

The Role of Extra-Ovarian Estrogen in Regulating Metabolism in Female Non-Human  
Primates

By

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**ABSTRACT**

Estradiol supports female metabolic function in multiple animal research models, including, mice, rats, and sheep. Estradiol action is mediated through estrogen receptors, predominately, estrogen receptor alpha (ER $\alpha$ ) and estrogen receptor beta (ER $\beta$ ). ER $\alpha$  has been shown extensively to be the predominant estrogen receptor regulating estradiol action in rodents by way of ovarian synthesized estradiol. While ovariectomy (OVX) in rodents enables obesity, OVX in nonhuman primates inconsistently alters weight gain, highlighting the possibility of estradiol produced outside the ovaries as an important site of estradiol synthesis signaling through ER $\alpha$  in nonhuman primates. We hypothesized that extra-ovarian estradiol provides key support through hypothalamic ER $\alpha$  for female metabolic function. We employed aromatase inhibition to eliminate extra-ovarian estradiol biosynthesis and elucidate the role of extra-ovarian estradiol in regulating energy balance in OVX adult female marmosets and rhesus macaques. Next, we employed RNAi technology to silence ER $\alpha$  in the mediobasal hypothalamus (MBH) of adult female OVX marmosets and ovary-intact rhesus macaques to assess the role of ER $\alpha$  in energy balance. Our findings provided the first evidence for estradiol's action through ER $\alpha$  contributing to energy homeostasis in a female nonhuman primate, and prompted speculation that extraovarian estradiol, and potentially, neuro-synthesized estrogens, may similarly regulate energy balance in other female primates, including humans.

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## **1. CHAPTER ONE: Literature Review: Estradiol Regulation of Metabolism**

### **1.1 Implications for Metabolic Dysfunction in Post-Menopausal Individuals**

Menopause, defined by menstruation cessation and a depletion of ovarian sex steroid hormones, increases the risk of metabolic dysfunction associated with reduced steroid actions in female adipose tissue, bone, cardiovascular tissues, pancreas, liver, and brain. Studies have shown, for example, an increased risk of metabolic syndrome by 60% in postmenopausal women<sup>1,2</sup>. Multiple investigations have attempted to dissociate effects of aging versus declining estradiol levels on adiposity, energy balance, cardiometabolic and brain health in menopausal women<sup>3,4,5</sup>. In general, these studies suggest that menopause per se is associated with increasing abdominal obesity and accompanying co-morbidities<sup>6</sup>. While findings on menopause-associated obesity parallel observations made in estrogen-depleted, ovariectomized (OVX) female rodents<sup>7</sup>, causal relationships between declining ovarian estradiol during menopause, and altered body composition and energy balance, have been difficult to confirm in women. Current clinical recommendations place hormonal replacement therapy (HRT), commonly a form of estrogen combined with a synthetic progestin, as the gold standard for treating symptoms associated with menopause<sup>8</sup>. There is conflicting evidence on the effectiveness of HRT in mitigating the adverse metabolic symptoms associated with the menopausal transition. Studies have demonstrated that both oral and transdermal estradiol therapy in postmenopausal women are associated with a reduction in central adiposity and an increase in lean body mass<sup>9,10</sup>, as well as reductions in insulin resistance and fasting glucose. More studies focused on glucoregulation have suggested estradiol-mediated increases in insulin sensitivity<sup>11</sup>, while others find no

improvement<sup>12,13</sup>. Studies on energy expenditure after commencing HRT have demonstrated increases and decreases in lipid oxidation and energy expenditure, respectively<sup>14</sup>. Selective estrogen receptor modulators (SERMs), compounds that interact with estrogen receptors in specific target tissues, have shown promise in improving osteoporosis and treatment of breast cancer<sup>15</sup>. Although SERMs can selectively engage target tissues, like systemically acting HRT, SERM use can cause adverse side effects, such as, increased risk for thromboembolic events<sup>16,17</sup>. The specific metabolic consequences of menopause and/or estrogen withdrawal in women remain unclear, due in part to the uncontrolled environmental, psychosocial, and genetic factors that predominate in most clinical studies.

## **1.2 Estradiol Production**

The major bioactive estrogen, estradiol, is a steroid hormone synthesized throughout the body by aromatase, a cytochrome P450 enzyme, encoded by the CYP19A1 gene<sup>18-21</sup>. While the main source of estradiol (E<sub>2</sub>) is the ovary, E<sub>2</sub> is also produced in low amounts in extra-ovarian sites including, adipose tissue, adrenal glands, skin, bone, pituitary gland, and multiple brain areas<sup>18-21</sup>.

## **1.3 Mechanisms of Estradiol Signaling**

E<sub>2</sub> can act through multiple receptors and mechanisms involving various signaling cascades to effect physiological action. E<sub>2</sub> crosses the plasma membrane and interacts with intracellular ER $\alpha$  and ER $\beta$  to exert direct genomic effects when binding to cognate DNA response element sequences, i.e., estrogen response elements (EREs) in the regulatory regions of target genes. E<sub>2</sub> can also exert genomic actions by complexing with other transcription factors that, in turn, bind to their own cognate response

elements. Alternatively, E<sub>2</sub> can exert non-genomic actions via interaction with the G-protein coupled estrogen receptor (GPER1) and/or ER $\alpha$  and ER $\beta$  complexed with anchoring proteins at the plasma membrane, thereby activating intracellular signaling cascades.

The classical mechanism of estrogen signaling is direct genomic signaling where ER $\alpha$  and ER $\beta$  nuclear receptors are ligand-activated transcription factors<sup>22,23</sup>. In this mechanism, nuclear and/or cytoplasmic bound ER $\alpha$  and ER $\beta$  undergo a conformational change inducing release from molecular chaperones and receptor dimerization<sup>24</sup>. After dimerization, the complex binds to genomic targets; chromatin at estrogen response element (ERE) sequences, enhancer regions by or within promoter regions and regions of target genes<sup>25</sup>. Research has identified 35% of genes acted on by E<sub>2</sub> do not contain ERE-like sequences<sup>26,27</sup>. In the absence of EREs, estrogen signaling can affect gene expression without binding directly to DNA, an “indirect genomic” action. In such mode of action, estrogen receptor complexes participate in protein-protein interactions with other transcriptional elements and response elements, like that of stimulating protein-1 (Sp-1), a crucial mediator of indirect genomic action that is enhanced by estrogen receptors<sup>28,29</sup>.

In contrast to the classic E<sub>2</sub> action of changing gene expression of target genes, non-genomic E<sub>2</sub> signaling consists of rapid intracellular changes via membrane associated receptors. Non-genomic estrogen signaling involves activation of signal-transduction mechanisms with the subsequent production of intracellular second messengers, such as cAMP and PKA, or regulation and protein-kinase signaling cascades. The activation of these membrane-initiated signaling cascades can result in

non-genomic changes in a host of cellular activities, as well as ultimately convey signals to the nucleus to evoke transcriptional responses<sup>30</sup>. GPER1 (G protein-coupled estrogen receptor 1), binds to estrogens, promoting estrogen-dependent activation of adenylyl cyclase and epidermal growth factor receptor (EGFR).

## **1.4 Steroid Hormones, Female Energy Homeostasis**

### *1.4.1 Female Rodent Models: Ovarian Hormones and Metabolism*

In many experimental animal models, ovarian E<sub>2</sub> regulates body weight and composition, energy balance, and glucoregulation<sup>31</sup>. In rodent models, ovarian E<sub>2</sub>-regulated control mechanisms are well established since OVX-mediated E<sub>2</sub> depletion reliably disrupts energy balance, and leads to increased body weight and visceral adiposity, reduced physical activity and energy expenditure<sup>32,33</sup>, and diminished insulin sensitivity. All these effects of metabolic dysfunction can be diminished or reversed by E<sub>2</sub> replacement<sup>34,35</sup>. Further, these E<sub>2</sub> actions are mostly mediated by estrogen receptor alpha (ER $\alpha$ ), as complete ER $\alpha$  knockout mice (ER $\alpha$ KO) exhibit increases in body weight and adiposity, and present energy metabolism phenotypes that largely mimic those observed in long-term OVX mice<sup>36</sup>. A similar marked degree of metabolic dysfunction is not found in ER $\beta$ KO or GPR30KO female mice<sup>37</sup>. Due to the vast research on the metabolic impact of ovarian E<sub>2</sub> via ER $\alpha$  in rodents, multiple avenues of E<sub>2</sub> regulated energy balance have been established. These include the direct actions of E<sub>2</sub> on metabolically active tissues such as adipose depots, liver, muscle, and pancreas<sup>38</sup>, E<sub>2</sub> activation of brown adipose tissue (BAT) via the sympathetic nervous system<sup>39</sup>, as well as E<sub>2</sub>-induced alterations in neural and peripheral control of food intake and energy expenditure.

#### 1.4.2. *E<sub>2</sub>-mediated Neural Regulation of Metabolic Function*

Rodent models of ovarian estradiol regulation have also highlighted the critical role of neural ER $\alpha$  signaling in mediobasal hypothalamic nuclei, the ventromedial nucleus (VMN) and the arcuate nucleus (ARC), to maintain energy homeostasis via the leptin-melanocortin system<sup>40,41</sup>. The leptin-melanocortin system is the predominant pathway regulating energy balance in mammals<sup>42</sup>. The central melanocortin system includes two distinct populations of neurons in the ARC—the proopiomelanocortin (POMC) and cocaine- and amphetamine-related transcript prepropeptide (CARTPT) expressing neurons and the agouti-related peptide (AgRP) and neuropeptide Y (NPY) expressing neurons<sup>43</sup>. Both populations of neurons have projections into the paraventricular nucleus (PVN) of the hypothalamus and facilitate downstream mechanisms to modulate energy balance. Leptin, an adipokine secreted by adipose tissue, conveys messages to the brain regarding the body's energy stores by binding to its receptors on the orexigenic neuronal populations, AgRP and NPY and the anorexigenic neuronal populations, POMC and CART<sup>44</sup>. Upon binding to POMC neurons, a cleavage product of the POMC transcript, alpha-MSH (alpha melanocyte stimulating hormone) is released and subsequently, activates MC4R, melanocortin 4 receptor, in the PVN (Figure 1). This reaction results in a satiety signal and reduces food intake. In opposition, AgRP competes with alpha-MSH to bind MC4R to increase food intake. Neuropeptide Y exerts its orexigenic effects via activation of NPY Y1 and Y5 receptors.

In addition to satiety signals, leptin activation of AgRP and POMC neurons through brain-derived neurotrophic factor (BDNF)-expressing neurons in the PVN, influences sympathetic innervation of white adipose tissue (WAT) and brown adipose tissue

(BAT)<sup>45</sup>. Leptin's ability to modify the sympathetic architecture of WAT and BAT and promote adipose thermogenesis when energy stores are in excess is an important downstream neuronal signaling mechanism for energy balance. Adipose depots, mainly BAT, are activated by sympathetic signals, such as, norepinephrine (NE) and epinephrine (E)- most notably, after cold exposure. NE and E signal through three G-protein coupled beta-adrenergic receptors located on brown adipocytes, termed 1 (ADRB1), 2 (ADRB2) and 3 (ADRB3). Upon stimulation by NE, in classical cold-induced thermogenesis in female rodents, beta-adrenergic receptors signal synthesis of cyclic AMP (cAMP), promoting cAMP driven protein kinase A (PKA) activation, stimulating lipolysis<sup>46</sup> (Figure 2). Lipolysis results in free fatty acid and glycerol release from the triglyceride lipid droplet and enhanced expression of uncoupling protein 1 (UCP1) mRNA. Increased UCP1 takes the energy stored in the mitochondrial electrochemical gradient and dissipates it as heat, independent of ATP synthesis, termed non-shivering thermogenesis.

E<sub>2</sub> modulates these neural MBH metabolic circuits in many ways; its actions can be dependent on the leptin-melanocortin system or act independently of leptin's actions altogether<sup>47</sup>. Like leptin, E<sub>2</sub> is a strong anorexigenic signal regulating energy balance by stimulating POMC and CART neurons and inhibiting NPY and AgRP neurons. It had been assumed that E<sub>2</sub>'s effect on energy balance acts in tandem with leptin's role via co-expression of leptin receptor (LepRb) and ER $\alpha$  in the mediobasal hypothalamus<sup>48</sup>, but recent research in rodents has shown co-expression is more pronounced in the preoptic area (POA) and is limited in the mediobasal hypothalamus. Quantification of LepRb and ER $\alpha$  colocalization throughout the murine hypothalamus has shown strong

colocalization absent in all areas aside from the POA and one month after OVX, co-expression tended to decline<sup>49</sup>. Further, STAT3 or ER $\alpha$  deletion from LepRb cells does not inhibit E<sub>2</sub>'s ability to manage body weight<sup>50</sup>. It has been shown that circulating levels of E<sub>2</sub> may enhance the anti-obesity effects of leptin and specifically modulate leptin sensitivity in certain metabolic tissues like adipose and skeletal muscle<sup>51</sup>. Taken together, E<sub>2</sub> mediation of energy homeostasis via ER $\alpha$  is not dependent on activation of LepRb on co-expressing cells and circulating levels of E<sub>2</sub> may act indirectly on the leptin-melanocortin pathway.

In addition to estrogenic effects on the leptin-melanocortin pathway of energy balance, E<sub>2</sub> and hypothalamic ER $\alpha$  have been well characterized as modulators of metabolism. It has been demonstrated that viral vector mediated *ESR1* gene silencing in the VMN of both female mice and rats largely recapitulates a metabolic phenotype observed in complete *ESR1*KO mice, including obesity, hyperphagia, impaired glucose tolerance and reduced energy expenditure<sup>52,53</sup> (Figure 2). It has also been demonstrated that ER $\alpha$  gene deletion in specific neuronal populations of the mediobasal hypothalamus have differing impacts on metabolism. Deletion of ER $\alpha$  in hypothalamic steroidogenic factor-1 (SF1) expressing neurons results in abdominal obesity and hypometabolism, not hyperphagia, while ER $\alpha$  deletion in POMC neurons leads to hyperphagia<sup>53</sup>.

Downstream of SF1 and POMC action, E<sub>2</sub> acts on more targets for metabolic regulation. Recent research has shown MC4R is a transcriptional target of ER $\alpha$  action<sup>54</sup>. Identification of a subset of E<sub>2</sub>-sensitive neurons in the ventrolateral ventromedial hypothalamic nucleus (VMHvl) with projections to the hippocampus and hindbrain

enables E<sub>2</sub> to modulate energy allocation. Upon E<sub>2</sub> influx, VMHvl neuronal MC4R signaling increases via recruitment of ER $\alpha$  to *MC4R*<sup>54</sup>.

The leptin-melanocortin system by nature is sensitive to changing caloric environments. Diet induced obesity (DIO) is a feeding regime commonly used in research to mimic human obesity. DIO can assert stress on energy balance mechanisms that can help elucidate signaling pathways. DIO rodents present with normal levels of leptin receptor but despite high levels of leptin, leptin fails to stimulate the leptin signaling cascade<sup>55</sup>. After switching to a reduced fat diet, leptin responsiveness of NPY/AgRP and POMC neurons could now be demonstrated, with mice regaining normal leptin sensitivity and glycemic control<sup>55</sup>, highlighting the importance of nutrient sensing in the leptin-melanocortin pathway. In the absence of obesity and diabetes, rodents fed a high fat/high calorie diet during pregnancy exhibited increased cytokine production in the hypothalamus and abnormal development of melanocortin pathways in fetal offspring<sup>56-60</sup>. Further, the fetal offspring, regardless of maternal obesity status, exhibit accelerated weight gain and adiposity paired with glucose intolerance<sup>56-60</sup>. Taken together, independent of maternal obesity, maternal diet can alter fetal development of melanocortin signaling, induce lipotoxicity and predispose female fetal offspring to adult metabolic dysfunction similar to the effects of obesity and glucose intolerance<sup>61</sup>.

DIO is critical in demonstrating the protective effects of E<sub>2</sub> against weight gain and disruption of energy homeostasis in rodents. OVX mice fed a high fat diet (HFD) and treated cyclicly with E<sub>2</sub>, displayed insulin sensitivity and no white adipose tissue (WAT) inflammation<sup>62</sup>. In contrast, OVX mice fed HFD and no E<sub>2</sub> replacement, presented with



insulin signaling dysfunction and low-grade inflammation in WAT. Importantly. Both animal groups attained similar body weights and demonstrated equivalent fat masses<sup>62</sup>. These results demonstrate that E<sub>2</sub> regulates glucose homeostasis independently of body weight and adiposity, highlighting the protective effect of E<sub>2</sub> against increased cardiovascular morbidities diabetic conversion associated with high fat/high caloric environments<sup>63</sup>.

In rodents, OVX induces rapid body weight gain and metabolic dysfunction, yet, in non-human primate models, OVX inconsistently alters body weight<sup>64-66</sup>. Additionally, while E<sub>2</sub> action via ER $\alpha$  and its regulation of energy balance is well characterized in rodent models, research is lacking in non-human primate (NHP) models. It remains crucial to identify whether similar mechanisms of E<sub>2</sub> signaling, and metabolic regulation apply to female NHPs and women.

#### *1.4.3 Female Primate Models: Ovarian Hormones and Metabolic Regulation*

Aspects of metabolism can fluctuate due to cyclic hormone secretions associated with female reproductive cycles. The degree and timing of hormonal regulation of metabolism is dependent on the type of reproductive cycle, thus eliciting species specific differences of hormonal control. Similar to rodents, some NHP species like lemurs, lorises and tarsiers, have an estrous cycle. New World Primates like marmosets (one model used in this thesis) and tamarins exhibit an ovarian cycle, while Old World Primates (i.e., rhesus macaques, model used in this thesis), great apes and humans share a menstrual cycle. It has been well established that food intake is reduced at the time of ovulation and increases during pregnancy in female animal models. Specifically, in a NHP model, decreased food intake has been observed in the peri-ovulatory phase

of the menstrual cycle in female rhesus macaques<sup>67,68</sup>. Such reduction of food intake during the peri-ovulatory period has also been observed in humans<sup>69-72</sup>. A well-supported hypothesis is the rise of E<sub>2</sub> in the circulation during this periovulatory period elicits alterations to hypothalamic signaling cascades involved in energy intake<sup>73</sup>.

As described in a previous section, OVX in rodents induces weight gain, adiposity, reduced energy expenditure and glucose intolerance. In NHP models, however, OVX-mediated E<sub>2</sub> depletion inconsistently increases body weight<sup>74-76</sup>. Despite the lack of a robust metabolic phenotype following OVX in NHPs, selective estrogen receptor modifiers (SERMs) can induce modest weight loss in OVX rhesus monkeys<sup>77</sup>. Interestingly, in OVX cynomolgus macaques, E<sub>2</sub> replacement therapy has no effect on body weight or insulin sensitivity in OVX cynomolgus macaques<sup>78</sup>. In addition to ovarian E<sub>2</sub>, E<sub>2</sub> synthesized in the brain can stimulate top-down neuronal signaling cascades and such neuro-E<sub>2</sub> is proposed to act as a neurotransmitter and neuromodulator<sup>79,80</sup>. Aromatase, the enzyme responsible for converting testosterone to E<sub>2</sub> has been documented to be present in rhesus macaque brain regions, including the hypothalamus<sup>81</sup>. We hypothesize that the failure for OVX to increase female NHP body weight is due to the residual actions of neuro-E<sub>2</sub> orchestrating energy balance in the hypothalamus of OVX primates. Rodent research models have provided comprehensive evidence supporting roles played by ovarian E<sub>2</sub> in female energy balance and body composition, yet there are few studies in female NHPs assessing ovarian and extra-ovarian E<sub>2</sub>, possibly neuro-E<sub>2</sub>, in regulating aspects of energy balance.

In addition to the hypoestrogenic environments induced by aromatase inhibiting drugs, like letrozole, hyperandrogenism is usually induced. In female rodent studies

treated with letrozole, animals present with hyperandrogenism and later metabolic phenotypes of dyslipidemia and insulin resistance by 5-weeks post-treatment start<sup>133</sup>. Further, in young female rhesus macaques, the combined treatments of hyperandrogenism and western-style diet contribute to a greater, more severe metabolic phenotype of weight gain and body fat than each treatment alone<sup>134</sup>. Moreover, the distribution of fat has been strongly correlated with androgen and estrogen hormonal ratios. Hyperandrogenism accompanies greater intra-abdominal fat mass in normal weight women with polycystic ovary syndrome (PCOS)<sup>135</sup>. Taken together, the balance between estrogens and androgens is crucial in maintaining energy balance in female rodents and primates.

## **1.5 Genetic and Hormonal Differences in NHP Research Models**

### *1.5.1. Obesity risk genes in rodent and human studies*

The etiology of obesity is complex and likely has many genetic, psychological, and environmental determinants. The traditional mechanism from which obesity arises involves a positive imbalance between food intake (energy consumption) and energy expenditure in favor. While caloric composition and intake, activity level and environmental alterations do contribute largely to the onset of obesity, genetic factors of obesity can contribute 40-70% of obesity variation in humans<sup>82</sup>. Although, genome-wide association scans (GWAS) have identified over 400 genes associated with type 2 diabetes, these genes have a low rate of predicting obesity. This may be due to the difficulty of discerning the effects of a gene or multiple gene mutations from other environmental or epigenetic factors in humans. Generally, genetic causes of obesity can be grouped into three categories<sup>83-86</sup>:

- 1) Monogenic obesity, a single gene mutation
- 2) Polygenic obesity, contribution of multiple genetic mutations or interactions
- 3) Syndromic obesity, neurodevelopmental and organ/system abnormalities

Rodent studies offer intricate, highly controlled investigations into how specific genetic modifications can contribute to the onset of obesity. Prior to the technological advancements that enable quick genetic investigation today, there had long been suspicion of genetic mutations of obesity. The parabiosis work of D.L. Coleman was the first to introduce a possible hormonal factor impacting satiety with a suspected genetic cause<sup>87</sup>, in which an obese mouse parabiosis paired with a normal partner was found to improve weight and satiety while a diabetic obese mouse parabiosis paired with a normal partner increased body weight and fat mass. Coleman concluded the obese mouse might not produce enough “satiety factor” while the diabetic mouse had sufficient satiety factor but could not respond to it due to a defective “satiety center”<sup>87</sup>. This led to the identification and genetic mapping of the mouse obese (*ob*) gene and the subsequent protein it encodes, leptin, which propelled research into identifying the role of leptin and the hereditary effects of obesity<sup>88</sup>. Rodent research confirmed the “satiety factor” and “satiety center” lacking in the parabiosis experiments were leptin and the leptin receptor, respectively<sup>89</sup>, leading to well established leptin deficient (*ob/ob*) and leptin receptor deficient (*db/db*) rodent models. Rodents with the genetic profile of *ob/ob* and *db/db* present with obesity, hyperphagia, reduced energy expenditure and insulin resistance<sup>90</sup>, and while these mutations in humans cause similar phenotypes, predominately, hyperphagia, the occurrence of such monogenic mutations are rare and only found in about 1%-5% of people with severe obesity<sup>91</sup>. A more common occurrence

of disrupted leptin action is via leptin resistance. The levels of leptin-receptor mRNA and protein are important, given that the basic metabolic effects of leptin depend primarily upon the number of available receptors, and it has been shown that circulating leptin levels correlated with leptin receptor levels<sup>136</sup>. Obesity leads to a decrease in the expression of both short and long isoforms of leptin receptors in the hypothalamus, fat, and muscles<sup>136</sup>.

The exploration into the *ob/ob* and *db/db* mouse phenotypes led to other discoveries of monogenic causes of obesity that can be generally categorized into three energy balance systems that become dysfunctional:

- 1) obesity via hypothalamic leptin-melanocortin system.
- 2) obesity via altered hypothalamic development and functions.
- 3) obesity via complex syndromes with unclear relationships to obesity.

The most characterized monogenic causes of obesity affect the hypothalamic leptin-melanocortin signaling pathway. As mentioned above, deleterious leptin and leptin receptor mutations have pronounced impacts on energy homeostasis. Additionally, genetic mutations of the associated genes, proopiomelanocortin (*POMC*), melanocortin receptor 4 (*MC4R*) and prohormone convertase 1/3 (*PC1/3*) in the leptin-melanocortin pathway result in disturbed energy intake and basal energy expenditure<sup>92</sup>. Obesity can also be a consequence of monogenic mutations associated with neurodevelopment. Three genes, *SIM1*, *BDNF* and *NTRK2* have been identified as having a critical role in hypothalamic development and compromising specific nucleotide variants induce hyperphagia and obesity. Although the etiology of obesity in these gene variant cases is not exactly known, it is posited that missing one copy of *Sim1* reduces paraventricular

nucleus (PVN) neuronal number. Since the PVN is the site of MC4R-expressing neurons, the impaired neuronal development within the PVN could result in a dysfunctional energy balance<sup>93,94</sup>. Acting downstream of MC4R are BDNF (brain-derived neurotrophic factor) and its receptor TRKB (tropomyosin-related kinase B) that are involved in proliferation, survival, and differentiation of neurons in neurodevelopment and plasticity, at least in adults. Haploinsufficiency of *BDNF* or *NTRK*, the gene that encodes for TRKB, also induce hyperphagia and obesity<sup>94-96</sup>. These single gene mutations in rodent models present with obesity, reduced energy expenditure and insulin resistance, and while they enable discovery of a gene's direct involvement in pathophysiology and are critical to identify therapies for monogenic causes of obesity in humans, the homogeneity of the inbred mouse lines are not representative of the genetic variation in humans and general obesity and may result in highly varied phenotypic outcomes.

More common are polygenic determinants of obesity in humans. Rodent models of DIO more closely mimic human obesity over monogenic models of obesity. Rodent models of polygenic obesity can therefore offer more clarity into gene-gene and gene-environment interactions due to the ability to control environmental contributions, like caloric content and consumption.

#### *1.5.2. Important differences in regulation of estradiol production and activity between female rodents, sheep and NHPs.*

Female reproductive cycles and their hormonal regulation differ among mammalian species. Rodents exhibit estrous cycles comprising four distinct phases, proestrus, estrus, metestrus, and diestrus. E<sub>2</sub> levels rise and eventually peak in the

proestrus phase. The estrus phase (ovulation) denotes day 1 of each estrous cycle. In contrast, certain New World Primates (NWP), such as marmoset and tamarin monkeys, exhibit ovarian cycles with two distinct phases, follicular and luteal. Cycle day 1 of each ovarian cycle denotes the onset of each follicular phase and is temporally distinct from ovulation that occurs on the last day of the follicular phase. Old World Primates (OWPs), such as rhesus macaques, great apes, and humans, however, exhibit menstrual cycles that have primate-typical follicular and luteal phases. The first day of menses denotes day 1 of each menstrual cycle and the first day of the follicular phase. Ovulation again occurs on the last day of the follicular phase. Both ovarian and menstrual cycles share similar endocrine dynamics with follicular and luteal phases separated by ovulation. The follicular phase involves rapid ovarian follicle growth stimulated by gonadotropins, FSH and LH (or analogous, chorionic gonadotropin, CG, in New World primates), that promote ovarian follicle growth resulting in a slow rise in circulating levels of  $E_2$ . Toward the end of the follicular phase, circulating  $E_2$  levels peak, inducing an LH (CG) surge, that ultimately triggers ovulation of a single dominant follicle (Old World and most New World primates) or 2-4 dominant follicles (certain New World primates, including marmoset and tamarin monkeys). Following ovulation, the single (or multiple) corpus luteum (lutea) produce(s) and release(s) progesterone and  $E_2$ , thus sustaining the luteal phase and implantation-receptive uterine endometrium.

The timing of pre-ovulatory  $E_2$  surges throughout mammalian reproductive cycles has been associated with aspects of energy balance such as body weight and food intake. In rodent estrous cycles, food intake increases during the metestrus-diestrus when  $E_2$  levels are lowest and decreases during the proestrus-estrus phase when  $E_2$

levels are the highest<sup>97</sup>. During hamster estrous cycles, body weight is lowest when circulating E<sub>2</sub> levels are the highest, body weight is the lowest<sup>98</sup>. Similar to rodents, NHPs and humans exhibit reduced food intake when high midcycle (pre-ovulatory) E<sub>2</sub> levels occur during ovarian and menstrual cycles<sup>99</sup>.

### *1.5.3. Important differences in regulation of energy homeostasis between rodents, sheep and NHPs.*

There are many components that contribute to the regulation of energy homeostasis in mammals, including sleep cycles. Metabolic syndrome is a major complication of sleep disorders, prompting the importance of the circadian rhythm in regulating metabolism<sup>100</sup>. In humans, this has been observed in the increased incidence of diabetes and obesity in shift workers as shift work interferes with the normal pattern of sleep<sup>101</sup>. Specifically, disruptions to sleep cycles have demonstrated increased postprandial circulating glucose, triglyceride, and insulin concentrations<sup>102-104</sup>. Adiposity and glucoregulatory dysfunction have also been observed in rodent models of circadian disruption<sup>105</sup>. An important component of metabolic regulation via the circadian rhythm is the hormone, melatonin. In contrast to rodents, sheep and humans produce melatonin in the rest phase, even during constant darkness<sup>106-108</sup>. Despite a similarity in circadian rhythm and melatonin regulation, sheep differ from humans in terms of alimentary organization, as they are ruminants. Their main source of energy is acquired by metabolizing cellulose to volatile fatty acid<sup>109</sup> and demonstrate lower basal glucose levels, reduced post-prandial fluctuations and lower insulin sensitivity compared to humans<sup>110,111</sup>. Moreover, disruption to circadian rhythms in sheep disrupted plasma glucose levels but without concomitant glucose tolerance or glucose-stimulated insulin



secretion dysfunction<sup>100</sup>. Taken together, although circadian rhythms are consistently important for metabolic balance across mammalian species, the specific mechanisms of regulations are different.

BAT thermogenesis is an additional regulator of energy homeostasis. BAT content is critical to the energy expenditure component adaptive thermogenesis. In rodents, there are defined depots of BAT that maintain throughout adulthood<sup>112</sup>. In large mammals, such as sheep, NHPs, and humans, brown adipocytes are mostly interspersed within WAT. Additionally, there are characteristically distinct beige adipocytes, that are interspersed in WAT depots<sup>112</sup>. Small BAT depots have also been identified in neck and supraclavicular region of adult humans and NHPs<sup>113-115,132</sup> and adult rhesus macaques<sup>131</sup>. In addition to BAT, skeletal muscle is an important tissue contributing to adaptive thermogenesis potential. While in rodents, the UCP1 independent pathways promoting mitochondrial uncoupling and futile calcium signaling are well researched, more investigation is needed in large mammals to elucidate these heat producing mechanisms<sup>112</sup>. It is speculated that subtle changes in skeletal muscle thermogenesis can produce large energy balance consequences due to the volume of skeletal muscle in large mammals, accounting for about 40% of body mass<sup>112</sup>.

## **1.6 Clinical Treatments for Estrogen-mediated Metabolic Dysfunction in Women**

Individuals experiencing symptoms of menopause, like hot flashes and night sweats, are typically prescribed hormonal replacement therapy (HRT) to alleviate symptoms and help reduce osteoporosis induced by declining circulating E<sub>2</sub> levels. There are two main types of systemic HRT, estrogen only therapies and combinations of estrogens and progestins, including synthetic forms of progesterone. Additionally,

there are varying delivery methods for HRT, such as oral formulations, transdermal creams, patches and gels, and vaginal preparations<sup>116</sup>. Although protection of bone integrity is a clear benefit of HRT<sup>117</sup>, increased risks of cancer of the breast and endometrium (if uterus is still present), blood clots, stroke, heart disease and dementia are consequences of considerable concern<sup>118</sup>. The literature investigating how HRT contributes to the risk of these diseases is not without inconsistencies. Many of these risks can be attributed to the timing of when HRT is started. There is an increased risk of dementia when HRT is started after mid-life, but a reduced risk if started before mid-life. In the same context, long-term HRT use<sup>119</sup> and chronic estrogen exposure are both associated with increased risk of breast cancer<sup>118</sup>. Considering the critical timing of starting HRT, new guidelines have recommended that HRT starts within 10 years of menopause should not exceed five years in duration and should not continue beyond the age of 60, unless with specific considerations<sup>120</sup>. Hot flashes and other symptoms associated with menopause may persist for 10-20 years from the start of menopause, but the risks associated with HRT greatly increase post-60 years of age<sup>120</sup>. Prior to the age of 60, hormonal combinations and doses for HRT are dependent on physiological condition and individualized risks for each patient.

Individuals who have breast cancer or a history of breast, ovarian, endometrial, or cervical cancer are not good candidates for HRT. Increased exposure to high levels of endogenous estrogens have been shown to increase breast cancer risk. Safe therapeutic interventions for breast cancer patients are incredibly important as many individuals prematurely enter menopause and experience menopausal symptoms due to cancer treatment<sup>121</sup>. Tamoxifen, a SERM, is a successful anti-cancer medication<sup>122</sup>

that blocks estrogen action by competitively binding to ER $\alpha$ , ER $\beta$  and GPER1. In contrast, letrozole, a non-steroidal aromatase inhibitor, blocks the conversion of testosterone to E<sub>2</sub> (as well as the conversion of androstenedione to E<sub>2</sub>) and reduces circulating E<sub>2</sub> (and E<sub>1</sub>) levels<sup>123</sup>. Both treatments have high efficacy in treating hormone-receptor-positive breast cancer in pre- and post-menopausal individuals, especially letrozole as adjuvant treatment, but both treatments present with varying safety profiles<sup>123</sup>. Compared to tamoxifen, letrozole use increased the chance of bone fracture, low-grade hypercholesterolemia, and cardiac events<sup>123</sup>. Additionally, there is evidence that both treatments can lead to weight gain and subsequent metabolic consequences<sup>124-129</sup>.

To treat estrogen-mediated metabolic dysfunction while reducing peripheral side effects, selective and accurate hormone therapies are necessary. One potential solution is a small-molecule bio precursor prodrug, 10 $\beta$ ,17 $\beta$ -dihydroxyestra-1,4-dien-3-one (DHED). The promise of this drug is immense as it converts to 17 $\beta$ -E<sub>2</sub> (or E<sub>2</sub>) in the brain but remains inert in the rest of the body and can thus be administered systemically. DHED was shown to have neuroprotective effects and to not induce proliferation of cancerous tissues in rodents<sup>130</sup>. DHED has the potential to mitigate metabolic consequences associated with disorders from depleted brain E<sub>2</sub> levels<sup>130</sup>, such as menopause related metabolic disorders.

## 1.7 Objectives and Importance

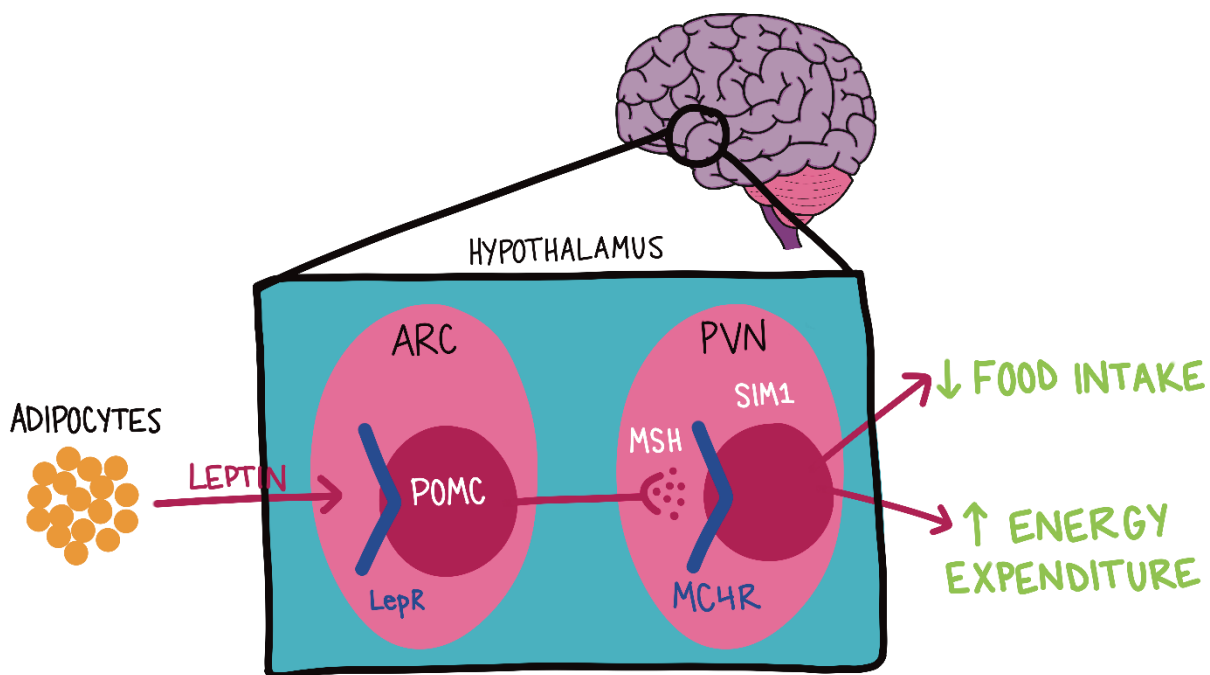
Activation of neural ER $\alpha$  orchestrates responses in multiple pathways that regulate metabolism in female rodents and sheep, yet we still do not fully understand ER $\alpha$  neuro-metabolic signaling in female NHPs and women. Understanding the mechanisms

through which E<sub>2</sub> mediates metabolic homeostasis via ER $\alpha$  will enable targeted therapeutic solutions to those experiencing menopausal- and cancer therapy-associated metabolic dysfunction.

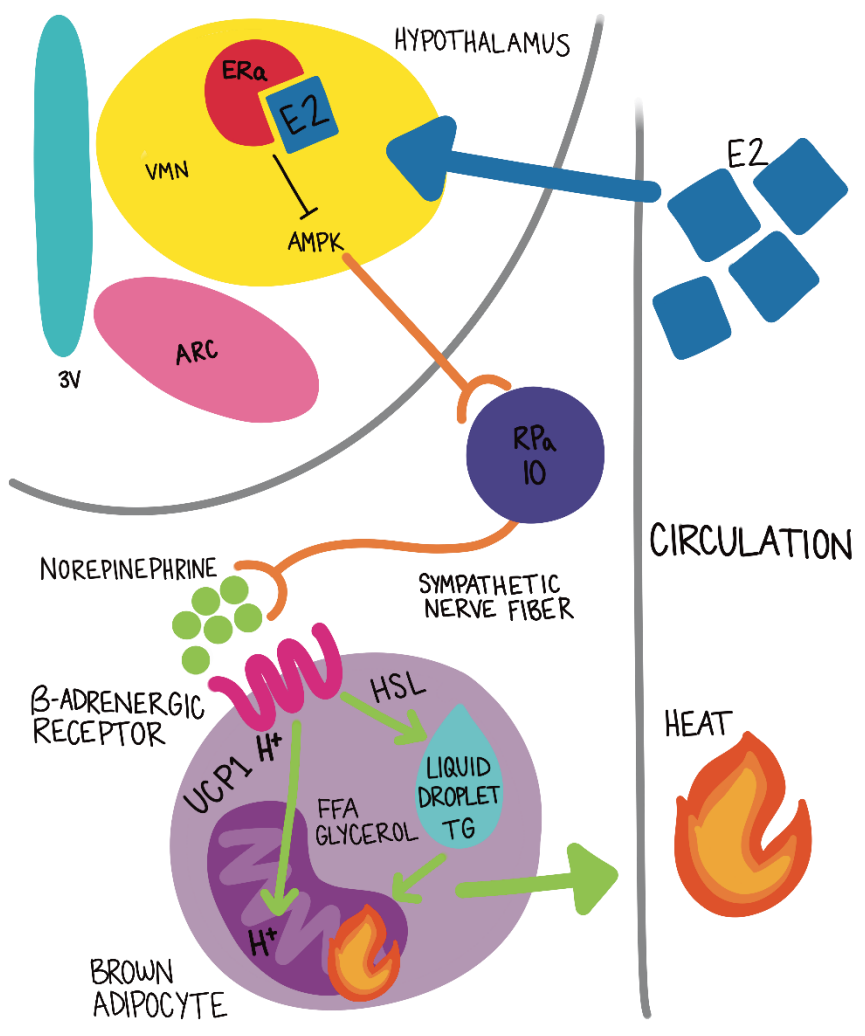
The objectives of this thesis were therefore to test the hypotheses that, 1) in a New World Primate model, the female common marmoset, extra-ovarian E<sub>2</sub> signaling through hypothalamic ER $\alpha$  regulates body weight, body composition, energy homeostasis and glucose homeostasis (**Chapter 2**), 2) in the absence of DIO, hypothalamic ER $\alpha$  signaling is critically important in regulating female energy homeostasis in rhesus macaques (**Chapter 3**), 3) extra-ovarian E<sub>2</sub> contributes to regulation of body weight, body composition, energy homeostasis and glucose homeostasis in female rhesus macaques (**Chapter 4**), and 4) concluding discussion revising consideration of E<sub>2</sub> regulation of female energy homeostasis in NHPs and likely women (**Chapter 5**).

This thesis is of clinical and biological importance because current modes of treating metabolic syndrome in menopausal women vary in effectiveness and possess co-morbidities. Moreover, extra-ovarian E<sub>2</sub> as a contributing source of estradiol maintaining energy homeostasis after the menopausal transition has never been investigated before in an NHP model. Identifying the source of E<sub>2</sub>, and the molecular pathways through which it regulates energy balance in female NHPs, will prompt a major reconceptualization of the fundamental control systems that govern metabolic function female NHPs and likely, women. Additionally, this understanding will provide novel therapeutic opportunities to target specific neural sites of action to combat menopausal associated metabolic syndromes.

## 1.8 Tables &amp; Figures



**Figure 1:** Depiction of leptin-melanocortin pathway. Leptin synthesized by adipose tissue activates leptin receptors (LepR) on POMC neurons in the arcuate nucleus (ARC) of the hypothalamus. This stimulates downstream activation of melanocortin 4 receptors (MC4R) in the paraventricular nucleus (PVN). Ultimately, influencing metabolic processes by decreased food intake and increased energy expenditure.



**Figure 2:** Depiction of estrogen receptor alpha (ERα) action via estradiol (E<sub>2</sub>) in the ventromedial nucleus (VMN) of the hypothalamus and downstream initiation of brown adipose tissue (BAT) thermogenesis.

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**2. CHAPTER TWO: Hypothalamic Estrogen Receptor Alpha, *but not extra-ovarian estradiol*, essential for Metabolic Regulation in Female Marmoset Monkeys**

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## 2.1 Abstract

Ovarian estradiol via estrogen receptor alpha (ER $\alpha$ ) regulates metabolic function in many female animal models. While ovariectomy (OVX) in rodents enables obesity, OVX in nonhuman primates inconsistently alters weight gain. We hypothesize that extra-ovarian estradiol provides key support for metabolism through hypothalamic ER $\alpha$ , and to test this idea, we completed two separate experiments. First, we employed aromatase inhibition to eliminate extra-ovarian estradiol biosynthesis and diet-induced obesity to enhance weight gain. Thirteen adult female marmosets were OVX and received (1) estradiol-containing capsules and daily oral treatments of vehicle (E2; n = 5); empty capsules and daily oral treatments of either (2) vehicle (VEH, 1 mL/kg, n = 4), or (3) letrozole (LET, 1 mg/kg, n = 4). VEH and LET females demonstrated increased % body weight gain (P = 0.01). Relative estradiol levels in peripheral serum were E2 >>> VEH > LET, while those in hypothalamus ranked E2 = VEH > LET, confirming inhibition of local hypothalamic estradiol synthesis by letrozole. Next, we utilized MRI-guided viral vector infusion to knockdown hypothalamic ER $\alpha$  expression in target regions of hypothalamic ventromedial (VMN) and arcuate (ARC) nuclei. Eleven adult female marmosets were OVX and received (1) OVX + estradiol replacement + scrambled virus shRNA (control; n=5) or (2) OVX + estradiol replacement + ER $\alpha$  gene silencing shRNA (ER $\alpha$ KD; n=4). ER $\alpha$  expressing cells were less numerous in the VMN in ER $\alpha$ KD compared to control females, while intensity of ER $\alpha$  expression in the ARC was less in ER $\alpha$ KD compared to control females. Our findings provide clear evidence for ovarian estradiol protecting female nonhuman primates from obesity and they also support the

notion that the majority of metabolically effective E<sub>2</sub> activity is mediated through ER $\alpha$  within the hypothalamic VMN and ARC.

