elevated adipose tissue inflammation and fibrosis in both males and females. Importantly, our novel viral-mediated knockdown of ER α expression reduces ER α in all adipose tissue cell types, including adipocytes, preadipocytes, immune cells and vascular cells. Therefore, in order to assess the role of ER α specifically in adipocytes, we created a mouse model in which we can knock down ER α selectively in adipocytes by generating a unique and novel animal model using the Cre-loxP system where we bred the $\text{ER}\alpha^{\text{lox/lox}}$ mouse [31] combined with the adipocytespecific Adiponectin-Cre transgenic model [32,33] to produce AdipoERa mice, or mice that specifically lack $ER\alpha$ only in adipocytes. Our data suggest that adipocyte ER α regulates adipocyte size, adipose tissue inflammation and fibrosis in both males and females. Finally, to rule out any contribution $ER\beta$ may have with respect to the metabolic phenotypes observed, we bred the AdipoER α mouse into the total body β ERKO mouse resulting in a mouse model that lacks both estrogen receptors on adipocytes. Our data suggest that knockdown of adipocyte ER α in the context of the BERKO mouse increases inflammation and fibrosis indicating a role for ER β in the absence of adipocyte ER α .

2. MATERIAL AND METHODS

2.1. αERKO mice

Wild type (WT) and α ERKO male and female littermates were housed in a temperature-controlled environment in groups of two to five at 22– 24 °C using a 12 h light/12 h dark cycle (singly housed to measure food intake). The mice were fed standard chow (#2916, Harlan-Teklad, Madison, WI) and ad libitum water. Body weight was monitored weekly, mice (*n*=8) were sacrificed at 6 weeks of age, a time in which body weight did not differ, and a time in which the WT mice had gone through puberty as confirmed by observation of vaginal opening in the females. Care of all animals and procedures were approved by the UT Southwestern Medical Center in accordance with the Guide for Care and Use of Laboratory Animals of the National Institutes of Health. All WT females were sacrificed in proestrus cycle.

2.2. Gonadal fat depot-specific viral knockdown of $ER\alpha$

In vivo fat pad-specific injections were made in WT C57BI6 male and female mice (n=9) at 8 weeks of age. Mice were anesthetized with isoflurane (Aerrane, Baxter, IL), the ventral surface shaved and treated with betadine and alcohol $3 \times$. A bilateral ventral abdominal incision was made and the gonadal fat pads (peri-ovarian for females, and epididymal for males) were injected with an adeno-associated (AAV) $ER\alpha$ siRNA virus (on the right side) and with a control (scrambled sequence) siRNA (on the left side). Each virus was diluted using sterile saline and was injected in equal volumes of 4×10^{11} particles per fat pad. Approximately 5-7 injections were performed to dispense the virus across the pad. The peritoneum was closed with Vicryl suture (Ethicon) and the skin was closed with nylon suture and staples. Post-operative pain was managed using 0.5 mg/kg buprenorphine injection postsurgery and 2 mg/kg rimadyl tablets for the following 48 h. After the surgery the mice were monitored daily for body weight and food intake to assess recovery. The tissues were harvested at 3 weeks postinjection (sacrificed at 11 weeks of age). To illustrate the validity and specificity of the technique, the AAV ER α siRNA virus was tagged with green fluorescent protein (GFP) and viewed under GFP fluorescence.

2.3. Adipocyte-specific knockdown of $ER\alpha$

In order to generate mice lacking $ER\alpha$ specifically in adipocytes, we used Adiponectin promoter driven-Cre transgenic mice [33], and crossed them with mice carrying the loxP-flanked ER α allele (ER α ^{lox/lox}) [31] producing ER $\alpha^{lox/lox}$ /Adiponectin-Cre (AdipoER α) mice and their ER $\alpha^{lox/lox}$ (WT) littermates on a C57BI6 background. Mice were housed in a temperature-controlled environment in groups of two to five at 22-24 °C using a 12 h light/12 h dark cycle (singly housed to measure food intake). The mice were fed standard chow (#2916, Harlan-Teklad, Madison, WI) and ad libitum water. Care of all animals and procedures were approved by the UT Southwestern Medical Center in accordance with the Guide for Care and Use of Laboratory Animals of the National Institutes of Health. Body weight was measured weekly from group housed mice (n=10). Food intake was measured daily, and average daily food intake was calculated using data from at least 10 continuous days. Body composition was determined using quantitative magnetic resonance (QMR) (Bruker's Minispec MQ10, Houston, TX). Mice were sacrificed at 18 weeks of age.

2.4. Adipocyte-specific knockdown of ER α in ER β null mice (AdipoER $\alpha/$ β ERKO)

In order to determine the potential role of ER β in mediating the AdipoER α phenotype, we generated mice lacking ER α specifically in adipocytes (ER $\alpha^{lox/lox}$ /Adiponectin-Cre (AdipoER α)) mice and bred them to β ERKO mice on a C57BI6 background (n=9). Care of mice, food intake, body composition, fat distribution, tissue analysis, gene expression and histology were carried out as previously described.

2.5. Tissue mRNA analyses

Mice (number of mice and age for each group are noted in figure legends where appropriate) were euthanized with isoflurane (Aerrane, Baxter, IL), and the gonadal white adipose tissue (WAT or Gonadal-the



fat pad surrounding the ovaries and uterus for the females, and the epididymal for the males) was isolated, weighed, and snap-frozen in liquid nitrogen. Total RNA was isolated following tissue homogenization in Trizol (Invitrogen, Carlsbad, CA) using a TissueLyser (Qiagen, Valencia, CA) and isolated using the RNeasy RNA extraction kit (Qiagen). The quality and quantity of the RNA were determined by absorbance at 260/280 nm. cDNA was prepared by reverse transcribing 1.5 μ g RNA with SuperScript III reverse transcriptase (Invitrogen) and oligo(dT)₂₀ (Invitrogen). Quantitative PCR (qPCR) was performed using TaqMan primers for tumor necrosis factor- α (TNF α), toll-like receptor-4 (TLR-4), serum amyloid A-3 (SAA3), lysyl oxidase (LOX), collagen-6 (COL6), and EGF-like module-containing mucin-like hormone receptor-like 1 (F4/80). Beta-2-microglobulin was used as the housekeeping gene.

