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# Insights of the role of estrogen in obesity from two models of ERa deletion

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#### **Abstract**

Sex hormones play a pivotal role in physiology and disease. Estrogen, the female sex hormone, has been long implicated in having protective roles against obesity. However, the direct impact of estrogens in white adipose tissue (WAT) function and growth are not understood. Here, we show that deletion of estrogen receptor alpha (ERa) from adipocytes using Adiponectin-cre does not affect adipose mass in male or female mice under normal or high-fat diet (HFD) conditions. However, loss of ERa in adipocyte precursor cells (APs) via *PdgfRa-cre* leads to exacerbated obesity upon HFD feeding in both male and female mice, with subcutaneous adipose (SWAT)-specific expansion in male mice. Further characterization of these mice revealed infertility and increased plasma levels of sex hormones, including estradiol in female mice and androgens in male mice. These findings compromise the study of estrogen signaling within the adipocyte lineage using the PdgfRa-cre strain. However, AP transplant studies demonstrate that the increased AP hyperplasia in male SWAT upon PdgfRa-cre-mediated ablation of ERa is not driven by AP-intrinsic mechanisms, but are rather mediated by off-target effects. These data highlight the inherent difficulties in studying models that disrupt the intricate balance of sex hormones. Thus, better approaches are needed to study the cellular and molecular mechanisms of sex hormones in obesity and disease.

#### **Keywords**

obesity; adipogenesis;	adipose tissue;	estrogen signalin	g	

RSP and MSR designed experiments. RSP and NT performed experiments. RSP and NT analyzed data. RSP and MSR interpreted data and wrote the manuscript. RSP, NT, and MSR approved the final version of the manuscript.

Declaration of interest

RSP, NT, and MSR declare no conflicts of interest.

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#### Introduction

Obesity is defined as the excessive accumulation of WAT. The role of sex hormones in obesity and disease has been recognized for decades (Cooke and Naaz 2004; Jones, et al. 2000; Marin and Arver 1998; Seidell, et al. 1990). Sex hormones influence many aspects that can lead to obesity including food intake, energy expenditure, and the expansion of WAT (Dubuc 1985; Heine, et al. 2000; Krause, et al. 2021; Musatov, et al. 2007; Ramirez 1980; Wurtman and Baum 1980). In particular, observations in mice and humans have indicated that estrogen has an overall protective effect against obesity and metabolic disease (Andersson, et al. 1997; Stubbins, et al. 2012). After menopause, when estrogen levels in plasma decline, women are more prone to obesity and metabolic dysfunction (Aloia, et al. 1996; Andersson et al. 1997; Stubbins et al. 2012; Toth, et al. 2000). Estrogen replacement therapy is able to ameliorate these effects in humans (Andersson et al. 1997) and in ovariectomized (OVX) female mice (Stubbins et al. 2012). Furthermore, loss of function mutations in the aromatase gene, which is required for estrogen production (Meyer 1955; Meyer, et al. 1955), causes increased fat mass, hyperinsulinemia, elevated cholesterol, and fatty livers (Bilezikian, et al. 1998; Carani, et al. 1997; Conte, et al. 1994; Jones et al. 2000; Jones, et al. 2001; Morishima, et al. 1995). Similar effects are also seen upon treatment with aromatase inhibitors (Gibb, et al. 2016; Kauffman, et al. 2015). In addition, modulating the function of estrogen receptors impacts obesity and metabolism. Whole-body deletion of estrogen receptor alpha (ERa) but not ERB leads to obesity in mice (Heine et al. 2000; Ohlsson, et al. 2000) while deletion of GPR30 receptor protects female mice from obesity (Wang, et al. 2016). Thus, both ERa and GPR30 influence obesity but the direct role of estrogen signaling on adipose biology remains unknown.

There are two main mechanisms of fat mass expansion: hypertrophy (increase in size of mature adipocytes) and hyperplasia (increase in number of mature adipocytes). Estrogen has been implicated to play a role in both of these processes. Estrogens regulate hypertrophy by modulating lipolysis and lipogenesis of mature adipocytes (Gavin, et al. 2013; Gormsen, et al. 2012; Monjo, et al. 2005; Stubbins et al. 2012; Zang, et al. 2007), and hyperplasia via affecting the differentiation of adipocyte precursors (APs) (Jeffery, et al. 2016). As mature adipocytes are post-mitotic, the generation of new adipocytes requires the proliferation and differentiation of APs (Berry and Rodeheffer 2013; Cristancho and Lazar 2011; Rodeheffer, et al. 2008). Recent work from our lab has elucidated the role of sex differences in adipocyte hyperplasia (Jeffery, et al. 2015; Jeffery et al. 2016; Sebo and Rodeheffer 2021). Male mice undergo adipocyte hyperplasia in obesity specifically in visceral fat (VWAT), while female mice display hyperplasia in both VWAT and SWAT. However, OVX females have VWAT-specific AP proliferation on HFD, similar to males (Jeffery et al. 2016). Furthermore, treating males with estrogen when on a HFD increases SWAT AP proliferation (Jeffery et al. 2016). Therefore, estrogen appears to influence the sexual dimorphic distribution of WAT in obesity via adipocyte hyperplasia in SWAT, but the cellular and molecular mechanisms are not understood. In this study, we use targeted models of ERa deletion to study the role of estrogen signaling in the adipocyte cellular lineage in obesity.

#### **Materials and Methods**

#### Animals

The Institutional Animal Care and Use Committee (IACUC) at Yale University approved all animal studies. All animals were kept in temperature and humidity-controlled rooms on a 12-hour:12-hour light:dark cycle, with lights on from 7:00 a.m. to 7:00 p.m. All mice used for these studies were on the C57BL/6J genetic background. *Adipoq-cre* mice (stock no. 028020) and *PdgfRa-cre* mice (stock no. 013148) were purchased from Jackson Laboratories. Esr1<sup>fl</sup> mice were a generous gift from Dr. Sean Morrison (UT Southwestern, Dallas, TX, USA). All cre lines were crossed to the mTmG (stock no. 007676) mice purchased from Jackson Laboratories. Breeding was done in the Yale Animal Resource Center and mice were weaned at p21. Unless otherwise noted, mice were males and females 6–9 weeks of age at the start of experiments. VWAT refers to the perigonadal WAT and SWAT refers to the inguinal WAT in mice. Body composition measures were done with NMR using the Echo MRI whole body composition analyzer (Echo Medical System, Houston, TX). High-fat diet is from Research Diets (D12492). Standard diet is from Harlan Laboratories (2018S).