## 2.2 Introduction

Aromatase, a cytochrome P450 enzyme, encoded by the CYP19A1 gene, converts testosterone to E<sub>2</sub>, and androstenedione to estrone. While the ovaries are a major source of E<sub>2</sub>, the same hormone is also produced at aromatase-expressing extra-ovarian sites, including liver, breast, skin, bone, pituitary gland, and various brain regions<sup>1-4</sup>. Neural production of E<sub>2</sub> in particular has been identified in birds<sup>5-7</sup>, mice<sup>8</sup>, rats<sup>9-12</sup>, as well as in monkeys<sup>13-16</sup>. Brain aromatase is expressed at high levels in the medial basal hypothalamus, preoptic area, amygdala, and hippocampus, and has a higher affinity for androgen substrates than its ovarian counterpart<sup>8,17</sup>. A role for hypothalamic aromatase in regulating gonadotropin-releasing hormone (GnRH) in female rhesus monkeys was recently demonstrated to show that (1) E<sub>2</sub> is produced and released at detectable levels within OVX monkey pituitary stalk-median eminence (SME), (2) E<sub>2</sub> synthesis and release depends upon acute elevations in systemic E<sub>2</sub> initiating detectable aromatase activity in the SME, and (3) hypothalamic E<sub>2</sub> can rapidly stimulate GnRH release in the SME<sup>14,15,18</sup>. These studies, in addition to previous work in female marmoset monkeys<sup>16,19</sup>, provide evidence for extra-ovarian E<sub>2</sub> production in female nonhuman primates (NHPs) engaging neural action.

In many mammalian species, including rodents (rats, mice, guinea pigs), sheep, and NHPs, E<sub>2</sub> is a major regulator of female body weight regulation and insulin–glucose homeodynamics<sup>20-22</sup>, mediating most of its activity by binding to estrogen receptor alpha (ER $\alpha$ )<sup>23-25</sup>. Neural signaling by E<sub>2</sub> in the female hypothalamus regulates metabolic homeostasis<sup>26,27</sup>. Hypothalamic E<sub>2</sub> activity is mostly mediated by estrogen receptor alpha (ER $\alpha$  and its encoding gene, *ESR1*)<sup>28,29</sup>. ER $\alpha$  exerts descending control over

systemic organ systems via autonomic innervation<sup>30-32</sup> and intra-hypothalamic innervation<sup>33</sup>, including E<sub>2</sub>-induced alterations in neuroendocrine regulation of ovarian function and food intake and energy expenditure, producing secondary endocrinological and metabolic states.

E<sub>2</sub>-regulated metabolic control mechanisms are pronounced in female rodents, exemplified by OVX-mediated E<sub>2</sub> depletion reliably increasing body weight and visceral adiposity, reducing physical activity and energy expenditure, as well as diminishing glucose tolerance and insulin sensitivity<sup>34-38</sup>. Consistent with these findings, aromatase knockout female mice develop obesity with insulin resistance in the absence of endogenous E<sub>2</sub> synthesis<sup>39</sup>. All these effects are prevented or reversed by physiological E<sub>2</sub> replacement<sup>40,41</sup>. Furthermore, while intact female mice are resistant to high-fat diet-induced obesity (DIO) and its associated sequelae, OVX-mediated E<sub>2</sub> depletion abolishes this protection<sup>42,43</sup>. Virtually all of these E<sub>2</sub> activities are mediated by ER $\alpha$ , as ER $\alpha$  knockout adult mice (ER $\alpha$ KO) bearing null mutations of *ESR1* gene exhibit body weight, adiposity, and energy metabolism phenotypes that largely mimic those observed in long-term OVX adult mice<sup>23-25</sup>. In addition, loss of E<sub>2</sub> bioactivity in bone, particularly that mediated by ER $\alpha$ , reliably results in skeletal bone loss<sup>44</sup>.

Estrogen deficiency is of particular clinical relevance for women in their post-reproductive years (menopause)<sup>45</sup>, as well as for 80% of girls and young women who survive cancer yet live with premature ovarian insufficiency (premature menopause) as a result of alkylating agent and/or radiation cancer therapy<sup>46,47</sup>. Women with such estrogen deficiency experience compromised cardiometabolic dysfunction, obesity, and osteoporosis<sup>48</sup>. Hormonal therapies, including systemically administered E<sub>2</sub>, alleviate



clinical conditions to varying degrees<sup>49,50</sup>. Currently approved estrogen medications, however, increase the risk for harmful systemic side-effects<sup>51,52</sup>, including blood clots and cancer<sup>53,54</sup>. Estrogen treatments will therefore never realize their full potential until, perhaps, their actions are limited to the brain. In this regard, however, the extent to which E<sub>2</sub> regulates neural control systems in women is largely unknown.

With regard to the hypothalamic regulation of female NHP metabolic homeostasis, the VMN has been identified as a key regulatory site following electrolytic lesions of the female macaque hypothalamus<sup>54,55</sup>. When lesions included portions of the VMN, female macaques demonstrate increased weight gain. Stimulatory effects of E<sub>2</sub> on energy expenditure are transduced in ER $\alpha$ -expressing neurons within the VMN<sup>56</sup> and by non-classical ER $\alpha$  signaling<sup>57</sup> coupled to activation of PI3-kinase<sup>58</sup>. E<sub>2</sub> also regulates gene expression associated with regulation of food intake and energy expenditure in the female hypothalamus, largely through ER $\alpha$  activation<sup>59,60</sup>. Furthermore, it has been demonstrated that viral vector-mediated ER $\alpha$  gene silencing in the VMN of both female mice and rats largely recapitulates a metabolic phenotype observed in whole-body ER $\alpha$  knockout (ER $\alpha$ KO) mice, including obesity, hyperphagia, impaired glucose tolerance and reduced energy expenditure<sup>21-23</sup>. In addition, ER $\alpha$ -expressing neurons within the ARC have also been demonstrated to play key roles in regulating female metabolic homeostasis<sup>27</sup>.

In menopausal women, declining E<sub>2</sub> concentrations and progressive testosterone predominance are generally associated with increased abdominal fat mass and increased risk for impaired glucose metabolism<sup>61-63</sup>. Metabolic functions of E<sub>2</sub> in women have been difficult to define, however, partly due to logistical and ethical constraints in

designing definitive experiments with rigorous control. OVX-mediated E<sub>2</sub> depletion had small effects on female rhesus macaque body weight in one study<sup>64</sup>, and no effects on female body weight were observed in two studies of cynomolgus macaques<sup>65,66</sup>. While a putative selective estrogen receptor modifier (SERM) can promote weight loss in OVX rhesus monkeys<sup>67</sup>, E<sub>2</sub> replacement therapy has no effect on body weight in OVX cynomolgus macaques<sup>66,68</sup>. E<sub>2</sub> and SERM activity, however, both reliably maintain skeletal bone mass in female macaques<sup>69-71</sup>.

To address the different contributions of ovarian and extra-ovarian E<sub>2</sub> in a female NHP model and to elucidate the major neural receptor mechanism of E<sub>2</sub> signaling and its role in regulating female metabolic function, two studies were undertaken. First, an aromatase inhibitor, letrozole, was employed to diminish E<sub>2</sub> production in OVX female marmosets and thereby enable an investigation into whether extra-ovarian estradiol contributes to female weight gain in addition to ovarian E<sub>2</sub>. We hypothesize that an extra-ovarian source of E<sub>2</sub>, likely the hypothalamus, will be diminished by aromatase inhibition and subsequently will enhance weight gain and skeletal bone loss, compared with both E<sub>2</sub>-replaced female marmosets and those experiencing the loss of ovarian E<sub>2</sub> alone. Second, to investigate the role for ER $\alpha$ -expressing neurons in the hypothalamic VMN and ARC in regulating female metabolic homeostasis in a female NHP, we used MRI-guided, neural infusions of an adeno-associated virus, AAV8, into the adult female marmoset hypothalamic VMN and ARC in order to deliver a small hairpin RNA (shRNA) either encoding a scrambled virus control, with no known gene targets in this species, or an shRNA specifically designed to associate with *ESR1* mRNA, alone, to prevent translation of ER $\alpha$  protein. We hypothesize that like female rodents<sup>5,6</sup>, *ESR1* gene

silencing in both the VMN and ARC of female marmoset monkeys will disrupt metabolic homeostasis.

## **2.3 Methods**

*EXPERIMENT 1: Diminished E<sub>2</sub> production: ovarian (OVX) depletion of E<sub>2</sub> (VEH) versus extra-ovarian and ovarian (OVX + aromatase inhibitor, letrozole) depletion of E<sub>2</sub> (LET) versus Peri-Ovulatory E<sub>2</sub> Replacement (OVX + E<sub>2</sub>, E<sub>2</sub>)*

### *Animals*

Thirteen adult female common marmosets from the Wisconsin National Primate Research Center (WNPRC) colony were OVX and assigned to 1 of 3 treatment groups that were balanced by age and body weight at the onset of the study (Table 1), systemic mid-cycle, peri-ovulatory E<sub>2</sub> replaced (E<sub>2</sub>; n = 5), systemic E<sub>2</sub> depleted, OVX plus daily vehicle (VEH; n = 4), or extra-ovarian E<sub>2</sub> depletion, OVX plus daily letrozole administration (LET; n = 4). Systemic (mid-cycle, peri-ovulatory) E<sub>2</sub> replacement was achieved through subcutaneous E<sub>2</sub>-filled capsules that maintained a systemic level of E<sub>2</sub> (Table 2) mimicking mid-cycle, peri-ovulatory circulating E<sub>2</sub> levels. The latter were sufficient to maintain negative feedback regulation of circulating pituitary gonadotropin levels within the ovary intact female range (Table 2)<sup>19</sup>. To maintain constant E<sub>2</sub> levels, capsules were replaced every 3 months throughout the study<sup>72</sup>. VEH females received a daily oral 200 µL of 1 mL/kg Ensure© as vehicle control, while LET females were given daily oral 1 mg/kg of letrozole dissolved in 200 µL of vehicle, as previously determined<sup>19</sup>, and females in both groups received empty capsules when E<sub>2</sub> females received E<sub>2</sub>-filled capsules and replacements.

All females lived with a well-established male cage mate in 0.60 m × 0.91 m × 1.83 m enclosures and were maintained with 12-hour lighting (06:00 hours to 18:00 hours), ambient temperature of ~27°C and humidity of ~50%. The protocol for this study was approved by the Animal Care and Use Committee of the Office of the Vice Chancellor for Graduate Research and Education of the University of Wisconsin–Madison, an AAALAC-accredited program. Following baseline assessments, bilateral OVX was performed. Cloprostenol (Estrumate®, 0.75-1.50 µg intramuscular injection for 2 successive days approximately 11-60 days after ovulation), an analog of prostaglandin-F<sub>2</sub>-alpha, was administered prior to OVX to facilitate scheduling of OVX during the follicular phase<sup>19</sup>. Treatment onset (0 months) was the day of the OVX procedure.

#### *Hormone Assays*

Plasma samples and hypothalami were analyzed for several hormones. Hypothalami were dissected at necropsy and frozen at –80°C. They were subsequently thawed, transected along the midline, and separated into hemi-hypothalami. The right hemi-hypothalamus per monkey was divided into 50- to 75-mg aliquots. For steroid hormone analyses, plasma and hemi-hypothalami aliquots underwent extraction<sup>13</sup> and hemi-hypothalamic aliquots were recombined after extraction. Extracted samples were subsequently submitted for analysis on a QTRAP 5500 quadrupole linear ion trap mass spectrometer (AB Sciex) equipped with an atmospheric pressure chemical ionization source (liquid chromatography tandem mass spectrometry [LC-MS/MS])<sup>19</sup>.

#### *Diet-induced Obesity from Consumption-driven Weekly Increments in Daily Calories*

Animals were fed with Mazuri Callitrichid High Fiber Diet #5MI6 (Purina Mills International, St. Louis, MO), providing 53% carbohydrate, 20% protein, 6% fat, and 10% fiber by weight, with a metabolizable energy of 3.3 kcal/g (~61%, 23%, and 16% kcal from carbohydrate, protein, and fat, respectively)<sup>73</sup>. Following OVX at study onset, total daily calorie consumption by each male–female study pair was recorded. To achieve DIO, diet allotment for each male–female study pair was increased weekly by ~66 kcal per day (equivalent to 20 g diet wet weight/day) if the entire daily allotment had been consumed during at least 4 out the 7 previous days. Calorie increments and total calories consumed, however, remained comparable across the pairs between treatment groups (Figure 1).

#### *Assessment of Daily Calorie Consumption While Maintaining DIO*

At 6 to 7 months post-OVX, each female was singly housed within a marmoset housing room, but outside visual contact with her male pair mate, while still maintaining vocal and auditory contact. Each female's daily calorie allotment began at 50% of those provided when housed with their male pair mate. Dietary allotment was increased weekly by ~33 kcal per day (~10 g diet wet weight) only when the entire daily allotment was consumed during at least 4 out the 7 previous days. Calorie increments and total calories consumed, however, remained comparable across females between treatment groups (Figure 1). The total kilocalories consumed daily were recorded for each female for 8 weeks.

#### *Body Composition and Bone Mass*

Animals were weighed weekly. Area under the curve (AUC) assessment of weight parameters over time, calculated by the trapezoid rule, was employed to better detect

recurring differences in weight gain, as previously employed<sup>74</sup>. For the first study, at baseline and 5 to 6 months post-OVX and for the second study, at baseline and 3 and 6 months post-OVX, total body composition, as well as bone mineral content (BMC) and bone mineral density (BMD), were assessed by dual-energy X-ray absorptiometry (DXA, iDXA, GE/Lunar Corp., Madison, WI) on sedated animals. Fat, fat-free mass (FFM) (excluding bone), BMC, and BMD were determined for total body as well as previously validated body regions of interest, including abdomen, chest, thighs, lower legs, and arms<sup>75</sup>.

#### *Locomotor Activity*

At baseline and 5 to 6 months post-OVX, a small accelerometer (Actiwatch Mini, CamNtech Ltd, Cambridge UK) was added to each female's standard collar. Activity and intensity of movement were recorded over a ~2-4-week period after which the accelerometers were removed. For the most part, activity recorded represented whole-body movements, and not limb or head movements, alone, as previously validated in marmosets<sup>76</sup>. The accelerometer sampled such activity counts every 30 seconds and these data were averaged for every hour, day (during lights on), night (during lights off), morning (06:00-12:00 hours), afternoon (12:00-18:00 hours), and 24 hours. AUC activity values were also assessed to detect recurring differences in activity over time.

#### *Fasting Glucose and Oral Glucose Tolerance Test*

Fasting glucose and glucoregulation were assessed in overnight fasted, awake animals. Fasting glucose was determined at baseline and 6 months post-OVX, while glucoregulation was assessed by oral glucose tolerance test (OGTT) at 6 months following OVX. Following a baseline blood sample, animals were given an oral dose (5

mL/kg) of 40% sucrose. Blood samples were then collected at 15, 30, 60, and 120 minutes (2 hours) following sucrose administration and assessed for glucose, as previously validated for marmosets<sup>77</sup>. Glucose was measured by glucometer (Accu-Check Aviva, Roche Diagnostics, Indianapolis, IN). AUC glucose values during the OGTT were also assessed to detect between group differences in accumulating high levels of glucose over time.

### *Statistical Analysis*

Data were analyzed utilizing SPSS software (IBM, Armonk, NY). Hormone measures were analyzed with a 1-way analysis of variance (ANOVA), followed by Bonferroni multiple comparison post hoc tests. All other between group analyses were performed using a 2-way ANOVA for repeated measures. Nonparametric Spearman's correlation tests were utilized to identify relationships between hormone values and metabolic parameters. Spearman's rho coefficients are expressed as  $r_s$ . Statistical significance was determined at  $P \leq .05$ .

### *EXPERIMENT 2: Diminished hypothalamic E<sub>2</sub> activity: silencing RNA specific for ESR1 (shRNA-ESR1) targeted on the hypothalamic VMN and ARC*

#### *Animals, Ovariectomy and E<sub>2</sub> Replacement*

Nine adult female common marmosets (2-6 years of age) from the WNPRC colony, and already housed in well-established male-female pairs were evenly assigned into two groups based on age, body weight and body mass index (BMI) (Table 3): OVX + E2 replacement + scrambled virus shRNA (control; n=5) and OVX + E2 replacement + ER $\alpha$  gene silencing shRNA (ER $\alpha$ KD; n=4). Monkeys were maintained in these two groups for ~9 months in 0.60m x 0.91m x 1.83m enclosures under 12-h lighting,

ambient temperature of ~27°C and humidity of ~50%. The animal care and use program at the University of Wisconsin maintains a Public Health Services Assurance. The protocol for this study was approved by the Animal Care and Use Committee of the Office of the Vice Chancellor for Graduate Research and Education of the University of Wisconsin–Madison, an AAALAC-accredited program.

Following baseline assessments, bilateral OVX was performed in all females at least one month prior to neural surgery. Cloprostenol (Estrumate., 0.75-1.50 µg intramuscular injection for two successive days approximately 11-60 days after ovulation), an analog of prostaglandin-F<sub>2</sub>-alpha, was administered prior to OVX to facilitate scheduling of OVX during the follicular phase. Such cloprostenol administration is without subsequent effect on study outcomes<sup>19,78</sup>. All females received subcutaneous, E2-filled silastic capsules immediately following OVX. Silastic capsules were removed and replaced at 3-month intervals post-OVX to approximate mid-cycle, peri-ovulatory systemic E2 levels and to maintain negative feedback regulation of circulating pituitary gonadotropin levels within the range of ovary intact female marmosets<sup>19,78</sup> (Table 2).

#### *Neural Infusion Surgery and ER $\alpha$ Knockdown*

OVX adult female marmosets received bilateral stereotaxic injections into the hypothalamic VMN and ARC of adeno-associated virus 8 (AAV8) expressing shRNA targeting *ESR1* for *in vivo* RNAi, and thus diminishing *ESR1* translation into ER $\alpha$  (ERaKD, n=4), or a control scrambled shRNA (Control, n=5). Treatment onset (0 months) was the day of neural infusion surgery.

Since accurate, within-brain placement of viral vector was essential for gene silencing efficacy and specificity, we used a single, presurgical, neuro-anatomical MRI



scan to refine the unique locations of each VMN and ARC within each marmoset, refined and adapted from previous studies<sup>79</sup>. To accomplish this, each female marmoset was food deprived overnight, sedated with an intramuscular (IM) injection of ketamine (10 mg), anesthesia was continued with an IM injection of meloxicam (2 mg), 2 ml isotonic saline was injected subcutaneously (SC) to provide fluid administration and atropine (0.016 mg) was administered IM, followed by placement of the marmoset into an MRI-compatible stereotaxic frame. Lateral placements of both left and right earbars, as well as vertical placements of palate and left and right eye bars, were all noted. Three-dimensional coordinate locations of both left and right VMN and ARC were subsequently obtained from a T1 MRI anatomical scan adapted for marmosets<sup>80</sup> in a next-generation 3T MRI system MR750 (GE Healthcare, Waukesha, WI) and performed at least 3 days prior to viral vector infusion (Figure 2).

Each presurgical scan employed a 3-inch surface coil and comprised 3D T1-weighted (T1W) images with an inversion-recovery (IR) prepped, fast gradient-echo (IR-fGRE) sequence adapted for marmosets<sup>80</sup>. Scanning parameters included an inversion time (TI) of 450 ms, repetition time (TR) of 6.8 ms, echo time (TE) of 2.9 ms, and receiver bandwidth of 62.5 kHz. The matrix size was 256 × 256 × 128 (X × Y × Z). The 3D T1W scans produced 248 coronal 0.3-mm-thick contiguous slices. Using coronal slices incorporating the hypothalamus from each monkey's presurgical scan, we utilized contrast-agent filled earbars in the stereotaxic frame to provide a rostral-caudal (RC) zero plane (Z coordinates), the middle of the 3<sup>rd</sup> ventricle at the level of the mediobasal hypothalamus to generate a medial zero plane (X coordinates), and the farthest lateral point of each earbar to provide a dorso-ventral zero plane for right and left sides

separately (X coordinates). Based on these 3D parameter sets, target coordinates for entry into the skull, as well as for rostral, medial-rostral, medial-caudal and caudal left and right VMN, and analogous locations for both left and right ARC, were estimated based on marmoset neuroanatomy atlas coordinates<sup>79</sup>.

For viral vector infusion surgery, each marmoset was sedated and prepped as described for their presurgical MRI scan. In addition, each monkey was intubated, and the anesthesia was continued under isoflurane (1-2%). Each marmoset was re-positioned into similar stereotaxic frame parameters as noted during their presurgical MRI scan. Vital signs (heart rate, blood oxygen, respiration, and temperature) were regularly monitored during the procedure and the animals were wrapped for warmth. A series of small-volume, subcutaneous (SC) injections of lidocaine (total, 1 mg) were given along the anticipated incision site to provide local anesthesia. A ~30mm incision was made on the top of the animal's head. Stereotaxic coordinates, estimated from our pre-surgery anatomical MRI scan, allowed calculation for the insertion sites of a 22-gauge guide cannula housing a 28-gauge stylet to target the VMN and ARC in the hypothalamus. The estimated midline location above the medial sites for both VMN and ARC was marked on the skull to enable excision of a ~10 mm diameter portion of skull using a surgical bone drill. A micromanipulator, with cannula holder and guide cannula with screw-in solid stylet, were placed on the stereotaxic frame to confirm midline of the brain from the medial location of the superior sagittal sinus, and presurgical scan estimated target coordinates for left VMN and ARC sites were revised accordingly. A 27g needle, customized into a miniature scalpel blade, was used to open a ~5mm incision in the dura mater vertically above the anticipated locations of the VMN and ARC

infusion sites on the left side of the marmoset brain, and avoiding the left lateral extent of the superior sagittal sinus.

At a trajectory angle of 0° (vertical), a 22-gauge guide cannula housing a sterile 28-gauge stylet was lowered into the medial-rostral estimated extent of the left ARC, as modified for marmosets<sup>82,84</sup>. The stylet was then replaced by a 28-gauge infusion cannula attached to a Hamilton syringe controlled by a programmable infusion pump. A solution of viral vector (~2 x 10<sup>10</sup> packaged genomic particles of AAV8 in 3.0 µL of sterile saline) mixed with gadolinium-containing Multihance MRI contrast agent (2 mM gadobenate dimeglumine, Bracco Diagnostics Inc.), was infused at 0.2 µL per minute. The AAV8 virus utilized in this study was shown to readily infect neurons<sup>83</sup> and was without notable inflammation in NHP brain<sup>84</sup>. To minimize dorsal tracking of viral particles and contrast agent when the infusion cannula was raised, there was a 5-min delay following cessation of infusion before infusion cannula removal alone to permit its flushing of sufficient fluid while outside the brain to confirm infusion integrity of the system. The stylet was repositioned inside the guide cannula. Both guide cannula and stylet were then gradually removed from the brain. Superficial skin sutures were placed using 4/0 vicryl suture to temporarily close the incision site during transfer to the MRI scanning room while the marmoset remained in the stereotaxic frame.

The contrast agent remaining at the medial-rostral ARC infusion site was used to confirm accuracy of the infusion site location from a mid-surgery, an ~23-min MRI T1 scan obtained approximately 10-20 min following the first infusion on the left side. When necessary, presurgical scan estimated target coordinates for left VMN and ARC sites were revised. Guide cannula, stylet and infusion cannula insertion, infusion and removal

procedures were then repeatedly performed to enable AAV8 infusion into rostral, medial-rostral, medial-caudal, and caudal VMN and ARC target sites on the left side of the brain.

Following all left side infusions, a 27g needle customized into a miniature scalpel blade, was used to open a ~5mm incision in the dura mater vertically above the anticipated locations of the VMN and ARC infusion sites on the right side of the marmoset brain, avoiding the right lateral extent of the superior sagittal sinus. A similar series of procedures, infusions and MRI scan were performed on the right side of the brain analogous to those performed on the left. When necessary, adjustments to presurgical estimates for rostral, medial-rostral, medial-caudal, and caudal right VMN and ARC target sites were made following the 2<sup>nd</sup> within-surgery MRI scan.

*Immunohistochemical processing of marmoset brain for visualizing and quantifying green fluorescent protein (GFP) and ER $\alpha$  protein expression*

Following upper body perfusion with 4% paraformaldehyde (PFA) (pH 7.6), the brain was removed, post-fixed at 4C for an additional 12-16h and cryoprotected in graded (10-30%) sucrose/PBS (pH 7.2) solutions. Before final storage at 4C in a sodium azide-30% sucrose-4% PFA solution, each post-fixed, brain was placed in a customized plastic mold designed to accommodate the dorsal surface of an adult marmoset brain and expose the ventral surfaces landmarks, enabling placement of mold-guided, microtome blades cutting the brain into coronal plane portions. The medial rostral-caudal portion (~2-3 mm), including the hypothalamus, was separated immediately rostral to the optic chiasm to immediately caudal to the mammillary bodies. This medial rostral-caudal portion of whole-brain was cryosectioned into consecutive 40 $\mu$ m frozen sections in the

coronal plane to include the rostral-caudal extent of the hypothalamus from the rostral extent of the mPOA to the caudal extent of the ARC.

Immunohistochemistry (IHC) procedures for GFP and ER $\alpha$  were performed and adapted for the marmoset monkey<sup>85</sup>. Cryo-sections of 40  $\mu$ m, at regular intervals, were immunostained for ER $\alpha$  using the mouse monoclonal antibody, 6F11 (Thermo Fisher Scientific, Hampton, NH)<sup>20</sup>. DAB-staining was subsequently used to reveal specific primary antibody binding. Sections close by to those selected for ER $\alpha$  immunostaining were stained for GFP using the rabbit polyclonal antibody, AB3080 (MilliporeSigma, Burlington, NH)<sup>86</sup>.

*Confirmation of neural targeting accuracy from immunohistochemical analyses of GFP expression in the mediobasal hypothalamus*

Digital images from coronal brain sections immunostained for GFP were obtained with a Nikon Microphot-FA microscope using a QImaging Retiga 200R CCD camera and Nikon NIS Elements software. Most images for analysis were obtained using a 4X objective. Each tissue section picture was the composite of several images stitched together by the Nikon system during the photo acquisition process. This resulted in a series of coronal images through the hypothalamus from the rostral POA at the level of the optic chiasm to the mammillary bodies in the caudal hypothalamus. As illustrated by a coronal section from approximately the mid-rostral-caudal extents of VMN and ARC within the hypothalamus (Figure 3), abundant GFP-immunoreactive (ir) labeling was demonstrated in VMN and ARC, and in portions of the immediately ventral PIT and immediately lateral optic tracts (Figure 3).

### *Analysis of Neural Targeting of ER $\alpha$ Gene Silencing*

ER $\alpha$ -immunoreactive (ir) labeling in the POA, VMN and ARC, employing measures of cell counts, cell density per unit area (number of cells/mm<sup>2</sup>), and labeling intensity (pixel intensity of individual cells). Labeled cells were detected using the NIS Elements AR Dark Spot Detection feature. The contrast threshold in the detection algorithm was determined for each case by adjusting cell counts derived from the spot detection algorithm to match cell counts made by three observers on identical images of tissue. Observer counts differed by < 15%, and NIS-derived cell counts fell within 5% of the average counts of all observers. The NIS Elements AR software calculated the mean pixel intensity of each identified cell, and we used this to derive the intensity of ER $\alpha$  staining. The average pixel values of the observer-derived threshold cells determined the low (lighter) end of the intensity staining range. The high (darker end) of the range was determined by the average pixel intensities of the darkest 2% of labelled cells, typically in the POA or pituitary, which were not targeted by the infusion. The staining intensity of individual cells was normalized (from 0 to 1) within the light to dark intensity range.

The number of labeled ER $\alpha$  cells was quantified in terms of cell density per unit area. This was determined by counting the number of immunostained neurons within regions of interest (ROIs) of 1000 $\mu$ m X 500 $\mu$ m ellipses (areas of 0.395 mm<sup>2</sup>) positioned within POA, VMN and ARC, in addition to ROIs within ventral VMN (VMNv) and dorsal ARC (ARCd) for all nine female marmosets. For each hypothalamic ROI within each female, we used the average of the three most intensely staining areas (comprising most 'dark' or most 'dark' and 'light' cells) as the density measure for that ROI. One exception, however, involved the rostral POA, the area with typically the most ER $\alpha$  expression, that

was lost during processing in one control female, The POA ROI density measures from this control female were thus obtained from a typically more demure, ER $\alpha$  immunostained portion of the POA. Since this POA ROI density value for ER $\alpha$  was not an outlier in the context of the remaining eight females, it was retained for quantitative analyses.

*Technical note regarding effects of focus plane on cell quantification.*

Cells appear most darkly labeled when the microscope objective is focused at the tissue depth that contains the cell. When the focal plane deviates from the depth of the cell, the cell blurs, and its staining darkness declines. To minimize this confound, images were collected when the microscope objective was focused on the middle depth of the tissue. While this will lead to underestimating the darkness of label of a cell located above or below the middle of the tissue and possibly prevent its detection, we found this effect to be small when using a 4X objective with its relatively large depth of field. To determine the effect of focus on estimates of staining intensity and cell count, we systematically varied focus across the depth of a tissue section and collected images through a large group of labeled cells. We applied the NIS analysis to the images and found that an image captured with a 4X objective focused in the center depth of the tissue provides a reliable estimate of the number of cells and their staining intensity. On average, mid-depth focus captured >95% of cells that were identified at all the focal positions. The additional cells were stained lightly, near detection threshold. To examine the effects on estimates of staining intensity, we tracked the mean pixel intensity of all identified cells across the depths of focus. The average difference between the estimate of staining darkness at mid-depth focus and the maximum darkness at any depth, normalized to the full range of staining intensities, was  $\sim 0.021 \pm 0.027$  std.

### *Hormone Values*

Plasma samples and hypothalami were analyzed for several hormones as described above in Experiment 1.

### *Diet-induced Obesity from Consumption-driven Weekly Increments in Daily Calories*

Animals were fed with Mazuri Callitrichid High Fiber Diet #5MI6 (Purina Mills International, St. Louis, MO), providing 53% carbohydrate, 20% protein, 6% fat, and 10% fiber by weight, with a metabolizable energy of 3.3 kcal/g (~61%, 23%, and 16% kcal from carbohydrate, protein, and fat, respectively)<sup>73</sup>. Following OVX at study onset, total daily calorie consumption by each male–female study pair was recorded, and DIO was achieved as described above in Experiment 1.

### *Assessment of Daily Calorie Consumption While Maintaining DIO*

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### *Body Composition and Bone Mass*

Animals were weighed weekly. Area under the curve (AUC) assessment of weight parameters over time, calculated by the trapezoid rule, was employed to better detect



recurring differences in weight gain, as previously employed<sup>74</sup>. At baseline, 3 and 6 months post-OVX, total body composition, as well as bone mineral content (BMC) and bone mineral density (BMD), were assessed by dual-energy X-ray absorptiometry (DXA, iDXA, GE/Lunar Corp., Madison, WI) on sedated animals as described in Experiment One.

#### *Locomotor Activity*

At baseline, 3 and 6 months post-OVX, a small accelerometer (Actiwatch Mini, CamNtech Ltd, Cambridge UK) was added to each female's standard collar. Activity and intensity of movement were recorded over a ~2-4-week period after which the accelerometers were removed. Activity was measured and analyzed as described in Experiment One.

#### *Fasting Glucose and Oral Glucose Tolerance Test*

Fasting glucose and glucoregulation were assessed in over- night fasted, awake animals. Fasting glucose was determined at baseline and 6 months post-OVX, while glucoregulation was assessed by oral glucose tolerance test (OGTT) at 6 months following OVX. Following a baseline blood sample, animals were given an oral dose (5 mL/kg) of 40% sucrose. Blood samples were then collected at 15, 30, 60, and 120 minutes (2 hours) following sucrose administration and assessed for glucose, as previously validated for marmosets<sup>77</sup>. Glucose was measured by glucometer (Accu-Check Aviva, Roche Diagnostics, Indianapolis, IN). AUC glucose values during the OGTT were also assessed to detect between group differences in accumulating high levels of glucose over time.

### *Statistical Analysis*

Data were analyzed utilizing SPSS software (IBM, Armonk, NY). Hormone measures were analyzed with a 1-way analysis of variance (ANOVA), followed by Bonferroni multiple comparison post hoc tests. All other between group analyses were performed using a 2-way ANOVA for repeated measures. Steroid and CG hormone data were log transformed, and behavioral data were transformed to arcsin, prior to ANOVA or correlation tests. Nonparametric Spearman's correlation tests were utilized to identify relationships between hormone values and metabolic parameters. Spearman's rho coefficients are expressed as  $r_s$ . Statistical significance was determined at  $P \leq 0.05$ .

## **2.4 Results**

### *EXPERIMENT 1*

#### *Circulating E<sub>2</sub> and Pituitary GC Levels Confirm Anticipated Estrogen Status*

As expected, E2 females alone exhibited circulating E<sub>2</sub> levels approximating those of mid-cycle, ovary intact female marmosets (Table 2). Greatly diminished circulating levels of E<sub>2</sub> were found in both VEH and LET females, with LET females demonstrating the more extreme E<sub>2</sub> depletion (Table 2). It was not surprising to find elevated circulating chorionic gonadotropin (CG) levels in VEH ( $P < 0.08$ , trend) and LET ( $P < 0.05$ ) compared with E<sub>2</sub> females, reflecting insufficient circulating E<sub>2</sub> for maintenance of negative feedback regulation of pituitary CG release in estradiol-depleted female groups (Table 2). CG levels in LET females, however, were ~35% greater than in VEH females.

#### *Circulating Steroid Hormones and Hypothalamic Steroid Content*

Eliminating ovarian E<sub>2</sub> (VEH) or both ovarian and extra-ovarian E<sub>2</sub> (LET) produced notable changes in hypothalamic estrogen and androgen content, as well as in circulating steroid hormone levels, when compared with females in the E<sub>2</sub> group. Hypothalamic E<sub>2</sub> content was diminished in LET females, alone, when compared with both E<sub>2</sub> (P = 0.005) and VEH (P = 0.013) female groups (Table 4). Notably, comparable hypothalamic E<sub>2</sub> content was maintained between E<sub>2</sub> and VEH groups despite the large systemic deficit in circulating E<sub>2</sub> levels (P = 0.001) exhibited by VEH females that greatly increased the ratio of hypothalamic E<sub>2</sub> (pg/g wet weight): systemic circulation E<sub>2</sub> (pg/mL) from approximately 0.5 (E<sub>2</sub> females) to 31.2 (VEH). Circulating levels of E<sub>2</sub>, however, tended (P = 0.08) to be lower in LET than in VEH females (Table 2), maintaining a high ratio of hypothalamic E<sub>2</sub>: systemic circulation E<sub>2</sub> of approximately 23.3 (LET).

Hypothalamic androgen content, in contrast to that for E<sub>2</sub>, was generally increased in the absence of ovarian E<sub>2</sub> levels (VEH) and was not further augmented by additional depletion of extra-ovarian E<sub>2</sub> (LET) (Table 4). Hypothalamic androstenedione content was elevated in both VEH (P = 0.03) and LET (P = 0.05) compared with E<sub>2</sub> (Table 34) females, resulting in hypothalamic: systemic ratios for androstenedione of 0.2 (E<sub>2</sub>), 1.3 (VEH), and 1.5 (LET). Hypothalamic dehydroepiandrosterone (DHEA) content, however, was only elevated in VEH (P = 0.04), but not LET, females when compared with those in the E<sub>2</sub> group, resulting in hypothalamic: systemic ratios for DHEA of 0.3 (E<sub>2</sub>), 0.1 (VEH) and 0.05 (LET). Despite demonstrating a similar overall pattern, hypothalamic testosterone values remained comparable across all female groups (Table 4) with relative consistency of hypothalamic vs circulating testosterone ratios between female groups of 0.2 (E<sub>2</sub>), 0.1 (VEH), and 0.2 (LET). Changes in circulating

androgen levels (Table 2), however, did not mirror those found in hypothalamic content (Table 4). Circulating levels of DHEA, alone, increased in VEH ( $P < .001$ ) and LET (Table 2;  $P = .001$ ) groups compared with those in E2 females. Circulating levels of androstenedione and testosterone (Table 2) remained comparable across all female groups.

While circulating levels of progestins, progesterone, and 17OHP4, and their ratios, remained comparable across female groups (Table 2), hypothalamic content of 17OHP4 more than doubled in VEH compared with E<sub>2</sub> and LET female groups (Table 4), thus increasing the hypothalamic to circulating 17OHP4 ratio to 10.9 (VEH) in comparison with 2.6 (E<sub>2</sub>) and 6.4 (LET). Hypothalamic progesterone content, and its hypothalamic to circulating ratio, demonstrated more modest increases following estrogen depletion (E<sub>2</sub>, 1.0; VEH, 2.7; LET 2.8).

With regard to ratios of hypothalamic steroid hormones, selected to quantify aspects of androgen biosynthesis (Table 4), the ratio of androstenedione: DHEA was higher in both VEH ( $P = 0.05$ ) and LET ( $P = 0.005$ ) than in E<sub>2</sub> females. There were no effects of eliminating ovarian E<sub>2</sub> (VEH) or both ovarian and extra-ovarian E<sub>2</sub> (LET) on the ratios of hypothalamic testosterone: androstenedione or 17OHP4: progesterone (Table 4), respectively. The ratio of hypothalamic E<sub>2</sub>: testosterone, in contrast, was decreased ( $P = 0.002$ ) in both VEH and LET compared with E<sub>2</sub>, as well as in LET compared with VEH females ( $P = 0.05$ ).

#### *Calorie Consumption, Body Weight, and Body Composition*

Only male–female pairs from the LET female group consumed more calories ( $P = 0.02$ ) per day than male–female pairs from the E<sub>2</sub> female group during weekly

increments of daily calories from months 1 to 5 following OVX (Figure 1). When females were separated from their male pair mates at 6 to 7 months following OVX for individual behavioral and calorie intake assessments, total daily calories provided to all 3 female groups were comparable, with female consuming ~80% of calories provided (Figure 1). Females from both E<sub>2</sub> depleted, VEH (P < 0.001) and LET (P < 0.001) groups, however, consumed more calories, corrected for FFM, than E<sub>2</sub> group females (Figure 1).

While female body weight increased ~5% to 10% during the DIO study regimen irrespective of treatment group (P < 0.016) (Figure 7) the AUC % body weight increases from baseline, with or without correction for FFM (Figure 7) was greater (P = 0.02) in both E<sub>2</sub>-depleted VEH and LET females compared with those in the E<sub>2</sub> group. At both baseline and 6 months following OVX, all female marmosets were obese (>14% body fat), with mean fat-to-lean mass ratios in each treatment group exceeding 0.3 (Table 5). While there were no obvious differences in DXA-determined fat mass 6 months following OVX in any female group for total body or previously validated body regions of interest, including abdomen and hips/thighs, in contrast, total body (P = 0.014), abdominal (P = 0.002), and upper leg (P = 0.025) FFM increased ~5% to 10% during the DIO study regimen irrespective of treatment group (Table 5). Increased abdominal FFM, however, was greater (P = 0.041) in VEH than in both E<sub>2</sub> and LET female groups. FFM increases in all female groups combined were not correlated with either circulating or hypothalamic steroid hormone levels or ratios, including those for E<sub>2</sub>.

Higher hypothalamic, but not circulating, androstenedione values and the androstenedione to DHEA ratio predicted greater AUC % body weight gain, with (androstenedione:  $r^2 = 0.55$ , P = 0.006; androstenedione to DHEA ratio:  $r^2 = 0.44$ , P =

0.019) or without (androstenedione:  $r^2 = 0.53$ ,  $P = .0007$ ; androstenedione to DHEA ratio:  $r^2 = 0.41$ ,  $P = 0.025$ ) correction for FFM, when all female groups were combined. In contrast, circulating but not hypothalamic, androstenedione to DHEA ratio predicted ( $r^2 = 0.45$ ,  $P = 0.011$ ) greater AUC calories consumed corrected for FFM. No correlations were found between measures of body weight or calories consumed and the remaining hypothalamic or circulating steroid hormone values or ratios, including those for estradiol.

Both total body BMC and BMD, and the same bone parameters in previously validated body regions of interest, were comparable across all 3 female groups. No bone parameter was diminished by depletion of either ovarian  $E_2$  or ovarian and extra-ovarian  $E_2$ .

#### *Locomotor Activity*

Activity collar assessments of female locomotion were obtained at both baseline (while pair housed with their male cage mate) and 6 months following OVX (when singly housed) in  $E_2$  depleted groups, alone. AUC locomotor activity was greater during the daytime in VEH than in LET females at both baseline ( $P = 0.001$ ) and 6 months following OVX ( $P = 0.01$ ) (Figure 8A, 8B). In contrast, during the resting hours of nighttime, there were no differences in locomotor activity between female groups. When comparing the AUC % change in locomotory activity, however, from baseline to 6 months following OVX, LET females became slightly more active than VEH during both daytime ( $P = 0.057$ ) and nighttime ( $P = 0.01$ ) (Figure 8C). Increased activity in LET females was notable late in the day and throughout much of the night.

#### *Fasting Glucose*

By 6 months following OVX, in all female groups combined, DIO induced a trend ( $P = 0.055$ ) toward increased fasting glucose levels from baseline (Table 6) irrespective of circulating  $E_2$  levels. OGTT 2-hour glucose, similar to fasting glucose, revealed no between group differences. Glycogenic hepatopathy was observed in 20% (1/5) of  $E_2$  females, 50% (2/4) of VEH females, and 75% (3/4) of LET females during post-necropsy histopathological hepatic assessment.

## *EXPERIMENT 2*

### *Targeted gene knockdown of $ER\alpha$ within VMN and ARC regions of interest (ROIs) of the mediobasal hypothalamus in $ER\alpha$ KD female marmosets*

Quantitative assessment of  $ER\alpha$  immunopositive cells within the mediobasal hypothalamus of OVX adult female marmosets demonstrated clear bilateral knockdown of  $ER\alpha$  protein expression within the targeted VMN and ARC, but not POA or pituitary, in  $ER\alpha$ KD compared to control female marmosets. Intra-hypothalamic infusion of AAV8 comprising shRNA specific for *ESR1* into  $ER\alpha$ KD female marmosets resulted in fewer numbers of cells expressing  $ER\alpha$  within ROIs incorporating the VMN, or portions of it, in comparison to control females ( $p=0.013-0.046$ ; Figure 9). In contrast, ROIs incorporating POA, ARC or portions of ARC, exhibited comparable numbers of  $ER\alpha$  immunopositive cells in  $ER\alpha$ KD compared to control females ( $p=0.069-0.561$ , Figure 9). Intensity of immunopositive  $ER\alpha$  cells, however, was diminished in  $ER\alpha$ KD compared to control females within the entire ARC ( $p=0.049$ ), as well as within its dorsal extent alone ( $p=0.021$ ). Intensity of the immunohistochemical stain for  $ER\alpha$  expression, nevertheless, remained comparable between the two female groups when POA and VMN ROIs were examined (Figure 9).

Not surprisingly, therefore, the product of ER $\alpha$  immunopositive cell density x ER $\alpha$  immunopositive intensity (D x I) was diminished in ER $\alpha$ KD compared to control females within the VMN and dorsal extent of the ARC ( $p=0.013-0.041$ ; Figure 9), but not within the POA. The combined ROI for VMN and ARC exhibited fewer ER $\alpha$  expressing cells ( $p=0.020$ ) and diminished immunopositive cell density x intensity for ER $\alpha$  expression in ER $\alpha$ KD compared to control females.

Positive correlations between ER $\alpha$  immunopositive cell numbers and intensity were identified in ER $\alpha$ KD, but not control females. Numbers of ER $\alpha$  immunopositive cells in ER $\alpha$ KD females positively correlated between VMN and ARC ROIs, alone, whereas intensity of ER $\alpha$  immunopositive expression values yielded positive correlates between VMN and ARC ROIs, as well as between POA and ARC ROIs. There were no between ROI correlations for ER $\alpha$  immunopositive cell density x intensity values. The finding of multiple positive correlations between ER $\alpha$  immunopositive values in VMN and ARC of ER $\alpha$ KD females, alone, highly suggests predictable degrees of ER $\alpha$  knockdown will be quantified in ARC from quantitation of ER $\alpha$  immunopositive cells in the VMN, alone. Thus, following intra-hypothalamic infusion of AAV8 comprising shRNA specific for *ESR1*, neuroanatomical locations for assignment of any change in female marmoset functionality are limited to the combined VMN-ARC and not to either VMN or ARC, separately.

Individual marmoset variation in the rostral-caudal extent from POA to ARC of ER $\alpha$  immunopositive cells within the mediobasal hypothalamus is illustrated by representative examples from a control (cj1) and an ER $\alpha$ KD (cj2) female, (Figures 5), respectively. In considering the control female, ER $\alpha$  immunopositive cells cluster peri-ventricularly within



the rostral POA, assuming a more dorsal and lateral position in the caudal POA. Distribution of VMN ER $\alpha$  immunopositive neurons in the VMN, are more dorsal to those in the ARC, and extend less caudally than those in ARC. Compared to the control female, the mediobasal hypothalamus of the ER $\alpha$ KD female exhibits fewer ER $\alpha$  immunopositive cells, almost all of which comprise 'light' immunostaining. In all female marmosets in which the pituitary remained intact during processing (ER $\alpha$ KD, n=3; C, n=1), the anterior pituitary exhibited a high proportion of 'dark', ER $\alpha$  immunopositive cells and no obvious between female group differences (Figures 5). All control females exhibited a similar pattern of ER $\alpha$  immunopositive cells, with highest immunopositive cell density and intensity in the POA and lowest in VMN.