2.6. Hematoxylin and eosin staining

As above, gonadal WAT was isolated, weighed, and fixed with 10% formalin for overnight and then stored in 50% ethanol. The fixed fat pads were sent to Richardson Molecular Pathology Core at UT South-western Medical Center, where the tissues were embedded with paraffin, sectioned and stained with haematoxylin and eosin (H&E). Images were viewed under rhodamine fluorescence and imaged using Leica DM2000 compound epifluorescence microscope equipped with an Optronics Microfire Color CCD Camera and analyzed for adipocyte area using NIH ImageJ software. Eight hundred to 1000 cells from each sample were included in the analysis. (See figure legends for details of each cohort).

2.7. ER α immunohistochemistry

To determine presence of ER α , sections (n=4) were de-paraffinized in xylene followed by decreasing concentrations of ethanol. Sections were microwaved in citric acid buffer for antigen retrieval. They were then blocked for endogenous peroxidases using 3% H₂O₂ in PBS. Tissues were also blocked for avidin and biotin (Vector SP-2001) as well as protein (10% NGS S-1000, Vector). After blocking, sections were stained with rabbit polyclonal anti-ER α primary antibody (1:400) (Santa Cruz) followed by biotinylated anti-rabbit antibody (Vector). Following incubation in HRP polymer (PK-6101) the secondary antibody (1:800) was detected using Vector Immpact DAB kit (SK-4105) and counterstained with hematoxylin. All images were obtained using Nikon Coolscope.

2.8. Adipocyte/SV fractionation and isolation of nuclei

To confirm knockdown of ER α specifically from adipocytes, adipose tissue was excised as previously described and adipocytes were isolated from the stromal vascular tissue (SV) using the collagenase fractionation method [34]. Specifically, isolated adipocytes were resuspended in cold nuclei extraction buffer (320 mM sucrose, 5 mM MgCl₂, 10 mM HEPES, 1% Triton-X at pH7.4) at 10⁶ cells/ml. The cells were vortexed gently for 10 s and incubated on ice for 10 min. Nuclei were pelleted at 2000 × *g*, and washed twice with nuclei wash buffer (320 mM sucrose, 5 mM MgCl₂, 10 mM HEPES at pH7.4). Nuclei in nuclei wash buffer was allowed to attach to silicone-coated slides for 1 h at 4 °C. Then nuclei were fixed in 3.7% formaldehyde in nuclei wash buffer for 10 min and subjected to BrdU analyses using BrdU Flow Kit (BD Biosciences). (See figure legends for details of each cohort).

2.9. Ovariectomy and estrogen replacement

Ovariectomy (OVX) and sham surgeries were performed in 12-week-old female mice as previously described [10]. In each study animal, a pellet of estradiol-17 β (0.03 mg per pellet for a 60 day release, (0.5 μ g/day), a dose in which plasma levels of estrogens mirror those of intact females in proestrus [35]), or placebo control was administered subcutaneously. Food intake, body weight, and body adiposity were tracked as previously described. Mice were sacrificed 4 weeks post-surgery.

2.10. Oral glucose tolerance test (OGTT)

Mice were fasted for 3 h (starting at 8 a.m.) prior to administration of glucose (2.5 g/kg body weight) by gavage. At the indicated time points, venous blood samples were collected in heparin-coated capillary tubes from the tail vein. Mice did not have access to food throughout the experiment. (See figure legends for details of each cohort).

2.11. Hormone measurements

For all measurements, plasma was obtained from trunk blood from mice fasted 3–4 h. Estradiol-17 β (Invitrogen) was assayed by enzyme-linked immunosorbent assay kits. (See figure legends for details of each cohort).

2.12. Statistics

The data are presented as mean \pm SEM. After confirming normal distribution of data, comparisons between two genotypes were made by the unpaired two-tailed Student's *t*-test; repeated-measures ANOVA was used to compare changes over time between two genotypes. *P* < 0.05 is considered to be statistically significant. Significance between WT and AdipoER α measurements is denoted by a (*) and significance between male and female, or VEH and estradiol-17 β within a genotype, is denoted by a (#).

3. **RESULTS**

3.1. Adipose tissue function (inflammation/fibrosis) in α ERKO mice

To determine the role of ER α in modulating adipocyte size, inflammation, and fibrosis, WT and littermate a ERKO weight-matched female and male mice (Figure 1A and B) were compared with respect to adipose tissue histology, inflammation, and fibrosis. Despite body weights being equivalent, a ERKO males and females have increased body fat as measured by NMR (Figure 1C and D). *a*ERKO mice have significantly larger adipocytes when compared to WT mice, as shown by representative histological images (Figure 1E) and adipocyte size calculations (Figure 1F), consistent with previous reports from Cooke et al. [30]. Enlarged adipocytes can be associated with increased macrophage infiltration into the fat pad [36]. To investigate if $ER\alpha$ in gonadal adipose tissue directly regulates macrophage infiltration, a marker for macrophage expression, F4/80, was measured. Figure 1G shows that in α ERKO mice there is an increase in F4/80 expression in both males and females over WT mice that reached statistical significance in the males. To test for adipose tissue inflammation in *α*ERKO mice, gene expression for the inflammatory markers serum amyloid A3 (SAA3), Toll-like receptor 4 (TLR4) and tumor necrosis factor- α (TNF α) was measured and compared between α ERKO and WT mice. Figure 1H demonstrates that α ERKO results in an up-regulation of SAA3 and TLR4 in both sexes and TNF α in males. Adipose tissue inflammation has often been correlated with enhanced adipose tissue fibrosis; therefore, to test the effects of α ERKO on adipose tissue fibrosis, we analyzed gene expression levels of collagen VI (COL6) and lysyl oxidase (LOX). Data