#### RNA extraction and cDNA synthesis

For gene expression analysis, whole tissues were collected and stored at –80C until RNA extraction was performed. For adipocyte fractionation, WAT was collected and digested with 0.8mg/mL collagenase type 2 for 45 – 60 min in a shaking water bath at 37C. Samples were passed through 40um filter and centrifuged 300 g for 3 minutes. Stromal vascular fraction (SVF) was removed using oral gavage needle and adipocyte layer was left intact. Both SVF and adipocytes were rinsed with DPBS (Life Technologies) and resuspended in Trizol reagent. RNA was isolated using Direct-zol RNA Miniprep Kit (Zymo Research), according to manufacturers' instructions. RNA was quantified by a nanodrop spectrophotometer (Thermo Fisher Scientific) and single stranded cDNA was synthesized from total RNA using the High-Capacity cDNA reverse Transcription Kit (Applied Biosystems, Life Technologies) according to the manufacturer's instructions.

#### **Quantitative real-time PCR**

Quantitative RT-PCR (qPCR) was performed on the cDNA using the Sybr green method of quantification on a Roche Lightcycler 480 using a SYBR FAST quantitative PCR kit (Kapa Biosystems; KK4611). Gene expression was analyzed for each sample in triplicate using the forward and reverse primers indicated on Table 1. Housekeeping genes were tbk1 or b-actin as deemed appropriate. For each experiment, cDNA samples were pooled and a standard curve was generated to quantify relative mRNA transcript levels. To confirm effective adipocyte fractionation, expression of *adipoq* and *pdgfra* were assessed in SVF and adipocyte fractions.

#### **BrdU treatments**

For BrdU experiments, BrdU (US Biological, B2850) was given in the drinking water at 0.8 mg/mL for experiments lasting one week or less and 0.4 mg/mL for experiments lasting more than one week. BrdU water was replenished every 48 hours.

#### **Confocal microscopy**

Adipose tissue was collected, embedded in paraffin, and stained as previously described (Holtrup, et al. 2017; Jeffery et al. 2015; Jeffery et al. 2016; Sebo and Rodeheffer 2021). For adipocyte nuclei analysis, 20–30 images for every tissue section were acquired at 40X with a Leica TCS SP5 confocal microscope. Quantification of BrdU in adipocyte nuclei was done as previously described (Jeffery et al. 2015). At least 50 adipocyte nuclei were scored for each animal.

For adipocyte diameter measurements, the area of each adipocyte (in square pixels) was measured using Cell Profiler. The diameter of each adipocyte was calculated using the measured area, assuming each adipocyte is a perfect circle. At least 200 adipocytes were measured for each animal.

For whole mount microscopy, tissues were dissected and cut into ~1.5×1.5 cm pieces. Samples were subsequently mounted onto microscope slides with Fluoromount-G (SouthernBiotech, 0100–01) and imaged at 20X with a Leica TCS SP5 confocal microscope.

#### Flow cytometry

Flow cytometry was performed as described previously (Jeffery et al. 2015) for BrdU analysis with the following antibodies: CD45 APC-eFluor 780 (eBioscience; 47–0451–80) at 1:1,000, CD31 PE-Cy7 (eBioscience, 25–0311–82), at 1:500, CD29 Alexa Fluor 700 (BioLegend, 102218) at 1:400 and Sca-1 V500 (BD Horizon, 561228) at 1:300. Cells were washed and then fixed and permeabilized using Phosflow lyse/fix and Perm Buffer III (BD Biosciences) according to the manufacturer's recommendations. Cells were then treated with DNase (deoxyribonuclease I; Worthington; × 200 units/ml) in DPBS (Sigma; with calcium chloride and magnesium chloride) for 2-hrs at 37C and then washed in HBSS with 3% BSA. Cells were then stained with anti-BrdU antibody (Alexa Fluor 647; Phoenix Flow Systems; AX647) at 1:30 in HBSS with 3% BSA overnight in the dark at 4C. Cells were then washed in HBSS with 3% BSA and incubated with CD34 Brilliant Violet 421 (BioLegend 230 119321) at 1:400 and CD24 PerCP-Cyanine 5.5 (eBioscience, 45–0242–80) at 1:250. Following antibody incubation, samples were washed and analyzed on a BD LSRII analyzer. Data analysis was performed using BD FACS Diva software (BD Biosciences).

#### Hormone quantification

Plasma 17-β estradiol was measured using Cayman's ELISA assay (No. 501890). All other sex hormones were measured with LC-MS/MS by OpAns LLC.

#### **Transplant Assay**

For cell transplant assays, APs were isolated and transplanted in WAT as previously described (Jeffery et al. 2016; Rodeheffer et al. 2008). For mTmG samples, NucRed<sup>TM</sup>

Live 647 ReadyProbes<sup>™</sup> Reagent (Thermo Fischer, R37106) was used as live/dead stain at 1:600 dilution. VWAT and SWAT of AP-ERαKO animals were pooled and GFP expression confirmed via FACS. Recipient male C57Bl/6J mice were anesthetized with isoflurane and surgeries performed using sterile technique. 0.5−1 million ERαKO AP cells were re-suspended in 15uL of PBS and injected into the left SWAT of 4−5-week-old congenic wildtype mice. A control PBS injection was done in the right SWAT. Mice were allowed to recover for 2 weeks, then placed on a HFD and treated with BrdU for 1 week. Left and right SWAT tissues were collected and analyzed separately via flow cytometry for incorporation with BrdU. ERαKO APs were identified by GFP fluorescence. Results were counted only for transplants in which more than 100 individual GFP-positive donor AP cells were recovered in the recipient SWAT.

#### **Statistical Analysis**

Statistical analyses are described in each figure legend. All tests were performed using GraphPad Prism version 9.0. Data are presented as mean  $\pm$  s.e.m. and p<0.05 was considered statistically significant. A minimum of 5 animals were used for each experiment, unless statistical significance was reached with fewer animals. Sample size is indicated in each figure legend. Experiments were not blinded, as genotypes of mice were known prior to analysis.