*Subtle changes in calorie consumption, but not in body weight and body composition, following ER $\alpha$  knockdown in the mediobasal hypothalamus*

Male-female pairs from the ER $\alpha$ KD female group consumed fewer calories ( $p=0.02$ ) than control male-female pairs during weekly increments of daily calories from months 1-7 following OVX (Figure 6A). When females were separated from their male pair mates after seven months following OVX for individual behavioral and calorie intake assessments, total daily calories provided to both female groups were comparable, with singly housed females from both groups consuming ~80% of calories provided (Figure 6B). ER $\alpha$ KD ( $p=0.001$ ) females, however, consumed fewer calories (AUC kcal corrected for fat-free mass,  $1004 \pm 12.50$  kcal/g fat free mass\*56days) in comparison to control females ( $1161 \pm 5.83$  kcal/g fat free mass\* 56days).

In contrast, female body weight, as well as AUC % body weight increased from baseline and AUC % body weight increase from baseline corrected for fat free mass all

demonstrated comparable increments during the study irrespective of female group (Figure 10). There were no obvious changes in DXA-determined body composition between baseline and 3- or 6-months following ER $\alpha$  knockdown, including fat and fat-free mass for total body or previously validated body regions of interest, including abdomen and hips/thighs. There were also no between female group differences in fat mass and fat-free mass percent change between any of the DXA timepoints (Figure 11).

*Diminished locomotor activity following ER $\alpha$  knockdown in the mediobasal hypothalamus*

Activity collar assessments of female locomotion were obtained at both 3 and 6 months post-AAV8 intra-hypothalamic infusion (while pair housed with their male cagemate). AUC locomotor activity was less during the daytime in ER $\alpha$ KD compared to control females at both 3 ( $p=0.001$ ) and 6 months following ER $\alpha$  knockdown ( $p=0.01$ ) (Figure 12). In contrast, during the resting hours of nighttime, there were no differences in locomotor activity between female groups. When comparing the AUC % change in locomotory activity, however, from 3 to 6 months following ER $\alpha$  knockdown, ER $\alpha$ KD females became slightly less active than controls during both daytime ( $p=0.057$ ) and nighttime ( $p=0.01$ ) (Figure 12). Decreased activity in ER $\alpha$ KD females was notable late in the day and throughout much of the night.

*Fasting glucose*

Fasting glucose levels remained comparable across female groups at baseline and at 6 months following ER $\alpha$  knockdown, as was 2h-glucose derived from an oral glucose tolerance test (OGTT) performed at 6 months following ER $\alpha$  knockdown (Figure

13). AUC glucose derived from the 6-month OGTT, however, revealed hyperglycemia in ER $\alpha$ KD females, alone (Figure 13).

## 2.5 Discussion

In Experiment 1, ovarian and extra-ovarian E<sub>2</sub> depletion of adult female marmoset monkeys, in contrast to depletion of ovarian E<sub>2</sub>, alone, demonstrated little functional contribution provided by extra-ovarian E<sub>2</sub> in support of female metabolic homeostasis. Ovarian E<sub>2</sub> depletion, in addition to ovarian and extra-ovarian E<sub>2</sub> depletion, nevertheless, enhanced DIO calorie consumption and weight gain in comparison to E<sub>2</sub> replete females, illustrating the contribution of E<sub>2</sub> toward protecting female marmoset monkeys from DIO-associated metabolic dysfunction. DIO, in contrast and regardless of E<sub>2</sub> depletion, induced modest glucose intolerance, possibly due to impaired hepatic glucose metabolism related to glycogenic hepatopathy observed in all 3 female groups. Prior to Experiment 2, hypothalamic neuronal receptor mechanisms governing E<sub>2</sub> regulation of female NHP metabolic function were previously unknown. The results of Experiment 2, however, provide little evidence that ER $\alpha$  is a key hypothalamic receptor mechanism for E<sub>2</sub> regulation of female metabolic homeostasis in NHPs, and likely women. Experiment 2 was modeled after published gene silencing approaches examining ER $\alpha$ 's role in discrete nuclei of the hypothalamus in female rats<sup>6</sup>. Compared with approaches employed with female rats, we employed MRI-guidance in order to refine individual targeting of the VMN and ARC, as NHPs have notable individual variations in neuroanatomical locations<sup>80,85</sup>. Based on IHC analysis of ER $\alpha$  expression in marmoset hypothalamic mPOA, VMN and ARC, ER $\alpha$  expression was significantly diminished in the VMN of females receiving ER $\alpha$  silencing shRNA. ER $\alpha$  expression in the ARC also appears diminished, but less so in the

mPOA. ARC and mPOA deficits in ER $\alpha$  expression, however, were less consistent than in the VMN. The results Experiments 2 and 2 taken together confirm hypothalamic ER $\alpha$  as a contributor to female marmoset metabolism and that systemic aromatase inhibition does not exaggerate the metabolic dysfunction of ovarian E<sub>2</sub> depletion, alone.

Endogenous estrogenic source(s) beyond the ovaries include a variety of organ systems as well as the brain<sup>7,8,11,12,17,88</sup>. This is of considerable importance to clinical management of women's health. For example, the oral, nonsteroidal aromatase inhibitor letrozole, employed in the current NHP study, is widely used in clinical practice, including minimizing recurrence of estrogen receptor positive breast cancer following surgical intervention<sup>89,90</sup> and enabling menopausal hormone therapy<sup>91,92</sup>. In addition, as found in this NHP study, aromatase inhibition treatment of women with breast cancer enables weight gain (including increased adiposity), perturbing female metabolic homeostasis<sup>93,94</sup>. In the present NHP study, however, DIO rather than aromatase inhibition, increased glucose intolerance. Women with naturally occurring gene variants in CYP19A1, while exceedingly rare, present with varying degrees of systemic estrogen depletion accompanied by overweight or obesity, impaired glucoregulation, borderline hyperlipidemia, and osteopenia/osteoporosis<sup>95</sup>. Unlike the present study, however, genetically determined CYP19A1 deficiency manifests E<sub>2</sub> depletion through all developmental stages, resulting in widespread organ system abnormalities that are absent from our adult-onset E<sub>2</sub> depletion.

As might be expected, systemically administered E<sub>2</sub> alleviates metabolic<sup>96</sup> dysfunction in women, but to varying degrees<sup>97</sup>, while also increasing the risk for harmful side-effects, including cardiovascular disease and cancer<sup>89,97</sup>. SERMs,

nonsteroidal compounds that interact with estrogen receptors, and display distinct differences in degree of agonism vs antagonism action at estrogen receptors in target tissues, show efficacy for osteoporosis and breast cancer<sup>98</sup>. SERMs, however, all carry safety risks, most notably venous thromboembolic events. Treatments that avoid systemic estrogenic activity by delivering bioactive E<sub>2</sub> to the brain, alone, thus have tremendous potential to alleviate female metabolic dysfunction, as indicated by the findings from our current female NHP study. In this regard a synthetically derived, inactive precursor of E<sub>2</sub>, when administered orally to female rodents, is only metabolized to a bioactive estrogen, in this case E<sub>2</sub>, within the central nervous system following transport across the blood–brain barrier<sup>99</sup>, providing amelioration to hypothalamically driven hot flashes without estrogenic effect on systemic estrogen-responsive organs and tissues, such as the uterus.

#### *E<sub>2</sub>, Body Fat, and Glucoregulation*

Attempts have been made in a variety of studies to dissociate effects of normal aging vs declining E<sub>2</sub> levels on adiposity, energy balance, and cardiometabolic health in postmenopausal women<sup>100-105</sup>. In general, these studies support the idea that menopause per se is associated with increasing abdominal obesity and that visceral fat accumulation may, in part, be secondary to an acceleration of aging-related decline in fat oxidation and metabolic energy expenditure<sup>62</sup>. While these changes parallel those observed in OVX rodents<sup>106</sup>, a causal relationship between declining ovarian E<sub>2</sub> in menopause and altered body composition and energy balance has been difficult to confirm<sup>107,108</sup>. Some randomized controlled studies have demonstrated that both oral and transdermal E<sub>2</sub> therapy in postmenopausal women are associated with a reduction

in central adiposity and an increase in lean body mass<sup>109,110</sup>, as well as reductions in insulin resistance and fasting glucose, new-onset type 2 diabetes, blood lipids, blood pressure, adhesion molecules, and procoagulant factors<sup>111</sup>. Of the few studies focusing on energy expenditure during menopausal hormone replacement therapy comprising a variety of estrogenic formulations, some demonstrate increases in lipid oxidation and energy expenditure<sup>112,113</sup>, while others reveal acute decreases in lipid oxidation and energy expenditure<sup>114</sup>. There are similarly conflicting data on the effects of hormone replacement therapy on insulin sensitivity, with some suggesting beneficial effects<sup>111</sup>, while others find no consistent improvement<sup>115-117</sup>.

In the first female marmoset monkey study completed here, mean baseline body weights in each female group (~400 g) were typical for this colony<sup>73,118,119</sup>, while baseline total body fat exceeded 14% of body mass in all females. Body fat in excess of 14% body mass is considered obese for this laboratory NHP<sup>120</sup>, but is typical for this colony<sup>75</sup>. By 9 months following OVX, however, ovarian as well as ovarian and extra-ovarian E<sub>2</sub> depletion resulted in DIO relative weight gain, with and without correction for FFM, as well as increased calorie consumption, in excess of the gains made by E<sub>2</sub>-replete females. E<sub>2</sub>, therefore, diminishes increased calorie consumption and weight gain in female marmosets in the context of DIO, as has previously been reported in mice<sup>20,21,121</sup>. There were, nevertheless, no changes in DXA- determined total, or depot-specific, body fat associated with either E<sub>2</sub> depletion. Such relatively modest fat accumulation in female marmosets may thus occur across a variety of specific depots.

Increased DXA-determined FFM, particularly in the abdomen and upper legs, unexpectedly contributed to combined E<sub>2</sub> depletion and DIO-induced weight gain in

female marmosets. While neither treatment has previously been associated with gain in FFM, E<sub>2</sub> depletion in female marmosets also induced hyperandrogenism. Induction of hyperandrogenism in female NHPs<sup>122</sup> and female-to-male transition<sup>123</sup> is associated with increased FFM, particularly in upper legs<sup>123</sup>, and with androgen-receptor activity in NHPs<sup>122</sup>, but in the present study no correlations between circulating or hypothalamic androgen concentrations or androgenic hormone ratios were associated with increases in FFM.

The actions of E<sub>2</sub> via ER $\alpha$  on adiposity may occur directly in white adipose tissue, liver, muscle, and/or pancreas<sup>34,38</sup>, as well as in hypothalamic neurons expressing ER $\alpha$ <sup>59,56</sup>. The latter exert descending control over systemic organ systems via autonomic innervation<sup>124-126</sup>, including E<sub>2</sub>-induced alterations in food intake and energy expenditure, producing secondary metabolic states, or by a combination of these. Stimulatory effects of E<sub>2</sub> on energy expenditure are transduced in ER $\alpha$  expressing neurons of the VMN of the hypothalamus<sup>56</sup> by nonclassical ER $\alpha$  signaling<sup>59</sup> coupled to activation of PI3-kinase<sup>9</sup>. E<sub>2</sub> also regulates gene expression associated with regulation of food intake and energy expenditure in the hypothalamus, largely through ER $\alpha$  activation<sup>21</sup>. Furthermore, a study by Musatov et al<sup>56</sup> demonstrated that viral vector-mediated ER $\alpha$  gene silencing in the VMN of both female mice and rats largely recapitulates a metabolic phenotype observed in whole-body ER $\alpha$ KO mice, including obesity, hyperphagia, impaired glucose tolerance, and reduced energy expenditure<sup>22-24</sup>. In the current female NHP study, however, and in contrast to female rodents, we found no evidence of impaired glucoregulation at 6 months following E<sub>2</sub> depletion. We nevertheless identified a trend toward DIO-associated glucose intolerance in all female

groups combined, likely the result of insufficient increase in compensatory pancreatic beta cell insulin release to accommodate DIO-induced systemic insulin resistance. Such DIO-associated impairments of glucoregulatory function in the present study are reminiscent of glucoregulatory dysfunction reported in a previous female marmoset study employing a glucose-enriched DIO<sup>127</sup>. Therefore, while E<sub>2</sub> ameliorated DIO increased energy intake and relative weight gain, E<sub>2</sub> may have failed to ameliorate pancreatic beta cell insulin decompensation due to weight gain–mediated systemic insulin resistance<sup>127</sup>. It is unclear whether the 50% to 75% incidence of glycogenic hepatopathy (excessive accumulation of glycogen in hepatocytes) found in E<sub>2</sub>-depleted female marmosets in the present study represents an increase above a previously reported incidence of 34% in laboratory housed female marmosets<sup>128</sup>. Glycogenic hepatopathy can indicate chronic recurrence of hyperglycemic episodes<sup>129</sup>.

Since E<sub>2</sub> depletion enables maturation of the female marmoset's androgenic zona reticularis in the adrenal cortex<sup>130</sup>, and aromatase inhibition commonly results in androgenic precursor excess<sup>19,131</sup>, the positive correlations between hypothalamic androstenedione and androstenedione to DHEA ratio with weight gain may simply represent biomarkers for the degree and duration of E<sub>2</sub> depletion rather than androgenic effects per se. Long-term testosterone treatment, however, increases body weight in ovary intact pre-pubertal<sup>29</sup> and adolescent–young adult<sup>30</sup> as well as ovariectomized adult<sup>122</sup> female rhesus macaques, without glucoregulatory impairment. Furthermore, addition of DIO to testosterone treatment of adolescent female rhesus macaques exaggerates weight gain and induces glucoregulatory impairments accompanying insulin resistance<sup>30</sup>. The absence of glucoregulatory impairments in our E<sub>2</sub>-depleted and



hyperandrogenic OVX female marmosets would be consistent with the requirement for E<sub>2</sub> activity, likely in the liver, to complete androgen-mediated glucoregulatory dysfunction, as found in organ-selective androgen receptor knockout female mouse models<sup>31</sup>.

#### *Skeletal Bone Mass Maintained Independently of E<sub>2</sub> in Female Marmosets*

In almost all female mammals with regular, frequent ovarian cycles, a reduction in circulating estrogen concentrations, either spontaneous or experimentally induced, leads to a reduction in bone mass. This has been demonstrated in numerous primate species and occurs in as little as 3 months in rhesus macaques<sup>32,60,134-136</sup>. Therefore, it is surprising that we found no deficit in bone mass or bone density of either the total body or lumbar spine associated with systemic or systemic and hypothalamic E<sub>2</sub> deficiency, though, importantly, estrogen and aromatase activity within the bone microenvironment<sup>4</sup> were not assessed. There are several potential reasons for this finding.

Evidence for estrogen control of bone mass in common marmosets is controversial. In several experiments, we were unable to find any evidence of bone loss following estrogen depletion due to either social subordination or OVX in marmosets<sup>118</sup>. Similar to the conclusions of Kraynak and colleagues<sup>75</sup> that primates evolved metabolic control systems regulated by extra-ovarian E<sub>2</sub> or that are generally less subject to E<sub>2</sub> regulation, a conclusion arising from this study was that brain-derived estrogen may be enough to maintain bone mass even in the context of circulating E<sub>2</sub> deficiency. There have been some limited reports of bone degeneration in marmosets. For example, Seidlová-Wuttke and colleagues<sup>137</sup> found some evidence of bone loss in

orchidectomized male marmosets though the level of bone loss only reached WHO criteria for osteoporosis in 2 older animals (10 and 11 years of age). It is challenging, however, to separate potential hormonal causes from overall gastrointestinal health issues, nutritional deficits, and vitamin D malabsorption<sup>138,139</sup>. While marmosets may represent an intriguing model of estrogen deficiency bone loss, some caution must be taken particularly given their high circulating vitamin D levels and associated end organ resistance related to the overexpression of vitamin D response element binding proteins<sup>140,141</sup>. In addition, such end organ resistance may be evidence of a broader resistance to select actions of steroid hormones. Bone in marmosets may therefore have evolved to be less subject to estrogen regulation.

#### *Applications to Understanding Metabolic Dysfunction in Women*

Our experiments are the first to investigate the metabolic consequence of knocking down hypothalamic ER $\alpha$  expression in a nonhuman primate model and subsequently, investigate the main source of E<sub>2</sub> synthesis in regulating female metabolism via ER $\alpha$ . Our investigations provided evidence that in adult female marmoset monkeys, mediobasal hypothalamic ER $\alpha$  is an important receptor in maintaining female energy balance and extra-ovarian E<sub>2</sub> is not crucial to maintain ER $\alpha$  metabolic signaling cascades. It is important to note that marmoset monkeys are not the most similar NHP research model to humans, in contrast to say the rhesus monkey, a more physiological, neuroendocrinological, and genetically similar NHP to humans. To further determine how ER $\alpha$  dictates energy balance via ovarian and extra-ovarian sources of E<sub>2</sub>, it would be of great benefit to perform hypothalamic ER $\alpha$  knockdown and use aromatase inhibition in adult female rhesus monkeys to elucidate possible neuro-E<sub>2</sub> mechanisms

likely relevant in human metabolic function. Identifying the predominate estrogen receptor and tissue from which E<sub>2</sub> is synthesized, potentially, the brain, and their roles in regulating female metabolism is critical understanding to create therapeutic targets that discretely act on a site while reducing peripheral side effects.

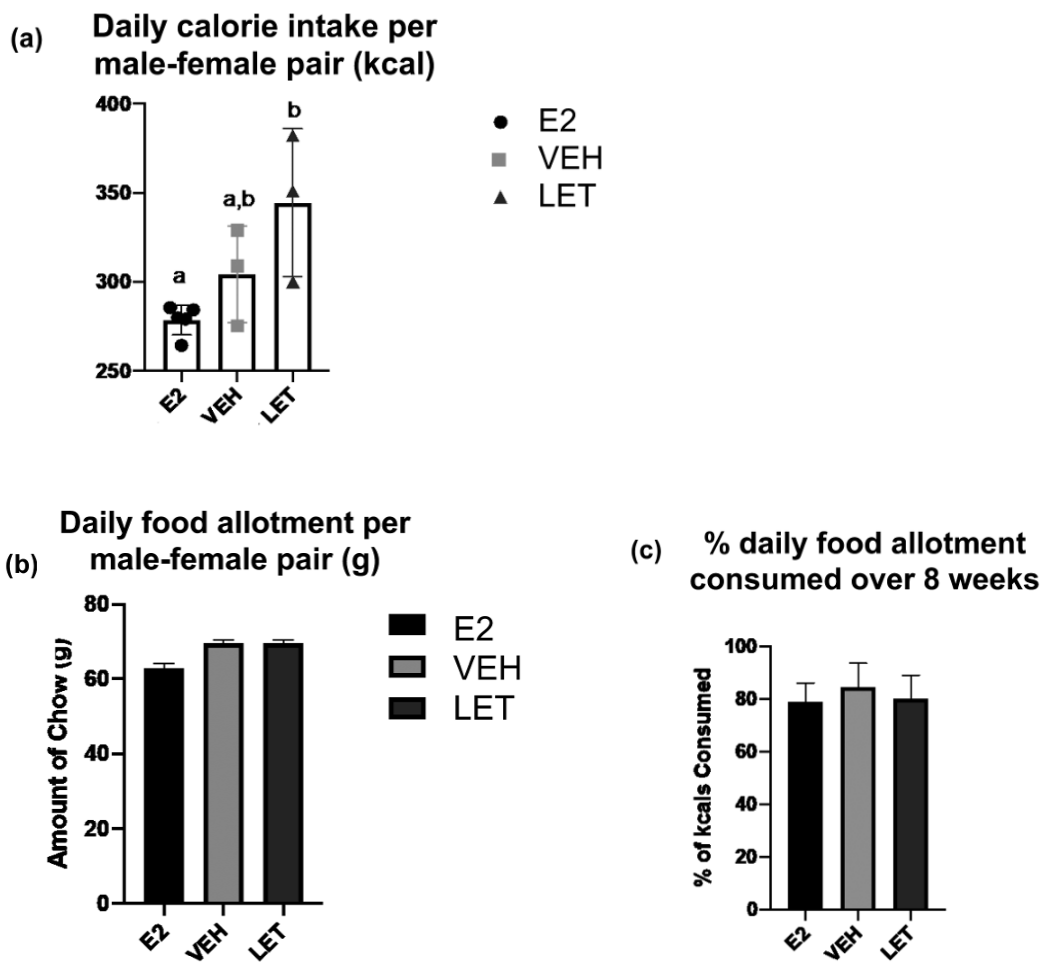
## 2.6 Tables & Figures

Parameter	E2 (n = 5)	VEH (n = 4)	LET (n = 4)	P value
Age at baseline (years)	3.1 ± 0.5	3.2 ± 0.3	2.9 ± 0.1	.854
Body weight at baseline (g)	402 ± 26	396 ± 35	382 ± 12	.863

**Table 1:** Age, body weight and uterine characteristics (mean ± SEM) of E2, VEH, and LET ovariectomized adult female marmoset monkeys.

Circulating hormone	E2 (n = 5)	VEH (n = 4)	LET (n = 4)	P value
Estradiol (pg/mL)	1174.8 ± 214.0	24.7 ± 11.9 <sup>a</sup>	3.0 ± 0.1 <sup>a,b</sup>	.001
Estrone (pg/mL)	ND	ND	ND	ND
Progesterone (ng/mL)	1.32 ± 0.40	0.94 ± 0.33	0.47 ± 0.30	.118
17-OHP <sub>4</sub> (ng/mL)	3.37 ± 1.89	2.19 ± 0.36	2.00 ± 0.36	.743
DHEA (ng/mL)	0.15 ± 0.02	1.93 ± 0.76 <sup>a</sup>	1.80 ± 0.79 <sup>a</sup>	.001
Androstenedione (ng/mL)	3.22 ± 0.37	7.16 ± 3.49	3.23 ± 2.13	.468
Testosterone (ng/mL)	0.18 ± 0.03	1.11 ± 0.62	0.42 ± 0.24	.518
CG (ng/mL)	2.70 ± 0.40	6.31 ± 2.22 <sup>c</sup>	8.50 ± 1.52 <sup>d</sup>	.014
<b>Hormone ratio</b>				
Estradiol:Testosterone	8.3 ± 1.9	0.3 ± 0.3 <sup>a</sup>	0.02 ± 0.01 <sup>a</sup>	.001
17OHP <sub>4</sub> :Progesterone	3.3 ± 1.8	2.8 ± 1.5	6.0 ± 2.1	.482
Androstenedione:DHEA	24.2 ± 4.6	4.1 ± 2.4 <sup>a</sup>	5.2 ± 2.0 <sup>d</sup>	.004
Testosterone:Androstenedione	0.06 ± 0.01	3.87 ± 2.15	3.32 ± 2.19	.225

**Table 2:** Circulating levels (mean± SEM) of sex steroid and gonadotropic hormones in ovariectomized female marmosets receiving E<sub>2</sub> replacement (E2), empty capsules (VEH) or the aromatase inhibitor, letrozole (LET) at 6 months following OVX at 09:00 hours, immediately prior to necropsy.

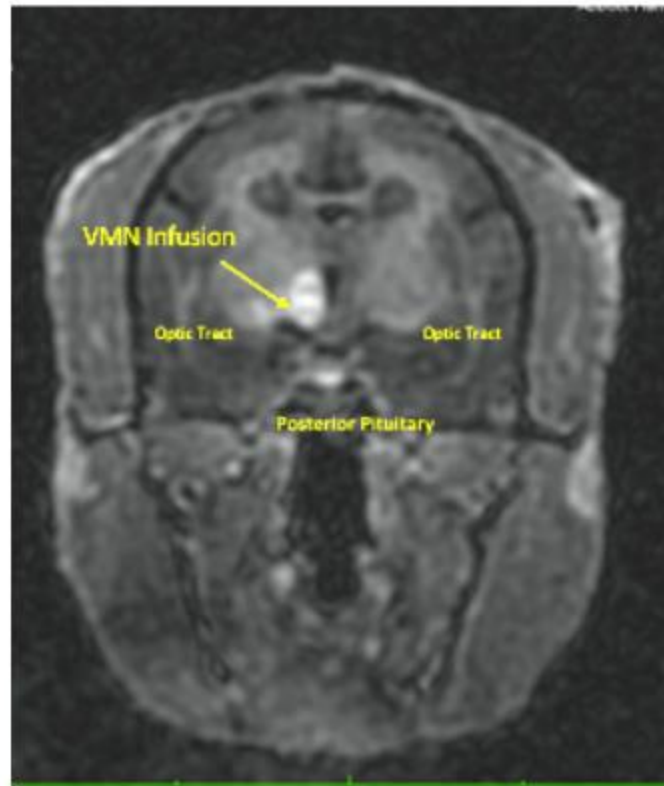


**Figure 1.** (A) Calories consumed (mean  $\pm$  SEM) by both males and females combined in the male–female pairs (symbols illustrate individual male–female pair consumption) comprising the treatment groups E2 (black circles), VEH (light gray squares), and LET (dark gray triangles) from ovariectomy (OVX) until 5 months (months) after OVX, (B) amount of chow (mean  $\pm$  SEM wet weight) provided to singly housed females during months 6 to 7 following OVX (E2 black bar, VEH light gray bar, LET dark gray bar), and (C) % calories consumed (mean  $\pm$  SEM) by singly housed females during months 6 to 7 following OVX (E2 black bar, VEH light gray bar, LET dark gray bar).

	Age (years)	Body Weight (g)
Scrambled Controls	3.44 ± 0.55	440.5 ± 24.3
ER $\alpha$ Gene Silenced	3.32 ± 0.31	412.4 ± 32.9

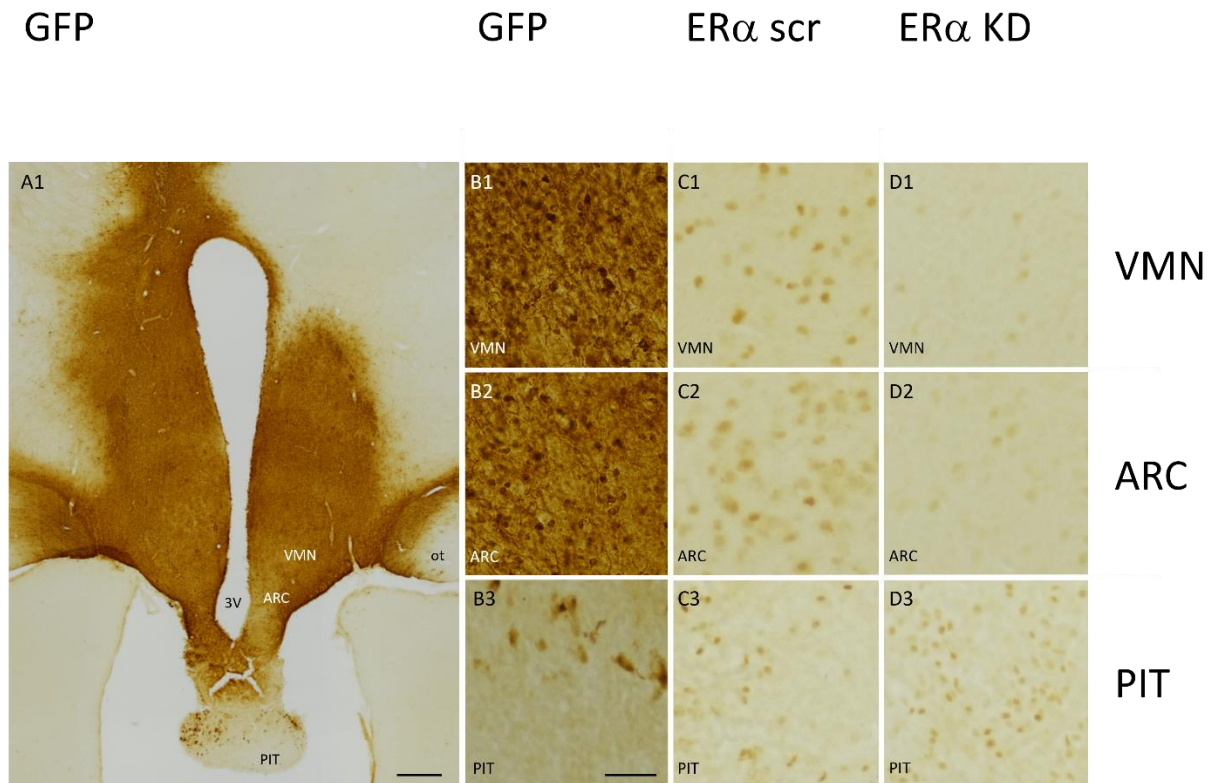
**Table 3:** Comparable ages and body weights among all female marmoset groups

(Scrambled controls, n=4; ER $\alpha$ KD, n=4) at study baseline.

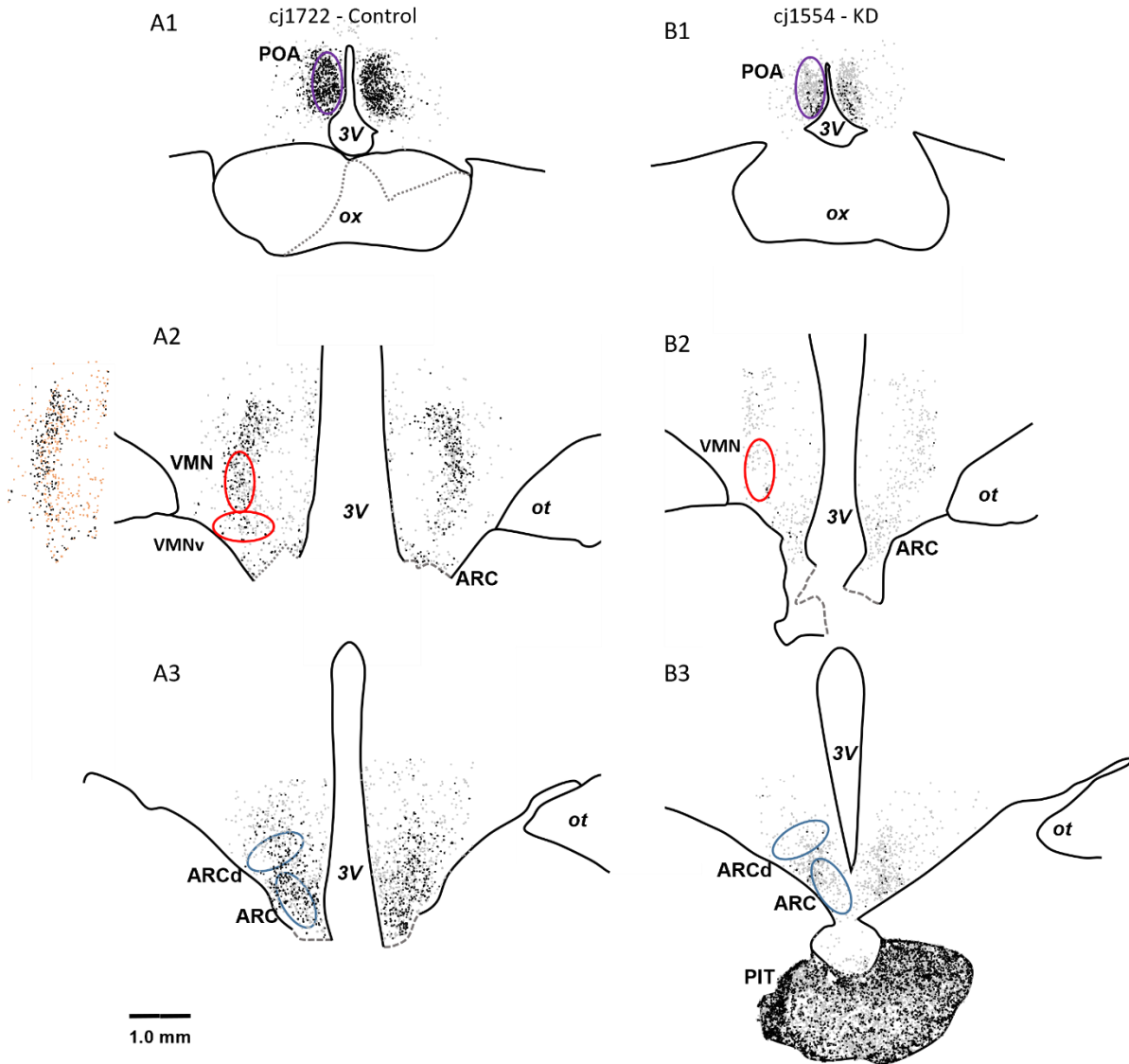


**Figure 2:** Use of MRI and Multihance, a contrast agent, allows successful targeting of the neural infusion of the viral vector encapsulated shRNA to the VMN and ARC of the hypothalamus. MRI images are taken prior to surgery to generate approximate targeting coordinates and following infusions (shown here) to confirm accurate targeting of the VMN.



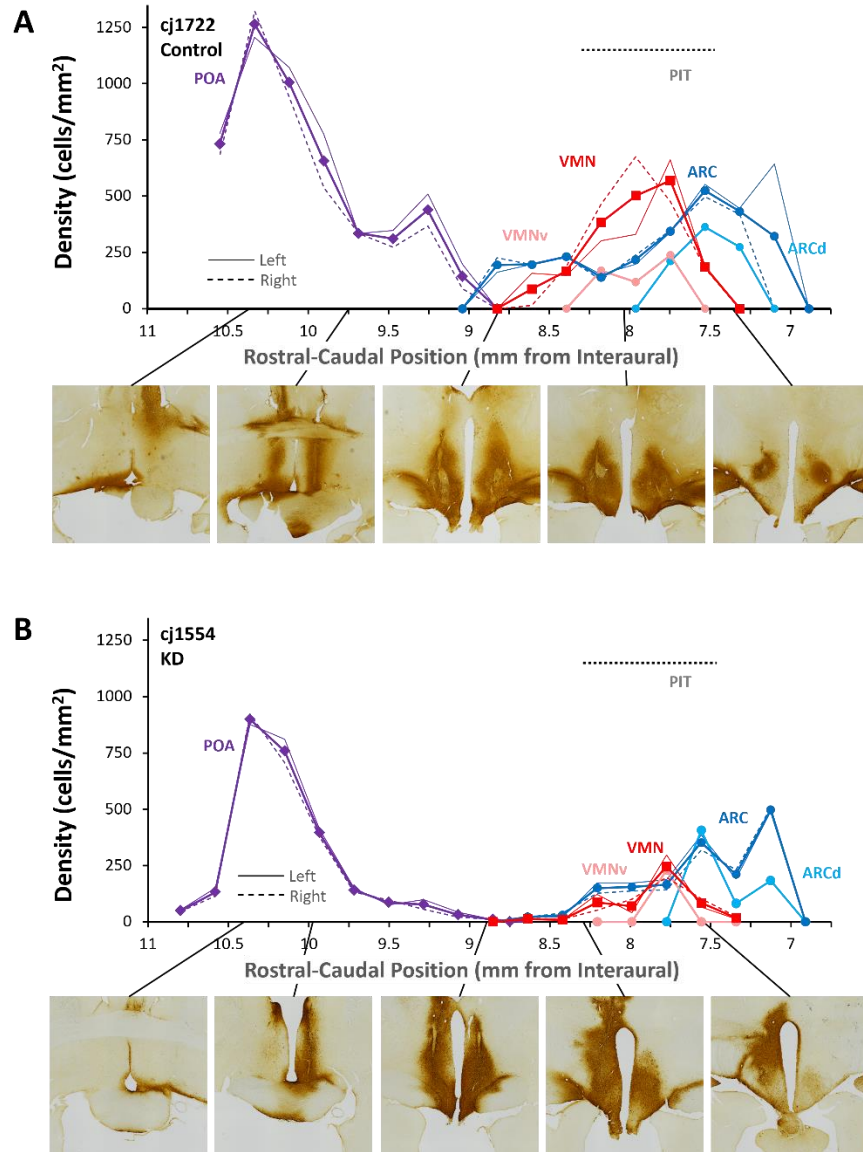


**Figure 3:** ER $\alpha$  gene silencing shRNA effectively ablates ER $\alpha$  in the VMN and ARC of female marmoset monkeys. 40x image of GFP in left most panel (A); panel (B) demonstrates expression of GFP (a/b: MAB3580) in an animal receiving the ER $\alpha$  gene silencing viral vector in the VMN, ARC and PIT, demonstrating infection of cells throughout the VMN and ARC, while the panel (C) illustrates 6F11 staining in a scrambled control animal and panel (D) illustrates an absence of ER $\alpha$  immunoreactivity in the VMN and ARC, with robust staining in the pituitary gland (D3).



**Figure 4:** Distribution of ER $\alpha$  labeled cells in the hypothalamus. Representative sections are shown from the rostral (top) to caudal (bottom) hypothalamus for an ER $\alpha$ KD (right) and scrambled control (left) monkey. Black dots indicate the location of darkly stained cells. Dotted lines indicate damaged tissue. Black dots indicate the location of darkly stained cells (upper half of the ER $\alpha$  staining intensity range for that animal). Gray dots indicate the location of the lightly stained cells (lower half of the

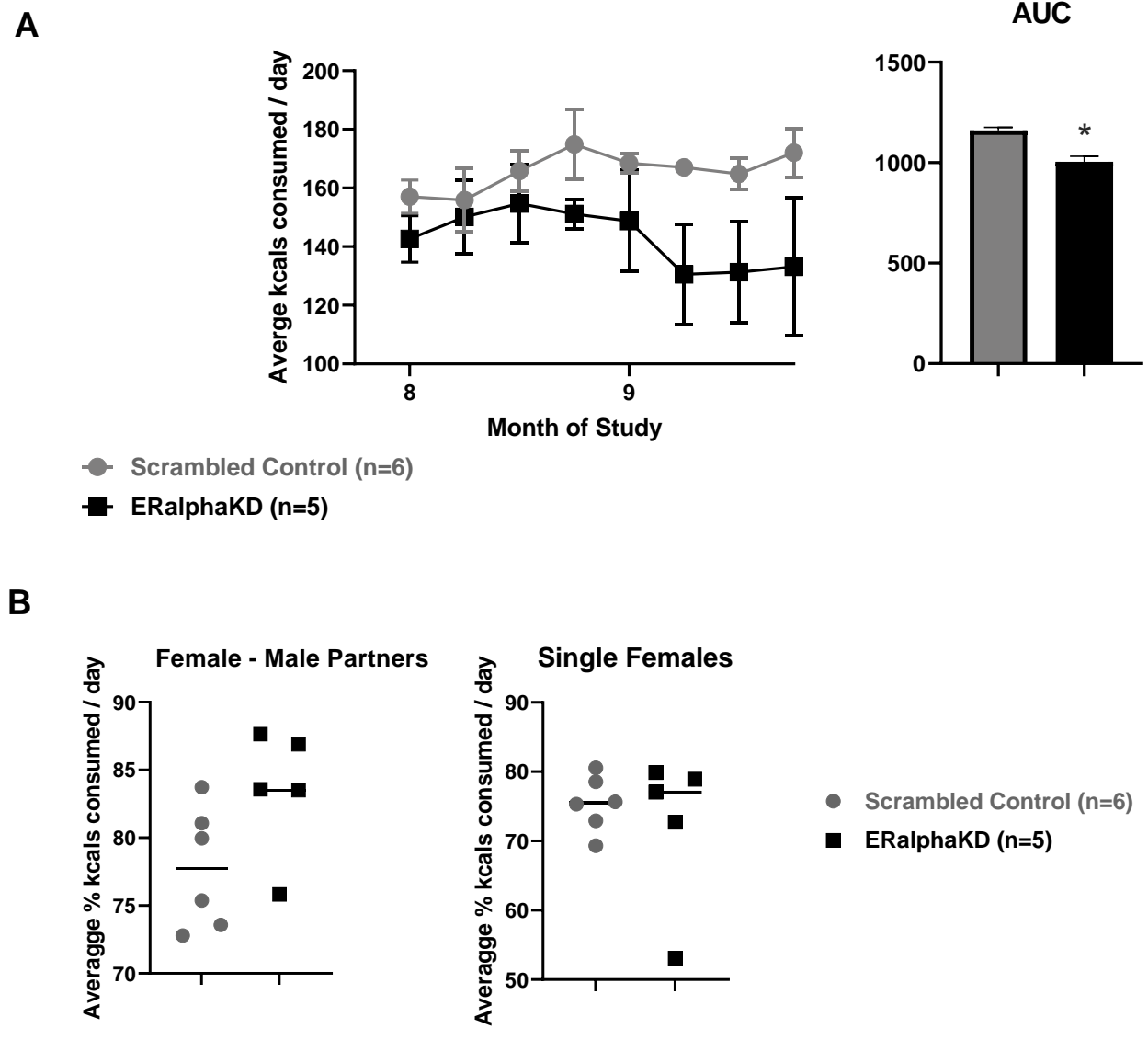
intensity range). Abbreviations: ox: optic chiasm; ac: anterior commissure; opt: optic tract; 3V: third ventricle



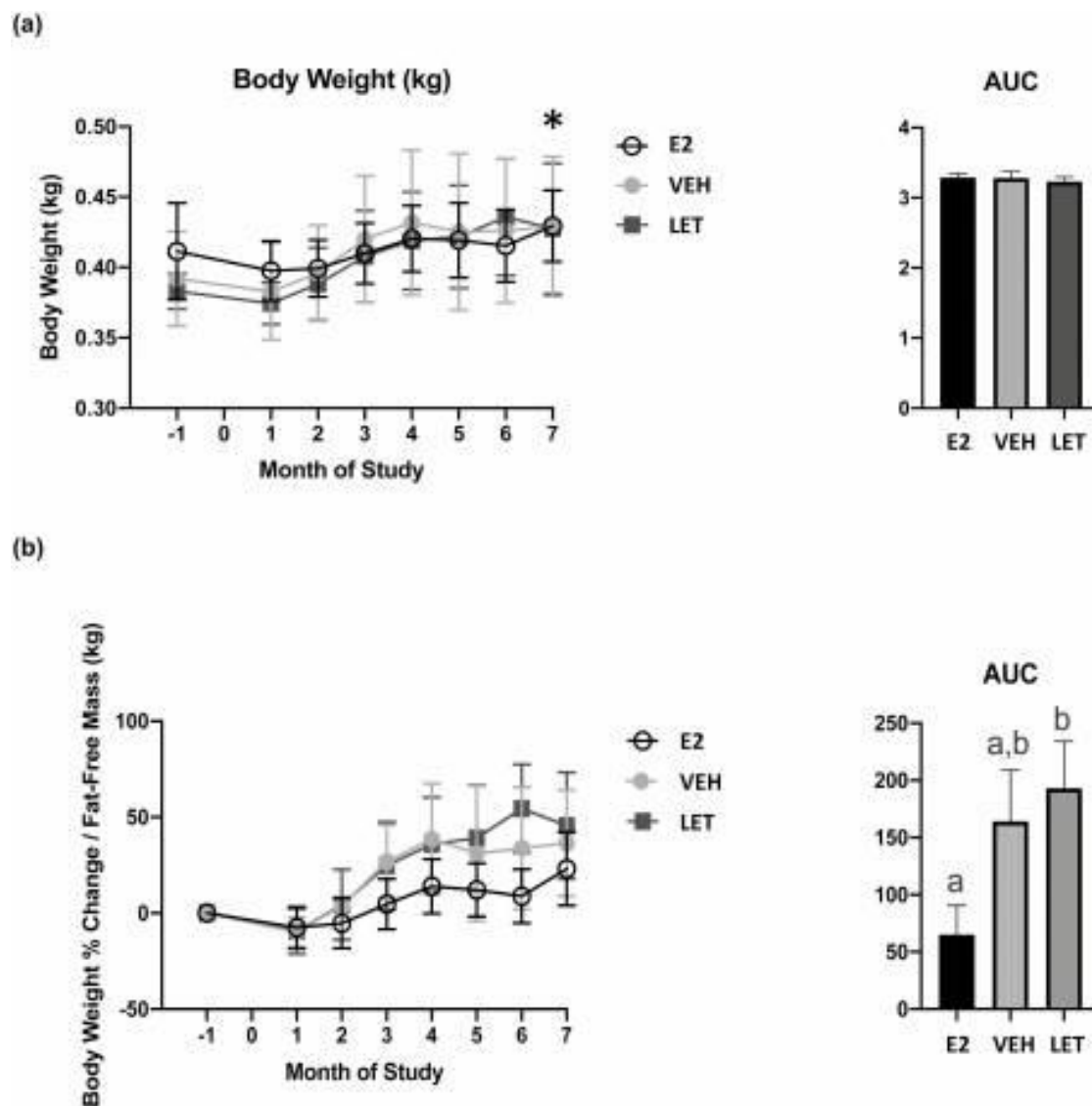
**Figure 5:** ER $\alpha$  positive cells were quantified in coronal brain sections from the most rostral extent of quantification (mid-optic chiasm) through the most caudal extent (caudal sections of the ARC including portions of the mammillary bodies). Thick, solid, purple lines denote numbers of ER $\alpha$  positive cells in the POA, red lines=VMN, and blue lines=ARC, and denote the summation of the right and left side of each nuclei. Thin lines represent the left (solid) and the right (dotted) cell counts at each location. Each plot represents one female (Scrambled control (A), ER $\alpha$  knockdown (B)).