Figure 1: Wild type vs α ERKO adipose tissue inflammation and fibrosis. (A and B) Weekly body weight was measured in singly housed female and male mice on normal chow (n=8/genotype). (C and D) Body composition was measured by NMR in 6-week-old female and male mice (n=8/genotype). (E) Representative photomicrographs of H&E staining of gonadal WAT from 6-week-old male and female mice. (F) Analysis of adipocyte cell size in both male and female of gonadal adipose tissue from 6-week-old mice (n=8/genotype). (G) Messenger RNA levels of F4/80 were quantified using qPCR whole adipose tissue from 6-week-old chow-fed females and males (n=8/genotype). (H and I) Messenger RNA levels were quantified using qPCR of whole adipose tissue from 6-week-old chow-fed females are presented as mean \pm SEM, and *P<0.05 between WT and α ERKO.



demonstrate a significant increase in the expression of LOX in both the female and male α ERKO adipose tissues when compared to WT control mice (Figure 1I). Further, female α ERKO mice have upregulated COL6 as compared to WT controls. Together these data suggest α ERKO results in fibrotic and inflamed adipose tissue associated with elevated macrophage infiltration, consistent with unhealthy adipose tissue, in both males and females, indicating that ER α is critical in maintaining adipocyte size, adipose tissue inflammation, and fibrosis in both males and females.

3.2. Gonadal fat depot-specific viral knockdown of $ER\alpha$

To begin to determine the adipose tissue-specific role of ER α in adult animals, we generated a novel technique that allows us to reduce the expression of ER α in a fat pad specific method. Importantly, this technique allows us to compare, within the same animal with the same hormonal milieu, the influence of adipose tissue $ER\alpha$. To illustrate the selectivity of the viral technique, Figure 2A demonstrates that only in the AAVsiRNA $ER\alpha$ GFP-injected pad (L Gon) is GFP visible (the tissue from each view has been outlined for reference), whereas the fat pad injected with the control AAVsiRNA/non-GFP virus is not fluroescent. Importantly, only trace amounts of signal are detected in the contralateral pad (R Gon), adjacent retroperitoneal pad (L Ret) and liver (Liv), which is most likely the result of auto-fluroescence. The effectiveness of the siRNA in gonadal adipose tissue was further determined using gPCR for ER α following injections of AAV-ER α siRNA (ER α siRNA) in the left gonadal pad or the control AAV-siRNA-scrambled sequence (control) in the right gonadal pad. These data suggest a \sim 50% reduction in ER α expression, as compared to control/contralateral fat pad (which is normalized to 1), in both males and females (Figure 2B). Fat pads that received the ER α siRNA are significantly heavier at 3 weeks post-injection than the pads injected with the control virus (regardless of sex) (Figure 2C). Consistent with *a*ERKO adipose tissue, adipocyte morphology analyses demonstrate the pad with reduced ER α has enlarged adipocytes (Figure 2D and E). Additionally, Figure 2F shows that reduced adipose tissue levels of $ER\alpha$ result in a 4-6 fold increase in F4/80 expression in both males and females over control injected pads, and this is associated with an upregulation of SAA3, TNF α , and TLR4 (Figure 2G). To test the effects of reduced adipose tissue $ER\alpha$ on fibrosis, we analyzed gene expression levels of COL6 and LOX. Consistent with α ERKO, adipose tissue specific reductions in ER α increase expression of LOX; however only a slight trend was observed in COL6 levels (Figure 2H). Together these data suggest that decreased adipose tissue $ER\alpha$ results in fibrotic and inflamed adipose tissue associated with elevated macrophage expression, consistent with unhealthy adipose tissue, in both males and females.

3.3. Adipocyte specific knockdown of ER α by generation of an ER α /Adiponectin Cre mouse

To determine the specific effect of ER α on adipocytes, we generated mice with adipocyte-specific knockdown of ER α (AdipoER α -ER α ^{lox/lox} + AdipoCRE) and control littermates (WT-ER α ^{lox/lox}). With this model, we demonstrate about ~60% knockdown of ER α in both male and female AdipoER α mice as compared to their WT littermates (Figure 3A). As previously indicated, females have higher ER α expression in the visceral depot than do males [10], and based on our previous work we know that knockdown of ER α from adipose tissue in females results in adipose tissue that is more 'male-like'. Here, we specifically chose an AdipoCRE line that results in downregulation of ER α expression in females to levels similar to those measured in males (Figure 3A–shown by comparing AdipoER α female to male WT levels). Adipocyte-specific

ER α gene expression was determined by RNA expression of collagenase-mediated isolation of adipocytes from the stromal vascular fraction. Consistently, representative gonadal adipose tissue sections stained for ER α demonstrate significantly fewer ER α immunoreactive adipocytes in the AdipoER α when compared to their littermate WT controls (Figure 3B). By generating mice with a 'physiological' knockdown of ER α we are able to metabolically determine the functional role of ER α specifically in adipocytes.

Male and female AdipoER α are born in normal Mendelian ratio and are viable. At weaning, WT and AdipoER α males and females do not differ in body weight or circulating estradiol 17ß levels (mice were sacrificed at 4 weeks of age in the proestrus stage of their cycle for comparison; WT 52.7 ± 8.2 vs AdipoER α 48.67 \pm 5.9 pg/mL estradiol 17 β however, at about 12 weeks of age, AdipoER α females begin to gain more weight when compared to WT females (Figure 3C). Male WT and AdipoER α mice do not differ in body weight through 25+ weeks of age (Figure 3D), or testosterone levels (WT 55.4 \pm 12.8 vs AdipoER α 57.49 \pm 14.9 pg/mL). Despite the findings that female AdipoER α mice weigh more, food intake does not differ between genotypes of either sex (Figure 3E). Nuclear magnetic resonance (NMR) imaging demonstrates that the increased body weight in AdipoER α females is due to increased fat mass, observed both in measurements of total grams of fat mass as well as percent of body weight (Figure 3F) while AdipoER α males do not differ significantly in any measure of body composition (Figure 3G). Moreover, female AdipoER α mice have an almost 2-fold increase in gonadal adipose tissue weight as compared to WT littermates (Figure 3H), equaling gonadal adipose tissue weights observed in males. These data suggest that adipocyte-specific knockdown of $ER\alpha$ does not alter body adiposity in males, dissimilar to knocking down ER α in the whole body or selectively in the adipose tissue depot (Figures 1D and 2C). In females however, reduced adipocyte ER α results in an increased fat mass, an increase that is largely due to increased gonadal adipose tissue, correlating to whole adipose body and tissue-selective knock down of ER α (Figures 1C and 2C).