#### Results

#### Adipocyte Estrogen Receptor alpha (ERa) signaling does not contribute to obesity

Estrogen signaling plays an important role in both female and male physiology, as both sexes of whole-body ERa-KO and Aromatase-KO mice have significant WAT accumulation via both hypertrophy and hyperplasia (Heine et al. 2000; Jones et al. 2000; Jones et al. 2001). To determine if estrogen affects mature adipocyte function directly, we crossed adipocyte-specific Adiponectin-cre mice (Eguchi, et al. 2011) to Esr1<sup>fl/fl</sup> mice (Feng, et al. 2007) to create Adi-ERa-KO mice (Figure 1A). To confirm the knockout model is effective, we isolated mature adipocytes and stromal vascular fraction (SVF) in WAT and performed quantitative PCR. After confirming effective adipocyte isolation by enrichment of adipoq (Supp. Figure 1A, 1C), we found a 90% reduction in ERa transcripts in adipocytes from both VWAT and SWAT in males and females (Figure 1B-C) with no impact on ERa expression in SVF (Supp. Figure 1B, 1D), demonstrating efficient and specific knockout of ERα. To determine if loss of ERα affects WAT accumulation, we fed Adi-ERα-KO mice and control littermates (ERa f/f) standard diet (SD) or HFD for 8 weeks and measured body composition via echo-MRI throughout. Adi-ERa-KO females had no differences in body weight accumulation in either diet (Figure 1D-E) but had a reduction in total fat mass after a HFD (Supp. Figure 2A–B). Adi-ERα-KO males on a HFD do not show differences in body composition (Figure 1F–G, Supp. Figure 3A–B), but Adi-ERa-KOs are slightly heavier than controls on a SD (Figure 1F). This increase is not due to differences in fat mass or lean mass (Figure 1G, Supp. Figure 3A–B). Together, these data suggest that adipocyte ERa signaling does not contribute to the obesity phenotypes seen in whole body ERa-KO males and females.

#### Adipocyte ERa signaling modestly impacts WAT accumulation in female mice

Even though there are no differences in body composition overall in Adi-ERα-KO mice, we assessed if differences in WAT distribution were present. After 8 weeks of HFD, both Adi-ERα-KO and control females accumulate more WAT than females on SD (Figure 2A). However, we found a significant reduction in VWAT mass in Adi-ERα-KO females on a HFD and a trend towards a reduction in SWAT (Figure 2A). Of note, there were no significant differences between liver or intrascapular brown adipose (iBAT) weights (Supp Figure 2C–D). In male Adi-ERα-KO mice, there are no differences in VWAT or SWAT accumulation on either diet (Supp. Figure 3C) or liver and iBAT weights (Supp. Figure 3D–E). These data indicate that adipocyte ERα does not play a role in male WAT expansion and only a minor role in female WAT expansion.

To determine if the reduction of VWAT in Adi-ERα-KO females on a HFD is due to changes in adipocyte hyperplasia, we employed a BrdU pulse-chase assay (Figure 2B) to measure the formation of new adipocytes, as described previously (Jeffery et al. 2015; Jeffery et al. 2016). After staining with caveolin to label the adipocyte plasma membrane, BrdU, and DAPI (Figure 2C), we found no differences in BrdU incorporation into mature adipocyte nuclei of female Adi-ERα-KO mice on a HFD (Figure 2D). To determine if adipocyte hypertrophy is affected in this model, we measured the diameter of adipocytes in WAT. Although there are no differences on the average adipocyte size between the HFD groups (Figure 2E), we found a slight VWAT-specific shift towards smaller adipocytes in Adi-ERα-KO females (Figure 2F, Supp. Figure 2E). Of note, VWAT adipocytes were larger than SWAT adipocytes in both groups (Figure 2E). These data indicate that lack of ERα signaling in adipocytes has a small effect on adipocyte size an does not affect adipocyte formation in female mice on HFD. Overall, the findings of the hyperplasia and hypertrophy assays indicate adipocyte-intrinsic ERα signaling does not significantly impact fat mass accumulation, even in obesity, in males or females.

# Ablation of $\mathsf{ER}\alpha$ in adipocyte precursors changes obesogenic AP proliferation patterns in mice

As loss of ERα in adipocytes does not significantly impact obesity in mice, we next focused on a potential role of ERα in APs in obesity. These cells do not express *adipoq* and therefore are not targeted in the Adi-ERα-KO mouse model (Berry and Rodeheffer 2013; Lee, et al. 2012). Thus, we deleted ERα in APs by crossing the *PdgfRa-cre* strain (Figure 3A) (Berry and Rodeheffer 2013) to *Esr1*<sup>fl/fl</sup> mice (Feng et al. 2007) to create the AP-ERα-KO mice. It is important to note that the mature adipocytes in this model also lack ERα. When quantifying the expression of ERα, we found the mice have a significant reduction of ERα in WAT from both male and female mice (Figure 3B). Furthermore, ERβ remains unchanged and lowly expressed in WAT (Supp. Figure 4A). To test obesogenic AP proliferation, we used flow-cytometry to measure BrdU incorporation into AP cells after 1 week of SD or HFD (Berry and Rodeheffer 2013; Jeffery et al. 2015; Jeffery et al. 2016; Rodeheffer et al. 2008). In this assay, we found female AP-ERα-KO still have HFD-induced AP proliferation, but, the APs in the AP-ERα-KO females have a more consistent proliferative response to HFD that is similar to the maximal response observed in control female mice (Figure 3C). Male AP-ERα-KO mice display normal HFD-induced VWAT

AP proliferation, however, unlike controls, these male mice also have increased SWAT AP proliferation on a HFD (Figure 3D). Thus, male AP-ER $\alpha$ -KO have a female pattern of HFD-induced AP proliferation. These data indicate that deleting ER $\alpha$  with PdgfRa-cre affects AP proliferation in male mice upon HFD feeding.

# Ablation of ER $\alpha$ in adipocyte precursors causes increased fat mass accumulation upon HFD feeding

As AP-ER $\alpha$ -KO females have HFD-induced AP proliferation, we next assessed if this proliferation is indeed adipogenic. AP-ER $\alpha$ -KO females and controls were fed a SD or HFD for 8 weeks and body composition was assessed weekly. Interestingly, AP-ER $\alpha$ -KO females gained significant more weight on the HFD compared to controls (Figure 4A) with increased fat and lean mass accumulation (Figure 4B). Although no changes in VWAT and SWAT mass, and thus fat distribution, were found, AP-ER $\alpha$ -KO females consistently gain fat mass on a HFD (Figure 4C). There are no differences in WAT accumulation on a SD (Figure 4C), indicating these phenotypes are HFD-specific.