Hypothalamic hormone	E2 (n = 5)	VEH (n = 4)	LET (n = 4)	P value
Estradiol (pg/mg)	0.56 ± 0.16	0.77 ± 0.33	0.07 ± 0.04 <sup>a,b</sup>	.004
Estrone (pg/mg)	0.26 ± 0.10	0.32 ± 0.15	0.10 ± 0.04	.154
Progesterone (pg/mg)	1.30 ± 0.25	2.55 ± 0.76	1.32 ± 0.40	.266
17-OHP <sub>4</sub> (pg/mg)	8.80 ± 1.46	23.83 ± 5.17 <sup>a</sup>	12.75 ± 2.11	.042
DHEA (pg/mg)	0.04 ± 0.01	0.20 ± 0.06 <sup>a</sup>	0.09 ± 0.02	.010
Androstenedione (pg/mg)	0.75 ± 0.04	9.34 ± 5.32 <sup>a</sup>	4.88 ± 1.75 <sup>c</sup>	.008
Testosterone (pg/mg)	0.04 ± 0.01	0.15 ± 0.03 <sup>a</sup>	0.08 ± 0.03	.019
<b>Hypothalamic hormone ratio</b>				
Estradiol:Testosterone	127.4 ± 63.2	62.9 ± 0.3 <sup>a</sup>	0.02 ± 0.01 <sup>a,b</sup>	.002
17OHP <sub>4</sub> :Progesterone	6.7 ± 0.8	9.8 ± 1.9	10.9 ± 2.0	.215
Androstenedione:DHEA	17.7 ± 1.2	38.9 ± 11.3	53.3 ± 9.9 <sup>a</sup>	.009
Testosterone:Androstenedione	0.06 ± 0.01	0.03 ± 0.01 <sup>a</sup>	0.02 ± 0.01 <sup>a</sup>	.003

**Table 4:** Hypothalamic levels (mean ± SEM) of sex steroid hormones in OVX female marmosets receiving E<sub>2</sub> replacement (E2), empty capsules (VEH), or the aromatase inhibitor letrozole (LET) at 6 months following OVX.



**Figure 6.** Average calories consumed while females singly housed (A) with corresponding AUC. Average percent of kcal consumed in female-male pairs and singly housed females (B). Plots expressed as mean  $\pm$  SEM. Significance assessed via RM-ANOVA and student's t-test.  $P^* < 0.05$ .



**Figure 7.** (A) Monthly body weights (mean  $\pm$  SEM) of adult female marmoset monkeys from baseline until 7 months following ovariectomy, and (B) AUC body weights (mean  $\pm$  SEM) incorporating all increments in body weight across the entire study, in E2 (open circles and black bars), VEH (light gray circles and bars), and LET (dark gray squares and bars) female groups.

Body region of interest	DXA parameter	Treatment group	Baseline	6 months	Change from baseline
Total body	Fat mass (g)	E2	121 ± 17	106 ± 19	-15 ± 18
		VEH	106 ± 22	156 ± 37	38 ± 33
		LET	113 ± 23	126 ± 35	5 ± 45
	Fat free mass (FFM) <sup>a</sup> (g)	E2	254 ± 15	270 ± 12	16 ± 7
		VEH	231 ± 11	251 ± 14	5 ± 11
		LET	234 ± 16	249 ± 7	22 ± 5
	Fat/FFM ratio	E2	0.5 ± 0.1	0.4 ± 0.1	-0.1 ± 0.1
		VEH	0.5 ± 0.1	0.6 ± 0.1	0.1 ± 0.1
		LET	0.5 ± 0.1	0.5 ± 0.1	-0.1 ± 0.1
	% fat	E2	31.6 ± 2.0	27.6 ± 3.8	-4.1 ± 3.2
		VEH	30.9 ± 4.2	36.6 ± 5.6	5.3 ± 6.3
		LET	32.3 ± 5.8	31.9 ± 6.3	-3.4 ± 7.1
% FFM	E2	68.4 ± 2.0	72.4 ± 3.8	4.1 ± 3.2	
	VEH	69.1 ± 4.2	63.4 ± 5.6	-5.3 ± 6.3	
	LET	67.7 ± 5.8	68.1 ± 6.3	3.4 ± 7.1	
Abdomen	Fat mass (g)	E2	27 ± 5	22 ± 6	-5 ± 7
		VEH	24 ± 8	43 ± 15	15 ± 14
		LET	23 ± 6	31 ± 13	19 ± 29
	FFM <sup>b</sup> (g)	E2	80 ± 9	85 ± 8	5 ± 3
		VEH	71 ± 8	92 ± 8	16 ± 3 <sup>c</sup>
		LET	73 ± 5	81 ± 6	9 ± 5
	Fat/FFM ratio	E2	0.3 ± 0.1	0.3 ± 0.1	-0.1 ± 0.1
		VEH	0.3 ± 0.1	0.4 ± 0.2	0.1 ± 0.2
		LET	0.3 ± 0.1	0.4 ± 0.1	0.1 ± 0.2
	% Fat	E2	24 ± 2	20 ± 4	-5 ± 3
		VEH	24 ± 5	29 ± 7	4 ± 8
		LET	23 ± 5	24 ± 7	-3 ± 6
% FFM	E2	76 ± 2	80 ± 4	5 ± 3	
	VEH	76 ± 5	71 ± 7	-4 ± 1	
	LET	77 ± 5	76 ± 7	2 ± 6	
Chest	Fat (g)	E2	37 ± 9	34 ± 7	-4 ± 6
		VEH	32 ± 11	48 ± 14	12 ± 10
		LET	35 ± 10	42 ± 14	1 ± 12

**Table 5A:** Mean (+/- SEM) regional body composition parameters as determined by dual x-ray absorptiometry (DXA) in E2, VEH and LET female groups of marmosets at baseline and after six months of treatment (continued in Figure B).

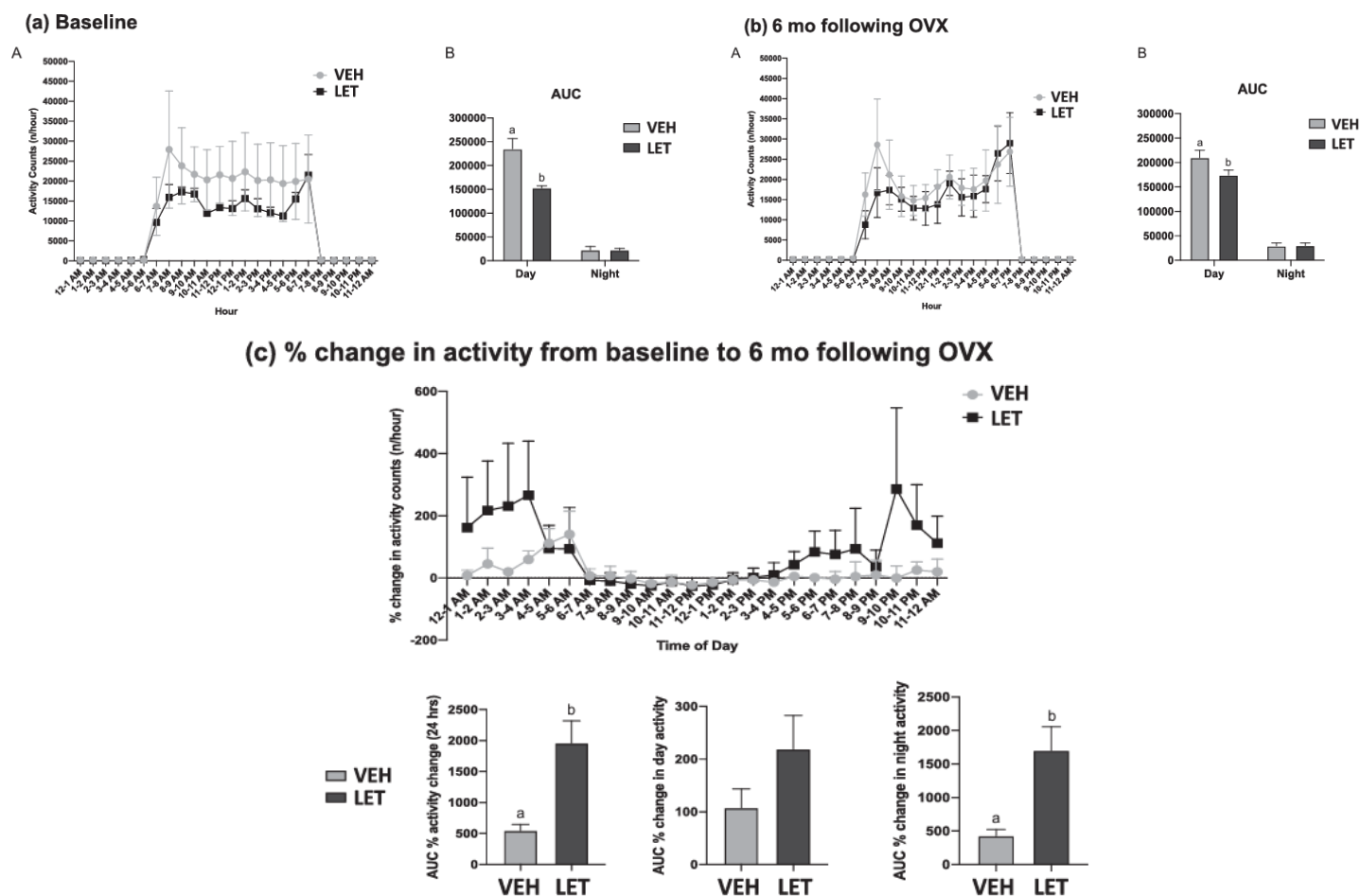


Body region of interest	DXA parameter	Treatment group	Baseline	6 months	Change from baseline
Chest (contd.)	FFM <sup>c</sup> (g)	E2	57 ± 4	61 ± 4	4 ± 2
		VEH	50 ± 4	44 ± 5	-8 ± 4
		LET	52 ± 8	52 ± 8	5 ± 7
	Fat/FFM ratio	E2	0.7 ± 0.1	0.6 ± 0.1	-0.1 ± 0.1
		VEH	0.7 ± 0.3	1.2 ± 0.4	0.5 ± 0.4
		LET	0.8 ± 0.3	1.0 ± 0.4	0.1 ± 0.5
	% Fat	E2	38 ± 5	35 ± 6	-3 ± 4
		VEH	37 ± 10	48 ± 10	10 ± 7
		LET	40 ± 11	43 ± 12	-4 ± 9
	% FFM	E2	63 ± 5	65 ± 6	3 ± 4
		VEH	63 ± 10	52 ± 11	-10 ± 7
		LET	60 ± 11	57 ± 12	4 ± 9
Upper legs	Fat mass (g)	E2	14 ± 1	12 ± 2	-1 ± 2
		VEH	12 ± 1	16 ± 2	3 ± 3
		LET	13 ± 2	11 ± 3	-1 ± 6
	FFM <sup>d</sup> (g)	E2	48 ± 3	50 ± 2	2 ± 1
		VEH	42 ± 3	48 ± 3	2 ± 3
		LET	44 ± 3	45 ± 2	5 ± 4
	Fat/FFM ratio	E2	0.3 ± 0.1	0.2 ± 0.1	-0.1 ± 0.1
		VEH	0.3 ± 0.1	0.3 ± 0.1	0.1 ± 0.1
		LET	0.3 ± 0.1	0.2 ± 0.1	-0.1 ± 0.1
	% Fat	E2	22 ± 1	19 ± 3	-3 ± 3
		VEH	21 ± 2	24 ± 2	2 ± 5
		LET	23 ± 4	19 ± 3	-5 ± 7
% FFM	E2	78 ± 1	81 ± 3	3 ± 3	
	VEH	79 ± 2	76 ± 2	-2 ± 5	
	LET	77 ± 4	81 ± 3	5 ± 7	
Lower legs	Fat mass (g)	E2	13 ± 1	12 ± 2	-2 ± 1
		VEH	12 ± 1	14 ± 2	-1 ± 2
		LET	12 ± 1	11 ± 2	-1 ± 1
	FFM (g)	E2	20 ± 1	23 ± 2	3 ± 2
		VEH	21 ± 2	19 ± 1	-2 ± 2
		LET	18 ± 1	22 ± 1	4 ± 3
	Fat/FFM ratio	E2	0.7 ± 0.1	0.5 ± 0.1	-0.1 ± 0.1
		VEH	0.6 ± 0.1	0.7 ± 0.1	0.1 ± 0.2
		LET	0.7 ± 0.1	0.5 ± 0.1	-0.2 ± 0.1

**Table 5B:** Mean (+/- SEM) regional body composition parameters as determined by dual x-ray absorptiometry (DXA) in E2, VEH and LET female groups of marmosets at baseline and after six months of treatment (Continued in Figure 5C).

Body region of interest	DXA parameter	Treatment group	Baseline	6 months	Change from baseline
Lower legs (contd.)	% Fat	E2	41 ± 2	34 ± 4	-6 ± 4
		VEH	37 ± 1	41 ± 4	1 ± 4
		LET	40 ± 4	32 ± 3	-7 ± 1
	% FFM	E2	59 ± 2	66 ± 4	6 ± 4
		VEH	63 ± 1	59 ± 4	-1 ± 4
		LET	60 ± 4	68 ± 3	7 ± 1
Arms	Fat mass (g)	E2	14 ± 2	12 ± 1	-2 ± 1
		VEH	12 ± 1	14 ± 2	2 ± 3
		LET	13 ± 2	13 ± 2	0 ± 3
	FFM (g)	E2	27 ± 1	28 ± 1	1 ± 1
		VEH	26 ± 2	27 ± 2	0 ± 2
		LET	26 ± 2	26 ± 2	0 ± 2
	Fat/FFM ratio	E2	0.5 ± 0.1	0.4 ± 0.1	-0.1 ± 1
		VEH	0.5 ± 0.1	0.6 ± 0.1	0.1 ± 0.2
		LET	0.5 ± 0.1	0.5 ± 0.1	0 ± 0.2
	% Fat	E2	34 ± 2	30 ± 3	-2 ± 1
		VEH	31 ± 2	35 ± 4	2 ± 3
		LET	33 ± 5	33 ± 3	0 ± 3
	% FFM	E2	66 ± 2	70 ± 3	4 ± 3
		VEH	68 ± 2	65 ± 4	-4 ± 7
		LET	67 ± 5	67 ± 3	1 ± 6
Abdomen/Upper legs ratio	Fat	E2	1.9 ± 0.2	1.7 ± 0.1	-0.2 ± 0.2
		VEH	1.5 ± 0.3	2.2 ± 0.6	0.6 ± 0.5
		LET	2.2 ± 0.6	2.4 ± 0.5	0.1 ± 0.1

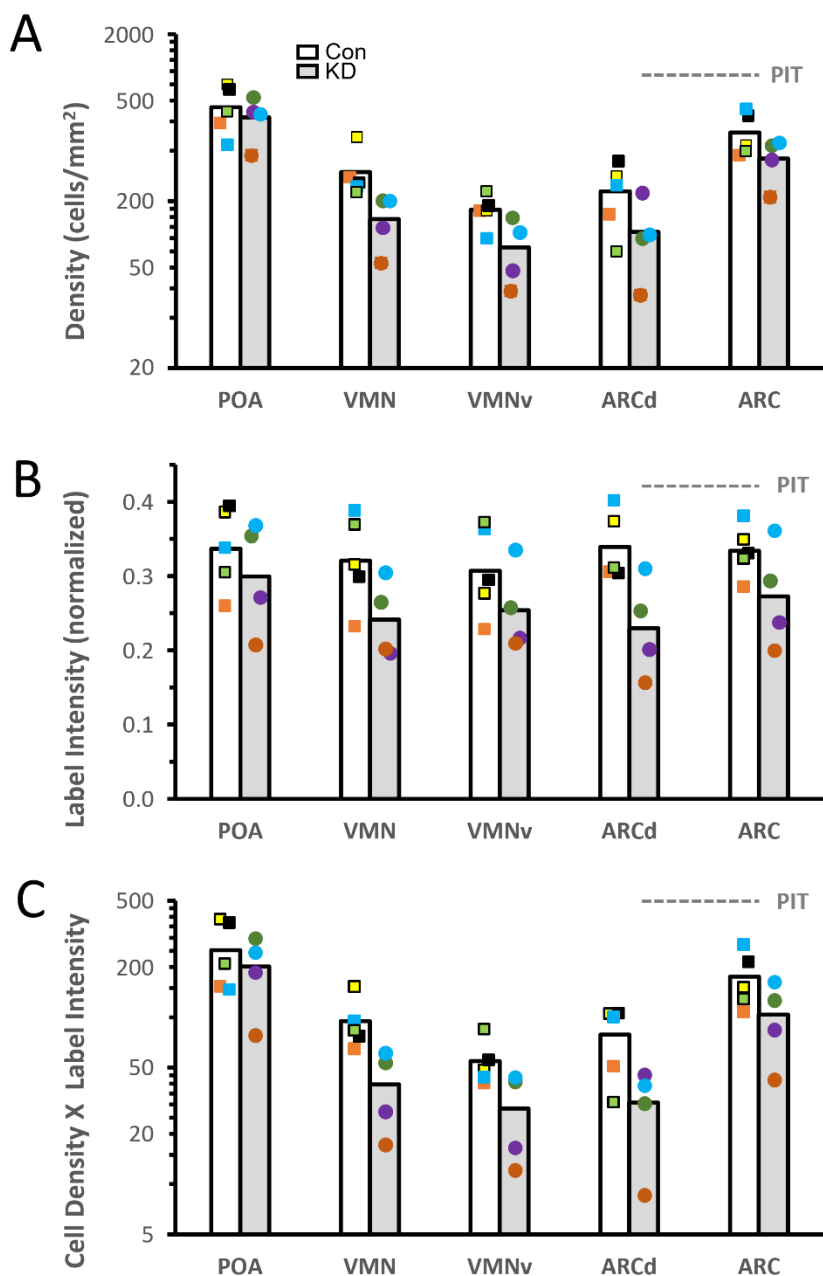
**Table 5C:** Mean (+/- SEM) regional body composition parameters as determined by dual x-ray absorptiometry (DXA) in E2, VEH and LET female groups of marmosets at baseline and after six months of treatment.



**Figure 8.** Actical collar determined body motion activity over 21 consecutive days (mean  $\pm$  SEM) in adult female marmosets in VEH (light gray circles and bars) and LET (dark gray squares and bars) groups (a) at baseline illustrating (A) hourly activity during 12 hour daytime and 12 hour nighttime and (B) AUC activity for daytime and nighttime, a, b:  $P = 0.001$ , at (b) 6 months (months) following OVX illustrating (A) hourly activity during 12 hour daytime and 12 hour nighttime and (B) AUC activity for daytime and nighttime, a, b:  $P = 0.01$ , and (c) % change in activity from baseline to 6 months following OVX during 12 hour daytime and 12 hour nighttime and AUC % activity change for 24 hours/day, daytime only and nighttime only, a, b:  $P = 0.01$ .

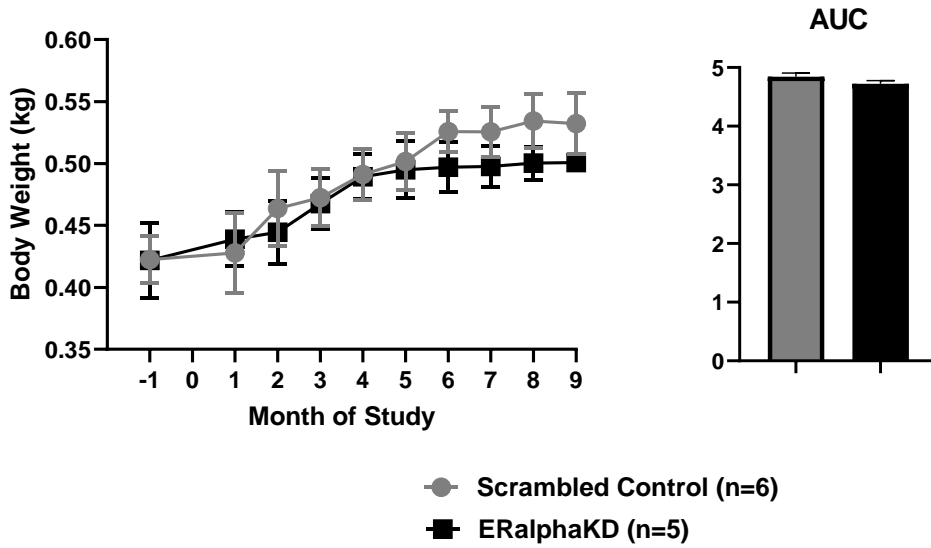
Parameter	E2	VEH	LET	P value
<b>Baseline</b>				
Basal glucose (mg/dL)	114 ± 3	119 ± 3	111 ± 6	—
<b>6 months after ovariectomy</b>				
Basal glucose (mg/dL) <sup>a</sup>	120 ± 11	127 ± 9	137 ± 7	.149
AUC OGTT glucose (mg/dL*120 min)	21251 ± 2386	19380 ± 1548	20843 ± 4829	.555
2 hour OGTT glucose (mg/dL)	130 ± 16	128 ± 13	180 ± 48	.440

**Table 6:** Mean ( $\pm$ SEM) basal glucose at baseline and 6 months after ovariectomy, as well as oral glucose tolerance test (OGTT) derived glucose values at 6 months after ovariectomy in E2, VEH and LET females. Abbreviations: AUC, area under the curve; OGTT, oral glucose tolerance test. <sup>a</sup>P = .055, 6 mo > baseline, all female groups combined. SI unit conversion : glucose  $\times$  0.0551 mmol/L.

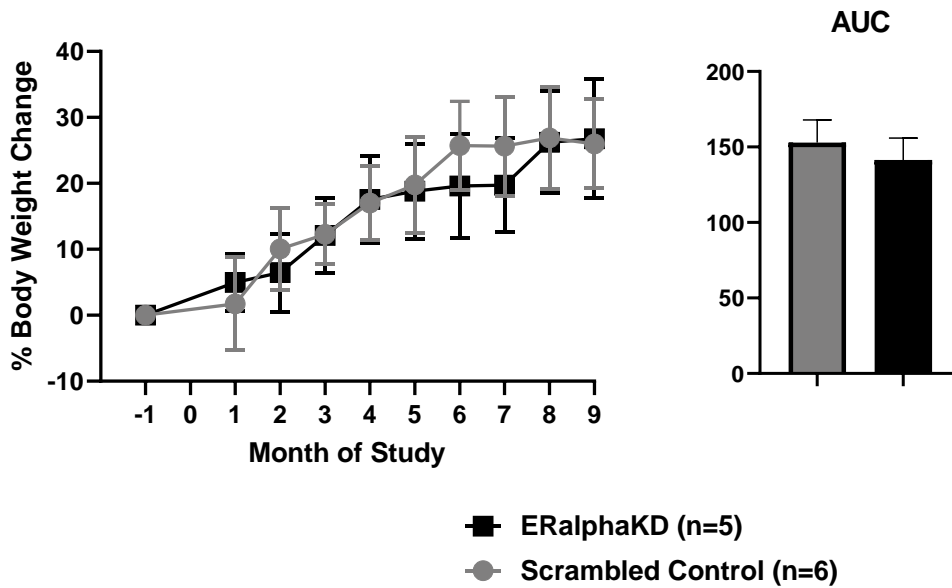


**Figure 9:** (A) The number of cells per unit area, or the ER $\alpha$  cell density is not statistically different between the groups in any of the hypothalamic nuclei analyzed. (B) ER $\alpha$  staining intensity in the VMN, but not the POA and ARC nuclei of the hypothalamus, is markedly less in the ER $\alpha$  gene silenced females (\* $p=0.04$ ) compared with control females. (C) density \* intensity in hypothalamic nuclei.

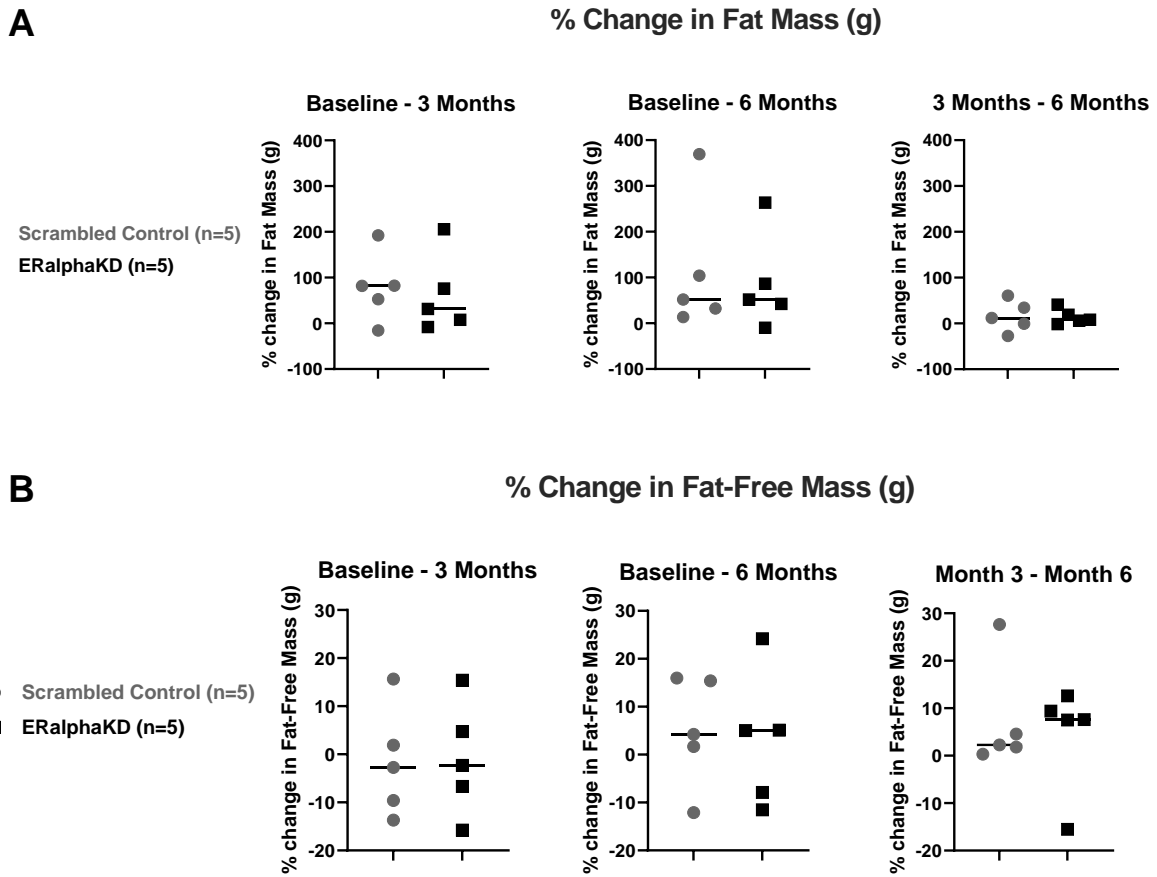
A



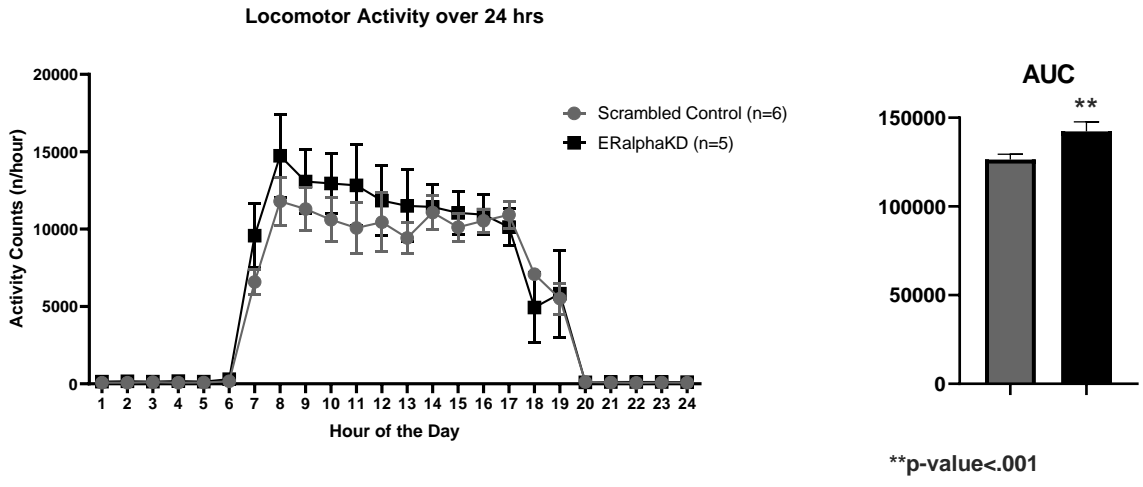
B



**Figure 10:** Body weight (kg) over duration of study with corresponding AUC bar graph (A) and body weight percent change from baseline weight over duration of study with corresponding AUC bar graph (B). Plots expressed as mean  $\pm$  SEM. Significance assessed via RM-ANOVA and student's t-test.

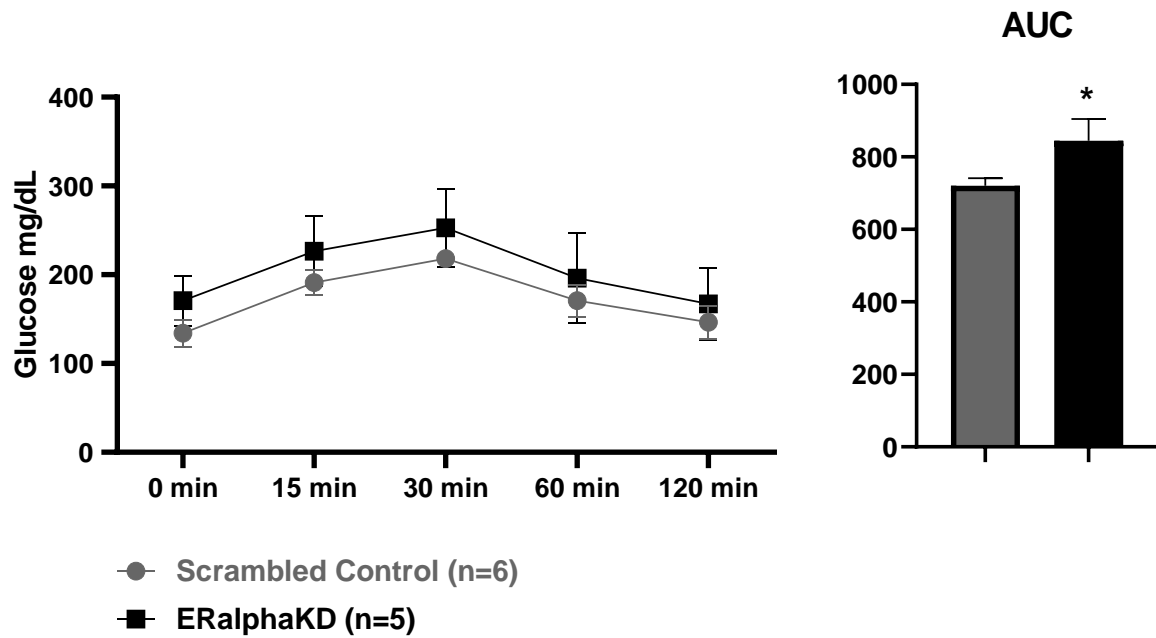


**Figure 11:** Percent change in fat mass (A) and fat-free mass (B) between DXA scans at baseline, 3-months, and 6-months post-treatment onset. Plots expressed as mean  $\pm$  SEM. Significance assessed via student's t-test.



**Figure 12:** Locomotor activity over 24-hours averaged over the duration of the activity assessment with corresponding AUC histogram. Plots expressed as mean  $\pm$  SEM. Significance assessed via RM-ANOVA and student's t-test. P\*\*<0.001.





**Figure 13:** Glucose levels during oral glucose tolerance test at 6-months post-treatment onset. Plots expressed as mean  $\pm$  SEM. Significance assessed via RM-ANOVA and student's t-test.  $P^* < 0.05$ .

## 2.7 References

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### **3. CHAPTER THREE: Hypothalamic *ESR1* Knockdown Induces Weight Gain and Reduces Morning Activity in Adult Female Rhesus Macaques**

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Contribution to this manuscript: Designed/carried out experiments, analyzed data, and prepared the manuscript for this thesis and subsequent publication]

### 3.1 Abstract

Estradiol ( $E_2$ ), the major bioactive estrogen, has been well documented as a major regulator of female metabolism in rodents. Estrogen receptor alpha ( $ER\alpha$ ) in the ventromedial nucleus (VMN) and arcuate nucleus (ARC) of the mediobasal hypothalamus (MBH) is the predominate estrogen receptor mediating female rodent metabolic homeostasis. In primates, however, it remains unknown if hypothalamic  $ER\alpha$  similarly mediates  $E_2$  metabolic actions. We hypothesized that reducing expression of  $ER\alpha$  in the VMN and ARC of female rhesus macaques would induce metabolic dysfunction by way of increased body weight, hyperphagia, decreased energy expenditure and insulin insensitivity. Adult female rhesus macaques were randomly assigned to one of two conditions: (1) control, scrambled shRNA (control,  $n=5$ ) and (2) selective  $ER\alpha$  protein knockdown with *ESR1* gene silencing shRNA ( $ER\alpha$ KD,  $n=6$ ). MRI-guided neural viral vector infusion surgery enabled delivery of scrambled or gene silencing shRNA into our target regions, VMN and ARC. Neurohistochemical visualization of green fluorescent protein (GFP) infected neurons confirmed accuracy of targeting as well as quantification of hypothalamic density and intensity of  $ER\alpha$  protein expression at study completion.  $ER\alpha$  expression was successfully diminished in our target regions of interest (ROIs); VMN ROIs of VMNv ( $p=0.03$ ) and VMNd1 ( $p=0.05$ ) and in ARC ROIs of ARC ( $p=0.02$ ) and ARCd ( $p=0.02$ ). The  $ER\alpha$ KD group presented with increased cumulative body weight % change (AUC,  $p=0.02$ ) and decreased morning locomotion between 0600h-0900h ( $p<0.01$ ). In both control and  $ER\alpha$ KD female groups,  $ER\alpha$  density in the ARC trended to be negatively correlated with body weight % change AUC ( $p=0.08$ ). Together, our findings highlight

ER $\alpha$  activation within hypothalamic nuclei as crucial for regulating metabolic homeostasis in female NHPs, and likely, women.

### 3.2 Introduction

The major bioactive estrogen, E<sub>2</sub>, is synthesized throughout the body by aromatase, a cytochrome P450 enzyme, encoded by the CYP19A1 gene<sup>1-4</sup>. While the main source of E<sub>2</sub> is synthesized in the ovaries, E<sub>2</sub> is also produced in extra-ovarian sites including adipose tissue, adrenal glands, skin, bone, pituitary gland, and multiple brain areas<sup>1-4</sup>. Rodent research has shown estrogen receptor alpha (ER $\alpha$ ), encoded by the gene, *ESR1*, to be the predominant estrogen receptor involved in female energy balance<sup>5-8</sup>, yet this has not yet been determined in nonhuman primates (NHPs). In female rodents, E<sub>2</sub> activates ER $\alpha$  in both the ventromedial nucleus (VMN) and arcuate nucleus (ARC) of the hypothalamus to initiate downstream neuroendocrine metabolic effects. In many experimental animal models, ovarian E<sub>2</sub> regulates body weight and composition, energy balance, and glucoregulation<sup>9</sup>. In rodent models, ovarian E<sub>2</sub>-regulated control mechanisms are well established since OVX mediated E<sub>2</sub> depletion reliably disrupts energy balance, including, increased body weight and visceral adiposity, reduced physical activity and energy expenditure<sup>10,11</sup>, and diminished insulin sensitivity<sup>10</sup>. All of these effects on metabolic function can be diminished or reversed by E<sub>2</sub> replacement<sup>12,13</sup>. Further, these E<sub>2</sub> actions are mediated by ER $\alpha$ , as complete knockout of *ESR1* in female mice (ESR1KO) increases body weight, adiposity and diminishes energy metabolism producing a phenotype that largely mimics that observed in long-term OVX mice<sup>14</sup>. Resulting from the vast research on the metabolic impact of ovarian E<sub>2</sub> via ER $\alpha$  in rodents, multiple mechanisms have been identified by which E<sub>2</sub> regulates of female energy balance. These include direct actions of E<sub>2</sub> on metabolically active tissues such as adipose tissue, liver, muscle, and pancreas<sup>15</sup>, E<sub>2</sub> activation of brown fat (BAT) via the

sympathetic nervous system (SNS)<sup>16</sup> and E<sub>2</sub>-induced alterations in neural and peripheral control of food intake and energy expenditure. Rodent models of ovarian E<sub>2</sub> regulation have also highlighted the critical role of neural ER $\alpha$  signaling in hypothalamic nuclei, VMN and ARC, to maintain energy homeostasis<sup>8,17</sup>. Importantly, one study demonstrated that viral vector mediated *ESR1* gene silencing in the VMN of both female mice and rats largely recapitulates a metabolic phenotype observed in complete ER $\alpha$  knockout mice, including obesity, hyperphagia, impaired glucose tolerance and reduced energy expenditure<sup>8</sup>. It remains unknown whether the same mechanisms of E<sub>2</sub> signaling, and metabolic regulation apply to female NHPs and women. Human studies suggest declining serum E<sub>2</sub> levels during the menopausal transition are associated with reduced energy expenditure and activity, increased central adiposity and insulin resistance<sup>12,13,18-20</sup>, but not all studies are consistent in these regards. In additional recently completed studies, we examined the role of ER $\alpha$  in the regulation of metabolic function in a female new-world, NHP, the common marmoset monkey (Chapter 2). Using a viral vector containing either a scrambled shRNA without known gene targets (control treatment) or gene silencing shRNA targeting *ESR1*, we assessed the metabolic effects of diminishing ER $\alpha$  protein expression in the mediobasal hypothalamus (MBH) of adult female marmoset monkeys. Our results suggested hyperglycemia and a reduction in daily caloric intake, despite increased body weight, induced by reduced ER $\alpha$  expression in VMN and ARC neurons in the MBH (Chapter 2), as found in female mice<sup>8</sup>. Taken together, our controlled NHP studies provide the first evidence that hypothalamic ER $\alpha$  signaling, in a diet-induced obesity (DIO) environment, is critically important in regulating energy balance in female NHPs. Moreover, recent research from our laboratory has shown that in female marmoset



monkeys, depletion of ovarian or both ovarian and extra-ovarian E<sub>2</sub> production removes protection from diet-induced obesity (DIO) and increased daily calorie consumption, suggesting that E<sub>2</sub> plays an important role in metabolic regulation in female NHPs (Chaper 2), and thus, possibly likely women<sup>21</sup>.

These findings reveal the importance of E<sub>2</sub> produced by the ovary, alone, as the primary organ source for E<sub>2</sub> activating hypothalamic ER $\alpha$  in at least one female primate NHP. Being able to replicate these findings that E<sub>2</sub> signaling hypothalamic ER $\alpha$  mediates similar regulation of metabolic homeostasis in a female NHP model with greater genetic, developmental, and physiological similarities to humans, such as rhesus macaques, will therefore be crucial to understanding how E<sub>2</sub> regulates female energy homeostasis and neuroendocrine-related functions in humans.

### **3.3 Methods**

#### *Animals*

Eleven adult female rhesus macaques (6-11 years of age) from the Wisconsin National Primate Research Center colony were randomized based on age, body weight and body mass index (BMI) into two groups: 1) estrogen receptor knockdown (ERaKD; n=6) in which monkeys received a viral vector infusion of a gene silencing shRNA targeting *ESR1* in the VMN and ARC of the hypothalamus and 2) scrambled control (scrambled control, SC; n=5) in which monkeys received a scrambled shRNA instead (Table 1). Monkeys were maintained in these two groups for ~34 months, pair-housed, in enclosures under 12-h lighting, ambient temperature of ~27°C and humidity of ~50%. (Figure 1). The protocol for this study was approved by the Animal Care and Use Committee of the Office of the Vice Chancellor for Graduate Research and Education of

the University of Wisconsin–Madison, an AAALAC-accredited program. Animals were fed chow twice a day, between 0700h-0900h and 1200h-1400h (Teklad Global 20% Protein Primate Diet, ENVIGO, Madison, WI). The calorie distribution of the biscuits was 29% protein, 14% fat and 57% carbohydrates with an energy density of 2.8 kcal/g. Animals were supplemented with daily fruit enrichment (<100 kcals), weekly tactile enrichments (<100 kcals), and weekly destructible enrichments (<100 kcals).

#### *Neural Infusion Surgery and Knockdown of ER $\alpha$*

OVX adult female rhesus monkeys received bilateral stereotaxic injections into the hypothalamic VMN and ARC of adeno-associated virus 8 (AAV8) expressing shRNA targeting *ESR1* for *in vivo* RNAi, and thus diminishing *ESR1* translation into ER $\alpha$  (ER $\alpha$ KD, n=6), or a control scrambled shRNA (Control, n=5). Treatment onset (0 months) was the day of neural infusion surgery.

Since accurate, within-brain placement of viral vector was essential for gene silencing efficacy and specificity, we used a single, presurgical, neuro-anatomical MRI scan to refine the unique locations of each VMN and ARC within each rhesus macaque, refined and adapted from<sup>22</sup>. To accomplish this, each female rhesus macaque was food deprived overnight, sedated with an intramuscular (IM) injection of ketamine (10 mg), anesthesia was continued with an IM injection of meloxicam (2 mg), 10 ml isotonic saline was injected subcutaneously (SC) to provide fluid administration and atropine (0.218 mg) was administered IM, followed by placement of the rhesus macaque into an MRI-compatible stereotaxic frame. Lateral placements of both left and right earbars, as well as vertical placements of palate and left and right eye bars, were all noted. Three-dimensional coordinate locations of both left and right VMN and ARC were subsequently

obtained from an ~23 min, T1 MRI anatomical scan adapted for rhesus monkeys<sup>23</sup> in a next-generation 3T MRI system MR750 (GE Healthcare, Waukesha, WI) and performed at least 3 days prior to viral vector infusion (Figure 2).

Each presurgical scan employed a 3-inch surface coil and comprised 3D T1-weighted (T1W) images with an inversion-recovery (IR) prepped, fast gradient-echo (IR-fGRE) sequence adapted for rhesus monkeys<sup>23</sup>. Scanning parameters included an inversion time (TI) of 450 ms, repetition time (TR) of 6.8 ms, echo time (TE) of 2.9 ms, and receiver bandwidth of 62.5 kHz. The matrix size was 256 × 256 × 128 (X × Y × Z). The 3D T1W scans produced 248 coronal 0.3-mm-thick contiguous slices. Using coronal slices incorporating the hypothalamus from each monkey's presurgical scan, we utilized contrast-agent filled earbars in the stereotaxic frame to provide a rostral-caudal (RC) zero plane (Z coordinates), the middle of the 3<sup>rd</sup> ventricle at the level of the mediobasal hypothalamus to generate a medial zero plane (X coordinates), and the farthest lateral point of each earbar to provide a dorso-ventral zero plane for right and left sides separately (X coordinates). Based on these 3D parameter sets, target coordinates for entry into the skull, as well as for rostral, medial-rostral, medial-caudal and caudal left and right VMN, and analogous locations for both left and right ARC, were estimated based on marmoset neuroanatomy atlas coordinates<sup>23</sup>.

For viral vector infusion surgery, each rhesus macaque was sedated and prepped as described for their presurgical MRI scan. In addition, each monkey received buprenorphine (0.072 mg) and cefazolin (174.9 mg) and was intubated with anesthesia continuing under isoflurane (1-2%). Each rhesus macaque was re-positioned into similar stereotaxic frame parameters as noted during for their presurgical MRI scan. Vital signs

(heart rate, blood oxygen, respiration, and temperature) were regularly monitored during the procedure and the animals were wrapped for warmth. A series of small-volume, subcutaneous (SC) injections of lidocaine (6 mg, total) were given along the anticipated incision site to provide local anesthesia. A ~60mm incision was made on the top of the animal's head. Stereotaxic coordinates, estimated from our pre-surgery anatomical MRI scan, allowed calculation for the insertion sites of a 22-gauge guide cannula housing a 28-gauge stylet to target the VMN and ARC in the hypothalamus. The estimated midline location between the medial sites for both VMN and ARC was marked on the skull to enable excision of a ~15 mm diameter portion of skull using a surgical bone drill. A micromanipulator, with cannula holder and guide cannula with screw-in solid stylet, were placed on the stereotaxic frame to confirm midline of the brain from the medial location of the superior sagittal sinus, and presurgical scan estimated target coordinates for left VMN and ARC sites were revised accordingly. A 27g needle, customized into a miniature scalpel blade, was used to open a ~10mm incision in the dura mater about 5-7mm to the right of the mid-sagittal sinus, lateral to potential hypothalamic locations on the right side of the rhesus monkey brain, to permit a 4-degree-to-the-vertical angled approach.