In order to determine the metabolic consequences of reduced adipocyte $ER\alpha$, oral glucose tolerance tests (OGTT) were performed on male and female WT and AdipoER α mice. In the females, we observe no statistical difference in glucose clearance with an OGTT (Figure 3I). This was surprising given their substantial increase in gonadal adipose tissue mass. The area-under-the-curve calculations demonstrate a slight impairment in glucose disposal in the female AdipoER α mice (Figure 3K); however, these differences do not attain statistical significance. More surprisingly the male AdipoER α mice, which do not differ in body weight or adiposity, show a profound compromise in alucose clearance (Figure 3J), as demonstrated by a near doubling in the area-under-the-curve measurement for AdipoER α males as compared to WT males (Figure 3K). These data suggest that, despite no changes in adiposity, reduced adipocyte $ER\alpha$ in males results in a profound metabolic compromise, further suggesting that reductions of $ER\alpha$ to near ablation levels result in unhealthy adipose tissue. AdipoER α females are able to maintain glucose clearance even though they have male-patterned adiposity.

Expansion of adipose tissues can occur through increased triglyceride storage as previously discussed, thus we analyzed adipocyte size with H&E staining. Our data demonstrate that both male and female AdipoER α mice have enlarged gonadal adipocytes as compared to their WT littermates (Figure 4A) and calculation of average adipocyte area further demonstrates this (Figure 4B). AdipoER α males have an approximate doubling in average adipocyte size, while female AdipoER α mice have 3 times larger gonadal adipocytes as compared to their WT



Figure 2: Viral mediated knockdown of ER α in visceral adipose tissue regulates adipocyte size and gene function. Injection methods were determined by using the average surface area and shape of the gonadal adipose tissue. The right gonadal pad that was injected with the control virus is set to 1, and data are normalized to the control-injected pad. (A) Representative images of green fluorescent protein (GFP) in the pad injected with the AAVsiRNA ER α demonstrate that *only* in the AAVsiRNA ER α GFP-injected pad (L Gon) is GFP visible (the tissue from each view has been outlined for reference), whereas the fat pad injected with the control AAVsiRNA/non-GFP virus is not. Representative photomicrographs of H&E staining of the Gonadal control and siRNA pad from both the males and females at 3 weeks post-injection (*n*=9). (B) Messenger RNA levels of ER α were quantified using qPCR in whole adipose tissue at 3 weeks post-injection of both males and females (*n*=9). (C) Gonadal pad weights of both the control and the ER α siRNA of both males and females were weighed at the time of sacrifice at 3 weeks post-injection (*n*=9). (D) Representative photomicrographs of H&E staining of gonadal WAT from male and female mice. (E) Analysis of adipocyte cell size in both male and female of gonadal adipose tissue (*n*=9). (G and H) Messenger RNA levels of indicated genes were quantified using qPCR in whole adipose tissue at 3 weeks post-injection of both male and female mice (*n*=9). Data are presented as mean \pm SEM, and **P* < 0.05 between the ER α sIRNA injected pad and the pad injected with the control virus.

littermates. Males have significantly larger gonadal adipocytes than females; however, our data demonstrate that when ER α levels in a female are reduced to male levels, the resulting adipocyte size is far larger than that observed in WT males.

To assess the macrophage content of adipose tissues in male and female WT and AdipoER α mice, F4/80 was measured (Figure 4C). These data demonstrate that even though the female AdipoER α mice have enlarged adipocytes, the macrophage content of their gonadal adipose





Figure 3: Adipocyte ER α regulates body weight adipocyte distribution. (A) Messenger RNA levels of ER α were quantified using qPCR in collagenase-isolated adipocytes from 18-week-old chow fed females and males (n= 10/genotype). (B) Immunohistochemistry for ER α from male and female gonadal adipose tissue from WT and AdipoER α mice demonstrating significantly fewer ER α immunoreactive positive adipocytes in the AdipoER α mice relative to WT. Arrows point to ER α positive adipocyte nuclei, whereas the arrowhead demonstrates non-adipocyte nuclei positively stained for ER α demonstrating efficacy of the staining. (C and D) Weekly body weight was measured in singly housed female and male mice on normal chow (n= 10/genotype). (E) Weekly food intake was measured in singly housed female and male mice on normal chow (n= 6/genotype). (F and G) Body composition was measured by NMR in 15-week-old male and female mice (n= 10/genotype). (H) Gonadal fat pad weights in 18 week old females and males (n= 10/genotype). (I and J) Crait glucose tolerance tests (CGTT) were performed in 15-week-old females and males (n= 10/genotype). (K) Area under the curve (AUC) from the OGTT's was calculated. Data are presented as meant \pm SEM, and *P < 0.05 between WT and AdipoER α mice.



Figure 4: Adipocyte ER α regulates adipocyte size and gene function. (A) Representative photomicrographs of H&E staining of gonadal WAT from 18-week-old male and female mice. (B) Analysis of adipocyte cell size in both male and female of gonadal adipose tissue from 18-week-old mice (n=10/genotype). (C) Messenger RNA levels of F4/80 were quantified using qPCR in collagenase-isolated adipocyte adipose tissue from 18-week-old chow-fed females and males (n=10/genotype). (D) Messenger RNA levels of inflammatory genes were quantified using qPCR in collagenase-isolated adipocytes form 18-week-old chow-fed females and males (n=10/genotype). (D) Messenger RNA levels of inflammatory genes were quantified using qPCR in collagenase-isolated adipocytes form 18-week-old chow-fed females and males (n=10/genotype). (D) Messenger RNA levels of bitrosis genes were quantified using qPCR in collagenase-isolated adipocytes form 18-week-old chow-fed females and males (n=10/genotype). Data are presented as mean \pm SEM, *P < 0.05 between WT and AdipoER α mice, and P < 0.05 between male and female mice of like genotype.

tissue does not differ from their WT littermates. In contrast, AdipoER α males have much higher levels of macrophage infiltration, which correlates with enlarged unhealthy adipocytes.