Next, we determined if the differences in fat mass in AP-ERa-KOs were caused by increased hypertrophy or hyperplasia using the BrdU pulse-chase assay (figure 2B) to quantify the formation of newly formed adipocytes (Figure 4D). Interestingly, we find a trend towards more hyperplasia in VWAT and SWAT of AP-ERa-KO females on HFD (Figure 4E). When adipocyte size was measured, AP-ERa-KO females on a HFD showed a trend towards larger adipocytes in both VWAT and SWAT, but this result did not reach statistical significance (Figure 4F). The weights of liver and iBAT were not altered in AP-ERa-KO females (Supp. Figure 5A–B). These data indicate that deleting ERa using *PdgfRa-cre* promotes obesity in female mice by affecting both adipocyte hyperplasia and hypertrophy.

As male AP-ERα-KO mice exhibited increased AP proliferation in SWAT, we determined if this led to long-term obesity with increased SWAT expansion. We fed AP-ERα-KO males SD or HFD for 8 weeks and found a significant increase in body weight on HFD, but not SD (Figure 5A). Similar to female AP-ERα-KO, this is the result of an increase in both fat mass and lean mass accumulation (Figure 5B). In line with increased SWAT AP proliferation, AP-ERα-KO males accumulate more SWAT on a HFD (Figure 5C). When we measured hyperplasia using the BrdU pulse-chase assay (Figure 5D), we found male AP-ERα-KO trend towards more adipogenesis in VWAT (Figure 5E) and significant increase in obesogenic hyperplasia in SWAT (Figure 5F). This is in contrast to control males where there is no hyperplasia in SWAT (Figure 5F), as has been previously reported (Jeffery et al. 2015). Furthermore, the size of SWAT adipocytes on a HFD is not changed in AP-ERα-KO males, suggesting the increase in SWAT mass is driven by adipocyte hyperplasia specifically (Supp. Figure 6A).

While there are overt phenotypes in the SWAT of male AP-ERα-KO mice, changes in VWAT accumulation were more subtle. These mice maintained the obesogenic VWAT expansion and adipocyte hyperplasia observed in control mice fed a HFD (Figure 5C, 5E). On a SD, AP-ERα-KO males have increased VWAT mass compared to controls (Figure 5C). Interestingly, this is due to larger adipocytes and not an increase in adipocyte

hyperplasia (Figure 5E, Supp. Figure 6A). In contrast, no significant changes in mass, size, or hyperplasia were seen in SWAT of AP-ERα-KO males on SD (Figure 5C, 5E–F, Supp. Figure 6A). Of note, no changes in iBAT mass were observed (Supp. Figure 6B), but AP-ERα-KO males had larger livers on a HFD compared to controls (Supp. Figure 6C). Together, these data demonstrate that AP-ERα-KO males on a HFD have a more feminized pattern of obesity, displaying increased AP proliferation and hyperplasia in SWAT.

#### AP-ERa-KO mice have altered sex hormone levels

Many mouse models of hormone receptor knockouts or inhibition via drug treatments result in excessive production of sex hormones, with several ERa knockout models having increased estrogen in plasma (Curtis Hewitt, et al. 2000; Gustafsson, et al. 2016; Kauffman et al. 2015; Xu, et al. 2011). Thus, we measured 17-β estradiol in plasma of AP-ERα-KO mice and found significantly higher levels in female KOs compared to controls (Figure 6A). In line with this finding, the expression of enzymes that produce and metabolize estrogens are significantly enriched in the gonads of AP-ERa-KO female mice (Figure 6B). Interestingly, AP-ERα-KO females also have elevated testosterone and androstenedione levels (Figure 6C, Supp. Figure 7A). Although to a lesser extent than females, males are capable of producing estrogen in the testis, brain, skin, and adipose tissue (Baird, et al. 1973; Liu, et al. 2013; MacDonald, et al. 1979; Ohlsson, et al. 2017; Schneider, et al. 1979). However, when we measured 17-β estradiol in male plasma, all but one sample fell below the detection limit (Supp Figure 7B), however, we do find increased testosterone and androstenedione in the plasma of AP-ERa-KO males (Figure 6C, Supp. Figure 7A). These data suggest that AP-ERa-KO mice have compensatory mechanisms that lead to elevated levels of multiple sex hormones, thus making it a difficult model to assess the cellular mechanism of ERα in APs as both androgens and estrogens have been implicated to play a role in hyperplasia and fat mass regulation (Jeffery et al. 2016; Sebo and Rodeheffer 2021).

Although *PdgfRa-cre* effectively targets the adipocyte lineage in adipose tissue (Berry and Rodeheffer 2013), its expression is not restricted to APs or WAT (Roesch, et al. 2008). Thus, we assessed other sites where ERα expression may be affected in the AP-ERα-KO mice using the dual fluorescent reporter mTmG (Muzumdar, et al. 2007). Upon expression of cre recombinase, cells switch from expressing plasma membrane-targeted Tomato (red fluorescence) to expressing plasma membrane-targeted GFP. Therefore, GFP fluorescence can be used to trace cre recombinase activity. Next, we extensively characterized WAT as well as other tissues from male and female AP-ERα-KO mice using whole-mount confocal microscopy as done previously (Berry and Rodeheffer 2013; Jeffery, et al. 2014; Sebo, et al. 2018). As anticipated, we found GFP expression from VWAT, SWAT, and iBAT but not in liver or muscle (Supp. Figure 8A). When other tissues were imaged, we found GFP expression in the brain and gonadal tissues of both male and female mice (Figure 6D). While it has been published that a population of glial cells express PdgfRα (Roesch et al. 2008), we did not anticipate to see GFP expression in the gonads of males and females.

Due to the important role of estrogens in the female reproductive system, we looked further into the physiological effects of ERa loss via *PdgfRa-cre* on the gonads. Gross morphology of the reproductive tract showed atrophied uterus and hemorrhagic ovaries, consistent with

previous models of ER $\alpha$  deletion (Figure 6D, Supp. Figure 9A–B) (Antonson, et al. 2014; Lubahn, et al. 1993). Next, we confirmed reduced ER $\alpha$  transcripts by qPCR in the female gonads (Supp. Figure 9C). Of note, we found no changes in ER $\beta$  transcripts. Based on these results, we assessed if there were disturbances in the estrus cycle of AP-ER $\alpha$ -KO females. Interestingly, AP-ER $\alpha$ -KO females do not cycle and are infertile (Supp. Figure 9D–E) similar to whole-body ER $\alpha$ -KOs (Heine et al. 2000).