At a trajectory angle of 4° (offset from vertical, 0°), with the flaps of the dura held apart, a 22-gauge guide cannula housing a sterile 28-gauge stylet was lowered into the medial-rostral estimated extent of the right ARC<sup>24,25</sup>. The stylet was then replaced by a 28-gauge infusion cannula attached to a Hamilton syringe controlled by a programmable infusion pump. A solution of viral vector (~3.06 x 10<sup>10</sup> packaged genomic particles of AAV8 in 12.0 µL of sterile saline) mixed with gadolinium-containing Multihance MRI

contrast agent (2 mM gadobenate dimeglumine, Bracco Diagnostics Inc.), was infused at 2  $\mu$ L per minute for a total of six minutes. The AAV8 virus utilized in this study was shown to readily infect neurons<sup>26</sup> and was without notable inflammation in NHP brain<sup>24</sup>. To minimize dorsal tracking of viral particles and contrast agent when the guide and infusion cannulae were raised, there was a 5-min delay following cessation of infusion before the cannulae were raised approximately 1 mm to the VMN infusion site, all at a 4 degree-angle-to-the vertical. Viral vector infusion and delayed raising of the cannula were then performed at the VMN, as undertaken at the ARC. The infusion cannula alone was raised and removed from the guide cannula to permit its flushing of sufficient fluid while outside the brain to confirm infusion integrity of the system. The stylet was re-positioned inside the guide cannula. Both guide cannula and stylet were initially raised 0.5 mm, pausing for 30 sec to minimize dorsal tracking of viral particles and contrast agent, and then both were gradually removed from the brain. Superficial skin sutures were placed using 4/0 vicryl suture to temporarily close the incision site during transfer to the MRI scanning room while the rhesus macaque remained in the stereotaxic frame.

The contrast agent remaining at the right medial ARC and VMN infusion sites was used to confirm accuracy of the infusion site location from a mid-surgery, ~23-min MRI T1 scan obtained approximately 20 min following the first infusion on the right side. When necessary, presurgical scan estimated target coordinates for right VMN and ARC sites were revised. Guide cannula, stylet and infusion cannula insertion, infusion and removal procedures were then repeatedly performed to enable AAV8 infusion into rostral, medial-rostral, medial-caudal and caudal VMN and ARC target sites on the right side of the brain.

Following all right side infusions, a 27g needle customized into a miniature scalpel blade, was used to open a ~10mm incision in the dura mater about 5-7mm to the left of the mid-sagittal sinus, lateral to potential hypothalamic locations on the left side of the rhesus monkey brain, to permit a 4-degree-to-the-vertical angled approach. A similar series of procedures, infusions, and MRI scan were performed on the left side of the brain analogous to those performed on the right. When necessary, adjustments to presurgical estimates for rostral, medial-rostral, medial-caudal and caudal left VMN and ARC target sites were made following the 2<sup>nd</sup> within-surgery MRI scan.

*Efficacy of RNAi activity for shRNA specific to ESR1 and absence of RNAi in control scramble sequence shRNA*

Two viruses were designed for this experiment, 1) ER $\alpha$ 34 shRNA driven by the H1 promoter and 2) ER $\alpha$ 78 shRNA driven by the U6 promoter. Vector Biolabs performed all recombinant AAV8 production, and the Dharmacon shRNA design center online was employed to design the shRNAs. Candidate shRNAs were only advanced to the next round if the chosen sequence was identical in marmoset and rhesus monkeys, human and mouse. Furthermore, candidate shRNAs were required to target a region of ER $\alpha$  that was common to all isoforms. Candidates were further scrutinized by examining potential off-target effects using BLAST analysis against the database for rhesus monkeys.

We obtained the AAV-H1-EGFP (green fluorescent reporter gene) or AAV-U6-RFP (red fluorescent reporter gene) plasmids from Vector Biolabs that were pre-designed, and then cloned in the shRNA sequence that we designed. Once the shRNA was chosen, and cloning completed, we transfected the shRNA expressing plasmids into MCF7 cells (a human breast cancer cell line with estrogen, progesterone, and glucocorticoid reporters<sup>27</sup>

at two different doses compared to mock transfected cells and ER $\alpha$  protein expression assessed by western blot quantified the ability of each shRNA to knockdown ER $\alpha$ . The pAAV-H1-ER $\alpha$ 34-EGFP plasmid was used to transfect MCF7 cells to validate the knockdown. Three candidate shRNAs were screened: ER $\alpha$ 12, ER $\alpha$ 34, ER $\alpha$ 56. After initial screening all three shRNAs, ER $\alpha$ 34 and ER $\alpha$ 56 were more successful than ER $\alpha$ 12 (validation blots, Figure 3) so ER $\alpha$ 34 and ER $\alpha$ 56 were compared in the follow up experiment.

The ER $\alpha$ 78 shRNA sequence mimicked that of the ER $\alpha$ 34 shRNA, except the leading nucleotide was changed from "A" to "G" to make it more compatible and efficient when employing the U6 promoter. Additionally, the pAAV-H1-ER $\alpha$ 34-EGFP engaged the GFP reporter for identifying hypothalamic areas infected during subsequent neuroimmunohistochemical morphological analyses, and the pAAV-U6-ER $\alpha$ 78-RFP engaged the RFP reporter for subsequent identification of infected hypothalamic areas. The oligos used for subcloning the shRNAs into the pAAV-H1-GFP and pAAV-U6-RFP plasmids are illustrated in Figure 4.

*Immunohistochemical processing of rhesus monkey brain for visualizing and quantifying green or red fluorescent protein (GFP or RFP) and ER $\alpha$  protein expression*

Following upper body perfusion with 4% paraformaldehyde (PFA) (pH 7.6) at necropsy, each female rhesus macaque brain was removed, post-fixed at 4C for an additional 12-16h and cryoprotected in graded (10-30%) sucrose/PBS (pH 7.2) solutions. Each post-fixed, dehydrated brain was placed in a customized plastic mold designed to accommodate the dorsal surface of an adult rhesus brain and expose the ventral surfaces landmarks, enabling placement of mold-guided, microtome blades cutting the brain into

coronal plane portions. The medial rostral-caudal portion (~8-10 mm), including the hypothalamus, was separated, comprising immediately rostral to the optic chiasm to immediately caudal to the mammillary bodies. This medial rostral-caudal portion of whole-brain was cryosectioned into consecutive 40 $\mu$ m frozen sections in the coronal plane to include the rostral-caudal extent of the hypothalamus from the rostral extent of the mPOA to the caudal extent of the ARC.

Immunohistochemistry (IHC) procedures for ER $\alpha$ , GFP (control monkeys) and RFP (ER $\alpha$ KD monkeys). Cryo-sections of 40  $\mu$ m, at regular intervals, were immunostained for ER $\alpha$  using the mouse monoclonal antibody, 6F11 (Thermo Fisher Scientific, Hampton, NH)<sup>29</sup>. DAB-staining was subsequently used to detect specific primary antibody binding. Sections adjacent to those selected for ER $\alpha$  immunostaining were stained for GFP using the rabbit polyclonal antibody, AB3080 (MilliporeSigma, Burlington, NH)<sup>30</sup> (control monkeys) or for RFP using the polyclonal antibody, R10367 (Invitrogen, Waltham, MA) (ER $\alpha$ KD monkeys).

*Confirmation of neural targeting accuracy from immunohistochemical analyses of GFP or RFP expression in the mediobasal hypothalamus*

Digital images from coronal brain sections immunostained for GFP and RFP were obtained with a Nikon Eclipse Ti microscope and Nikon NIS Elements software. Images were acquired of the hypothalamus and peri-hypothalamic areas from the rostral preoptic area to the caudal arcuate nucleus and mammillary bodies at 4x objective. Each hypothalamic section image was the composite of several images stitched together by the Nikon system during the photo acquisition process. As illustrated by a coronal section from approximately the mid-rostral-caudal extents of VMN and ARC within the



hypothalamus (Figure 5), abundant GFP-immunoreactive (ir) labeling was demonstrated in VMN (Figure 5B) and ARC (Figure 5B) from a control monkey, and in VMN and ARC (Figure 5C) from an ER $\alpha$ KD monkey. Selected images from each rostral-caudal series of coronal sections of GFP-ir labeling, including the POA, VMN and ARC, for each rhesus macaque were positioned at appropriate rostral-caudal locations associated with quantification of ER $\alpha$  immunostaining, as illustrated in Figure 5 for a representative control (Figure 5A) and ER $\alpha$ KD female rhesus monkeys (Figure 5C) to confirm AAV8 infection in both VMN and ARC.

#### *Analysis of Neural Targeting of ER $\alpha$ Gene Silencing*

Digital images from sections immunostained for ER $\alpha$  were obtained as described above for GFP and RFP imaging and analyzed with NIS Elements Advanced Research (AR) image analysis software (ver. 5.02) was then used to quantify the amount of ER $\alpha$ -immunoreactive (ir) labeling in the POA, VMN and ARC, employing measures of cell counts, cell density per unit area (number of cells/mm<sup>2</sup>), and labeling intensity (pixel intensity) of individual cells. Labeled cells were detected using the NIS Elements AR Dark Spot Detection feature. The contrast threshold in the detection algorithm was determined for each monkey by adjusting contrast settings in the spot detection algorithm to match cell counts made by three observers on a series of six cropped images of the hypothalamic section to be analyzed. Observer counts differed by less than 15 %, and NIS threshold values were adjusted so NIS-derived cell counts fell within 5% of the average counts of all observers. The NIS threshold value for labeled cells was then applied across all hypothalamic sections to derive cell counts for that monkey.

The NIS Elements AR software calculated the mean pixel intensity of each identified cell, and we used this to estimate the intensity of ER $\alpha$  staining. Threshold staining intensity was determined by the threshold process described above, and the threshold value was quantified as lowest 1% intensity of all stained cells. The darkest staining label was defined as the pixel intensity of the darkest 2% of ER $\alpha$  expressing cells; these were typically a subset of cells in the POA and anterior pituitary, which were distant from the infusion and served as a positive control within each monkey. This established a range from threshold to dark for each monkey. Staining of each labeled cell was normalized within this range, and staining intensity measures were calculated from the normalized intensity values.

Density of individual neuronal nuclear immunostaining for ER $\alpha$  in each of the hypothalamic areas was determined by counting the number of immunostained neurons within regions of interest (ROIs) of 1000 $\mu$ m X 500 $\mu$ m ellipses (areas of 0.395 mm<sup>2</sup>) positioned within POA, VMN and ARC. Multiple ROIs were used in the VMN (VMNv, VMN, VMNd1, VMNd2, ventral to dorsal) and ARC (ARC and ARCd) because of the broad extent of the ER $\alpha$  cells they contained (Figure 6). For each hypothalamic ROI within each female, we used the average of the three most densely-staining sections as an overall density measure for that ROI.

Quantitative analyses were performed on digital images obtained while the microscope lens was focused on the middle depth of each hypothalamic section. At that focal point, the depth of field with a 4X objective allowed visualization of labeled cells throughout the entire depth of the hypothalamic section (Chapter 2). To confirm validity of this strategy, we compared cell counts from images acquired when focus was on the

surface of the hypothalamic section, middle of the hypothalamic section and bottom of the hypothalamic section. Cell counts obtained from center-focused images varied less than 5% from those obtained from top-of-section or bottom-of-section focused images.

#### *Hormone and Metabolic Analyte Determinations*

Steroid hormone analysis was adapted from methods previously described<sup>31,32</sup>. Briefly, internal standard was added to rhesus serum samples (1mL) and then they were extracted twice using methyl tert butyl ether followed by dichloromethane. Samples were derivatized using dansyl chloride and then analyzed by LC-MS/MS (Sciex QTRAP 5500). Individual calibration curves were constructed for each analyte with at least 8 points. The linearity was  $r > 0.9990$  and the curve fit was linear with 1/x weighting. None of the compounds of interest were detected in blank or double blank samples. Inter-assay coefficient of variation was determined by 3 levels of human serum and ranged from 5.6-9.7%.

#### *Body Weight, Body Composition & Bone Mass*

Animals were weighed weekly. Incremental area under the curve (AUC) assessment of weight parameters over time, calculated by the trapezoid rule, was used to better detect recurring differences in weight gain. At ~18 months post-neurosurgery and ~28 months post- neurosurgery, total body composition was assessed by dual-energy X-ray absorptiometry (DXA, iDXA, GE/Lunar Corp., Madison, WI) on sedated animals. Monkeys received up to 7 mg/kg ketamine IM and up to 0.03 mg/kg dexmedetomidine IM, to be reversed on conclusion of the procedure by up to 0.3 mg/kg atipamezole IM. After placing the anesthetized monkey onto the scanner bed, DXA-compliant cushioning was used to position the monkey with arms and tail flat on the

scanner bed, clearly distinct from its trunk and other extremities. A DXA scan was then performed for approximately 30 minutes. The scanner room was maintained at 27-28°C during the performance of all scans to minimize the risk of hypothermia. Monitoring of anesthesia recovery was documented every 15 min until the monkey was sitting upright, then every 30 minutes until the animal was fully recovered from the anesthesia. Fat mass, fat-free mass, bone mineral content (BMC) and bone mineral density (BMD) were determined for total body as well as regions of interest, including abdomen, chest, thighs and extremities<sup>33</sup>. Body mass index was calculated using  $\frac{\text{Body Weight (kg)}}{\text{Crown Rump Length (m}^2\text{)}}$ <sup>34</sup>.

#### *Somatometric Measurements*

Somatometric parameters were measured using tape measure or calipers while monkeys were anesthetized for DXA imaging. Measurements included femur length; abdomen, chest, pelvis, arm, leg, and head circumferences; head diameter; and chest and abdomen skinfolds. Additionally, monkeys were placed on an osteometric board for measurement of crown rump and crown-heel lengths.

#### *Indirect calorimetry*

At ~28-months post treatment onset, monkeys were singly housed in a transparent metabolic chamber (75 cm x 75 cm x 80 cm (0.45 m<sup>3</sup>) with respiratory system (Sable Systems, North Las Vegas, NV) for 48 hours. While occupying the chamber, each monkey was able to see and hear its cagemate or another monkey residing in a standard cage or another metabolic chamber within the same room; no other monkeys resided in the room. Monkeys were fed chow, fruit enrichment (~1,000 total kcal a day) split between two feeding periods, 0700h-1000h and 1200h-1400h while in the chamber. Oxygen (O<sub>2</sub>) consumption and carbon dioxide (CO<sub>2</sub>) productions

were measured. As determined by behavioral observations in an acclimation period of 24 hours, such circumstances were not stressful for these monkeys. Room air was drawn through the chamber and flow rate was continuously monitored. Exhaust air was sampled for analysis of O<sub>2</sub>, and CO<sub>2</sub> content. Energy expenditure, respiratory quotient, daily, daytime and nighttime metabolic rate, and meal-induced thermogenesis, noted as the energy expenditure in the five hours post meal consumption, was determined from these data.

#### *Locomotor Activity Assessment*

At ~18 months and ~28 months post-treatment onset Actical accelerometers (Respironics, Murrysville, PA) were attached to a fabric collar and placed around each monkey's neck. Activity and intensity of movement were recorded over a ~4-week period after which the accelerometers were removed. Activity was recorded in counts (n), where one count represented body movements on an x, y, or z axis. The accelerometer sampled such activity counts every minute and these data were averaged for every hour (h), day (during lights on, 0600-1800 h), night (during lights off, 1800-0600 h), morning (0600–1200 h), afternoon (1200–1800 h), and over 24 hours. Incremental area under the curve activity values were also assessed to detect cumulative differences over the course of the study.

#### *Caloric Intake Assessment*

At ~28 months post-treatment onset, monkeys were singly housed 0700-1800 for 7-21 days. Monkeys were fed 9 biscuits of standard chow twice a day at 0700-0900 and 1200-1400. At each feeding, monkeys received on average ~423 kilocalories from chow. In addition, monkeys received daily fruit enrichment (i.e., pear, banana, green

pepper, kiwi and apple) totaling on average ~100 kilocalories per enrichment occurrence. Daily chow and enrichment were weighed prior to be given each day (Calories Offered). At 1800h, any chow or enrichment not consumed from the day was weighed (Calories Left) and returned to each monkey for the overnight period (1800h-0700h). Average calories (kcal) consumed was calculated by multiplying the weight (g) of chow consumed each day by the chow's energy density of 2.8 kcal/g and averaged over the 20-day food assessment period. Monkeys were repaired with their partner for the overnight period (1800h-0700h).

#### *Intravenous Glucose Tolerance Test (ivGTT)*

The ivGTT was conducted according to the Modified Minimal Model (MMM) protocol as adapted for rhesus monkeys<sup>35</sup>. A catheter was inserted through the saphenous or femoral vein and the tip positioned in the inferior vena cava. Four pretreatments ~1.5 ml blood samples were taken at approximately -15, -10, -5, and -1 minutes to establish fasted baseline values of insulin and glucose. A ~300 mg/kg glucose bolus was administered over about one minute starting at ~ 0 min, and ~1.5 ml blood samples were drawn at 2, 3, 4, 5, 6, 8, 10, 12, 14, 16, 19, 22, 23, 24, 25, 27, 30, 40, 50, 60, 70, 80, 90, 100, 120, 140, 160, and 180 minutes after glucose administration. Blood volume availability for individual animals was calculated by body weight (animal's body weight (kg) x 60 = estimated total volume of blood). If a monkey's blood volume availability did not meet the required volume for the full ivGTT (48 ml), samples at timepoints of 100, 120, 140 and/or 160 were omitted. Twenty minutes after glucose bolus administration, tolbutamide (~25mg/ml up to a maximum volume of ~5ml 1N sodium hydroxide, 0.1N hydrochloric acid in sterile saline solution) was administered

via the catheter to induce an additional increment in insulin secretion to enhance MMM calculations. Maintenance of catheter patency and fluid volume was accomplished with intravenous Plasma-Lyte A provided by an infusion pump hooked up to the indwelling catheter. Additional fluid replacement was provided by 1-2ml cannula line flushes following each blood sample withdrawal. The duration of anesthesia was approximately 240 minutes. Monkeys were maintained under warm heat lamps (with a 24-inch minimum space between the monkey and the heating lamp), a heated air warming unit and/or a water-circulating heating pad. During the procedure, each monkey's blood oxygen, heart rate and rectal temperature were continuously monitored and recorded every 15 minutes. After completion of the procedure, progress of anesthesia recovery was documented every 15 min until the monkey was sitting upright, then every 30 min until the monkey was fully recovered from the anesthesia.

### *Statistical Analysis*

Data collected was analyzed utilizing GraphPad PRISM and SPSS software. Metabolic parameters collected at both time points (~18 mo and ~28 mo) were compared by ANOVA with repeated measures (RM-ANOVA). Neuronal nuclear ER $\alpha$  immunostaining intensity, immunostained cell density within neurons, as well as metabolic, endocrine and somatometric parameters were analyzed using independent t-tests to compare means. Plots are expressed as means $\pm$ SEM. Relationships between hormones, metabolic parameters, and ER $\alpha$  immunostaining intensity and density were assessed via nonparametric Spearman's correlation tests. Statistical significance was determined as  $p \leq 0.05$  and trending towards significance was determined as  $0.05 < p < 0.1$ .

### 3.4 Results

#### *Targeted knockdown of ER $\alpha$ within VMN and ARC*

ER $\alpha$ KD monkeys typically demonstrated less ER $\alpha$  protein density and ER $\alpha$  density\*intensity in multiple VMN and ARC ROIs as determined by immunohistochemical analysis when compared to controls (Figure 7). In addition, ER $\alpha$  density was less in the VMN ROIs of VMNv ( $p=0.03$ ) and VMNd1 ( $p=0.05$ ) and ER $\alpha$  density\*intensity was less in the VMNv ( $p=0.05$ ) and tended to be less in the VMNd1 ( $p=0.06$ ) in ER $\alpha$ KD monkeys compared to controls. Additionally, there was reduced ER $\alpha$  density in the ARC ( $p=0.02$ ) and ARCd ( $p=0.02$ ) and reduced ER $\alpha$  density\*intensity in the ARC ( $p=0.02$ ) and ARCd ( $p=0.03$ ) in ER $\alpha$ KD compared to control monkeys.

Certain metabolic and hormone parameters involving all females combined across both groups correlated with degree of ER $\alpha$  knockdown in specific brain regions. In the ARC, there was a trend towards a negative correlation between body weight percent change AUC with ER $\alpha$  expression density (Figure 8A;  $R^2=0.33$ ,  $p=0.08$ ). In each of the VMNv, VMNd1, ARC, and ARCd, luteinizing hormone (LH) AUC between 18-30 months negatively correlated with ER $\alpha$  expression density (VMNv,  $R^2=0.58$ ,  $p=0.02$  (Figure 8B), VMNd1,  $R^2=0.42$ ,  $p=0.057$  (trend), ARC,  $R^2=0.75$ ,  $p=0.001$ , ARCd,  $R^2=0.52$ ,  $p=0.02$ ) and ER $\alpha$  expression density x intensity (VMNd1,  $R^2=0.37$ ,  $p=0.08$  (trend), ARC,  $R^2=0.63$ ,  $p=0.006$ , ARCd,  $R^2=0.48$ ,  $p=0.03$ ).

#### *Steroid Hormones*

There were no treatment differences observed in steroid hormone measurements of E<sub>2</sub>, estrone (E<sub>1</sub>), testosterone (T), androstenedione (A<sub>4</sub>), DHEA, progesterone (P<sub>4</sub>), 17-OHP, LH and FSH at baseline, 18-months, and 30-months post-treatment onset (Table



2). Area under the curve analyses of hormones revealed increased LH ( $p < 0.001$ ) and  $A_4$  ( $p = 0.03$ ) between 18-months and 30-months of the study.

### *Body Composition*

Weekly weight measurements were averaged for each month of the study. Body weight via AUC over the duration of the study was increased in ER $\alpha$ KD compared to control females ( $p = 0.01$ , Figure 9A). Body weight percent change AUC was also increased in the ER $\alpha$ KD compared to the control females ( $p = 0.02$ , Figure 9B). Although there was a cumulative increase in body weight % change over the course of the 34-month study in the ER $\alpha$ KD monkeys, DXA scans at 18-months and 28-months of the study indicated no differences between ER $\alpha$ KD and control females with regard to total fat mass (1<sup>st</sup> DXA,  $p = 0.39$ ; 2<sup>nd</sup> DXA,  $p = 0.38$ ) and total fat-free mass (1<sup>st</sup> DXA,  $p = .18$ ; 2<sup>nd</sup> DXA,  $p = 0.33$ ) (Figure 10). Additionally, percent change of fat mass ( $p = 0.45$ ) and fat-free mass ( $p = 0.85$ ) between the 18-month and 28-month DXA scans did not differ between female groups (Figure 9). There were also no differences in fat mass or fat-free mass in any defined region of interest (abdomen, extremities, thigh, or head) between female groups (Table 3).

### *Caloric Intake*

There were no differences between female groups in average kcals consumed ( $p = 0.29$ , Figure 11) or average kcals assumed when adjusted for kilogram of body weight ( $p = 0.12$ , Figure 11). There was a trend, however, towards reduced kcal consumed when adjusted for kilogram of FFM in the ER $\alpha$ KD compared to control females ( $p = .085$ , Figure 11).

### *Indirect Calorimetry*

The metabolic chamber analysis enabled investigation into energy expenditure (EE) per minute (EE/min) with EE values adjusted for kilogram of fat-free mass. Cumulative EE assessed via AUC analysis indicated there were no female group differences in EE over the course of the 48-hour period the monkeys occupied the metabolic chambers (Figure 12). More specifically, there was also no difference in averaged EE with or without being adjusted for fat-free mass for 24-hours ( $p=0.80$ , w/FFM adjustment,  $p=0.35$ , Figure 13) daytime EE, defined as 0600-1800 ( $p=0.53$ , w/FFM adjustment,  $p=0.31$ , Figure 13) and nighttime EE, defined as 1800-0600 ( $p=0.65$ , w/FFM adjustment,  $p=0.59$ , Figure 13) between female groups. Energy expended was calculated from the difference of averaged nighttime EE from daytime EE. Energy expended with ( $p=0.32$ , Figure 14) and without FFM ( $p=0.40$ , Figure 14) adjustment showed no between female group difference. Sleeping EE, defined as the lowest continuous 3-hour EE period during the night also showed no between female group difference ( $p=0.46$ , w/FFM adjustment,  $p=0.54$ , Figure 15).

### *Locomotor Activity*

Activity counts were averaged over 24-hrs, daytime (0600-1800), nighttime (1800-0600), morning (0600-1200) and afternoon (1200-1800) for the duration of the activity assessment at ~28-months post-treatment onset. There were no between group differences in 24-hr, day, night, morning, or afternoon periods (Figure 16). Although there was not a between group difference in the morning period (0600h-1200h), during the hours of 0600h-0900h, the ER $\alpha$ KD females exhibited reduced morning activity compared to the controls (0600h-0700h,  $p<0.001$ ; 0700h-0800h,  $p<0.001$ , 0800h-0900h,  $p<0.05$ ; Figure 17).

### *Glucoregulation via ivGTT*

Glucoregulatory values were calculated via the Bergman-Cobelli Minimal Model software. There were no treatment differences in insulin sensitivity ( $S_i$ ,  $p=0.45$ ), disposition index (DI,  $p=0.95$ ), acute insulin response (AIRg,  $p=0.27$ ), glucose effectiveness ( $S_G$ ,  $p=0.58$ ), and basal insulin ( $I_b$ ,  $p=0.68$ ). There was a trend towards reduced basal glucose ( $G_b$ ) in the ER $\alpha$ KD compared to control females ( $p=0.06$ , Figure 18).

### **3.5 Discussion**

Hypothalamic ER $\alpha$  has been well documented as the main estrogen receptor governing E $_2$ -dependent energy homeostatic mechanisms in female adult rodents, yet, prior to this study the role of ER $\alpha$  in a female adult NHP model with close physiological, reproductive, and genetic similarities to humans was unknown. Our results suggest that NHP hypothalamic ER $\alpha$  is an important neural target for E $_2$  to maintain energy balance in adult female rhesus macaques and likely, women.

In the present study, we demonstrated that mediobasal hypothalamic ER $\alpha$  mediates energy balance in terms of body weight regulation as well as locomotion activity in adult female rhesus macaques. The hypothalamic nuclei regions, VMN and ARC, targeted for MRI-guided viral vector infusion diminishing ER $\alpha$  protein expression were divided into multiple regions of interests (ROIs) for immunohistochemical (IHC) analysis. The ROIs VMNv, VMNd1, ARC and ARCd demonstrated significantly less ER $\alpha$  expression density and ER $\alpha$  density x intensity in ER $\alpha$ KD compared to control monkeys. Twelve months after neural ER $\alpha$  knockdown surgery, visually, ER $\alpha$ KD females began diverging in body weight percent increase compared to the controls.

Incremental AUC analyses were used to highlight the subtle cumulative weight gain induced by reduced hypothalamic ER $\alpha$  expression. We also demonstrated reduced ER $\alpha$  expression contributing to reduced locomotion in specific morning hours compared to control monkey activity. The positive energy balance of body weight gain and reduced morning locomotion resembles previous studies in rodents<sup>10,14</sup> and humans<sup>18</sup> that demonstrated menopause-associated body weight gain and reduced spontaneous activity. Similarly, to previous human studies<sup>18</sup>, we found no caloric intake differences in menopausal compared with premenopausal women despite weight gain and reduced spontaneous activity.

In contrast to our findings of body weight percent increase and reduced locomotor activity in specific morning hours, mice subjected to ER $\alpha$  knockdown in the VMN exhibit a more robust metabolic phenotype. ER $\alpha$ KD mice demonstrate increased body weight, reduced energy expenditure, hyperphagia and impaired glucose tolerance within 10 weeks compared to control mice<sup>8</sup>. The difference in severity of metabolic phenotype after ER $\alpha$  knockdown could be due to a more complete knockdown of ER $\alpha$  protein expression in the mice compared to the rhesus monkeys in our study<sup>8</sup>. Additionally, we saw similar contrasts in our recently completed adult female marmoset monkey ER $\alpha$  knockdown study. In ER $\alpha$ KD female marmosets, we observed reduced 24-hour locomotor activity and a trend towards impaired glucose tolerance. Due to the less robust metabolic phenotypes observed in our ER $\alpha$ KD female marmoset and rhesus monkey studies, we speculate that fewer metabolic processes are mediated by hypothalamic ER $\alpha$  within the VMN and ARC in adult female rhesus monkeys compared

to adult female mice. Perhaps, one metabolic process mediated by ER $\alpha$  and conserved across species is the regulation of locomotion.

Previously, our group investigated the metabolic consequence of silencing hypothalamic ER $\alpha$  expression in a marmoset monkey model and provided similar evidence that mediobasal hypothalamic ER $\alpha$  expression is crucial for maintenance of female energy homeostasis in female NHPs (Chapter 2). The female marmoset study, in contrast to the present study employing female rhesus macaques, diet-induced obesity (DIO). Estradiol and functional ER $\alpha$  have been shown in rodent models to have a protective effect against DIO attributed weight gain. We speculate that adoption of a DIO methodology in this rhesus monkey study, may have exaggerated the weight gains and reductions in morning activities in ER $\alpha$ KD compared to control females that we observed in the present study. Although the present rhesus monkey study approximated 3 years in duration, the menopausal transition in humans averages 3-4 years and can proceed over a decade<sup>36</sup>. It is plausible that the specific metabolic differences detected are only the first indications of metabolic dysfunction due to reduced ER $\alpha$  expression and disruption of E<sub>2</sub> mediated energy homeostatic mechanisms. Moreover, it is possible that ER $\alpha$  and E<sub>2</sub> independent mechanisms of energy balance can compensate in the absence of hypothalamic ER $\alpha$  action, such as increased signaling through leptin-mediated melanocortin pathways that are not all regulated by ER $\alpha$ . It had been assumed that E<sub>2</sub>'s effect on energy balance acts in tandem with leptin's role via co-expression of leptin receptor (LepRb) and ER $\alpha$  in the mediobasal hypothalamus, but recent research in rodents has shown co-expression is more pronounced in the preoptic area (POA) and is limited in the mediobasal

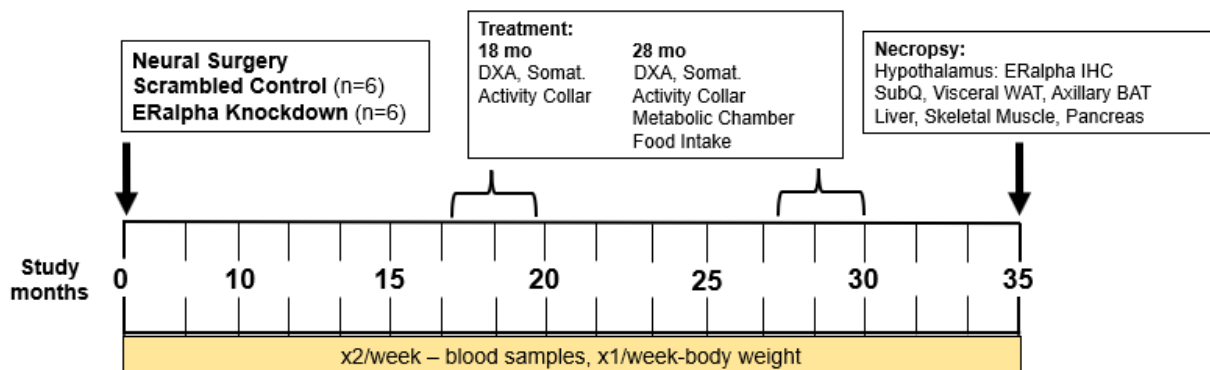
hypothalamus. Quantification of LepRb and ER $\alpha$  colocalization throughout the murine hypothalamus has revealed that robust colocalization does not occur in all areas aside from the POA and one month after OVX, co-expression tended to decline<sup>37</sup>. Further, STAT3 or ER $\alpha$  deletion from LepRb cells does not inhibit E<sub>2</sub>'s ability to regulate body weight<sup>38</sup>. It has been shown that circulating levels of E<sub>2</sub> may enhance the anti-obesity effects of leptin and specifically modulate leptin sensitivity in certain metabolic tissues like adipose and skeletal muscle<sup>39</sup>. Taken together, E<sub>2</sub> mediation of energy homeostasis via ER $\alpha$  is not dependent on activation of LepRb on co-expressing cells and circulating levels of E<sub>2</sub> may act indirectly on the leptin-melanocortin pathway and it can be posited that in the absence of ER $\alpha$  expression, leptin control of energy balance may partially compensate, enough, if not further stressed by DIO or a sedentary lifestyle.

In summary, partial silencing of ER $\alpha$  in the VMN and ARC of rhesus macaques results in significant body weight gain and a significant reduction in locomotion during the peak hours of physical activity. These findings replicate only some of much more extensive effects of similar treatments in rodents, a difference that may be attributable to technical limitations in deploying RNAi strategies in NHPs vs. rodents, or incompletely overlapping metabolic roles of ER $\alpha$  in these differing experimental models. Nevertheless, our findings confirm ER $\alpha$  as a crucial neural receptor of E<sub>2</sub> action in female NHPs and reveal the potential for therapies to engage this specific neural target as a way to combat obesity and E<sub>2</sub>-associated metabolic disorders.

### 3.6 Tables & Figures

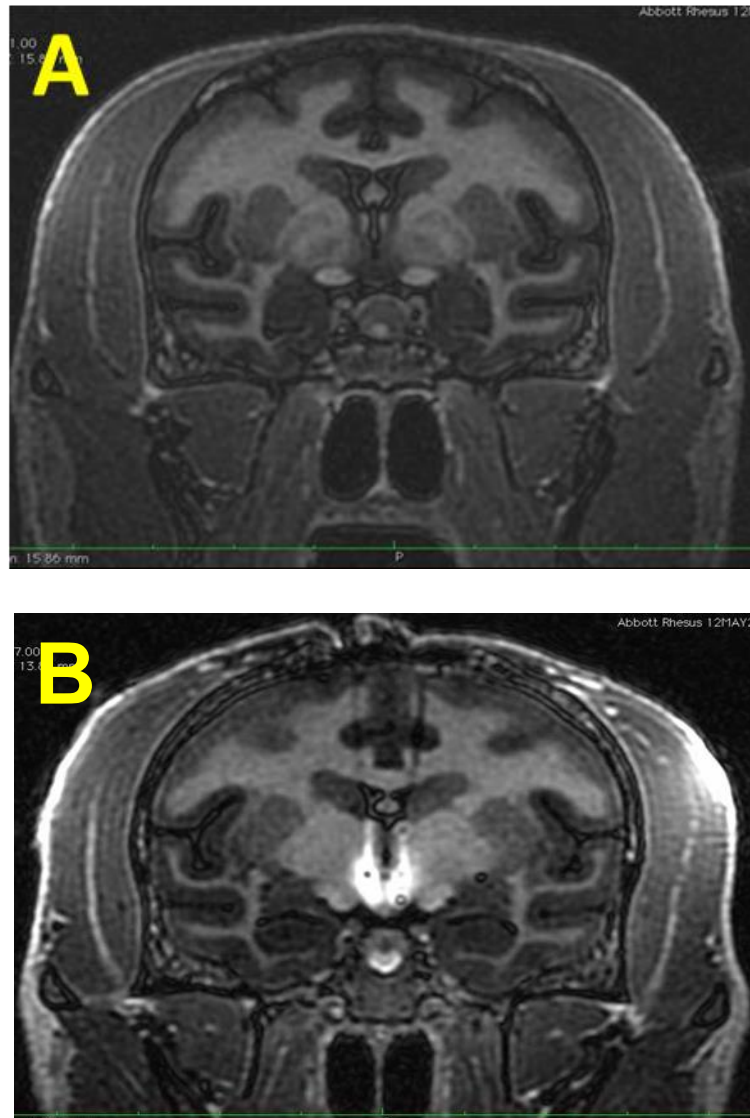
	Age (years)	Body Weight (kg)
Scrambled Controls (n=6)	10.75±1.16	7.71±1.05
ER $\alpha$ Knockdown (n=6)	11.26±0.77	8.35±0.51
p-value	0.71	0.56

**Table 1:** Comparable ages and body weights between both female rhesus treatment groups. Scrambled control (n=6) and ER $\alpha$  knockdown (n=6) at the start of the study. P-values calculated via student's t-test.



**Figure 1:** Timeline of treatment onset, metabolic procedures, and end of study.





**Figure 2:** (A) Coronal anatomical MRI images used to determine targeting of the MBH prior to neural surgery. (B) During surgery, infusions contained a contrast agent (white) to enable MRI imaging of infusion placement.



**Figure 3:** Western blots validated the knockdown of ER $\alpha$  protein expression for shRNA 34 and shRNA 56 employing antibody 6F11.

**A****ERa34 oligo 1**5'-GATCCA**CCTACTACCTGGAGAACGA**CTTCCTGTCA**TCGTTCTCCAGGTAGTAGG**TTTTTTGGAAA-3'**ERa34 oligo 2**5'-AGCTTTTCCAAAAA**CCTACTACCTGGAGAACGA**TGACAGGAAG**TCGTTCTCCAGGTAGTAGG**TG-3'Oligos 1 and 2 annealed:5'-GATCCA**CCTACTACCTGGAGAACGA**CTTCCTGTCA**TCGTTCTCCAGGTAGTAGG**TTTTTTGGAAA-3'  
3'-GT**GGATGATGGACCTCTTGCT**GAAGGACAGT**AGCAAGAGGTCCATCATCC**AAAAAACCTTTTCGA-5'

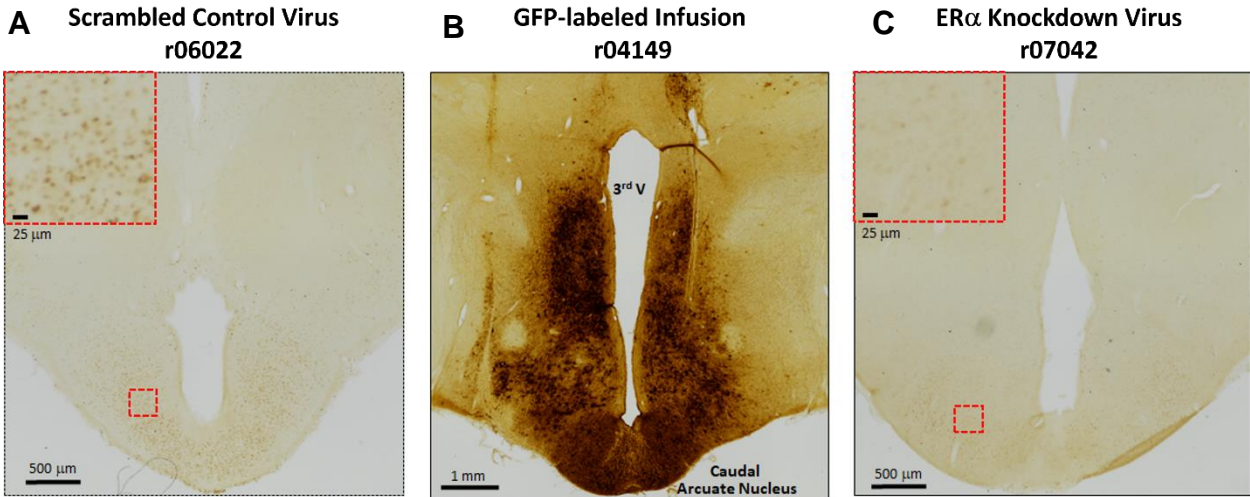
pAAV-H1-EGFP from VB: clone into 5' BamHI (GGATCC) and 3' HindIII (AAGCTT)

BamH1: GGATCC                      HindIII: AAGCTT  
          CCTAGG                        TTCGAA**B****ERa78 oligo 1**5'-GATCCG**CCTACTACCTGGAGAACGA**CTTCCTGTCA**TCGTTCTCCAGGTAGTAGG**TTTTTTGGAAA-3'**ERa78 oligo 2**5'-AGCTTTTCCAAAAA**CCTACTACCTGGAGAACGA**TGACAGGAAG**TCGTTCTCCAGGTAGTAGG**CG-3'Oligos 1 and 2 annealed:5'-GATCCG**CCTACTACCTGGAGAACGA**CTTCCTGTCA**TCGTTCTCCAGGTAGTAGG**TTTTTTGGAAA-3'  
3'-GC**GGATGATGGACCTCTTGCT**GAAGGACAGT**AGCAAGAGGTCCATCATCC**AAAAAACCTTTTCGA-5'

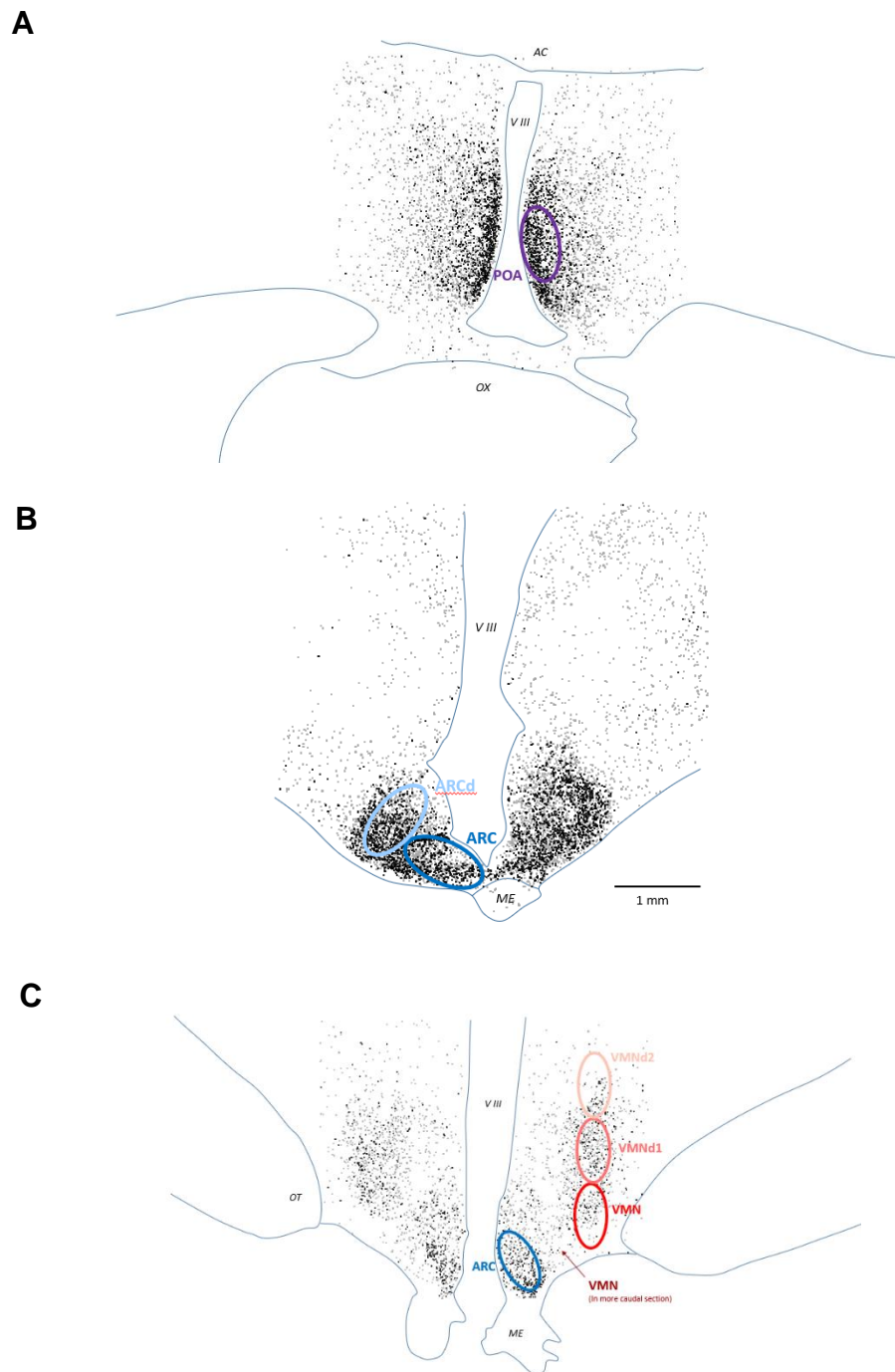
pAAV-H1-EGFP from VB: clone into 5' BamHI (GGATCC) and 3' HindIII (AAGCTT)

BamH1: GGATCC                      HindIII: AAGCTT  
          CCTAGG                        TTCGAA

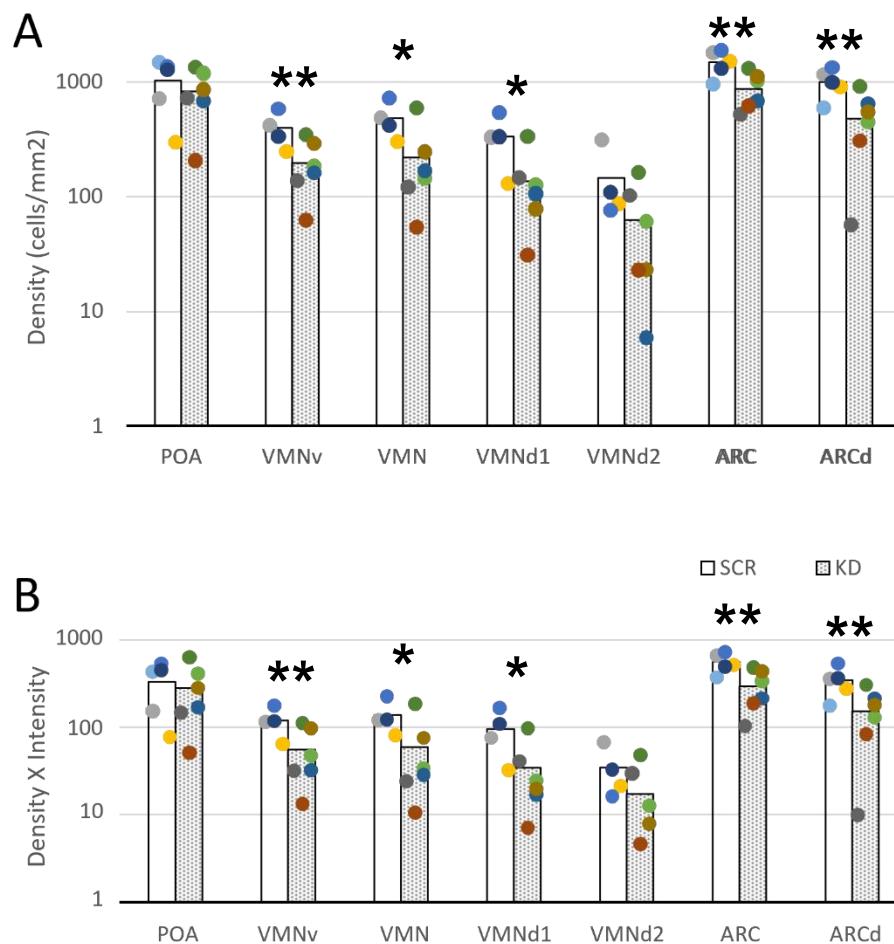
**Figure 4:** Oligo sequences for ERa34 (A) and ERa78 (B) shRNA candidates for virus production. Green highlighted is the shRNA portion. Bold in between highlighted regions are the loop.