To further measure the level of adipocyte inflammation, we isolated adipocytes, extracted RNA and performed qPCR for known inflammatory genes. Similar to the F4/80 data, there is no marked increase in inflammatory gene expression in female AdipoER α mice as compared to WT (Figure 4D). However, AdipoER α males show a substantial increase in inflammatory gene expression as compared to male WT levels. Similar to the inflammatory data, AdipoER α males show significant increases in expression of both LOX and COL6, while females show no apparent increase (Figure 4E).

Thus, these data together suggest that reduction of $ER\alpha$ levels in females to that of males results in male-like patterning in adiposity, without the metabolic complications of visceral adipose tissue deposition. By contrast, males do not have any alterations in adipose tissue

distribution when adipocyte ER α levels are reduced; however there are derangements in glucose homeostasis accompanied by elevated adipose tissue inflammation and fibrosis. These data demonstrate a sexual dimorphism with respect to the metabolic effects of gonadal adiposity and adipocyte ER α expression.

3.4. Adipocyte $\text{ER}\alpha$ is critical for the beneficial estrogenic effects on adipose tissue and metabolism in females

Despite having similar expression levels of ER α , AdipoER α females do not display the same metabolic consequences as WT males. AdipoER α females maintain normal ovarian production of estrogens and this may contribute to metabolic stability even in the absence of adipocyte ER α and the associated increase in gonadal adiposity. AdipoER α females cycle normally (data not shown); therefore, it could be hypothesized that the differential effects of adipocyte ER α in males and females could be levels of circulating estrogens. In order to determine the role of



adipocyte $ER\alpha$ in the capacity of circulating estrogens to regulate body adiposity and metabolic homeostasis, female WT and AdipoER α mice were ovariectomized (OVX) at 15 weeks of age and implanted with a subcutaneous pellet of estradiol-17 β or vehicle control (VEH). Postsurgical body weight demonstrates that OVX without supplementation of exogenous estrogens results in significant body weight gain in WT mice and with the addition of estradiol-17ß, the OVX-induced weight gain does not occur (Figure 5A). In the AdipoER α group, there is little to no observed estradiol-176 induced reduction in body weight gain (Figure 5B). Even more striking, the AdipoER α group that received VEH does not show the OVX-induced body weight gain, as seen in the WT group. Food intake data demonstrate that OVX WT VEH females eat significantly more as a per day average than WT mice receiving estradiol-17 β (Figure 5C); however this estrogenic reduction in food intake is not observed in the AdipoER α cohort. Further, OVX AdipoER α females eat significantly less than OVX WT females. OVX WT females treated with estradiol-17ß have decreased fat mass as compared to those treated with VEH (Figure 5D), while AdipoER α females show no effect of exogenous estrogens on body composition (Figure 5E). To this end, the weight of gonadal adipose tissues is decreased with estradiol-17ß administration in WT mice; however no such decrease is observed in AdipoER α mice when treated with exogenous estrogens (Figure 5F). Furthermore, our data demonstrate that the estrogenic reduction on body adiposity that occurs in WT females is dependent on adipocyte $ER\alpha$. Moreover, the OVX-induced weight gain is also dependent on adipocyte ER α , as OVX does not induce weight gain in AdipoER α females, thus suggesting that in the absence of circulating estrogens, adipocyte ER α may contribute to adipose tissue expansion.

In order to better understand the role of adipocyte ER α in the estrogenic regulation of metabolic homeostasis, WT and AdipoER α mice treated with VEH or estradiol-17 β were administered an OGTT. Data show that while exogenous estrogen improves glucose tolerance in WT OVX females (Figure 5G), it has little discernible effect in AdipoER α OVX females (Figure 5H). This is further demonstrated with area under the curve calculations; estradiol-17 β administration significantly reduces the area under the curve in WT OVX mice, however has no significant effect on the glucose curves in AdipoER α OVX females (Figure 5I).

Other data presented here demonstrate that in males and females, adipocyte ER α functions to maintain adipocyte size and lower levels of $ER\alpha$ results in enlarged adipocytes. To this end, OVX of WT female mice results in enlarged adipocytes and addition of exogenous estrogens results in a significant reduction in adipocyte area (Figure 6A and B), which is consistent with previous reports [37]. While there is a slight trend for a reduction in adipocyte area with estradiol-17ß treatment of AdipoER α OVX females, it does not attain significance. Interestingly, gonadal adipocytes from AdipoER α OVX mice treated with VEH are significantly smaller than those from WT OVX treated with VEH, thus suggesting that in the absence of exogenous estrogens, adjpocyte $ER\alpha$ acts adversely and enhances adipocyte size. Similar to trends in previous data. F4/80 expression is drastically decreased in AdipoER α OVX treated with VEH as compared to WT OVX treated with VEH, again suggesting that $ER\alpha$, in the absence of circulating estrogens, may have unfavorable effects on adipose tissue macrophage infiltration (Figure 6C). Administration of exogenous estradiol-17 β to WT OVX results in a decrease in the expression of the pro-inflammatory genes TNF α , TLR4 and SAA3 (Figure 6D); however, in AdipoER α OVX the addition of estradiol-17ß actually enhances expression of both TLR4 and SAA3 (there is no observable effect on TNF α expression). Markers of adipose tissue fibrosis, LOX and COL6, both decrease with administration of estradiol-17 β in WT OVX females; however no decrease is observed in AdipoER α females (Figure 6E). Interestingly, there is no improvement in fibrotic gene expression in AdipoER α OVX females over WT OVX females treated with VEH, as observed in other measures of adipocyte dysfunction. Taken together, these data suggest that adipocyte ER α is critical for the estrogen-induced improvements in adipocyte metabolism. In the absence of circulating estrogens, our data suggest that reduced adipocyte ER α expression helps counter the OVX-induced adipose tissue inflammation and fibrosis, thus improving metabolic homeostasis.