#### Increased SWAT hyperplasia in AP-ERa-KO males is due to AP-extrinsic mechanisms

Our lab has previously shown that microenvironment changes in adipose tissue influence the response of APs to HFD (Jeffery et al. 2016). Adipose tissue is a major source of estrogen production in humans (Hetemäki, et al. 2017; Nelson and Bulun 2001). There is some evidence that mice adipose can produce estrogen (Chow, et al. 2009; Polari, et al. 2015; Zhao, et al. 2009), although some groups report undetectable levels in mice WAT (Kim, et al. 2021). In addition, treating male mice with 17-β estradiol induces SWAT proliferation (Jeffery et al. 2016). To determine if the increased SWAT proliferation in AP-ERa-KO males is due to tissue microenvironment changes, we attempted to quantify sex hormones within the adipose tissue of these mice via several methods, including mass spectrometry, but our efforts were unsuccessful (data not shown). As an alternative approach, we employed an AP transplant assay to test if the proliferation of these ERα-KO cells was sustained in a wildtype tissue microenvironment. Briefly, we sorted GFP+ AP cells from AP-ERα-KO males and injected them into the SWAT of wildtype C57BL/6J males (Figure 6E). After 2 weeks of recovery, recipient mice were fed HFD and treated with BrdU for one week. AP proliferation was measured for endogenous cells (GFP-) and transplanted ERα-KO cells (GFP+) via flow cytometry. In contrast to the AP-ERa-KO mice, ERa-KO cells do not proliferate in the SWAT of wildtype male mice (Figure 6F). These data indicate that the increased AP proliferation, hyperplasia, and accumulation of SWAT in AP-ERα-KO male mice is driven by AP-extrinsic changes in the tissue microenvironment and not a lack of ERa function in APs.

#### **Discussion**

The expansion and distribution of adipose tissue is sexually dimorphic. Previous work in the lab has shown that sex hormones play a role in this process by influencing hyperplasia (Jeffery et al. 2016; Sebo and Rodeheffer 2021). Our attempt to elucidate the role of estrogen signaling in APs led to an important lesson in the use of hormone receptor knockout models. *PdgfRa-cre* is known be expressed in several non-adipose tissues. Here we find that it also marks female and male reproductive organs (Figure 6D, Supp. Figure 9C). This deletion of ERα in one or several of the target tissues leads to compensatory mechanisms that elevate sex hormones in the plasma of AP-ERα-KO mice, including estradiol in females (Figure 6A–C, Supp. Figure 7A–B), making this model unsuitable to study ERα signaling in APs. Despite having increased estrogen, an obesity-protecting hormone, AP-ERα-KO mice are more obese. This finding replicates many other ERα-KO models using cre promoters that target the brain (Xu et al. 2011), and whole-body ERα-KO (Curtis Hewitt et al. 2000; Gustafsson et al. 2016) where the mice are more obese despite the increased estrogen levels. This highlights that ERα mediates many of estrogens protective

roles against obesity. In addition, estrogen can still signal through ER $\beta$  and novel GPR30 receptors in these models. Thus, it is difficult to determine if the phenotypes we observe here are due to lack of ER $\alpha$  signaling in target cells or the result of alteration of sex hormones and their intermediates that can signal via ER $\alpha$ -independent mechanisms (Sebo and Rodeheffer 2021).

Nonetheless, our findings in AP-ERa-KO mice do support an important role of estrogens in obesity. The fact that AP-ERa-KO females are more likely to become obese than control females is of interest. Our lab has found that wildtype females have a large variation in hyperplasia (Jeffery et al. 2016) and obesity phenotypes overall (Hong, et al. 2009; Yang, et al. 2014). This indicates that estrogen is likely involved in the hyperplastic response of females to a HFD and this mechanism should be further explored. Perhaps even more important is the feminized pattern of obesity that we find in male AP-ERa-KO (Figure 3D, 5C). Although the increased expansion of SWAT in these mice occurs via hyperplasia and not hypertrophy (Figure 5F, Supp. Figure 6A), we find that lack of ERa in APs itself does not cause more proliferation (Figure 6E-F). This suggest that the hormonal milieu in SWAT of male AP-ERa-KOs provides the necessary cues for adipogenesis to occur in absence of ERa. Though our attempts to measure sex hormones within WAT were unsuccessful, the fact that wildtype males treated with estrogen have SWAT proliferation (Jeffery et al. 2016) suggests increased estrogen production in AP-ERa-KO is a potential mechanism to explain this phenotype, as free estradiol correlates to increased SWAT mass in men (Vermeulen, et al. 2002).

The AP-ERα-KO model also affects estrogen signaling in mature adipocytes, and although estrogens have overall protective effects against obesity, their effects in adipocyte biology are not well understood. This is the first study to address the role of adipocyte ERα in diet-induced obesity. Our data suggest that the extensive adipose tissue accumulation seen in whole body ERα-KO are not replicated in adipocyte-specific ERα-KO. Male Adi-ERα-KO mice do not show differences in fat mass accumulation (Figure 1F, Supp. Figure 3A, 3C). However, loss of adipocyte ERα in females affects VWAT expansion via a modest reduction in adipocyte size (Figure 2A, 2F) without affecting hyperplasia (Figure 2D). It is unclear if this is due to energy expenditure or food intake changes in Adi-ERα-KO females. Importantly, estrogen levels in the plasma of male and female Adi-ERα-KO mice are not different from controls. Thus, the absence of phenotypes in these mice is not explained by changes in systemic estrogen levels.

One way estrogen is believed to modulate obesity is by affecting adipocyte lipolysis and lipogenesis (Gormsen et al. 2012; Pedersen, et al. 2004). Our data demonstrates that this potential role could be mediated by ERa in female mice as some subtle differences in VWAT adipocyte sizes were present (Figure 2F). Of note, our findings in the Adi-ERa-KO mice stand in contrast to previous findings (Davis, et al. 2013). This could be due to the use of different *Adiponectin-cre* strains and their respective efficiencies as the strain used in this study has demonstrated higher recombination levels in adipocytes (Eguchi et al. 2011; Wang, et al. 2010). Overall, these data suggest that signaling of ERa in adipocytes is not a driver of the protective effects of estrogen in obesity and that there is potential

VWAT-specific role of ERa in regulating adipocyte size in females. We do not know how these findings will translate to postmenopausal adipocyte-ERa signaling or obesity in aging.