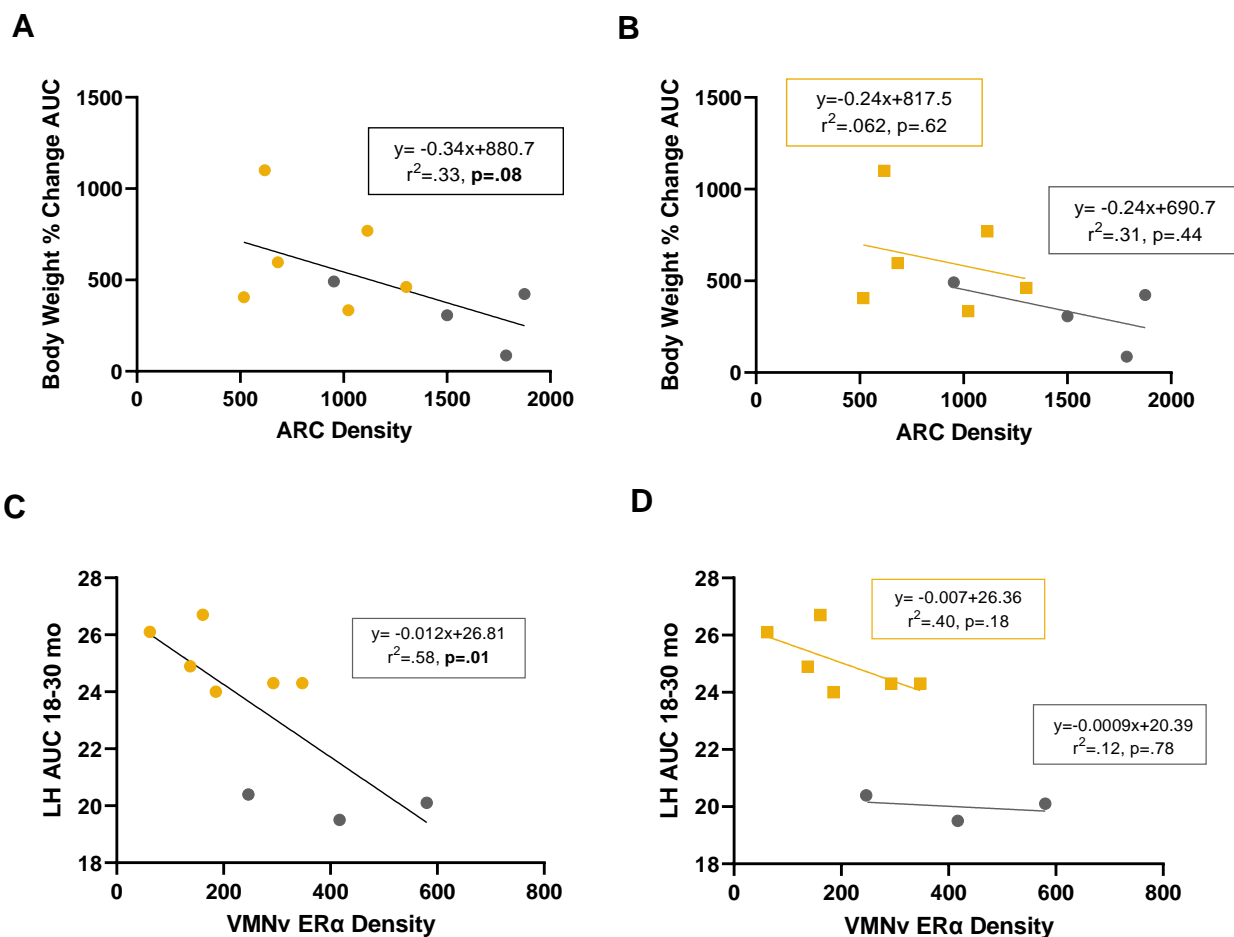
**Figure 5:** Adult rhesus macaque coronal sections from approximately the caudal extents of the ARC within the hypothalamus. Abundant GFP-immunoreactive (ir) labeling was demonstrated in ARC (B). ER $\alpha$  protein expression was abundant in scrambled control (A) and was less obvious in ER $\alpha$ KD females (C).



**Figure 6:** Regions of interest (ROIs) drawn for the POA (A, purple), ARC (B, blue) and VMN (C, red) ERα density and intensity analyses from a control monkey.



**Figure 7:** ER $\alpha$  density (A) and density\*intensity (B) for control (SCR, white bars) and ER $\alpha$ KD females (KD, shaded bars). Dots represent individual values for each monkey. Bars represent mean of each female group.  $p^* < 0.1$ ,  $p^{**} < 0.05$ .



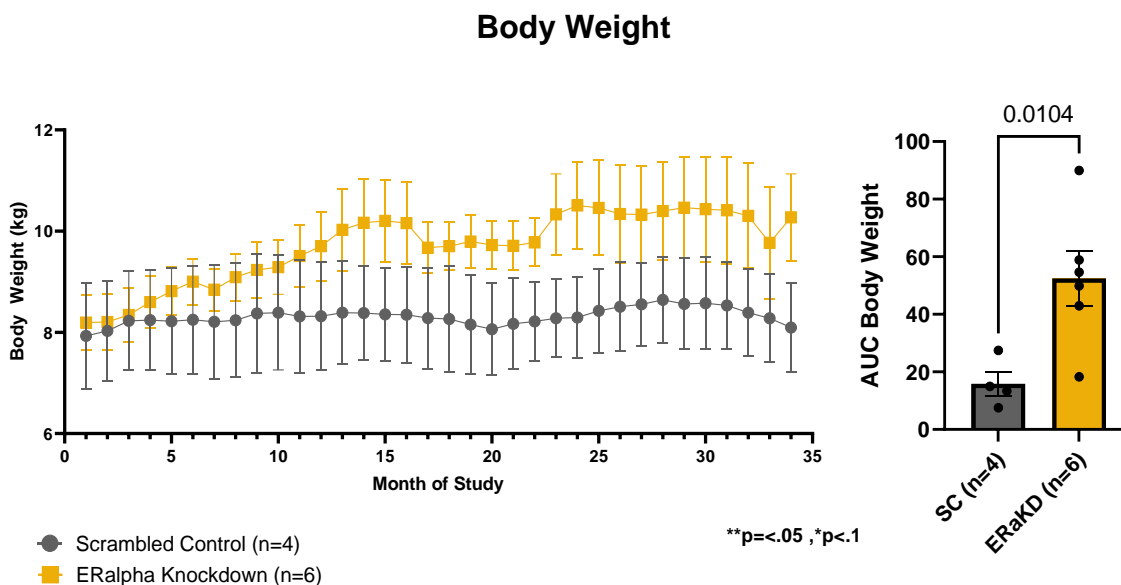
**Figure 8.** Associations between ARC ER $\alpha$  expression density and body weight percent change AUC for all females combined (A) and by female group (B). Relationships of VMNv ER $\alpha$  density and LH AUC 18-30 months for all females combined (C) and by female group (D). There was a trend ( $p=0.08$ ) towards a negative correlation between ARC ER $\alpha$  expression and body weight percent change AUC, all females combined (A) and a negative correlation ( $p=0.01$ ) between ARC ER $\alpha$  expression and LH AUC 18-30 months, all females combined (C). Scrambled control (grey) and ER $\alpha$ KD (yellow) females.

		Female Group		
		Scrambled Control (n=4)	ER $\alpha$ Knockdown (n=6)	p-value
<b>Estradiol (pg/mL)</b>	Baseline	64.85 $\pm$ 17.45	60.02 $\pm$ 2.91	0.80
	18-months	63.80 $\pm$ 16.97	59.50 $\pm$ 12.40	0.84
	30-months	77.70 $\pm$ 14.13	54.37 $\pm$ 5.25	0.20
<b>Estrone (pg/mL)</b>	Baseline	37.85 $\pm$ 12.10	23.80 $\pm$ 3.25	0.33
	18-months	12.48 $\pm$ 3.57	8.87 $\pm$ 1.75	0.41
	30-months	13.93 $\pm$ 6.10	13.05 $\pm$ 4.19	0.91
<b>Testosterone (ng/mL)</b>	Baseline	0.24 $\pm$ 0.03	0.20 $\pm$ 0.03	0.32
	18-months	0.16 $\pm$ 0.01	0.17 $\pm$ 0.02	0.64
	30-months	0.17 $\pm$ 0.01	0.18 $\pm$ 0.12	0.87
<b>Androstenedione (ng/mL)</b>	Baseline	0.44 $\pm$ 0.03	0.42 $\pm$ 0.05	0.80
	18-months	0.25 $\pm$ 0.03	0.31 $\pm$ 0.03	0.20
	30-months	0.33 $\pm$ 0.05	0.37 $\pm$ 0.04	0.50
<b>DHEA (ng/mL)</b>	Baseline	20.79 $\pm$ 2.85	14.84 $\pm$ 1.15	0.13
	18-months	12.60 $\pm$ 3.00	14.87 $\pm$ 2.89	0.60
	30-months	21.04 $\pm$ 4.13	20.70 $\pm$ 2.98	0.95
<b>LH (ng/mL)</b>	Baseline	1.63 $\pm$ 0.29	1.38 $\pm$ 0.07	0.48
	18-months	1.75 $\pm$ 0.29	1.82 $\pm$ 0.16	0.85
	30-months	1.60 $\pm$ 0.20	2.17 $\pm$ 0.26	0.12
<b>Progesterone (ng/mL)</b>	Baseline	0.24 $\pm$ 0.03	0.20 $\pm$ 0.03	0.32
	18-months	0.15 $\pm$ 0.03	0.15 $\pm$ 0.03	0.93
	30-months	0.17 $\pm$ 0.01	0.18 $\pm$ 0.01	0.87

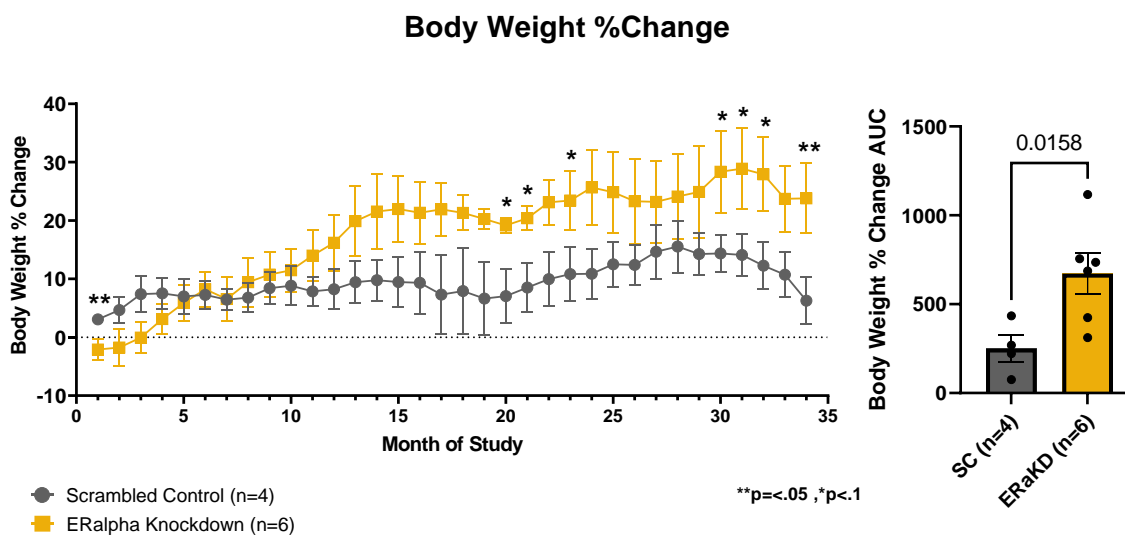
**Table 2:** Circulating steroid hormones during the follicular phase at baseline, 18-months, and 30-months post-treatment onset. Significance assessed via student's t-test. Values represented as mean  $\pm$  SEM.



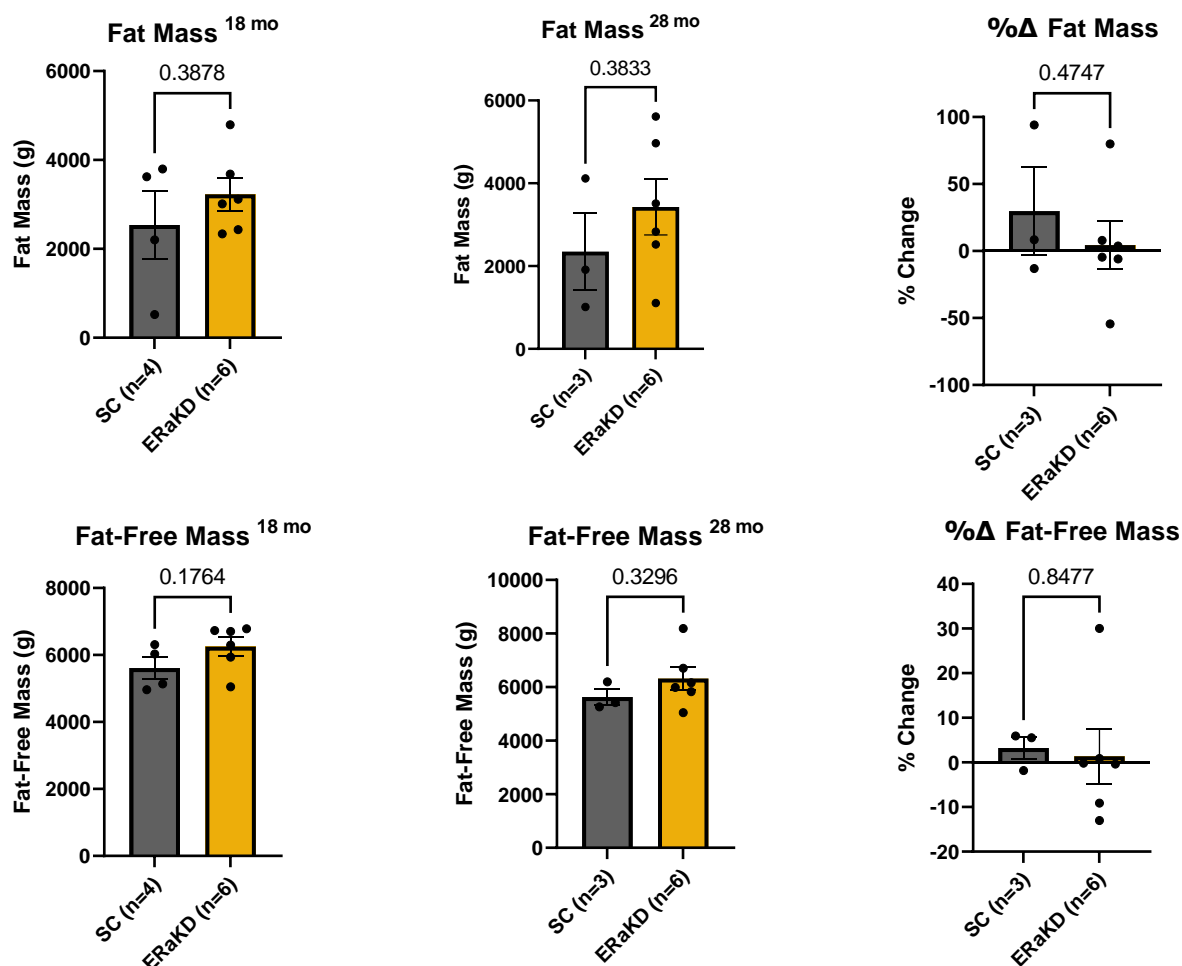
A



B



**Figure 9.** Body weight (A) and body weight percent change (B) over 34-months of the study with corresponding area under the curve (AUC) measurements. Asterisks represent months significant, ERαKD compared to control females from repeated-measures ANOVA time effect (\*p<0.1, \*\*p<0.05). Plots are expressed as means±/ SEM.



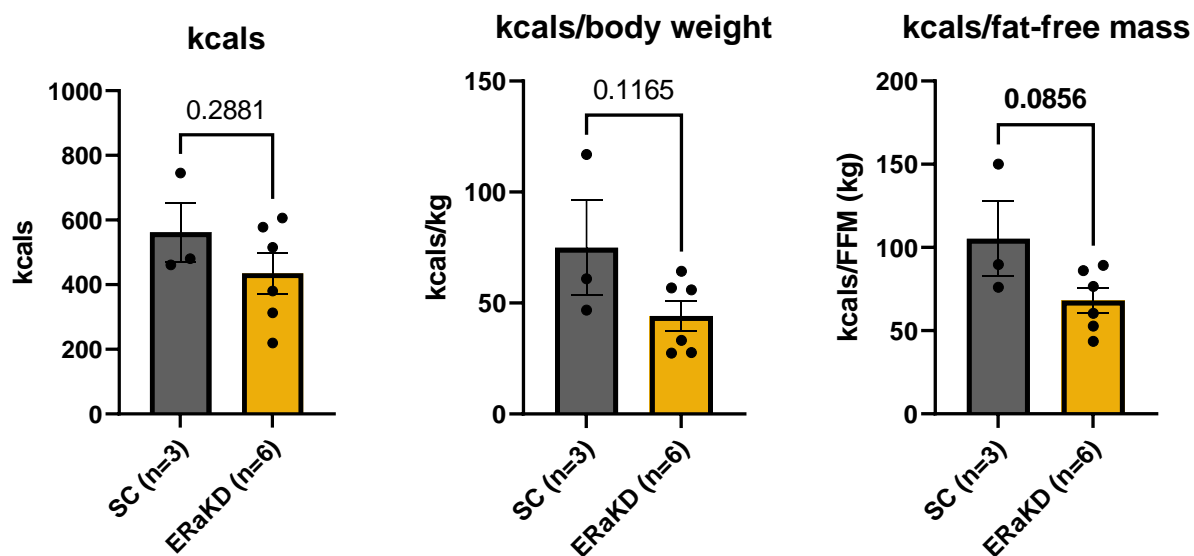
**Figure 10:** Total fat mass and fat-free mass from DXA scan at 18-months and 28-months of the study with corresponding percent change between the two scans.

There were no significant differences between the female groups. Significance assessed via student's t-test. Plots are expressed as means $\pm$ SEM.

		Treatment Group					
		Scrambled Control (n=4)		ER $\alpha$ Knockdown (n=6)		p-value	p-value
		Fat Mass (kg)	Fat-Free Mass (kg)	Fat Mass (kg)	Fat-Free Mass (kg)	Fat-Mass	Fat-Free Mass
<b>Abdomen</b>	18-mo	1.07 $\pm$ 0.38	<b>1.67<math>\pm</math>0.07</b>	1.33 $\pm$ 0.21	<b>2.04<math>\pm</math>0.11</b>	0.58	<b>0.02*</b>
	28-mo	0.87 $\pm$ 0.40	1.68 $\pm$ 0.07	1.32 $\pm$ 0.30	2.05 $\pm$ 0.23	0.41	0.17
<b>Upper Legs</b>	18-mo	0.30 $\pm$ 0.09	1.13 $\pm$ 0.17	0.40 $\pm$ 0.06	1.22 $\pm$ 0.07	0.38	0.63
	28-mo	0.29 $\pm$ 0.10	1.08 $\pm$ 0.21	0.38 $\pm$ 0.09	1.26 $\pm$ 0.11	0.54	0.50
<b>Chest</b>	18-mo	0.56 $\pm$ 0.18	1.12 $\pm$ 0.13	0.80 $\pm$ 0.09	1.21 $\pm$ 0.08	0.29	0.70
	28-mo	0.60 $\pm$ 0.27	1.09 $\pm$ 0.03	0.97 $\pm$ 0.18	1.23 $\pm$ 0.07	0.33	0.13
<b>Lower Legs</b>	18-mo	0.19 $\pm$ 0.04	0.53 $\pm$ 0.04	0.21 $\pm$ 0.02	0.52 $\pm$ 0.04	0.61	0.92
	28-mo	0.19 $\pm$ 0.03	0.58 $\pm$ 0.04	0.21 $\pm$ 0.03	0.55 $\pm$ 0.04	0.70	0.61
<b>Arms</b>	18-mo	0.29 $\pm$ 0.03	0.71 $\pm$ 0.01	0.33 $\pm$ 0.03	0.75 $\pm$ 0.04	0.63	0.39
	28-mo	0.25 $\pm$ 0.09	0.72 $\pm$ 0.02	0.37 $\pm$ 0.09	0.71 $\pm$ 0.04	0.42	0.80

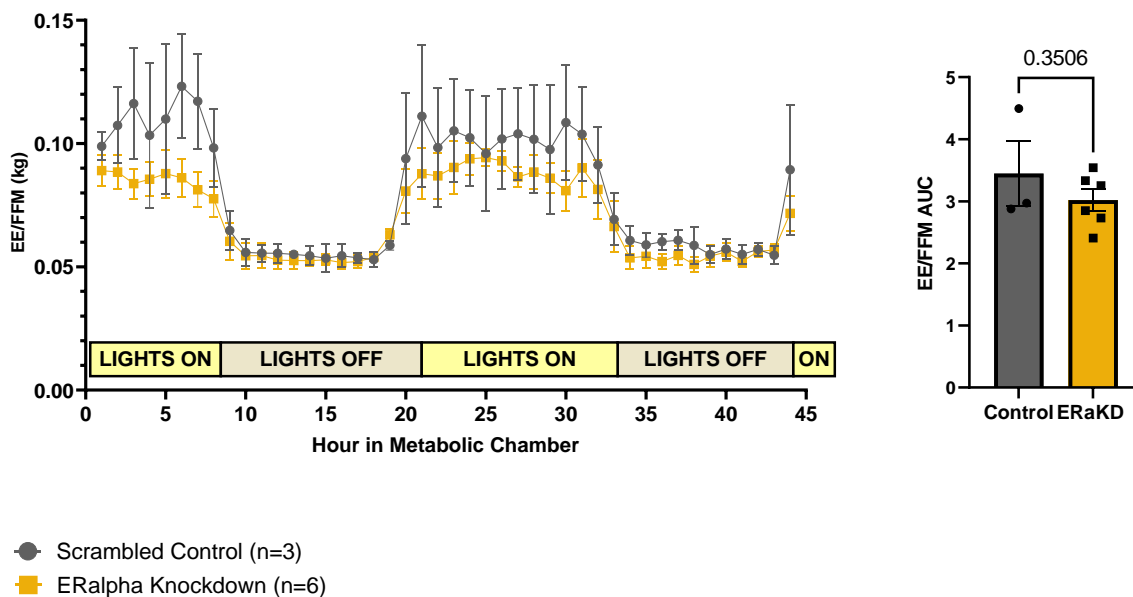
**Table 3:** Regions of interest analyzed at ~18-months and ~28-months post-treatment onset via DXA scan. Significance assessed via student's t-test ( $p^* < 0.05$ ) between ER $\alpha$ KD and control females. Values represented as mean  $\pm$  SEM.

## Average Daily kcals Consumed

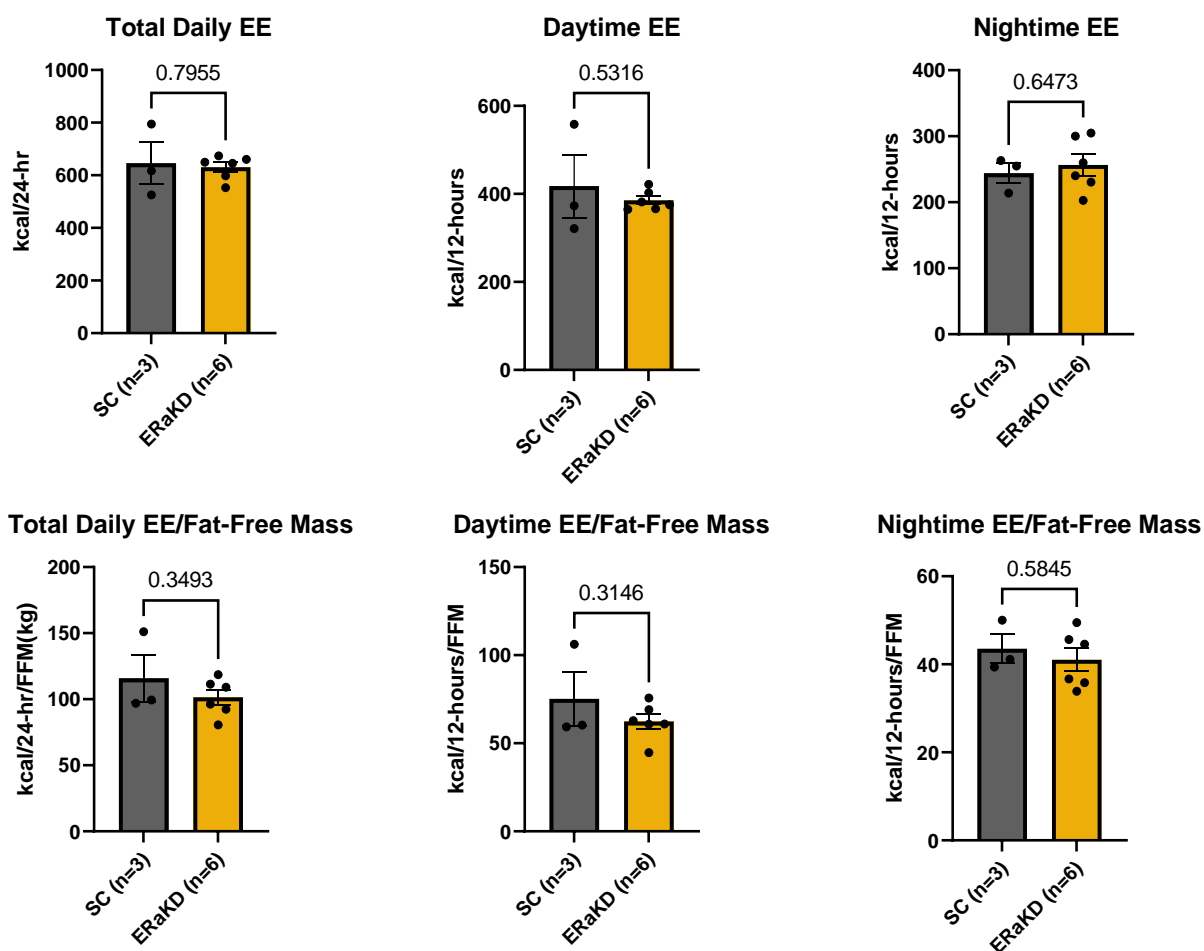


**Figure 11:** Average kcals consumed throughout the food assessment without adjustment and with body weight (kg) and fat-free mass (kg) adjustments. Trend ( $p=.08$ ) towards reduced kcals consumed when adjusted for kilogram of fat-free mass in ERαKD compared to control females. Significance assessed via student's t-test. Plots are expressed as means $\pm$ SEM.

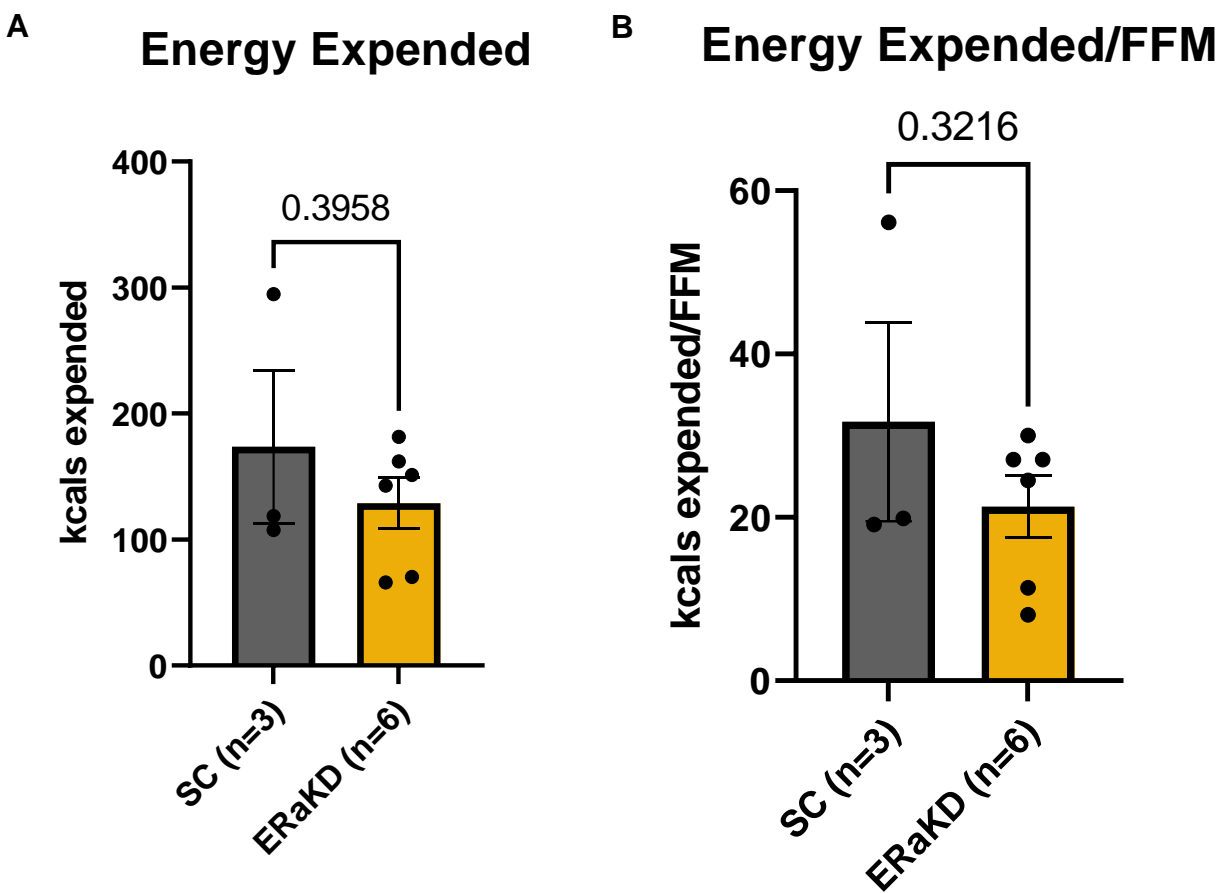
## Energy Expenditure Adjusted for Fat-Free Mass



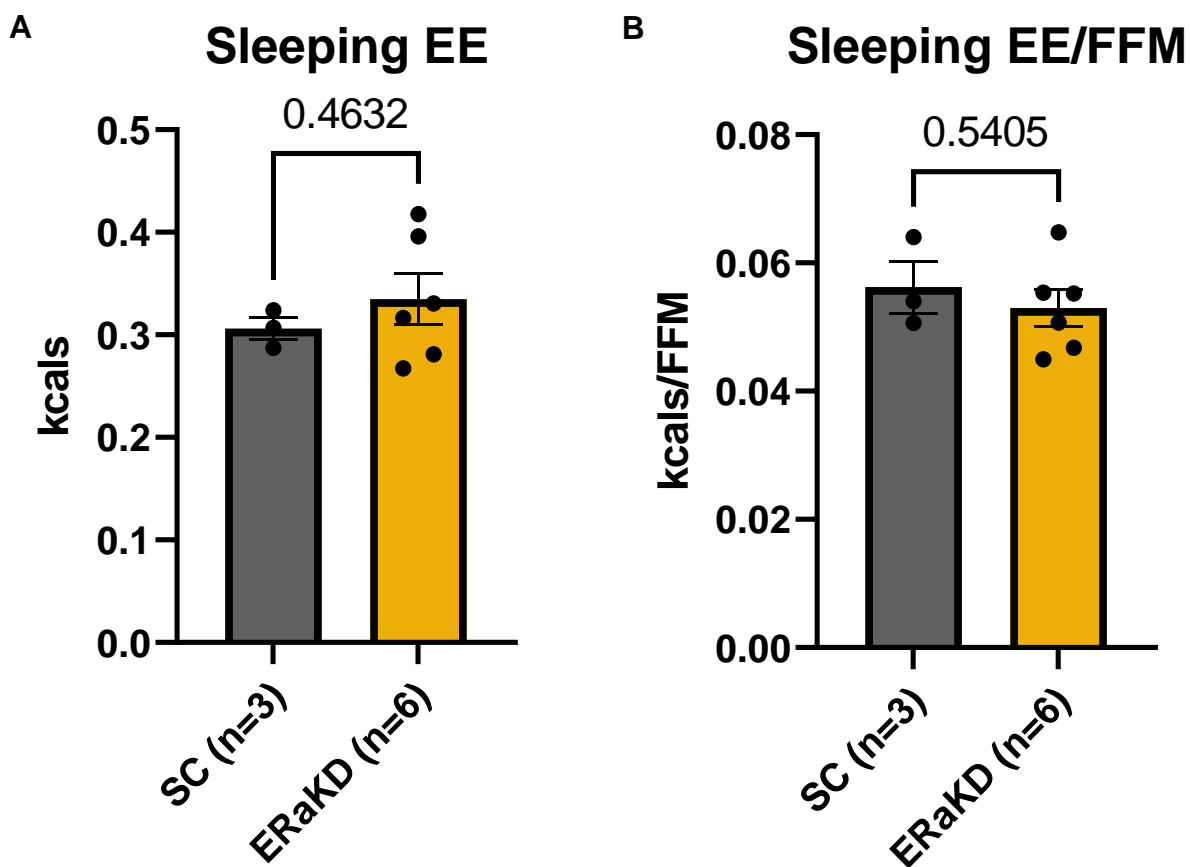
**Figure 12.** Averaged hourly energy expenditure (EE) adjusted for kilogram of fat-free mass (FFM) with corresponding area under the curve (AUC) measurement. There were no between female group differences. Significance assessed via student's t-test and RM-ANOVA. Plots are expressed as means $\pm$ SEM.



**Figure 13.** Summaries of energy expenditure over 24-hr, daytime (0600-1800) and nighttime (1800-0600) periods within the metabolic chamber with and without adjustments for fat-free mass (kg). There were no observed differences between female groups. Significance assessed via student's t-test. Plots are expressed as means $\pm$ SEM.

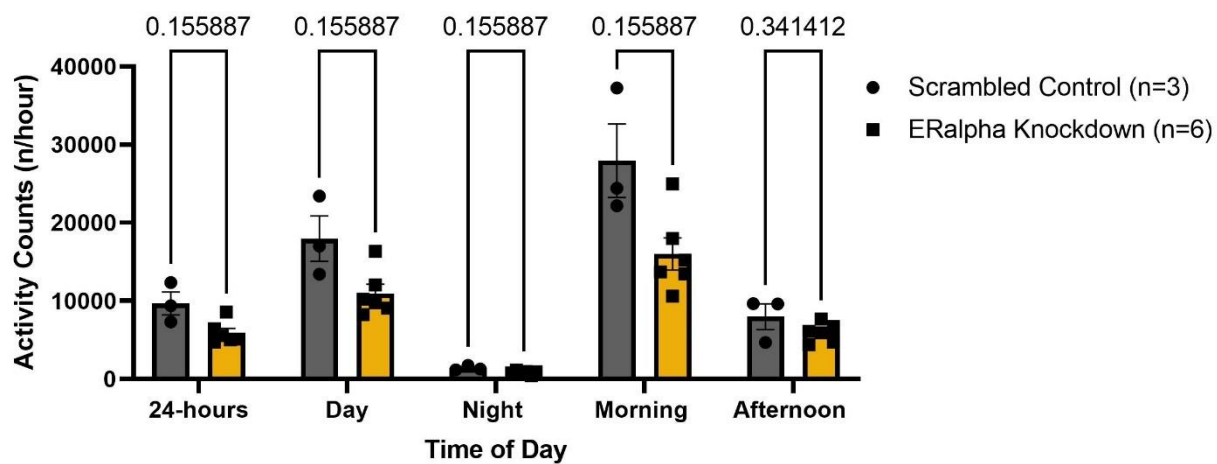


**Figure 14.** Average energy expenditure over duration in metabolic chamber with and without correction for fat-free mass (kg). No observed differences between female groups. Significance assessed via student's t-test. Plots are expressed as means $\pm$ SEM.

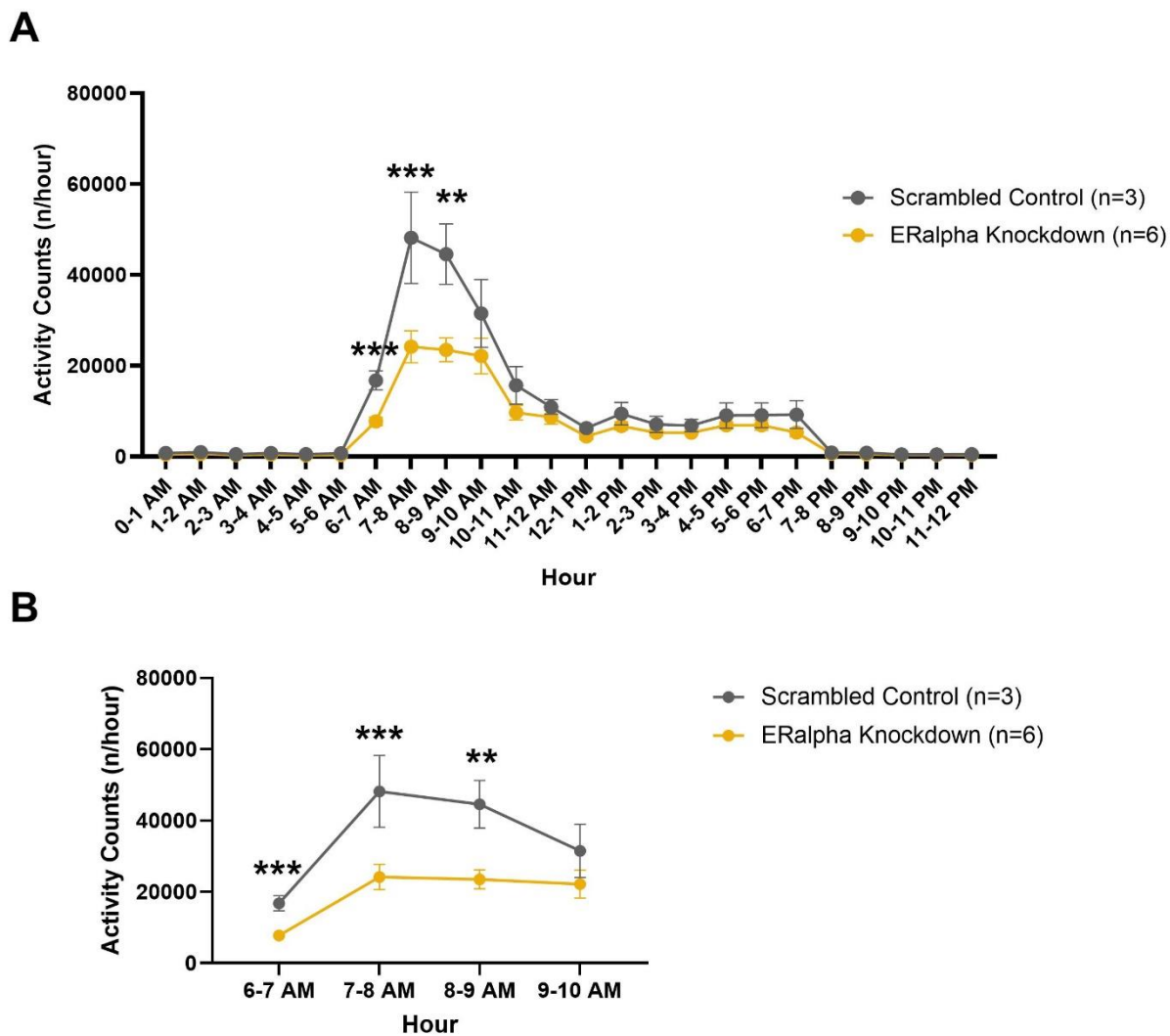


**Figure 15.** Average sleeping energy expenditure averaged over both nights in the metabolic chamber with and without adjustment for fat-free mass (kg). No observed differences between female groups. Significance assessed via student's t-test. Plots are expressed as means $\pm$ SEM.

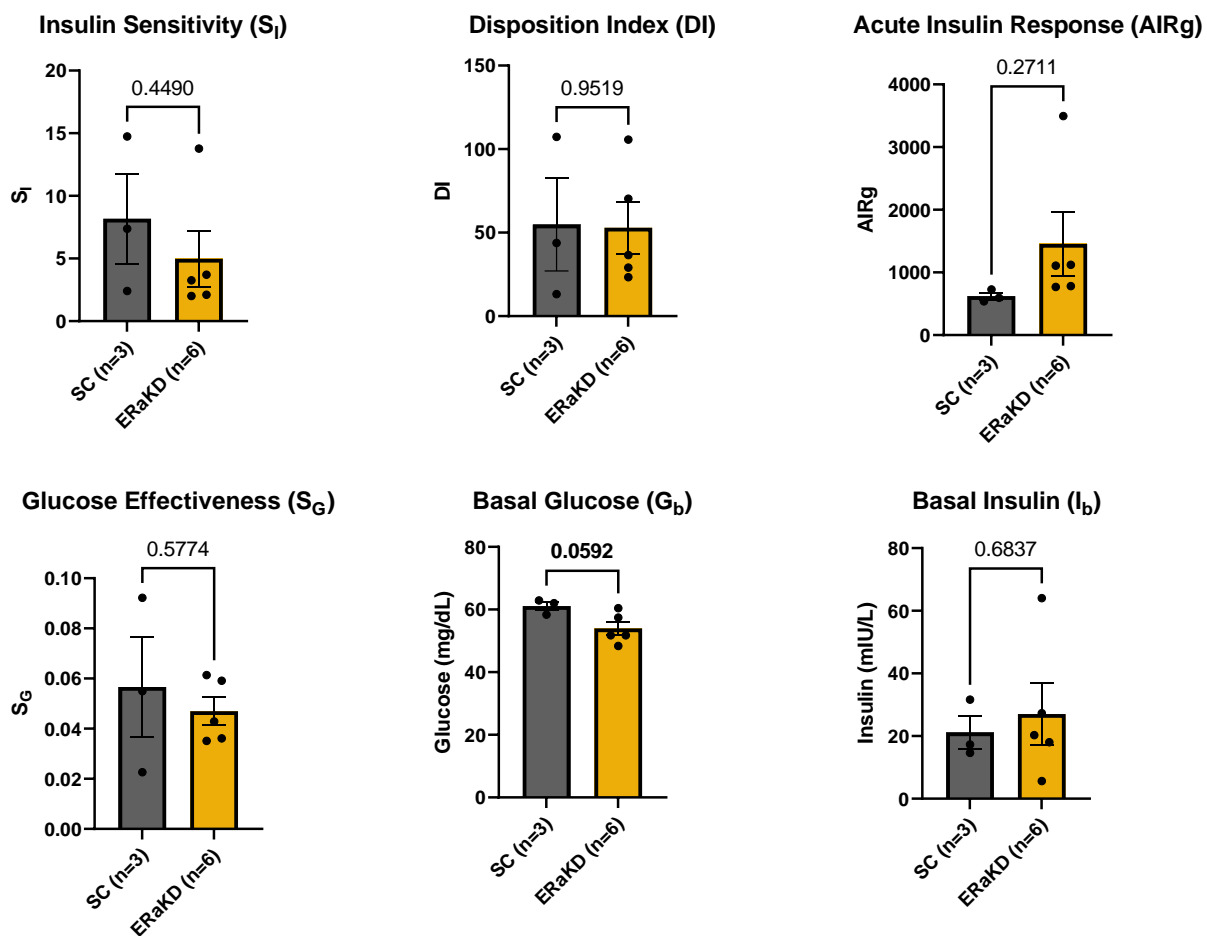




**Figure 16:** Activity counts averaged over 24-hours, Day (0600-1800), Night (1800-0600), Morning (0600-1200) and Afternoon (1200-1800). There were no observed differences between female groups. Significance assessed via student's t-test. Plots are expressed as means $\pm$ SEM.