3.5. Adipocyte-specific knockdown of ER α in ER β null mice (AdipoER $\alpha/$ β ERKO)

Estrogen receptor beta (ER β) is also expressed in adipose tissue [20] and following adipocyte specific knockdown of ER α there is a significant increase in the expression of ER β in both females and males (Figure 7A). Therefore, to understand the contribution of ER β on the outcomes of adipocyte-specific ER α knockdown, we generated mice with adipocyte-specific knockdown of ER α (AdipoER α) in the β ERKO background (AdipoER α/β ERKO). Similar to our previous results, we demonstrate about ~50% knockdown of ER α (Figure 7B) and 100% knockdown of ER β (Figure 7C) by RNA expression of collagenase-mediated isolation of adipocytes in both male and female AdipoER α/β ERKO mice as compared to their WT littermates.

Male and female AdipoER α/β ERKO are born in normal Mendelian ratio and are viable. At weaning, AdipoER α and AdipoER α/β ERKO males and females do not differ in body weight (Figure 7D and E) or Estradiol 17 β levels (mice sacrificed at 12 weeks of age in the proestrus stage of their cycle for comparison; AdipoER α 49.6 \pm 6.9 pg/mL vs AdipoER α/β ERKO 54.5 \pm 5.9). NMR imaging demonstrates that both AdipoER α/β ERKO males and females do not differ from AdipoER α of comparable sex on any measure of body composition (Figure 7F and G). These data suggest that lack of both ER β and adipocyte ER α does not enhance the already existing body weight phenotype of the AdipoER α mice.

In order to determine if there are additional metabolic consequences of reduced adipocyte ER α and ER β , oral glucose tolerance tests (OGTT) were performed on male and female WT and AdipoER α/β ERKO mice (cycle day was determined at the time of the test, but did not factor into the results). Here, we find AdipoER α/β ERKO females had reduced glucose clearance when compared to the AdipoER α females (Figure 7H). For the males, we do not see any difference from the already impaired glucose clearance previously observed in the AdipoER α (Figure 7I). We found that the gonadal adipocytes in both male and female AdipoER α/β ERKO mice are similar to those of AdipoER α mice (Figure 8A and B). Consistent with the changes in glucose clearance, we found increases in F4/80 (Figure 8C), markers of inflammation (Figure 8D), and fibrosis (Figure 8E) in both sexes when compared to the AdipoER α indicating that in the absence of adipocyte ER α , ER β plays a protective role in adipocyte fibrosis and inflammation.

4. **DISCUSSION**

Here we demonstrate the critical role of adipose tissue and adipocyte ER α in regulating adipocyte size, fibrosis, and inflammation. ER α is well established to regulate metabolic homeostasis and the ER α whole body knockout displays obesity and metabolic compromise [29]. Here we extend those findings to indicate they also have inflamed and fibrotic adipose tissue independent of differences in body weight. Our data demonstrate that knockdown of ER α in gonadal adipose tissues results in an enlarged depot with larger adipocytes and with increased



Figure 5: Adipocyte ER α regulates the capacity of estradiol-17 β to modulate body weight and adipose tissue distribution. All females were OVX and administered VEH or estradiol-17 β at 12 weeks and all post-sacrifice measurements were taken in 18-week-old mice, 42 days post-OVX (n=8/genotype). (A and B) Daily body weight was measured in singly housed female mice post-OVX on normal chow (n=8/genotype). (C) Daily food intake was measured in singly housed female mice post-OVX on normal chow (n=8/genotype). (C) Daily food intake was measured in singly housed female mice post-OVX on weight were weighed. Data presented as a fold change over VEH of pad weight mouse body weight. (G and H) Oral glucose tolerance test (OGTT) was performed in 16-week-old females (n=8/genotype). (D) Area under the curve (AUC) was calculated for the OGTTs. Data are presented as mean ± SEM, and *P < 0.05 between WT and AdipoER α mice and # P < 0.05 between VEH and 17 β -estradiol.

expression of a marker of macrophages, local inflammation, and fibrosis. Altering ER α levels in total gonadal adipose tissues has similar effects in males and females. These data demonstrate that many

adipose tissue effects observed in the total body ER α knockout can be attributed to lack of ER α selectively in adipose tissues. We further demonstrate that adipocyte ER α *per se* appears to be critical in





Figure 6: Adipocyte ERα regulates the capacity of estradiol-17β to modulate adipose tissue function, adipocyte size and glucose homeostasis. All females were OVX and administered VEH or Figure 0. Adjourds that globales the regulates the capacity of establish p to modulate adjourds that p at 12 weeks and all post-sacrifice measurements were taken in 18-week-old mice, 42 days post-OVX (n=8/genotype). (A) Representative photomicrographs of H&E staining of gonadal WAT. (B) Analysis of adjoocyte cell size in gonadal WAT from 18-week-old mice, 42 days post-OVX (n=8/genotype). (C) Messenger RNA levels of F4/80 were quantified using qPCR in collagenase-isolated from 18-week-old chow-fed females (n=8/genotype). (D and E) Messenger RNA levels of indicated genes were quantified using qPCR in Collagenase-isolated from 18-week-old chow fed females (n=8/genotype). (D and E) Messenger RNA levels of indicated genes were quantified using qPCR in Collagenase-isolated from 18-week-old chow fed females (n=8/genotype). Data are presented as mean ± SEM, and *P<0.05 between WT and AdipoER α mice and P<0.05 between VEH and 17 β -estradiol.

regulating adjpocyte size, inflammation, and fibrosis. We demonstrate this by using mice with selective knockdown of ER α in adipocytes, which results in significantly increased fibrosis and inflammation selectively in the males. Lastly, our data suggest that in the absence of adipocyte ER α , ER β provides a protective role in adipose tissue inflammation and fibrosis. Our data for the first time begin to delineate the roles of ER α and ER β in modulating adipose tissues.