Estrogens protective effects extend to other metabolic organs like the liver, as it has been shown to improve liver function and prevent hepatic steatosis. (Chow, et al. 2011; Shen and Shi 2015; Zhu, et al. 2013) The major estrogen receptor in the liver, ERa, has been shown to have oscillating activity in the liver correlating to the estrus cycle in females. (Villa, et al. 2012) This pulsatile activity during proestrus is sufficient to drive changes in gene expression to prevent lipid accumulation. This is mediated in part by ERa as liver-specific knockouts do not show these gene expression changes and both male and female liver-ERa-KOs have fatty livers. (Villa et al. 2012) In addition, OVX female mice are more prone to develop hepatic steatosis compared to sham females, but treatment with estrogen rescues this phenotype (Zhu et al. 2013) and the hepatic steatosis seen in Aromatase-KO males can be rescued by treatment with an ERa agonist. (Chow et al. 2011) Our data shows that male but not female AP-ERa-KOs had substantially larger livers on a HFD compared to controls (Supp. Figure 6C). This is not due targeting of ERa in the livers as PdgfRa-cre is does not label cells in the liver (Supp. Figure 8A). Interestingly, male Adi-ERa-KOs showed a trend to larger livers on a SD (Supp. Figure 3D). These data points at a potential role for estrogen in mediating organ crosstalk to promote lipid accumulation in the liver. Other possibilities are that this is driven by the distinct hormonal milieu present in AP-ERa-KO males or a product of the increased adipose tissue mass of HFD-fed AP-ERα-KO males.

Overall, these data provide important insights in the role of estrogens in diet-induced obesity. As obesity rates and its associated comorbidities continue to rise (Finkelstein, et al. 2012; Ward, et al. 2019) it is crucial to understand the mechanisms underlying this disease. Although it is evident that sex hormones are key players in the expansion of adipose, we must be cautious of conclusions derived from hormone receptor knockout mouse models as these have consistently shown to have compensatory mechanisms that elevate estrogen and other sex hormones (Curtis Hewitt et al. 2000; Gustafsson et al. 2016; Kauffman et al. 2015; Xu et al. 2011). It is important to note that the hyperplastic phenotypes seen in AP-ERα-KO mice are exclusively present in HFD conditions. Our findings suggest the hormonal milieu in AP-ERα-KO mice is not sufficient to drive hyperplastic obesity, and that a HFD is needed for this process. Identifying these nutrient signals and how estrogen and other hormones influence expansion of adipose is crucial to our understanding of the regulation of fat mass expansion in obesity.

### **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

### **Funding**

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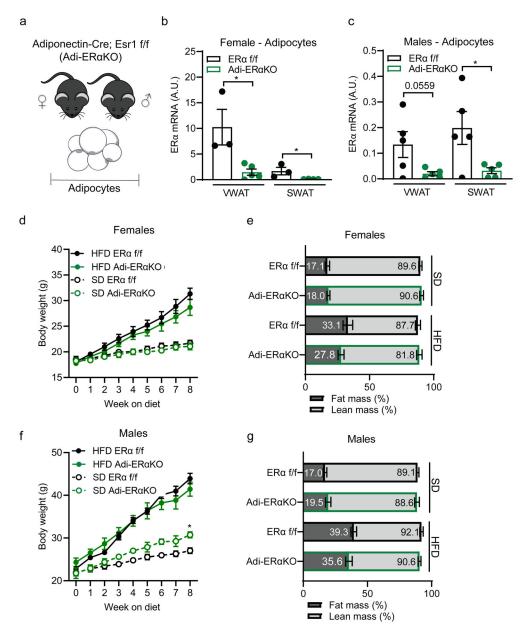


Figure 1. Adipocyte ERa deletion does not impact body composition.

- (A) Experimental mouse model to delete ERa in mature adipocytes using the *Adiponectin-cre*, *Esr1 fl/fl;mTmG* strain.
- **(B)** ERα expression in isolated mature adipocytes from Adi-ERα-KO females and controls. (n=3–5 mice per group)
- (C) ER $\alpha$  expression in isolated mature adipocytes from Adi-ER $\alpha$ -KO males and controls. (n=5 mice per group)
- (**D**) Body weight of female Adi-ER $\alpha$ -KO and controls during 8 weeks of SD or HFD. (n= 6–7 mice per group)
- (E) Body composition of female Adi-ER $\alpha$ -KO and controls after 8 weeks of SD and HFD. (n=6–7 mice per group)

(F) Body weight of male Adi-ER $\alpha$ -KO and controls during 8 weeks of SD and HFD. Comparison shown is between SD groups. (n= 5–6 mice per group)

(G) Body composition of male Adi-ER $\alpha$ -KO and controls after 8 weeks of SD and HFD. (n=5–6 mice per group)

Statistical significance determined by unpaired t-tests in panel B-C and two-way ANOVA with Tukey's test for panels C and E. \*p<0.05. Error bars represent mean  $\pm$  S.E.M. SD: standard diet, HFD: high-fat diet, VWAT: visceral fat, SWAT: subcutaneous fat. See also Figure S1.

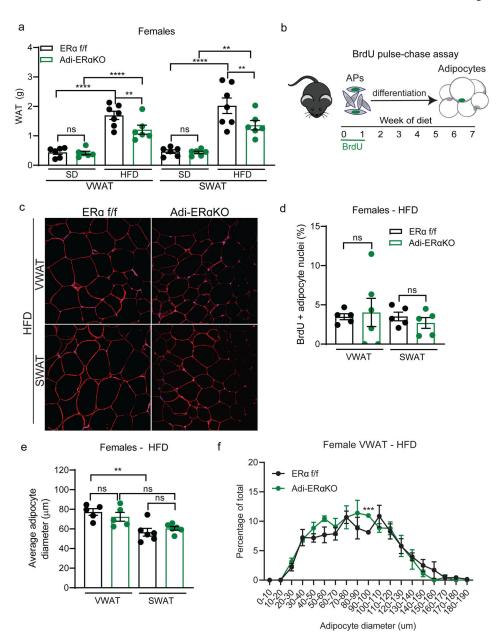
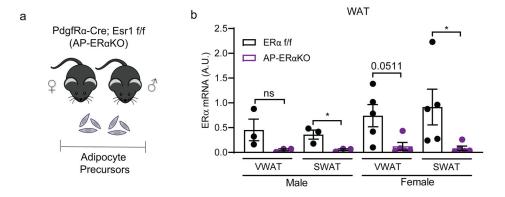


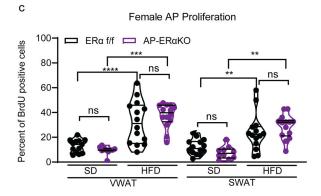
Figure 2. Adipocyte ER $\alpha$  deletion modestly impacts VWAT accumulation in HFD-fed female mice.