**Figure 17:** Activity counts (n) averaged per hour over the duration of the locomotor assessment (A) and activity counts over the morning period, 6 AM – 10 AM (B). Significance assessed via RM-ANOVA ( $p^* < 0.05$ ,  $p^{**} < 0.001$ ,  $p^{****} < 0.0001$ ) ER $\alpha$ KD compared to control females. Plots are expressed as means $\pm$ -SEM.



**Figure 18.** Summary of values computed from the Bergman Modified Minimal Model (MMM) for glucoregulation. No observed differences between female groups aside from a trend towards reduced basal glucose in ERαKD compared to control females ( $p=.0592$ ). Significance assessed via student's t-test. Plots are expressed as means $\pm$ SEM.

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**4. CHAPTER FOUR: Ovarian and Extra-Ovarian Estradiol Depletion Both  
Contribute to Bone Loss but not to Body Weight Gain or Diminished Energy  
Expenditure in Adult Female Rhesus Monkeys**

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[Manuscript is being prepared for submission to the Journal of the Endocrine Society.  
Contribution to this manuscript: Designed/carried out experiments, analyzed data, and  
prepared the manuscript for this thesis and subsequent publication]



#### 4.1 Abstract

Declining serum estradiol (E<sub>2</sub>) levels during the menopausal transition are associated with increased central adiposity and heightened risk for metabolic disease. Ovarian estradiol, E<sub>2</sub>, supports female metabolic function in rodents. While ovariectomy (OVX) in rodents enables obesity, OVX in nonhuman primates (NHPs) inconsistently alters weight gain. We therefore hypothesized that in female NHPs, extra-ovarian E<sub>2</sub> provides key support for metabolic homeostasis. To test this, we employed aromatase inhibition, via letrozole, to eliminate extra-ovarian E<sub>2</sub> biosynthesis. Nineteen adult female rhesus monkeys were OVX and received: (1) E<sub>2</sub>-containing capsules and daily oral treatments of letrozole (E<sub>2</sub>; n=5); empty capsules and daily oral treatments of either vehicle (VEH, 1ml vehicle/kg, n=7), or (3) letrozole (LET, 1 mg/kg in 1ml vehicle/kg, n=7). After 12 months, we demonstrated reduced percent change of bone mineral density in LET (p=0.005) and VEH (0.06, trend) females compared to E<sub>2</sub> females. We also demonstrated a trend (p=0.08) in reduced glucose effectiveness for VEH and LET compared to E<sub>2</sub> females at 12-months, despite no other differences in metabolic components such as body weight, body composition and locomotor activity. Our findings demonstrate that ovarian and ovarian + extra-ovarian E<sub>2</sub> depletion results in decreased bone mineral content and bone mineral density and contributes to reduced ability for glucose uptake independent of insulin action in adult female rhesus macaques.

## 4.2 Introduction

Women largely spend the last third of their lives in a post-menopausal state. Menopause is characterized as the cessation of menstrual cyclicity and ovarian estrogen production. The depletion of the ovarian follicular pool and consequent reduction of circulating estradiol ( $E_2$ ) during menopause has been suggested by some investigators to contribute to post-menopausal weight gain, reduced energy expenditure, osteoporosis, hot flashes, cardiovascular disease, and cognitive decline<sup>1-6</sup>. There is greater consensus supporting the idea that menopause is associated with an increase in visceral relative to subcutaneous adiposity, with or without an increase in overall body weight<sup>7</sup>. Women experiencing personally distressing symptoms of menopause are prescribed hormonal replacement therapy (HRT), comprising of estrogen only therapies and combinations of estrogens and progestins, including synthetic forms of progesterone, to alleviate symptoms and help reduce osteoporosis induced by declining circulating  $E_2$  levels. While HRT has been shown in some studies<sup>8-9</sup>, but not others<sup>10-12</sup>, to reduce weight gain and slow skeletal bone loss, HRT use has been greatly reduced since the WHO-sponsored studies originally reported small but significant increases in the risks of blood clots, dementia, and breast cancer<sup>13-14</sup>.

$E_2$  supports optimal female metabolic function in multiple female laboratory animals, including, mice, rats, and sheep<sup>15-17</sup>.  $E_2$  action is mediated through estrogen receptors, predominately, estrogen receptor alpha ( $ER\alpha$ ) and estrogen receptor beta ( $ER\beta$ ).  $ER\alpha$  has been shown extensively to be the predominant estrogen receptor regulating  $E_2$  action in rodents<sup>18</sup> and as we have recently shown,  $ER\alpha$ , is an important hypothalamic regulator of metabolic homeostasis in adult female rhesus macaques

(Chapter 3). Thus, while ovarian E<sub>2</sub> has been established as a major contributor to female metabolism in female rodents and sheep, its role in female NHPs is less well understood. While ovariectomy (OVX) in female rodents enables obesity, OVX in female NHPs inconsistently alters weight gain<sup>18-20</sup>, highlighting the possibility of E<sub>2</sub> produced outside the ovaries as an important site of E<sub>2</sub> synthesis signaling through ER $\alpha$  in female NHPs.

Aromatase, a cytochrome P450 enzyme, encoded by the CYP19A1 gene, converts testosterone (T) to E<sub>2</sub>, as well as androstenedione (A<sub>4</sub>) to estrone (E<sub>1</sub>). The ovaries serve as the major site of E<sub>2</sub> synthesis in female non-primates, NHPs and humans. E<sub>2</sub>, however, is also produced in various other tissues in which aromatase is expressed, such as adipose depots, adrenal glands, liver, skin, and in various brain regions. In particular, neural production of E<sub>2</sub> (neuro-E<sub>2</sub>) has been characterized in birds, rats, and in NHPs. Brain aromatase is expressed at its highest levels in the medial basal hypothalamus (MBH), preoptic area (POA) and amygdala in rodents, monkeys, and humans.

We hypothesize that the failure for OVX to increase female NHP body weight is due to the residual actions of neuro-E<sub>2</sub> orchestrating energy balance in the hypothalamus of OVX female NHPs. Rodent research models have provided comprehensive evidence supporting roles played by ovarian E<sub>2</sub> in female energy balance and body composition, yet there are few studies in female NHPs assessing ovarian and extra-ovarian E<sub>2</sub>, possibly neuro-E<sub>2</sub>, in regulating aspects of energy balance. To test this hypothesis, we employed the aromatase inhibitor, letrozole, to diminish systemic and central E<sub>2</sub> production in OVX adult female rhesus macaques to

enable investigation into extraovarian E<sub>2</sub> source(s) regulating of female metabolism in a NHP. We propose that an extra-ovarian source of E<sub>2</sub>, likely from the brain, will be greatly diminished or extinguished by body-wide aromatase inhibition and resulting in metabolic dysfunction via increased body weight, increased adiposity and reduced energy expenditure in female rhesus macaques compared with both E<sub>2</sub> replaced females and those experiencing depleted ovarian E<sub>2</sub>, alone.

### **4.3 Methods**

#### *Animals*

Nineteen adult female rhesus macaques (5-11 years of age) from the Wisconsin National Primate Research Center colony were all OVX assigned to one of three female groups so that age, body weight and body mass index (BMI) were comparable (Table 1): 1) OVX + letrozole + E<sub>2</sub> replacement via subcutaneous capsules (E<sub>2</sub>, n=5), 2) OVX + vehicle subcutaneous capsules (VEH, n=7) and 3) OVX + letrozole + vehicle capsules (LET, n=7) (Table 1). E<sub>2</sub> replacement was achieved through subcutaneous E<sub>2</sub>-filled capsules that maintained a sufficient systemic level of E<sub>2</sub> (40-100 pg/mL) to suppress gonadotropin secretion and resembled E<sub>2</sub> values observed during the early-to-late follicular of the menstrual cycle<sup>21</sup>. To maintain constant E<sub>2</sub> levels, capsules were replaced every 3 months throughout the study. OVX females received a daily oral dose of 1ml/kg low calorie pudding as a vehicle control. LET and E<sub>2</sub> females were given a daily oral dose of 1 mg/kg of letrozole dissolved in 1ml/kg low calorie pudding. Monkeys were maintained in these three groups for ~17 months, pair-housed, in enclosures under 12-h lighting, ambient temperature of ambient temperature of ~27°C and humidity of ~50%. (Figure 1). The protocol for this study was approved by the Animal Care and Use Committee of the

Office of the Vice Chancellor for Graduate Research and Education of the University of Wisconsin–Madison, an AAALAC-accredited program.

### *Diet*

Monkeys were fed chow twice a day, between 0700h-0900h and 1200h-1400h (Teklad Global 20% Protein Primate Diet, ENVIGO, Madison, WI). The calorie distribution of the biscuits was 29% protein, 14% fat and 57% carbohydrates with an energy density of 2.8 kcal/g. Monkeys were supplemented with daily fruit enrichment (<100 kcals), weekly tactile enrichments (<100 kcals), and weekly destructible enrichments (<100 kcals).

### *Hormone and Metabolic Analyte Determinations*

Steroid hormone analysis was adapted from methods previously described<sup>22,23</sup>. Briefly, internal standard was added to rhesus serum samples (1mL) and then they were extracted twice using methyl tert butyl ether followed by dichloromethane. Samples were derivatized using dansyl chloride and then analyzed by LC-MS/MS (Sciex QTRAP 5500). Individual calibration curves were constructed for each analyte with at least 8 points. The linearity was  $r > 0.9990$  and the curve fit was linear with 1/x weighting. None of the compounds of interest were detected in blank or double blank samples. Inter-assay coefficient of variation was determined by 3 levels of human serum and ranged from 5.6-9.7%.

Serum samples were analyzed for estradiol (25uL) and progesterone (20uL) using a cobas e411 analyzer equipped with ElectroChemiLuminescence technology (Roche, Basal, Switzerland) according to manufacturer instructions. Results were determined via a calibration curve which was instrument generated by 2-point calibration using traceable

standards and a master curve provided via the reagent barcode. Inter-assay coefficient of variation (CV) was determined by a pool of rhesus plasma. For estradiol, the limit of quantitation (LOQ) was 25 pg/mL, the intra-assay CV was 2.02%, and the inter-assay CV was 5.05%. For progesterone, the LOQ was 0.2 ng/mL, the intra-assay CV was 1.37%, and the inter-assay CV was 4.63%.

Metabolic peptides were measured with a Milliplex Non-Human Primate Metabolic Magnetic Bead Panel (NHPMHMAG-45K; EMD Millipore). The assay was run according to the kit insert using a BioPlex 200 (BioRad). The intra-assay CV was determined by two pools of rhesus serum and ranged from 2.2-9.3%.

#### *Hypothalamic Hormone Determinations*

The method for measurement of steroid hormone measurement in the hypothalamus was adapted from Bertin et al., 2015<sup>24</sup>. Briefly, up to 100 mg of brain tissue was homogenized in methanol. Internal standard was added, and the methanol was run through solid phase extraction using Oasis HLB cartridges (Waters Corporation). The eluate was derivatized and analyzed by LC-MS/MS (Sciex 6500+). Individual calibration curves were constructed for each analyte with at least 8 points. The linearity was  $r > 0.9990$  and the curve fit was linear with 1/x weighting. Inter-assay coefficient of variation was determined by two pools of spiked macaque cerebellum and ranged from 8.9-17.2%.

#### *Body Weight, Body Composition & Bone Mass*

All monkeys were weighed weekly. Incremental area under the curve (AUC) assessment of weight parameters over time, calculated by the trapezoid rule, was used to better detect recurring differences in weight gain, as previously employed by Risal

and colleagues (2019). At baseline, 6-months, and 12-months post-OVX, total body composition was assessed by dual-energy X-ray absorptiometry (DXA, iDXA, GE/Lunar Corp., Madison, WI) on sedated animals. Monkeys received up to 7 mg/kg ketamine IM and up to 0.03 mg/kg dexmedetomidine IM, and the latter was reversed on conclusion of the procedure by up to 0.3 mg/kg atipamezole IM. After placing the anesthetized animal onto the scanner bed, DXA-compliant cushioning was used to position the animal with arms and tail flat on the scanner bed, clearly distinct from the monkey's trunk and other extremities. A DXA scan was then performed for approximately 30 minutes. The scanner room was maintained at 27-28°C during the performance of all scans to minimize the risk of hypothermia. Monitoring of anesthesia recovery was documented every 15 min until the monkey was sitting upright, then every 30 minutes until it was fully recovered from the anesthesia. Fat mass, fat-free mass, bone mineral content (BMC) and bone mineral density (BMD) were determined for total body as well as regions of interest, including abdomen, chest, thighs and extremities. Body mass index was calculated using  $\frac{\text{Body Weight (kg)}}{\text{Crown Rump Length (m}^2\text{)}}^{25}$ .

### *Somatometric Measurements*

Somatometric parameters were measured using tape measure or calipers while monkeys were anesthetized for DXA imaging. Measurements included femur length; abdomen, chest, pelvis, arm, leg, and head circumferences; head diameter; and chest and abdomen skinfolds. Additionally, monkeys were placed on an osteometric board for measurement of crown rump and crown-heel lengths.

### *Caloric Intake Assessment*

At baseline, 6-months post-OVX and 12-months post-OVX, monkeys were singly housed 0700h-1800h for 7-21 days. Animals were fed 9 biscuits of standard chow twice a day at 0700h-0900h and 1200h-1400h. At each feeding, monkeys received on average ~423 kilocalories from chow. In addition, animals received daily fruit enrichment (i.e., pear, banana, green pepper, kiwi and apple) totaling on average ~100 kilocalories per enrichment occurrence. Daily chow and enrichment were weighed prior to being given each day (Calories Offered). At 1800h, any chow or enrichment not consumed from the day was weighed (Calories Left) and returned to each monkey for the overnight period (1800h-0700h). Calories Consumed was calculated by subtracting Calories Left from Calories Offered. The weight of chow of Calories Consumed was multiplied by the chow's energy density of 2.8 kcal/g to convert to calories. Monkeys were repaired with their partner for the overnight period (1800h-0700h).

### *Locomotor Activity Assessment*

At baseline, as well as at 6-months and 12-months post-OVX accelerometers were attached to a fabric collar and placed around each monkey's neck. Activity and intensity of movement were recorded over a ~4-week period after which the accelerometers were removed. Activity was recorded in counts (n), where one count represented body movements on an x, y, or z axis. The accelerometer sampled such activity counts every minute and these data were averaged for every hour (h), day (during lights on, 0600h-1800h), night (during lights off, 1800h-0600h), morning (0600h-1200h), afternoon (1200h-1800h), and over 24 hours. Incremental areas under the



curve activity values were also assessed to detect cumulative differences over the course of the study.

#### *Doubly Labeled Water (DLW) Procedures*

Total energy expenditure was measured during a 4-day period by the two-point doubly-labeled water (DLW) methodology<sup>26</sup>. Monkeys were injected IV with 1.0 g/kg of body weight of a premixed cocktail. The mixture comprised 0.147 g of 98 APE 18O water (Medical Isotopes, Inc., Pelham, NH), 0.083 g of at 99.9 APE deuterium oxide (2H<sub>2</sub>O, Cambridge Isotope Laboratories, Andover, MA), 0.650g of sterile water, and 0.125 g of 7.2% hypertonic NaCl to achieve physiological osmolarity. Blood samples were collected from non-anesthetized monkeys prior to (baseline), two hours, and four days after DLW injection. Immediately after blood collection, the samples were centrifuged for 10 min for serum separation. Serum was stored at -20°C in cryogenically stable tubes until analysis by isotope ratio mass spectrometry.

Water from serum samples was extracted by centrifugation (4°C, 30 minutes at 4,500 rpm, RCF 3,100 ×g, Centrifuge Marathon 21k/BR, Thermo Fisher Ltd) on regenerated cellulose filters (Amicron Ultra-4, Ultracel-30K, Merk Millipore Ltd). Isotope enrichments were determined as previously described (Blanc et al., 2003). Carbon dioxide production was calculated according to the equation proposed by Speakman et al. (2021):  $r\text{CO}_2 = 0.4554 * N * [(1.007 * k_o) - (1.043 * k_d)] * 22.26$ , in which total body water (N) was calculated by the average of dilution spaces of stable isotopes (2H<sub>2</sub>18O) obtained by the plateau method (Coward et al., 1990) using the 2-h post-dose sample and corrected for isotope exchange by the factors 1.043 and 1.007, respectively (Speakman, et al 2021). The observed isotope dilution space ratio was  $1.037 \pm 0.019$

(mean  $\pm$  SD). The isotope constant elimination rates ( $k_o$  and  $k_d$ ) were calculated by linear regression of the natural logarithm of isotope enrichment as a function of elapsed time from 2h samples. TEE was calculated by Wier's equation (1949) using a food quotient of 0.93 estimated from the animal's diet.

#### *Intravenous Glucose Tolerance Test (ivGTT)*

At baseline, 6-months post-OVX and 12-months post-OVX, an ivGTT was conducted according to the Modified Minimal Model (MMM) protocol as adapted for rhesus monkeys<sup>27</sup>. A catheter was inserted through the saphenous or femoral vein and the tip positioned in the inferior vena cava. Four pretreatments ~1.5 ml blood samples were taken at approximately -15, -10, -5, and -1 minutes to establish fasted baseline values of insulin and glucose. A ~300 mg/kg glucose bolus was administered over about one minute starting at ~ 0 min, and ~1.5 ml blood samples were drawn at 2, 3, 4, 5, 6, 8, 10, 12, 14, 16, 19, 22, 23, 24, 25, 27, 30, 40, 50, 60, 70, 80, 90, 100, 120, 140, 160, and 180 minutes after glucose administration. Blood volume availability for individual animals was calculated by body weight (animal's body weight (kg) x 60 = estimated total volume of blood). If a monkey's blood volume availability did not meet the required volume for the full ivGTT (48 ml), samples at timepoints of 100, 120, 140 and/or 160 were omitted. Twenty minutes after glucose bolus administration, tolbutamide (~25mg/ml up to a maximum volume of ~5ml 1N sodium hydroxide, 0.1N hydrochloric acid in sterile saline solution) was administered via the catheter to induce an additional increment in insulin secretion to enhance MMM calculations. Maintenance of catheter patency and fluid volume was accomplished with intravenous Plasma-Lyte A provided by an infusion pump hooked up to the indwelling catheter. Additional fluid replacement

was provided by 1-2ml cannula line flushes following each blood sample withdrawal. The duration of anesthesia was approximately 240 minutes. Monkeys were maintained under warm heat lamps (with a 24-inch minimum space between the animal and the heating lamp), a heated air warming unit and/or a water-circulating heating pad. During the procedure, the monkey's blood oxygen, heart rate and rectal temperature were continuously monitored and recorded every 15 minutes. After completion of the procedure, progress of anesthesia recovery was documented every 15 min until the monkey was sitting upright, then every 30 min until it was fully recovered from the anesthesia.

### *Statistical Analysis*

Data collected was analyzed utilizing GraphPad PRISM and SPSS software. Metabolic parameters collected at multiple timepoints were compared by ANOVA with repeated measures (RM-ANOVA). Metabolic parameters at individual timepoints between three treatment groups were assessed via one-way ANOVA. Plots are expressed as means $\pm$ SEM. Relationships between hormones, metabolic parameters and hypothalamic hormones were assessed via nonparametric Spearman's correlation tests. To determine cumulative differences over time, incremental area under the curve (AUC) was calculated via the trapezoid rule using GraphPad Prism software. Outliers were determined via Grubbs test at  $\alpha=0.05$  level. If outliers were determined, appropriate statistical tests were completed after removal of outlier(s). Data presented includes all monkey values, nevertheless. Normality was assessed via GraphPad software. If normality was not met,  $\log_{10}$  transformations were performed prior to statistical

analysis. Untransformed data, however, are reported. Statistical significance was determined as  $p < 0.05$  and trending towards significance was determined as  $0.05 < p < 0.1$ .

#### 4.4 Results

##### *Hypothalamic and Circulating Hormone Levels*

Circulating hormones values averaged over the duration of the study and compared between groups, revealed reductions in estrogens but increases in androgens in both the VEH and LET E<sub>2</sub>-depleted groups compared to the E2 group (Figure 2). In VEH and LET females, levels of both circulating estradiol (VEH,  $p < 0.0001$ ; LET,  $p < 0.0001$ ) and estrone (VEH,  $p = 0.02$ ; LET,  $p = 0.005$ ) were diminished in both E<sub>2</sub>-depleted groups compared to the E2 group. In contrast, VEH and LET females exhibited hyperandrogenism compared to E2 by way of increased circulating T (VEH,  $p = 0.002$ ; LET,  $p < 0.0001$ ) and DHEA (VEH,  $p = 0.008$ ; LET,  $p = 0.003$ ) compared to the E2 females. LET females also developed elevated androstenedione levels compared to both VEH ( $p = 0.09$ , trend) and E2 ( $p = 0.008$ ) females. There were no observed between female group differences in circulating progesterone levels (Figure 2). With regard to ratios between circulating steroid hormones, the T: E<sub>2</sub> ratio favored testosterone in LET ( $p = 0.0003$ ) and VEH (0.07) females compared to the E2 group. Additionally, the T: A<sub>4</sub> ratio trended to favor to testosterone in VEH ( $p = 0.08$ ) compared to E2 females (Figure 2).

Hypothalamic hormone content measured by LC-MS/MS from the right hemi-hypothalamus at 17-months from study onset revealed largely undetectable hypothalamic E<sub>2</sub> in both the VEH and LET groups, significantly reduced compared to E2 females (VEH,  $p < 0.0001$ ; LET,  $p = 0.0001$ , Figure 3A). E<sub>1</sub> was significantly reduced in

LET compared to E2 females ( $p=0.0002$ , Figure 3B). Hypothalamic content of T, DHEA, A<sub>4</sub>, progesterone, 17-OHP, and cortisol all showed no between female group differences (Figure 3C-H). Hypothalamic hormone ratios between T and E<sub>2</sub> and T and A<sub>4</sub> showed no between female group differences (Figure 4A-B). The hypothalamic ratio of E<sub>1</sub> to E<sub>2</sub>, however, was reduced in E2 compared to VEH females ( $p=0.04$ , Figure 4C).

#### *Circulating Metabolic Analytes*

At baseline pre-OVX, there were no between female group differences in circulating levels of leptin, glucagon, GIP, c-peptide, and amylin (Table 2). There was a trend towards increased GLP-1 in LET compared to E2 females ( $p=0.07$ ), but this trend disappeared at 6-months and 12-months post-OVX. At 6-months post-OVX, there was a trend towards reduced c-peptide in LET compared to E2 females ( $p=0.07$ ) and increased leptin in VEH compared to E2 females ( $p=0.08$ ), while there was no observed between group differences in any of the other metabolic analytes between female groups (Table 2). At 12-months post-OVX, no observed trends or differences between female groups remained for any metabolic analyte (Table 2).

#### *Body Weight, Body Composition and Bone Analysis*

Monthly body weights were calculated by averaging the weekly body weights for all animals in each treatment group. There were no differences in body weight or body weight percent change between the three female groups (Figure 5A, 5C). Additionally, there were no differences in AUC analyses performed on body weight and body weight percent change through 17 months of the study (Figure 5B, 5D). Body mass index (BMI) calculated at baseline, 6-months post-OVX and 12-months post-OVX (Figure 6A) and percent change from baseline to 12-months post-OVX (Figure 6B) showed no

difference between female groups. Body composition via DXA scans at baseline, 6-months post-OVX and 12-months post-OVX showed no between female group difference in total body fat mass or total body fat-free mass. Additionally, there were no between female group differences in calculated percent change of fat mass and fat-free mass between baseline and 12-months post-OVX (Figure 7). In all validated regions of interest (abdomen, thighs, extremities) there were no between female group differences in fat mass or fat-free mass (Table 3).

Bone mineral content (BMC) was reduced at 6-months and 12-months post-OVX in the LET females compared to their baseline values ( $p < 0.0001$ ;  $p = 0.0011$ , Figure 8A). In contrast, E2 females presented with increased BMC at 12-months post-OVX compared to their baseline values ( $p = 0.045$ , Figure 8A). Both VEH and LET females presented with negative percent change of BMC from baseline to 12-months post-OVX compared to E2 females (VEH vs E2,  $p = 0.007$ ; LET vs E2,  $p = 0.0002$ ; Figure 8B). Similarly with bone mineral density (BMD), LET females presented with reduced, while VEH females presented with a trend towards reduced percent change from baseline to 12-months post-OVX compared to E2 females (VEH vs E2,  $p = 0.06$ ; LET vs E2,  $p = 0.005$ ; Figure 8D). Only VEH females demonstrated a time effect of reduced BMD between 6-months and 12-months post-OVX ( $p = 0.03$ ; Figure 8B), in contrast to E2 and LET females.

#### *Caloric Intake*

Total daily calories (kcal) consumed were averaged between all the days of the caloric intake assessment at each timepoint. There were no differences in average calories consumed at baseline, 6-months post-OVX or 12-months post-OVX between

the three female groups (Figure 9A). Additionally, when daily average calories consumed were adjusted for body weight (kg) and fat-free mass (kg) for each animal at each timepoint, there were no observed between female group differences (Figure 9B).

#### *Locomotor Activity*

Locomotor activity data was calculated for a total of 24-hours and specific periods of the day. Total 24-hrs was an average of all the activity counts over 24-hrs averaged over the number of days the collar was worn. Daytime activity was the average of activity counts between 0600h-1800h, nighttime was 1800h-0600h, morning was 0600h-1200h and afternoon was 1200h-1800h. There were no observed between female group differences in total locomotor activity counts at baseline, pre-OVX and 12-months post-OVX (Figure 10, Figure 11). Additionally, there was no between female group differences in percent change of activity counts from baseline to 12-months post-OVX (Figure 12).

#### *Doubly labeled water (DLW) Energy Expenditure*

Total energy expenditure (TEE, kcal/d) was adjusted for fat-free mass (FFM, kg) at baseline, 6-months, and 12-months. There were no observed between female group differences in TEE adjusted for FFM at baseline, 6-months, or 12-months post-OVX (Figure 13A-C). Additionally, there was no observed between female group differences in percent change between baseline and 12 months in TEE adjusted for FFM (Figure 13D).

### *Glucoregulation via ivGTT*

Glucoregulatory values were calculated via the Bergman-Cobelli Modified Minimal Model for rhesus macaques (Table 4). There were no between female group differences at baseline (B) and 12-months post-OVX for insulin sensitivity (SI, baseline,  $p=0.38$ ; 12-mo,  $p=0.46$ ), disposition index (DI, baseline,  $p=0.86$ ; 12-mo,  $p=0.34$ ), acute insulin response (AIRg, baseline,  $p=0.88$ ; 12-mo,  $p=0.67$ ), basal glucose (Gb, baseline,  $p=0.62$ ; 12-mo,  $p=0.79$ ) and basal insulin (Ib, baseline,  $p=0.54$ ; 12-mo,  $p=0.30$ ). At 12-months post-OVX, there was a trend towards reduced glucose effectiveness (Sg) in LET females ( $p=0.07$ ) compared to E2 and VEH female groups, suggesting reduced ability for insulin-independent glucose uptake in LET females. Percent change between baseline and 12-months post-OVX revealed no between female group differences in any glucoregulatory measure (SI,  $p=0.11$ , AIRg,  $p=0.34$ , DI,  $p=0.59$ , Sg,  $p=0.37$ , Ib,  $p=0.42$  and Gb,  $p=0.39$ ).

### **4.5 Discussion**

In the present study, we comprehensively demonstrate that neither ovarian E<sub>2</sub> depletion nor ovarian + extra-ovarian E<sub>2</sub> depletion for 17-months elicits a disruption in energy balance in adult female rhesus macaques. We hypothesized that extra-ovarian E<sub>2</sub>, specifically neuro-E<sub>2</sub>, would be the source of E<sub>2</sub> signaling through hypothalamic ER $\alpha$  to maintain female metabolic homeostasis, but our findings reveal that neither long-term ovariectomy alone, nor ovariectomy and complete estrogen synthesis inhibition, produced any major alterations in body weight, body composition, adiposity, energy expenditure, food intake, or glucose homeostasis. Our observations suggest that



neither ovarian nor extra-ovarian estrogens are required to maintain metabolic homeostasis in a non-human primate under normal dietary conditions.

Our results surprisingly revealed that the hypothalamic from ovariectomized animals treated with either vehicle or letrozole contained largely both the VEH and LET females exhibited undetectable levels of circulating and hypothalamic E<sub>2</sub> content. E<sub>2</sub> females nevertheless demonstrated early to late follicular phase levels of circulating E<sub>2</sub> due to E<sub>2</sub> replacement and averaged 0.89 pg of E<sub>2</sub> in the right hemi-hypothalamus. It is interesting that both VEH and LET females exhibited comparable undetectable hypothalamic content of E<sub>2</sub> despite administration of high doses (1 mg/kg/day) of the aromatase inhibitor, letrozole, to LET females. Despite a highly sensitive limit of assay quantification for E<sub>2</sub> of 250 fg, it is possible that we were not able to identify aromatase inhibition driven differences as both systemic E<sub>2</sub> levels and hypothalamic E<sub>2</sub> content were below this limit. In addition, hypothalamic E<sub>2</sub> content in E<sub>2</sub> females could be derived from circulating E<sub>2</sub> crossing the blood brain barrier, highlighting the possibility that hypothalamic E<sub>2</sub> in adult female rhesus macaques may not be synthesized in the hypothalamus or is only synthesized for brief periods following acute elevations in systemic E<sub>2</sub> levels<sup>28</sup>.

In contrast to hypothalamic E<sub>2</sub> content, hypothalamic content of E<sub>1</sub> was reduced in LET, but not VEH female groups compared to the E<sub>2</sub> group. This latter finding clearly demonstrates that the adult female rhesus monkey hypothalamus does express aromatase since aromatase inhibition and OVX diminished hypothalamic E<sub>1</sub> content more than OVX alone. Since this effect of aromatase inhibition was only observed for the hypothalamic E<sub>1</sub> and not hypothalamic E<sub>2</sub>, content it is possible that hypothalamic

conversion of  $A_4$  to  $E_1$  was favored over hypothalamic conversion of  $A_4$  to T. This notion of neuro-steroidogenic pathways having a bias towards  $A_4$  and  $E_1$  production relative to T and  $E_2$  would be consistent with studies of bird brain neurosteroidogenesis<sup>29</sup>. Interestingly, hypothalamic  $A_4$  content was nearly undetectable in all female groups and there were no between female group differences of hypothalamic  $A_4$  and T. The greater hypothalamic content of  $E_1$  over  $E_2$  is also consistent with research identifying  $E_1$  as the predominate active estrogen in post-menopausal women.

In the systemic circulation of adult female rhesus monkeys in this study,  $E_1$ , like  $E_2$ , was below the limit of quantitation for most VEH and LET females. T and  $A_4$  were nevertheless both increased to a greater extent in LET females, compared to VEH and  $E_2$  females, strongly implicating aromatase inhibition induced hyperandrogenism. The contrast between systemic circulation hormone levels and hypothalamic hormone content could be due to tissue specific regulation of estrogens as circulating levels of  $E_2$  decline experimentally or naturally, with menopause. Investigation into the hormonal and genetic profiles of metabolically relevant tissues such as adipose and liver could reveal specific changes within a tissue.

With regard to  $E_2$  depletion association with impaired female metabolic function, we found a subtle trend in reduced glucose effectiveness in the VEH and LET group compared to the  $E_2$  group. This suggests that reduced ovarian as well as reduced ovarian + extra-ovarian  $E_2$  contributes to reduced ability for glucose uptake independent of insulin action. The menopausal transition in women lasts on average 4 years but can last up to a decade. During this period of vast hormonal and metabolic changes,  $E_2$  levels decline. Our study duration was only 17 months so while it exceeded a year, the

representative time compared to humans experiencing reduced E<sub>2</sub> is short. In the absence of body weight gain and lipid accumulation, it is possible we are highlighting only early onset disruptions of energy balance due to reduced estrogens. The reduced ability of glucose effectiveness could elicit pancreatic islets to compensate and if stressed by a Western style diet or sedentary lifestyle, could evolve into insulin intensity and subsequently, diabetes.

With regard to E<sub>2</sub> depletion and the development of osteoporosis, we provided evidence that ovarian and ovarian + extra-ovarian E<sub>2</sub> depletion results in decreased bone mineral content and bone mineral density. Percent change of bone mineral density from baseline to 12-months post-OVX trended towards greatest bone loss in LET females. Understanding the effects of aromatase inhibitor use and bone loss is incredibly important as aromatase inhibition is used as a frontline treatment for breast cancer patients<sup>30</sup>. Bone loss has been shown to occur rapidly in post-menopausal women with an estimated 10-20% of bone loss around the first 5 years<sup>31</sup> and occurs more severely in post-menopausal breast cancer patients receiving aromatase inhibition therapy<sup>32</sup>. In the present female monkey study, reductions in bone mineral content started to appear six-months post-treatment onset in LET females, confirming past literature of rapid deterioration of bone<sup>33</sup>. In contrast to other components of energy balance such as body weight and energy expenditure, our study confirms that bone metabolism appears sensitive to changes in circulating estrogens and is exaggerated by aromatase inhibition.

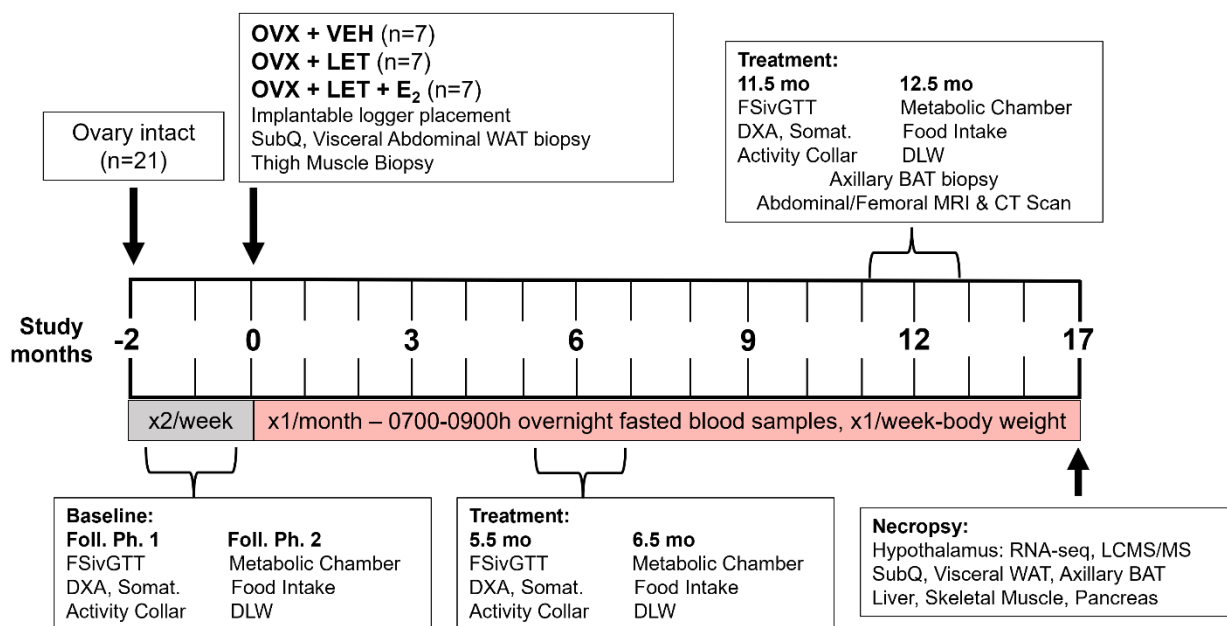
In summary, we demonstrated that OVX + aromatase inhibition exaggerates circulating hyperandrogenism and induces hypothalamic E<sub>1</sub>:E<sub>2</sub> ratios favorable to E<sub>1</sub>.

While all but one of these endocrine differences failed to diverge from just OVX alone, except for E<sub>1</sub>, only bone loss differed in phenotypic outcomes between female groups. Potentially, the subtle difference in reduced glucose effectiveness in the E<sub>2</sub> depleted groups is highlighting the first disruption to energy balance and unless compensated for effectively, and even without achieving a positive energy balance, could lead to diabetes.

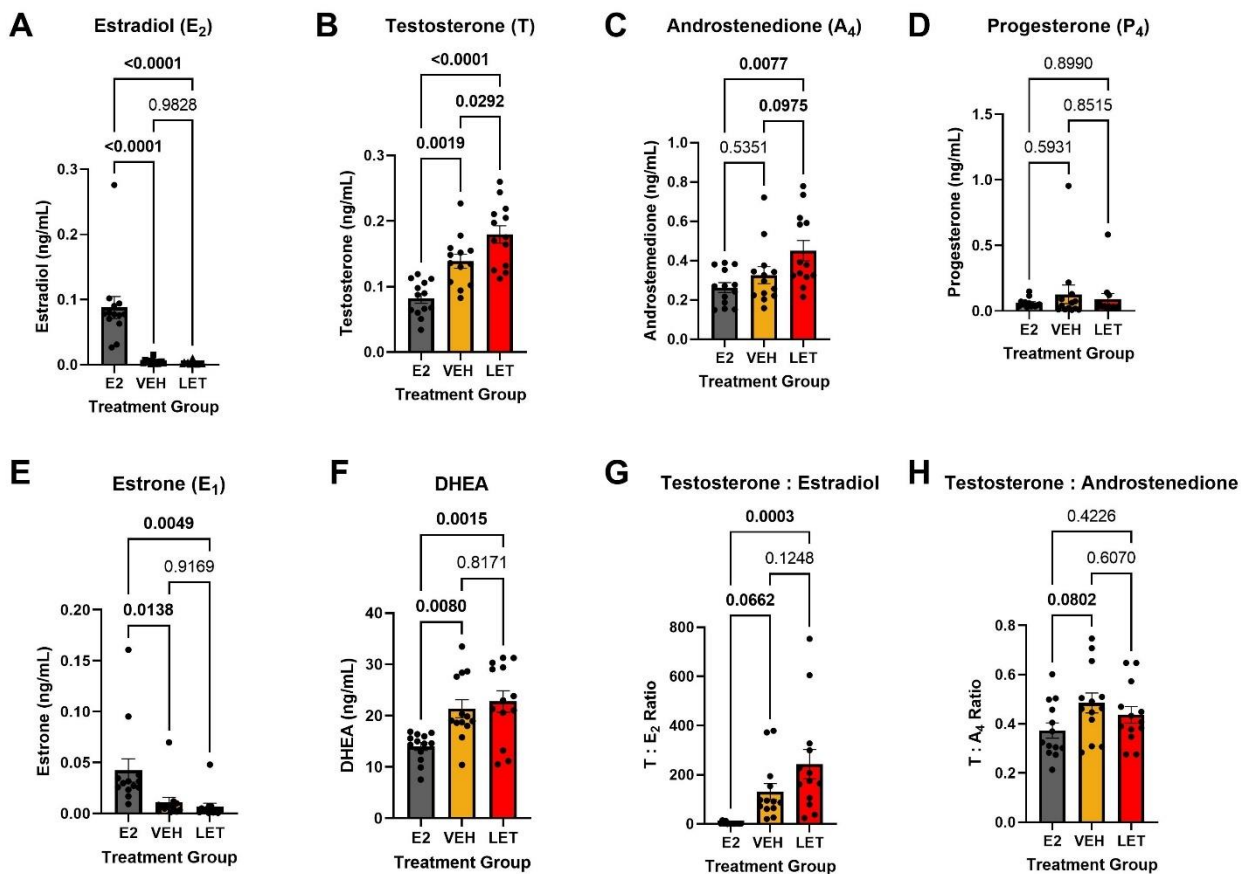
#### 4.6 Tables/Figures

	<b>Age (years)</b>	<b>Body Weight (kg)</b>
<b>E2 (n=7)</b>	7.40 ± 3.31	6.59 ± 1.10
<b>VEH (n=7)</b>	7.50 ± 2.10	7.00 ± 1.16
<b>LET (n=7)</b>	9.94 ± 1.79	7.06 ± 1.03
p-value	0.13	0.75

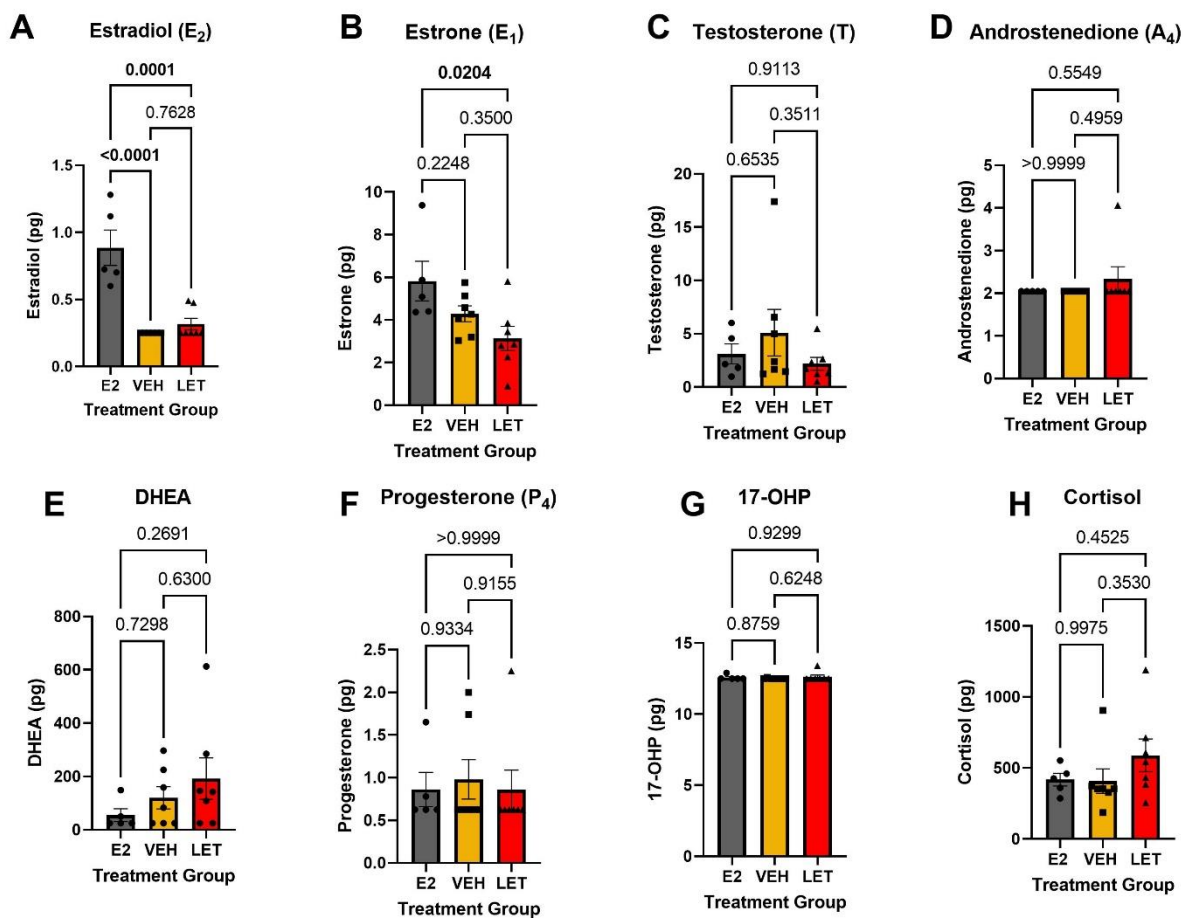
**Table 1:** Comparable ages and body weights between all three female rhesus treatment groups. OVX+LET+E2 (n=7), OVX+VEH (n=7) and OVX+LET (n=7) at the start of the study. Values expressed as mean ± SEM. P-values calculated via ordinary one-way ANOVA.