0

LOX wт

COL6

LOX

COL6

AdipoERα

4.1. αERKO mice have enlarged adipocytes with increased inflammation and fibrosis in both males and females

It has previously been demonstrated by Cooke et al. [30] that α ERKO mice have enlarged fat cells relative to WT/control mice. Here we confirm these data in weight-matched male and female mice, and extend these findings to demonstrate increased levels of inflammation and fibrosis in adipose tissue.

4.2. Viral-mediated knockdown of ER α in visceral adipose tissue results in enlarged adipocytes with increased inflammation and fibrosis in males and females

In order to address the role of $ER\alpha$ in individual adipose depots, without the complications of changing the metabolic or hormonal profile in vivo, our lab has developed a novel technique with which we can site specifically modulate gene expression in a fat pad specific way using adeno-associated viral (AAV) technology. We demonstrate for the first time that gonadal fat pad specific knockdown of ER α results in increased adipose tissue weight, adipocyte size, and increased markers of inflammation and fibrosis. The purpose of this experiment is to demonstrate, regardless of the hormonal milieu, that fat pad specific knockdown of ER α has a profound effect. This experimental paradigm was not conducted to determine the metabolic perturbations associated with fat pad knockdown of ER α , but rather to probe/interrogate the importance of gonadal adipose tissue $ER\alpha$ per se.



Figure 7: Contribution of ER β to the AdipoER α phenotype. (A) ER β is upregulated in gonadal adipose tissue in male and female AdipoER α mice (n=9/genotype, 18-week old at sacrifice). (B) ER α mRNA from isolated adipocytes is reduced to similar levels in AdipoER α and AdipoER α /BERKO mice (n=9/genotype). (C) ER β mRNA from isolated adipocytes is reduced to similar levels in AdipoER α and AdipoER α /BERKO mice (n=9-10/genotype). (D and E) Weekly body weight was measured in AdipoER α and AdipoER α /BERKO singly housed female and male mice on normal chow (n=6/genotype). (F and G) Body composition was measured by NMR in 15-week-old AdipoER α and AdipoER α /BERKO male and female mice (n=9-10/genotype). (H and I) Oral glucose tolerance tests (OGTT) were performed in 15-week-old AdipoER α /BERKO females and males (n=9-10/genotype).





Figure 8: (A) Representative photomicrographs of H&E staining of VISC WAT from 18-week-old male and female mice. (B) Analysis of adipocyte cell size in both AdipoER α and AdipoER α /βERKO male and female of gonadal adipose tissue from 18-week-old mice (n=9-10/genotype). (C) Messenger RNA levels of F4/80 were quantified using qPCR whole adipose tissue from 18-week-old AdipoER α and AdipoER α and AdipoER α /βERKO chow-fed females and males (n=9-10/genotype). (D) and E) Messenger RNA levels were quantified using qPCR of whole adipose tissue from 18-week-old AdipoER α and AdipoER α /βERKO chow-fed females and males (n=9-10/genotype). (D and E) Messenger RNA levels were quantified using qPCR of whole adipose tissue from 18-week-old AdipoER α and AdipoER α /βERKO chow-fed females and males (n=9-10/genotype). Data are presented as mean \pm SEM, and *P < 0.05 between AdipoER α and AdipoER α /βERKO.

4.3. Adipocyte-specific ER α regulates body adiposity, adipocyte inflammation, fibrosis and systemic glucose tolerance in a sexually dimorphic way

Adipose tissues are comprised of many different cell types, including adipocytes, inflammatory cells and preadipocytes. In order to understand the specific role of adipocyte ER α we used the adipocyte-specific Adiponectin-CRE crossed to the ER α -floxed mouse, thus altering ER α only in adipocytes. We chose to work with mouse lines where adipocyte ER α in female mice is reduced to levels seen in males. Our data demonstrate that reductions in adipocyte ER α levels result in increased body weight, adipose tissue mass, and adipocyte size in females. This increased adiposity, however, does not result in altered metabolic homeostasis, adipose tissue inflammation or fibrosis. Contrary to females, reduced adipocyte ER α in males does not alter body weight or adiposity; however, adipocyte size is enlarged and accompanied by increased adipose tissue inflammation and fibrosis. Furthermore, glucose tolerance is blunted in AdipoER α males, but not in females.

Despite enlarged adipose tissues, AdipoER α females remain protected from the obesity-associated co-morbidities seen in males. While males have increased levels of macrophage infiltration and elevated adipocyte inflammation and fibrosis, AdipoER α females display none of these characteristics. Interestingly, knockdown of whole adipose depot $ER\alpha$ in females results in an upregulation of macrophage infiltration, adipocyte inflammation and fibrosis, similar to male levels. Together these data suggest other adipose tissue cell types, and the ER α levels within those cell types regulate the obesity-associated co-morbidities in females. A recently published study demonstrates that knockout of macrophage $ER\alpha$ expression in females results in obesity and metabolic complications [38]. The discontinuity between the data in females with total depot knockdown and adipocyte-specific knockdown could be the fact that, unlike in the depot-specific knockdown, the macrophages in the AdipoER α females retain their ER α levels, thus reducing the inflammatory state. In addition, AdipoER α females do not have altered glucose clearance, while reducing adipocyte $ER\alpha$ in males results in profound

metabolic compromise. This could be in part due to changes in adipocyte metabolism and inflammatory state. Alternatively, estrogen signaling in the brain and other peripheral tissues may compensate to maintain systemic metabolic homeostasis in females.