- (A) VWAT and SWAT accumulation of Adi-ER $\alpha$ -KO and control females after 8 weeks of diet. (n=6–7 mice per group)
- (B) Experimental design for BrdU pulse-chase to measure adipocyte formation.
- (C) Representative confocal microscopy images of WAT from female Adi-ERa-KO and controls after 8 weeks of HFD. Tissue stained for caveolin (red), BrdU (green), and DAPI (blue) taken at 40X.
- (**D**) Quantification of BrdU in adipocyte nuclei of female Adi-ER $\alpha$ -KO mice and controls after 8 weeks of HFD. (n=5-6 mice per group)
- (E) Average adipocyte diameter ( $\mu m$ ) of female Adi-ER $\alpha$ -KO and controls after 8 weeks of HFD. (n=5–6 per group)

(F) Distribution of adipocyte diameters in VWAT of female Adi-ER $\alpha$ -KO and controls after 8 weeks of HFD. (n=5–6 per group)

Statistical significance determined by two-way ANOVA with Tukey's test for panels A, D, and E. Statistical significance determined by multiple unpaired t-tests for panel F. Error bars represent mean  $\pm$  S.E.M. \*\*p<0.01, \*\*\*p<0.001, \*\*\*\*p<0.0001. SD: standard diet, HFD: high-fat diet, VWAT: visceral fat, SWAT: subcutaneous fat, AP: adipocyte precursors. See also Figure S2–S3.





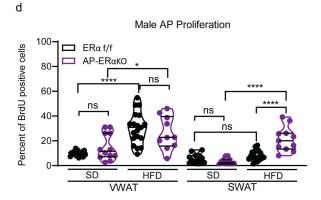


Figure 3: Lack of ERa in adipocyte precursors (APs) changes obesogenic proliferation patterns. (A) Experimental mouse model to delete ERa in APs using *PdgfRa-cre*; *Esr1 fl/fl*; *mTmG* strain

- (B) ERa expression in whole white adipose tissue (WAT) from AP-ERa-KOs and controls. (n=3–5 mice per group)
- (C) AP proliferation of female AP-ER $\alpha$ -KOs and controls after one week of SD or HFD. (n=10–14 mice per group)
- (**D**) AP proliferation of male AP-ER $\alpha$ -KOs and controls after one week of SD or HFD. (n=10–19 mice per group)

Statistical significance determined by unpaired t-tests for panel B. Statistical significance determined by two-way ANOVA with Tukey's test for panels C-D. In dot plots, error bars represent mean  $\pm$  S.E.M. In violin plots, mean is showed as filled line and quartiles as dotted

lines. \*p<0.05, \*\*p<0.01, \*\*\*p<0.001, \*\*\*\*p<0.0001. SD: standard diet, HFD: high-fat diet, VWAT: visceral fat, SWAT: subcutaneous fat, AP: adipocyte precursors. See also Figure S4.

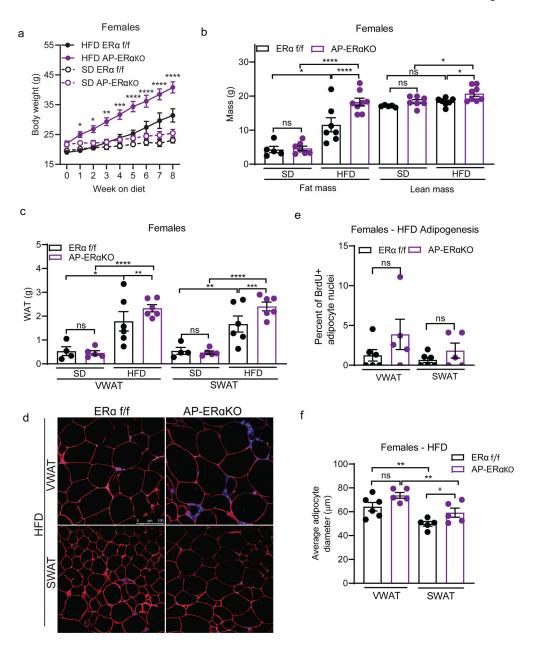


Figure 4: AP-ERa-KO female mice are more susceptible to diet-induced obesity.

- (A) Body weight of female AP-ERα-KOs and controls during 8 weeks of SD and HFD. Significance shown between HFD groups. (n=5–8 mice per group)
- (B) Total fat and lean mass of female AP-ER $\alpha$ -KOs and controls after 8 weeks of SD and HFD. (n=5–8 mice per group)
- (C) VWAT and SWAT accumulation of female AP-ER $\alpha$ -KOs and controls after 8 weeks of SD and HFD. (n=5–8 mice per group)
- (**D**) Representative confocal microscopy images of WAT from female AP-ERa-KO and controls after 8 weeks of HFD. Tissue stained for caveolin (red), BrdU (green), and DAPI (blue) taken at 40X. Scale bar is 100µm.

(E) Quantification of BrdU in adipocyte nuclei of female AP-ERα-KO mice and controls after 8 weeks of HFD. (n=5–6 mice per group)

(**F**) Average adipocyte diameter ( $\mu m$ ) of female AP-ER $\alpha$ -KO and controls after 8 weeks of HFD. (n=5–6 per group)

Statistical significance determined by two-way ANOVA with Tukey's test. Error bars represent mean  $\pm$  S.E.M. \*p<0.05, \*\*p<0.01, \*\*\*p<0.001, \*\*\*\*p<0.0001. SD: standard diet, HFD: high-fat diet, VWAT: visceral fat, SWAT: subcutaneous fat. See also Figure S5.