Given this possibility, we investigated the role of adipocyte ER α in the ability of estrogens to regulate both adipose tissue and systemic metabolism. Circulating estrogens are known to influence adipose tissue deposition and whole body metabolic homeostasis [19,22]. Additionally, the contribution of the several-fold increase in circulating E2 levels in α ERKO females [39] has been guestioned as to its contribution of obesity/adiposity phenotype. Specifically, circulating estrogens in the absence of functional ER α could have adverse effects on metabolic homeostasis. Our data demonstrate that OVX WT females respond metabolically to exogenous estrogens in a beneficial way, to include decreased adiposity and adipose tissue inflammation and fibrosis and improved glucose tolerance which may be mediated peripherally as well as centrally. OVX AdipoER α mice, however, seem largely unresponsive to estrogens with respect to body weight and adiposity. Moreover, the addition of estradiol-17 β to AdipoER α females results in increased fibrosis and inflammation, thus suggesting that estrogens can have a negative impact locally on adipose tissue with reduced levels of adipoctye ER α expression. As the OVX AdipoER α females express $ER\alpha$ to levels seen in males, we compared levels of adiposity and adipocyte gene expression between the two groups. Even with little circulating estrogens and low adipocyte $ER\alpha$ expression, OVX AdipoER α females retain better glucose clearance than did males. Similar to results described above, OVX AdipoER α females have far lower adipose tissue macrophage infiltration than WT males. Measures of adipocyte size as well as inflammation and fibrosis, however, seem

on par with levels in male adipocytes. These data provide further support that the levels of adipocyte ER α are critical in determining adipose tissue deposition and adipocyte size. However, once circulating estrogens (as seen in the intact AdipoER α females) are removed, adipocyte dysfunction appears.

Another intriguing observation is the apparent improvement of the OVX AdipoER α treated with VEH over the OVX WT treated with VEH. These data suggest that in the absence of circulating estrogens, adipocyte ER α has antagonistic actions with respect to adiposity and adipocyte function. Even more, this is also seen in the OGTT AUC, where there is a slight but significant improvement in the OVX AdipoER α VEH over OVX WT VEH. The estrogen-induced pathways that regulate metabolic function are not completely understood and it is likely that, in the absence of sufficient ligand, ER α may bind to alternative estrogen response elements (EREs) or other factors critical to adipocyte function and induce metabolically detrimental effects. In addition, ER α could also associate with an alternative group of signaling factors in the absence of ligand, thus inducing a differential signaling program. Either way these data demonstrate that the presence of ligand or receptor alone can be detrimental to metabolic function.

4.4. Adipocyte-specific $\text{ER}\alpha/\beta\text{ERKO}$ regulates body adiposity, adipocyte inflammation, fibrosis and systemic glucose tolerance in a sexually dimorphic way

The specific function of ER α in adipose tissue has previously been unknown. Additionally, the ratio of ER α /ER β could be important with respect to adipose tissue function. The absence of heterodimer formation by ER α and ER β [40] or other putative estrogen receptors [41] in total body α ERKO mice could play a role in the development of



Figure 9: Cartoon depiction of our results. (A) 'Healthy' female adipose tissue represented by small (expandable) adipocytes (that contain ER α), low levels of inflammation and fibrosis. (B) Reductions in adipocyte ER α as seen in males (relative to females) and AdipoER α females result in larger (expandable) adipocytes with low levels of inflammation and fibrosis. (C) Knockout of ER α (as seen in AdipoER α males) results in enlarged adipocytes that lack the ability to expand further due to increased levels of inflammation and fibrosis. (D) Knockout of ER α form total adipose tissues results in enlarged adipocytes that lack the ability to expand further due to increased levels of inflammation and fibrosis. (E) Reductions in adipocyte ER α and knockout of adipose tissue ER β result in enlarged to expand due to increased levels of inflammation and fibrosis.



the obese/adipose tissue phenotype. Interestingly, the double knockout of ER α and ER β does not seem to have metabolic effects over that of the $ER\alpha$ knockout alone [42], suggesting that $ER\beta$ functions may require the presence of ER α . In our AdipoER α model, adipocyte ER α is only downregulated ~60% and metabolically does not have a huge impact in females. It is possible that $ER\beta$ expression plays a protective role in the state of reduced ER α through enhanced estrogen signaling through ER β or other estrogen receptors [41] thereby mediating the effects on WAT. Therefore, to test the contribution of ER β to the AdipoER α phenotype, we crossed the AdipoER α mouse to the ER β total body knockout, creating a mouse that lacks $ER\beta$ in adipose tissue with reduced levels of adipocyte ER α . Our findings suggest that in the absence of adipocyte ER α , ER β plays a protective role in suppressing inflammation and fibrosis in the females. However, in the absence of both $ER\alpha$ and $ER\beta$ there is a significant increase in adipose tissue markers of fibrosis, inflammation, macrophage infiltration in both sexes. Interestingly, knocking out $ER\beta$ in female AdipoER α mice reduces glucose clearance while in males this effect is not above and beyond that of the AdipoER α male mice. Our data presented here begin to define the roles of ER α and ER β in modulating adipose tissue function. Additionally, our data suggest that ER β may serve a protective role in states of reduced ER α expression.

5. CONCLUSIONS

We demonstrate for the first time that adipose tissue ER α regulates body fat distribution, adipose tissue inflammation, and fibrosis. Our data further suggest that adjpocyte ER α is required for many of the metabolically beneficial effects of estrogens on metabolism. In males, extremely low levels of adipocyte $ER\alpha$ result in significant increases in adipose tissue fibrosis, with systemic (data not shown) as well as adipose tissue inflammation, which results in metabolic dysfunction (Figure 9). AdipoER α females retain modest levels of both adipose tissue $ER\alpha$ and $ER\beta$ which appears to facilitate adipose tissue expansion (as indicated by increased adipocyte size), reductions in markers of fibrosis, reductions in markers of systemic (data not shown) as well as adipose tissue inflammation, and increased adiposity. Therefore, our data are consistent with recent research focusing on inflammation and the metabolic syndrome. Importantly, our data suggest that the expandability of the fat cell suppresses adipose tissue and systemic inflammation and is 'protective' for the metabolic syndrome and that both ER α and ER β play critical roles in modulating the expandability of adipocytes. Together, our studies demonstrate for the first time the effects of adipocyte-specific $ER\alpha$ in males and females, and the requirement of adipocyte ER α for estrogen-mediated metabolic effects.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

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