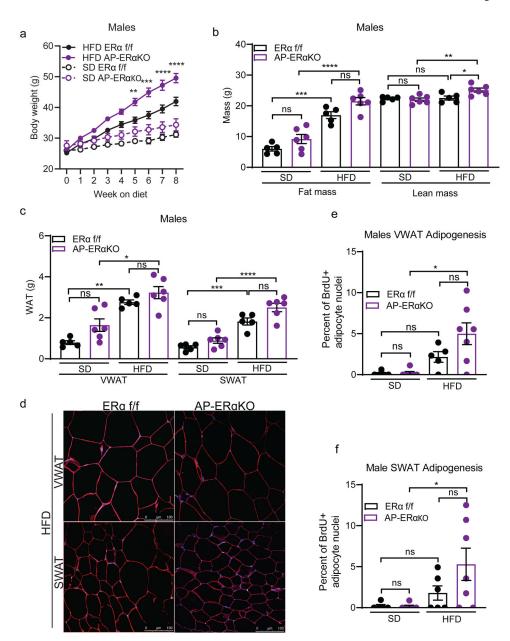


Figure 5: AP-ERa-KO male mice have diet-induced SWAT hyperplasia.

- (A) Body weight of male AP-ERa-KOs and controls during 8 weeks of SD and HFD. Significance shown between HFD groups. (n=5-6 mice per group)
- (B) Total fat and lean mass of male AP-ER $\alpha$ -KOs and controls after 8 weeks of SD and HFD. (n=5-6 mice per group)
- (C) VWAT and SWAT accumulation of male AP-ER $\alpha$ -KOs and controls after 8 weeks of SD and HFD. (n=5–6 mice per group)
- (**D**) Representative confocal microscopy images of WAT from male AP-ERa-KOs and controls after 8 weeks of HFD. Tissue stained for caveolin (red), BrdU (green), and DAPI (blue) taken at 40X. Scale bar is 100µm.

(E) Quantification of BrdU in adipocyte nuclei in VWAT of male AP-ERα-KOs and controls after 8 weeks of SD and HFD. (n=5–6 mice per group)

(F) Quantification of BrdU in adipocyte nuclei in SWAT of male AP-ER $\alpha$ -KOs and controls after 8 weeks of SD and HFD. (n=5-6 mice per group)

Statistical significance determined by two-way ANOVA with Tukey's test. Error bars represent mean  $\pm$  S.E.M. \*p<0.05, \*\*p<0.01, \*\*\*p<0.001, \*\*\*\*p<0.0001. SD: standard diet, HFD: high-fat diet, VWAT: visceral fat, SWAT: subcutaneous fat. See also Figure S6.

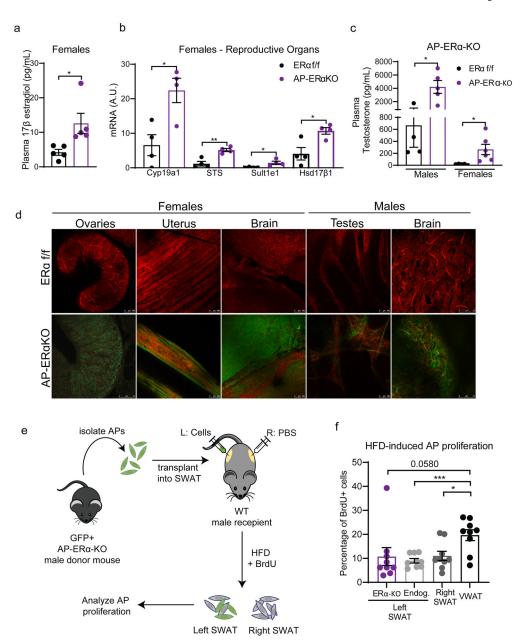


Figure 6: Off-target effects in AP-ERa-KO mice drive obesogenic hyperplasia.

- (A) Plasma levels of 17- $\beta$  estradiol in female AP-ER $\alpha$ -KOs and controls. (n=5 mice per group)
- **(B)** Expression of estrogen metabolizing genes in the reproductive organs of AP-ERα-KO and control females. (n=4 mice per group)
- (C) Plasma levels of testosterone in male and female AP-ERα-KOs and controls. (n=4–6 per group)
- **(D)** Whole-mount confocal images of tissues from *AP-ERa-KO*; *mTmG* mice taken at 20X. GFP fluorescence indicates presence of cre activity. Scale bar is 100μm. (n=1 mice per group)

(E) Experimental design of AP transplant experiment. AP cells lacking ER $\alpha$  were injected into the SWAT of wildtype C57BL/6J male mice. AP proliferation was measured after one week of HFD.

(F) One week AP proliferation of transplanted ER $\alpha$ -KO cells, endogenous cells from left SWAT, right SWAT (PBS injection), and VWAT of recipient males. (n=9 mice per group) Statistical significance determined by unpaired t-tests. Error bars represent mean  $\pm$  S.E.M. \* p<0.05, \*\*p<0.01, \*\*\*p<0.001. HFD: high-fat diet, VWAT: visceral fat, SWAT: subcutaneous fat. See also Figure S7–9.

Table 1.

## Primer sequences for gene expression analysis

Gene	Primer sequences
Esr1	5'-CCCGCCTTCTACAGGTCTAAT-3' 5'-CACACGGCACAGTAGCGAG-3'
Esr2	5'-ACACCTTGCCTGTAAACAGAGA-3' 5'-GCAGAAGTGAGCATCCCTCTT-3'
Сур19а1	5'-CCATGCCACTCCTGCTGAT-3' 5'-TTCCCAGACAGTAGCCAGGAC-3'
Sult1e1	5'-GAAGGTGATGTGGAAAAATGC-3' 5'-TTCTGGGAGATTCCTTCTCTTTT-3'
Sts	5'-CTCGCTGACATCATCCTCGC-3' 5'-GCGTTTTGTTGCCGTAGCAG-3'
Hsd17b1	5'-ACTTGGCTGTTCGCCTAGC-3' 5'-GAGGGCATCCTTGAGTCCTG-3'
Adipoq	5'-GGAGATGCAGGTCTTCTTGG-3' 5'-GCGATACACATAAGCGGCTTC-3'
PdgfRa	5'-AACGGAGGAGCTGCGGGGAA-3' 5'-CCCATAGCTCCTGAGACCTTCTCCT-3'
Tbk1	5'-AGGGCTTTGTGACGGGAACAG-3' 5'-GGCACCCGGTCAAATGAGA-3'
Actb	5'-GGCTGTATTCCCCTCCATCG-3' 5'-CCAGTTGGTAACAATGCCATGT-3'