**Figure 1:** Timeline of treatment onset, metabolic procedures, and end of study.

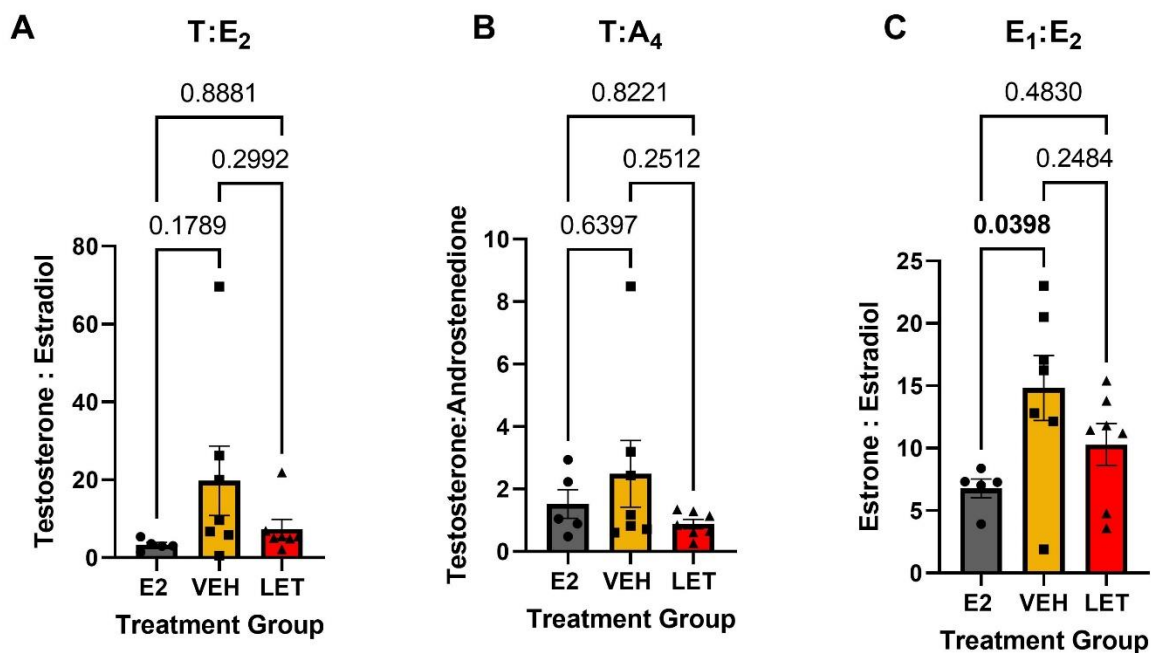


**Figure 2:** Circulating hormones averaged over 17 months of the study. Plots expressed as mean  $\pm$  SEM. Significance assessed via ordinary one-way ANOVA. VEH and LET group presented with reduced E<sub>1</sub> and E<sub>2</sub> and increased T, A<sub>4</sub> and DHEA compared to the E2 group. T : E<sub>2</sub> ratio was favored to T in VEH and LET group and T : A<sub>4</sub> ratio favored to T in VEH group. E2 (grey, n=5), VEH (yellow, n=7) and LET (red, n=7).



**Figure 3.** Hypothalamic steroid hormone content from the right hemi-hypothalamus 17-months following study onset. Individual values and mean  $\pm$  SEM values are illustrated. E2 (grey, n=5), VEH (yellow, n=7) and LET (red, n=7).

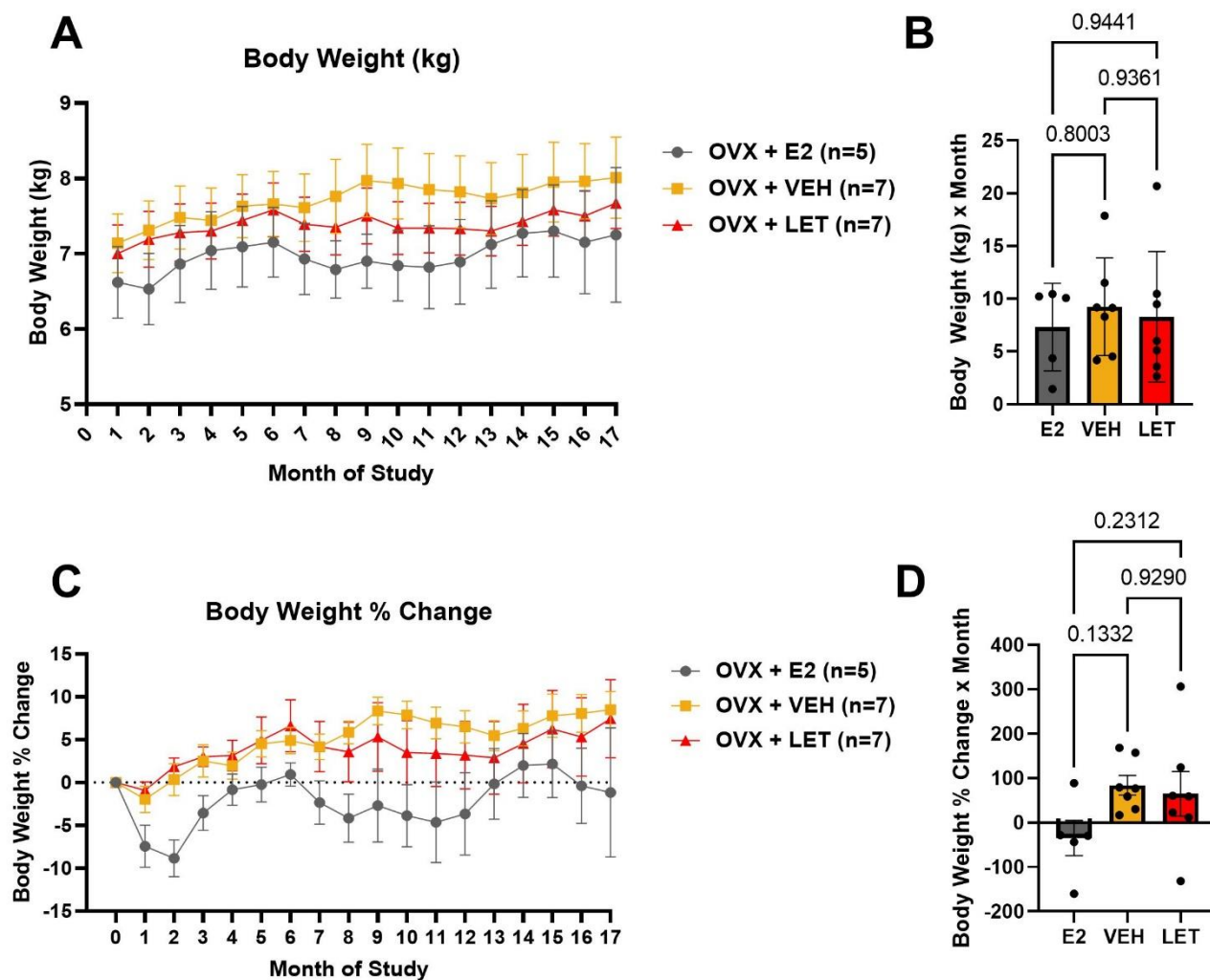




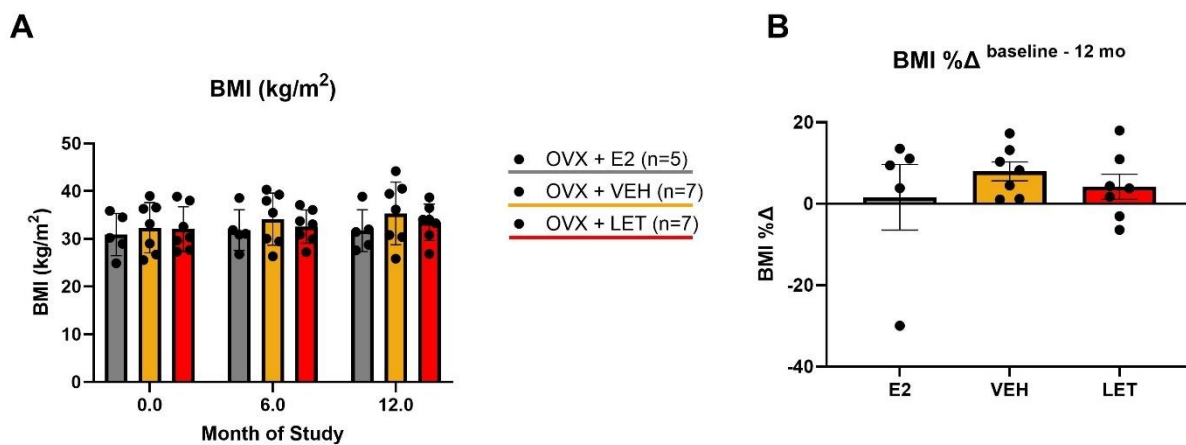
**Figure 4:** Ratio of hypothalamic hormones (T= testosterone, A<sub>4</sub>=androstenedione, E<sub>2</sub>= estradiol and E<sub>1</sub>= estrone). Plots expressed as mean  $\pm$  SEM. Significance assessed via ordinary one-way ANOVA. E2 (grey, n=5), VEH (yellow, n=7) and LET (red, n=7).

		Treatment Group			p-value
		E2	VEH	LET	
<b>Leptin</b>	Baseline	168.9±68.4	372.7±164.8	375.5±95.04	0.49
	6-months	<b>253.7±108.3</b>	<b>949.4±337.7</b>	<b>314.9±85.8</b>	<b>0.08</b>
	12-months	231.3±92.6	519.5±155.7	479.8±115.2	0.32
<b>Ghrelin</b>	Baseline	NA	NA	NA	NA
	6-months	20.15±6.43	13.93±0.21	18.27±2.15	0.40
	12-months	55.41±23.18	27.72±11.78	39.61±14.50	0.56
<b>GLP-1</b>	Baseline	<b>30.83±14.46</b>	<b>27.86±6.27</b>	<b>59.58±9.80</b>	<b>0.07</b>
	6-months	209.8±102.4	233.4±93.8	146.7±71.6	0.76
	12-months	30.37±3.58	44.14±7.12	47.73±14.50	0.53
<b>Glucagon</b>	Baseline	127.3±40.85	117.5±39.27	33.58±12.56	0.14
	6-months	36.60±8.95	37.48±7.00	44.63±18.59	0.11
	12-months	50.58±6.43	31.68±3.99	34.00±12.58	0.34
<b>GIP</b>	Baseline	49.00±10.8	32.04±5.3	41.25±7.4	0.32
	6-months	38.43±11.2	34.82±5.5	32.75±6.6	0.87
	12-months	52.50±12.0	59.00±5.6	70.69±13.9	0.52
<b>C-peptide</b>	Baseline	2002±637	1927±491	1655±183	0.86
	6-months	<b>963.8±218</b>	<b>1577±411</b>	<b>594.5±127</b>	<b>0.07</b>
	12-months	2091±315	2948±891	1754±161	0.34
<b>Amylin</b>	Baseline	3061±10.9	25.18±10.6	51.19±18.4	0.68
	6-months	13.72±0.00	19.26±5.54	54.22±40.50	0.71
	12-months	13.72±0.00	16.63±2.91	20.49±6.77	0.49

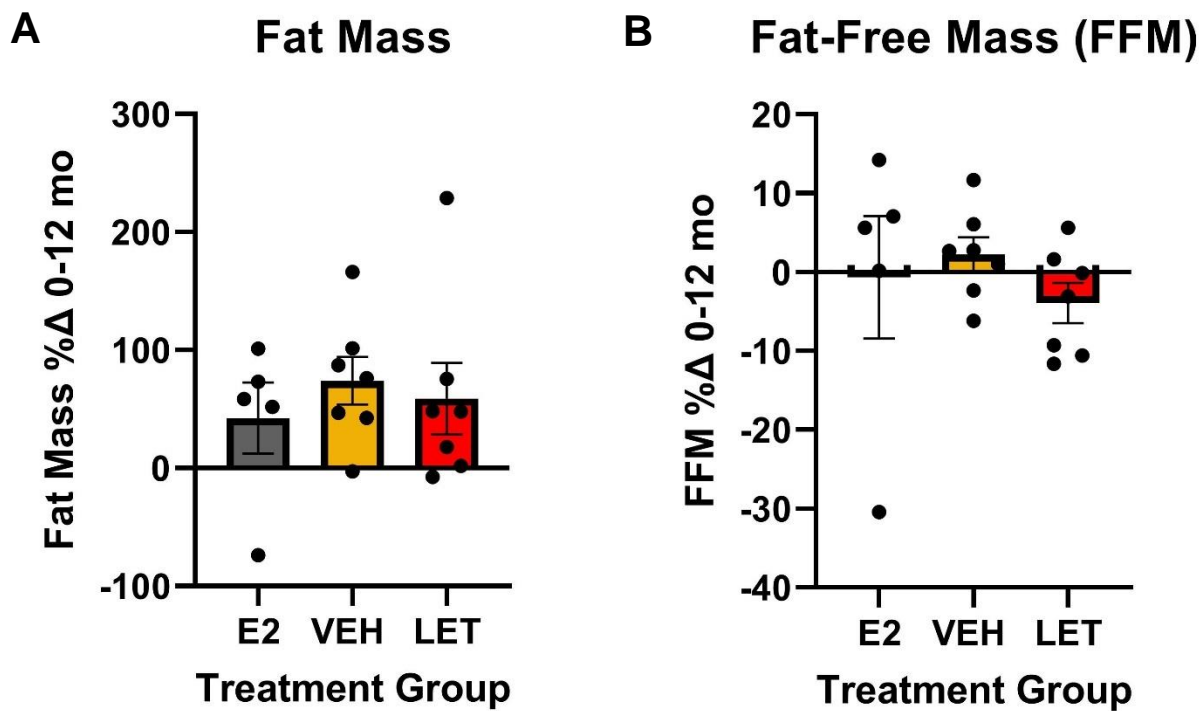
**Table 2:** Metabolic analytes measured at baseline pre-OVX, 6-months and 12-months post-OVX. Percent (%) change is the percentage change between baseline pre-OVX and 12-months post-OVX. Values expressed as mean ± SEM. Trending significance is represented by p-value <0.1.



**Figure 5:** Body weight (A) over 17 months of the study with corresponding AUC bar graph (B) and body weight percent change from month 0 (C) over 17 months with corresponding AUC bar graph (D). Plots are expressed as mean  $\pm$  SEM. Significance assessed via RM-ANOVA (A, C) and ordinary one-way ANOVA (B,D). E2 (grey, n=5), VEH (yellow, n=7) and LET (red, n=7).



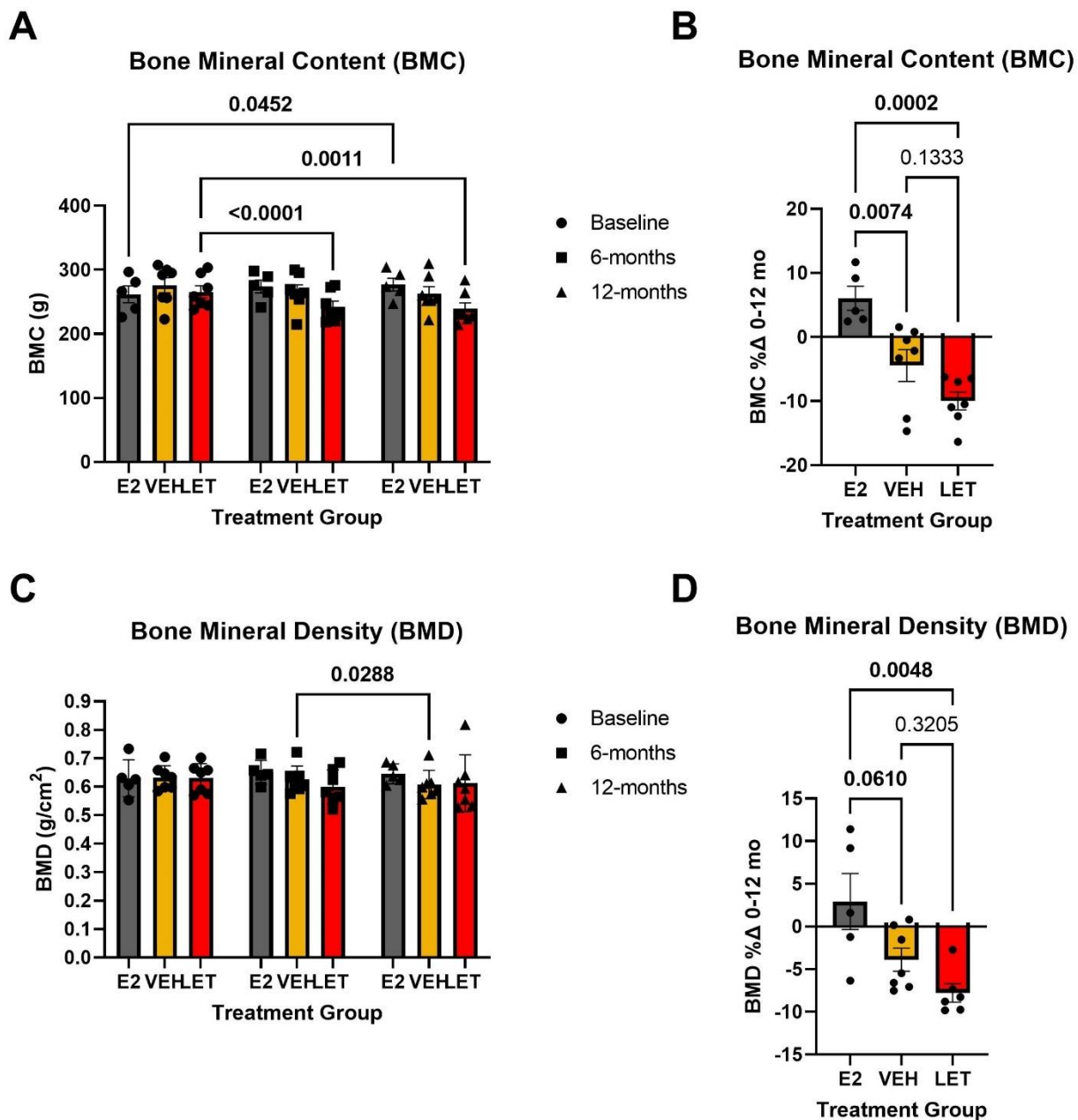
**Figure 6.** Body mass index (BMI) at baseline, 6-months and 12-months post-OVX (A) and BMI percent change between baseline and 12-months post-OVX (B). Plots are expressed as mean  $\pm$  SEM. Significance was assessed via RM-ANOVA (A) and ordinary one-way ANOVA (B). E2 (grey, n=5), VEH (yellow, n=7) and LET (red, n=7).



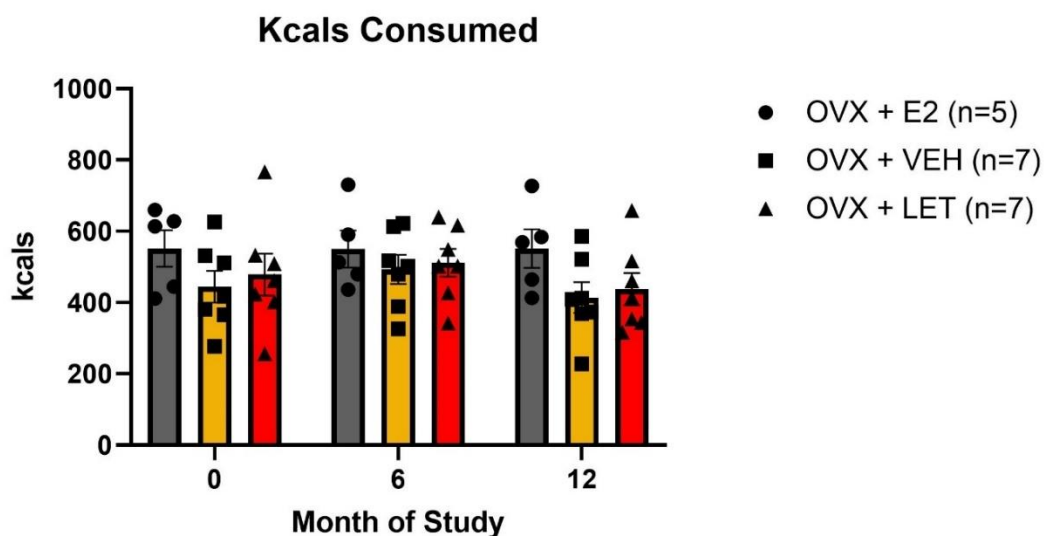
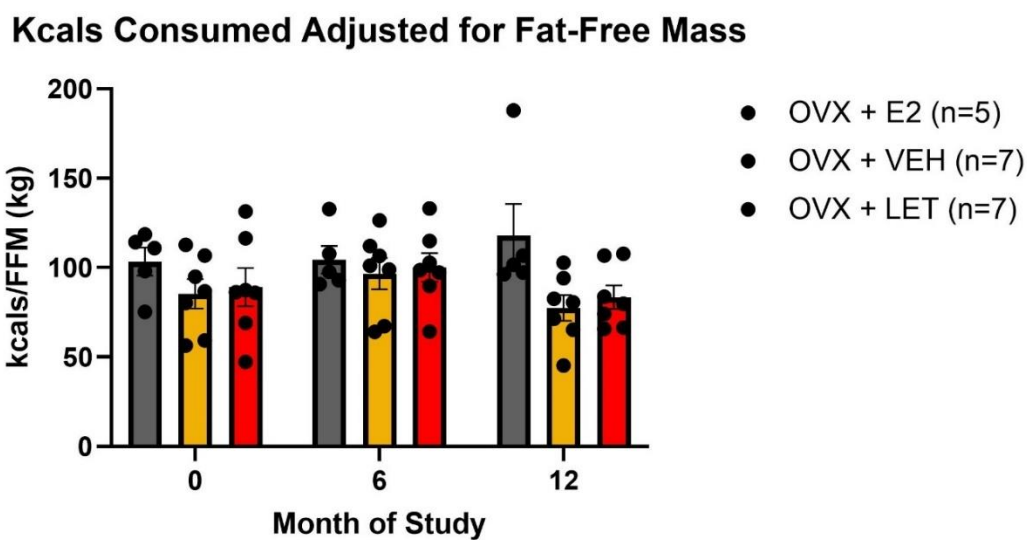
**Figure 7.** Fat mass and fat-free mass percent change from baseline to 12-months post-OVX. Plots are expressed as mean  $\pm$  SEM. Significance was assessed via ordinary one-way ANOVA. E2 (grey, n=5), VEH (yellow, n=7) and LET (red, n=7).

		Treatment Group			
<b>Fat-Mass (kg)</b>		E2	VEH	LET	p-value
<b>Abdomen</b>	Baseline	0.29±0.11	0.56±0.17	0.48±0.13	0.13
	6-months	0.34±0.11	0.86±0.19	0.78±0.14	
	12-months	0.38±0.16	0.98±0.20	0.82±0.14	
<b>Upper Legs</b>	Baseline	0.18±0.06	0.22±0.06	0.22±0.04	0.39
	6-months	0.21±0.04	0.28±0.04	0.29±0.04	
	12-months	0.02±0.03	0.29±0.05	0.27±0.04	
<b>Chest</b>	Baseline	0.23±0.08	0.33±0.09	0.29±0.08	0.27
	6-months	0.29±0.08	0.54±0.09	0.44±0.11	
	12-months	0.24±0.01	0.44±0.11	0.38±0.08	
<b>Lower Legs</b>	Baseline	0.12±0.02	0.13±0.02	0.13±0.02	0.91
	6-months	0.14±0.02	0.16±0.01	0.14±0.01	
	12-months	0.15±0.03	0.15±0.02	0.15±0.02	
<b>Arms</b>	Baseline	0.14±0.02	0.17±0.04	0.17±0.03	0.53
	6-months	0.16±0.02	0.22±0.04	0.23±0.04	
	12-months	0.16±0.02	0.20±0.03	0.21±0.03	
<b>Fat-Free Mass (kg)</b>					
<b>Abdomen</b>	Baseline	1.60±0.09	1.57±0.07	1.69±0.08	0.89
	6-months	1.51±0.09	1.54±0.09	1.54±0.07	
	12-months	1.76±1.24	1.68±0.07	1.68±0.07	
<b>Upper Legs</b>	Baseline	1.02±0.99	1.07±0.08	0.96±0.09	0.48
	6-months	1.00±0.86	1.09±0.07	0.99±0.09	
	12-months	0.80±0.21	1.01±0.03	0.92±0.07	
<b>Chest</b>	Baseline	1.06±0.07	1.02±0.06	1.15±0.05	0.49
	6-months	1.11±0.06	0.99±0.06	1.07±0.05	
	12-months	1.06±0.06	1.00±0.10	1.03±0.06	
<b>Lower Legs</b>	Baseline	0.55±0.05	0.48±0.03	0.47±0.02	0.51
	6-months	0.48±0.02	0.48±0.02	0.47±0.02	
	12-months	0.51±0.05	0.53±0.02	0.47±0.02	
<b>Arms</b>	Baseline	0.73±0.05	0.67±0.04	0.71±0.02	0.21
	6-months	0.73±0.03	0.62±0.04	0.62±0.03	
	12-months	0.71±0.04	0.66±0.02	0.64±0.02	

**Table 3.** Fat-mass (kg) and fat-free mass (kg) values from DXA scan ROIs (abdomen, upper legs, chest, lower legs, and arms). P-values represent treatment factor from RM-ANOVA over baseline, 6-months, and 12-months.

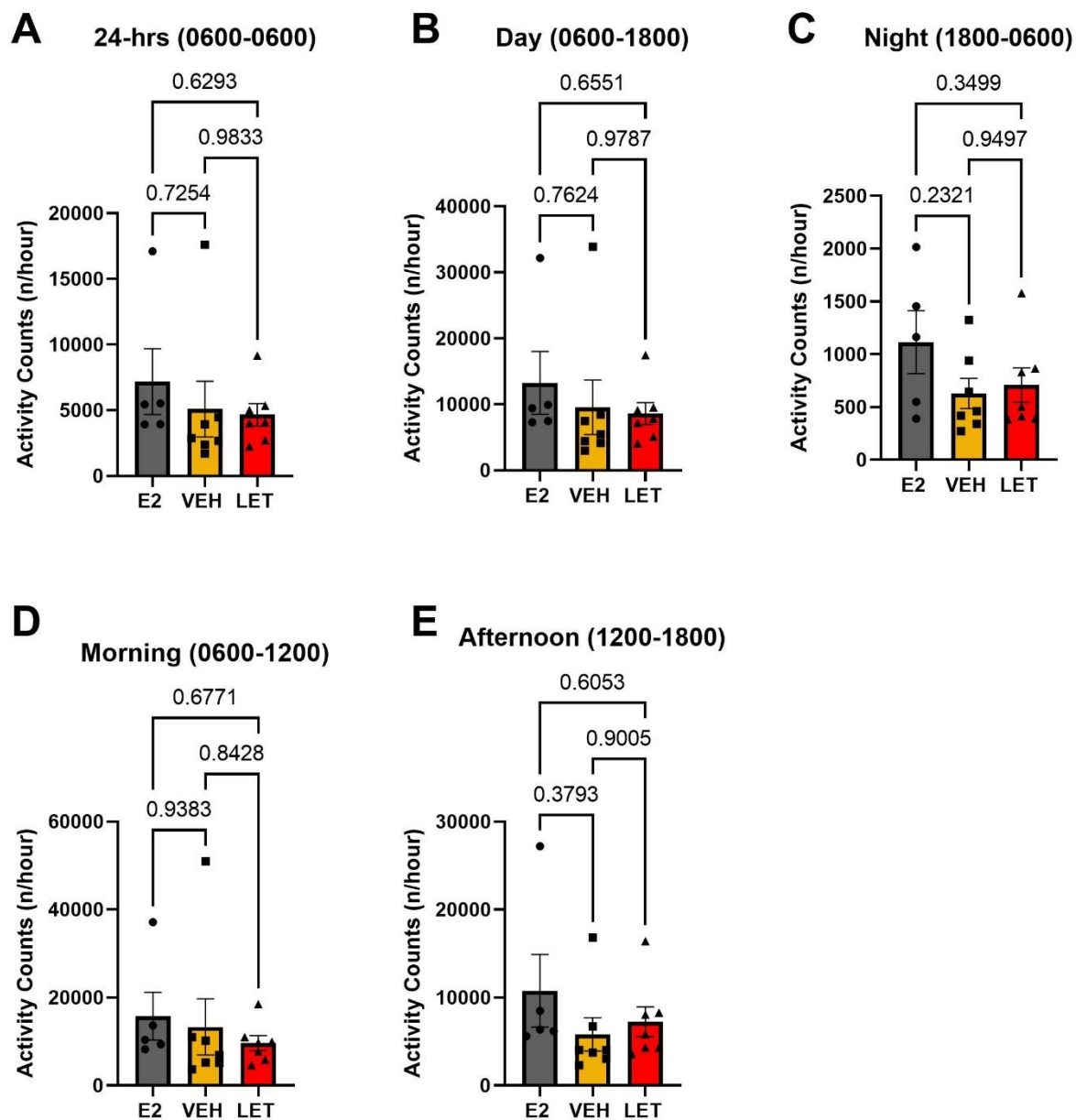


**Figure 8:** Bone mineral content (A) and bone mineral density (C) at baseline, 6-months, and 12-months post-OVX DXA scans and percent change BMC (B) and BMD (D) from baseline to 12-months post-OVX. Plots expressed as mean  $\pm$  SEM. Significance assessed via RM-ANOVA and ordinary one-way ANOVA. E2 (grey, n=5), VEH (yellow, n=7) and LET (red, n=7).

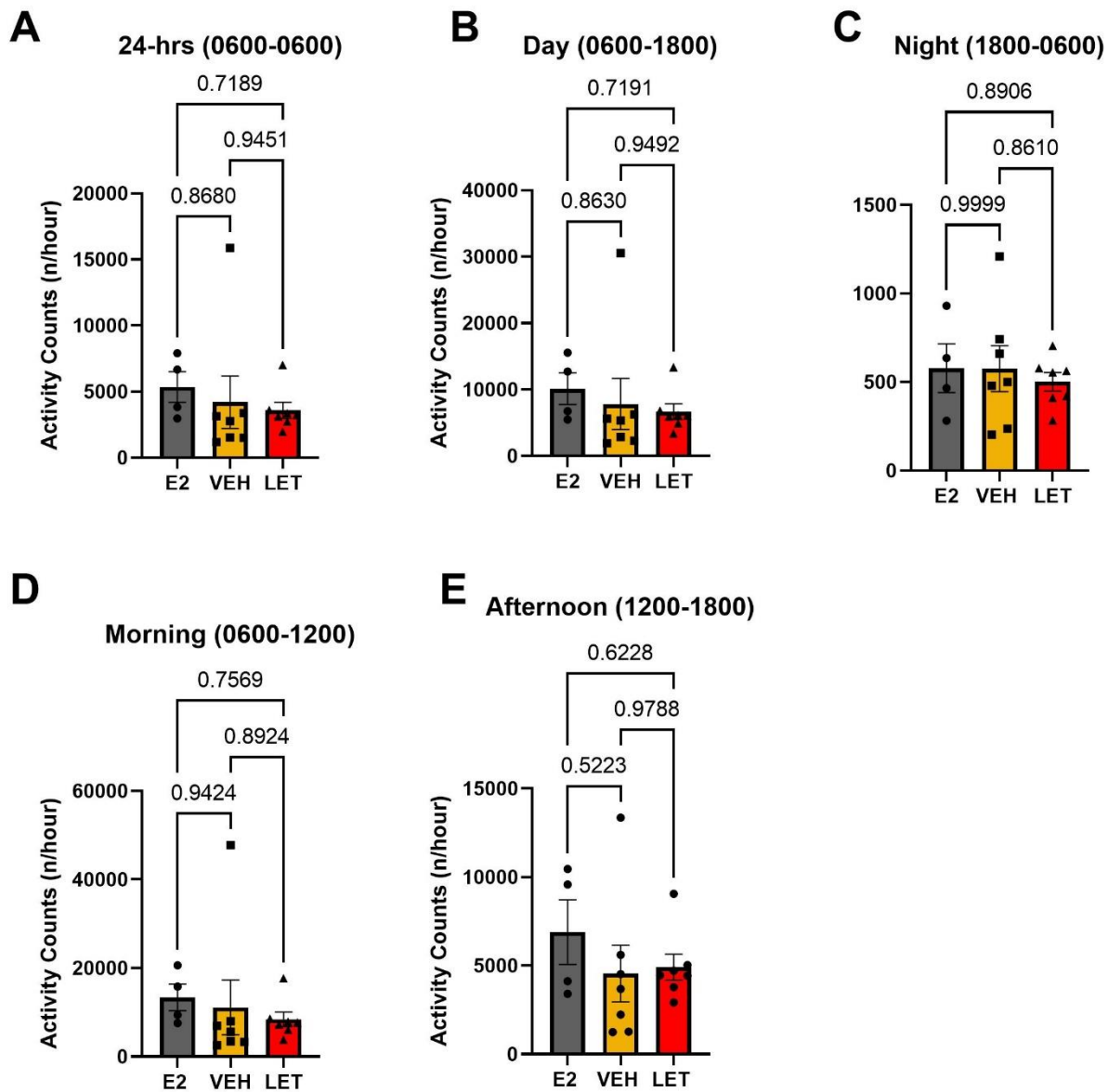
**A****B**

**Figure 9:** Calories (kcal) consumed (A) and Calories (kcal) consumed adjusted for fat-free mass (kg) averaged over the food assessment at baseline, 6-months, and 12-months post-OVX. Plots expressed as mean  $\pm$  SEM. Significance assessed via RM-ANOVA. E2 (grey, n=5), VEH (yellow, n=7) and LET (red, n=7).

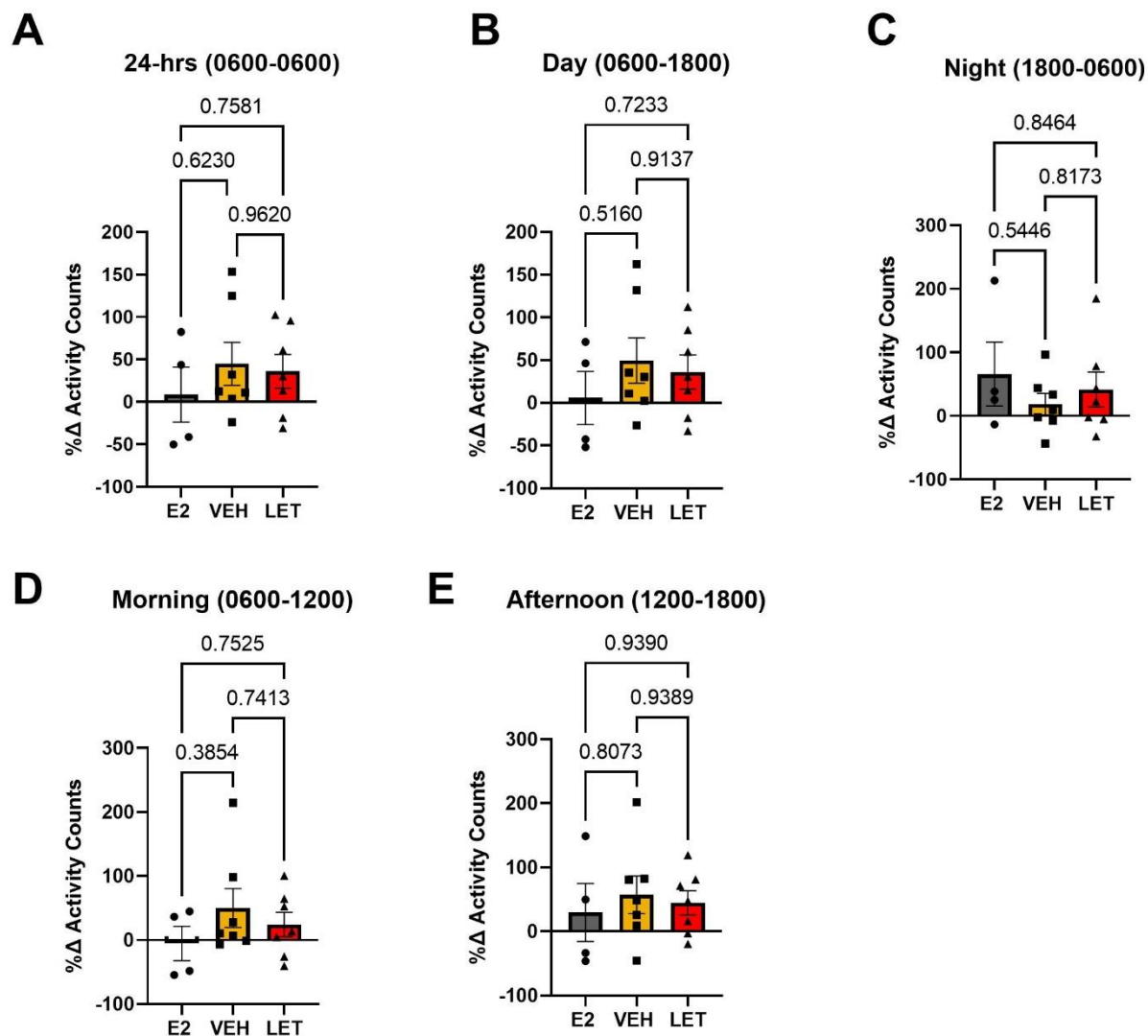




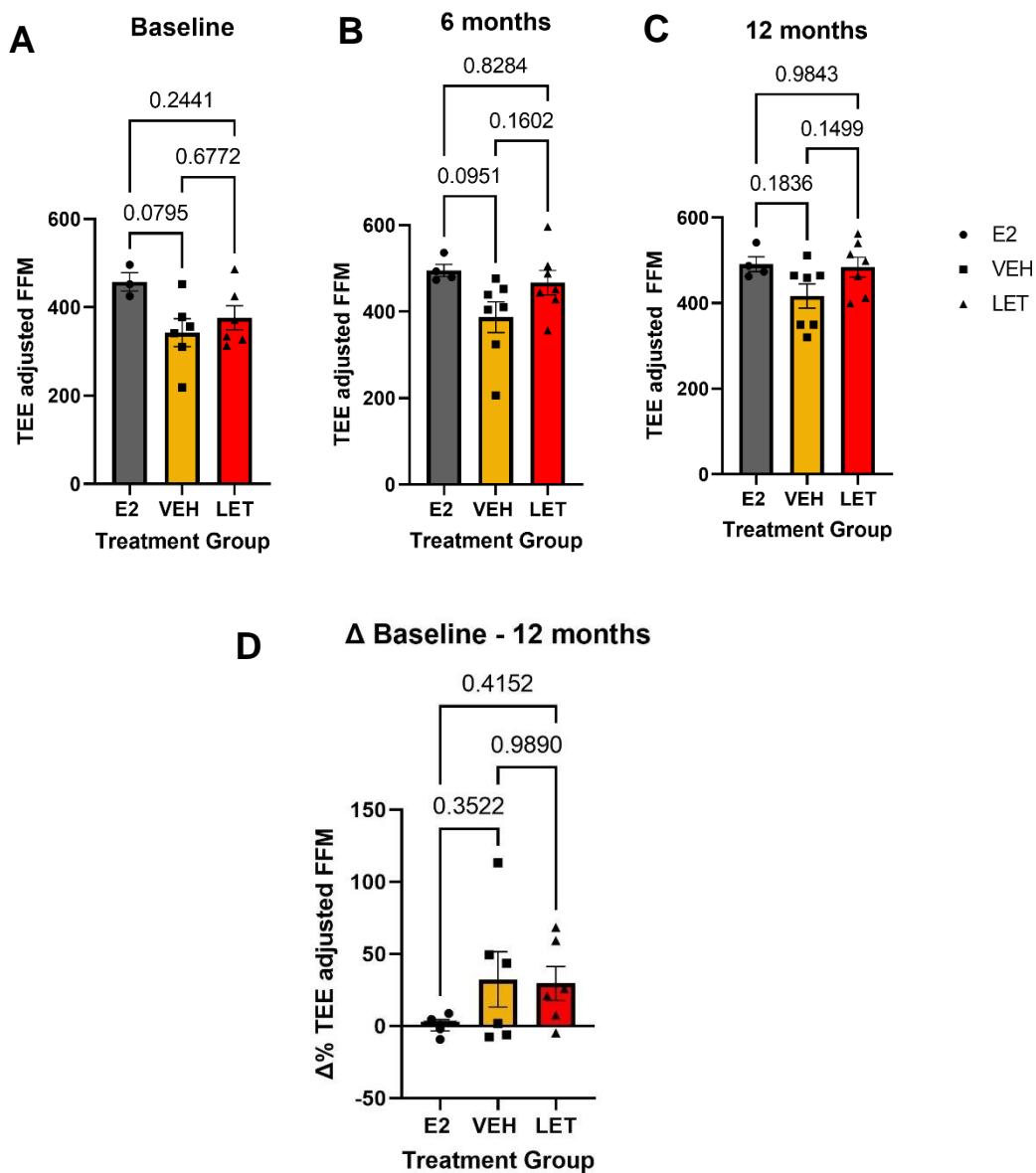
**Figure 10:** Activity counts averaged over 24-hrs (A), Day (B), Night (C), Morning (D) and Afternoon (E) at baseline. Plots expressed as mean  $\pm$  SEM. Significance assessed ordinary one-way ANOVA. E2 (grey, n=5), VEH (yellow, n=7) and LET (red, n=7).



**Figure 11:** Activity counts averaged over 24-hrs (A), Day (B), Night (C), Morning (D) and Afternoon (E) at 12-months post-OVX. Plots expressed as mean  $\pm$  SEM. Significance assessed ordinary one-way ANOVA. E2 (grey, n=5), VEH (yellow, n=7) and LET (red, n=7).



**Figure 12:** Percent change in activity over 24-hrs (A), Day (B), Night (C), Morning (D) and Afternoon (E) between baseline and 12-months post-OVX. Plots expressed as mean  $\pm$  SEM. Significance assessed ordinary one-way ANOVA. E2 (grey, n=5), VEH (yellow, n=7) and LET (red, n=7).



**Figure 13:** Total energy expenditure (TEE) adjusted for fat-free mass (kg) at baseline (A), 6-months (B) and 12-months (C). Percent change of TEE adjusted for FFM between baseline and 12 months (D). Plots expressed as mean  $\pm$  SEM. Significance assessed ordinary one-way ANOVA. E2 (grey, n=5), VEH (yellow, n=7) and LET (red, n=7).

		Treatment Group			p-value
		E2	VEH	LET	
<b>SI</b>	Baseline	4.95±1.04	7.18±1.21	5.83±0.93	0.38
	12-months	8.16±1.18	5.79±2.52	4.81±0.94	0.46
	% change	114.6±68.2	-11.39±40.8	-8.58±21.2	0.11
<b>AIRg</b>	Baseline	1243±394	1034±235	1103±240	0.88
	12-months	900±352	1300±318	1276±311	0.67
	% change	-18.73±16.0	62.46±47.3	42.65±31.1	0.34
<b>DI</b>	Baseline	40.83±5.30	44.61±3.08	43.13±5.34	0.86
	12-months	46.39±3.61	36.04±4.67	40.08±5.11	0.34
	% change	26.7±27.15	-13.21±17.16	31.61±47.2	0.59
<b>Sg</b>	Baseline	0.04±0.01	0.03±0.01	0.05±0.01	0.25
	<b>12-months</b>	<b>0.042±0.003</b>	<b>0.031±0.003</b>	<b>0.032±0.003</b>	<b>0.07</b>
	% change	40.62±39.3	24.92±37.1	-22.74±17.75	0.37
<b>Ib</b>	Baseline	22.06±5.79	31.36±11.67	18.70±4.73	0.54
	12-months	9.18±1.95	21.56±8.19	12.63±2.27	0.30
	% change	-45.99±15.1	-13.52±19.3	74.87±98.3	0.42
<b>Gb</b>	Baseline	55.92±2.56	58.26±2.61	54.46±3.13	0.62
	12-months	51.02±0.98	51.17±2.00	52.41±1.31	0.79
	% change	-8.34±3.00	-11.49±4.05	-1.62±6.76	0.39

**Table 4:** Values of glucoregulation calculated via the Bergman Minimal Model

program. Values represented as group mean ± SEM. **Bold** indicates trend towards significance,  $p < 0.1$ . SI=insulin sensitivity, AIRg=acute insulin response to glucose, DI=disposition index, Sg=glucose effectiveness, Ib= basal insulin, Gb=basal glucose, KG=X.

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## 5. CHAPTER FIVE: Future Directions

Willging MM, Levine JE, and Abbott DH

### *5.1 Menopausal and ovarian estrogen depletion and implications for female metabolic health*

Menopause is characterized by a cessation of menstrual cycles and a reduction in circulating levels of estradiol, E<sub>2</sub>. Individuals experiencing personally distressing symptoms of menopause, including hot flashes and night sweats, are typically prescribed hormonal replacement therapy (HRT, usually including synthetic or naturally occurring estrogens with or without accompanying progestins or progesterone) to alleviate symptoms and help reduce osteoporosis induced by declining circulating E<sub>2</sub> levels. Although protection of bone integrity is a clear benefit of HRT<sup>1</sup>, increased risks of cancer of the breast and endometrium, blood clots, stroke, heart disease and dementia are consequences of considerable concern<sup>2</sup>. Therefore, it is incredibly important to elucidate the mechanisms by which E<sub>2</sub> regulates energy homeostasis and how these mechanisms are altered in a menopausal environment.

Ovariectomy (OVX) during a female's reproductive years, while mimicking the ovarian E<sub>2</sub> depletion of menopause, occurs within a less aged bodily environment and may not completely represent the E<sub>2</sub> depletion of menopause in the context of aging. While OVX in rodents induces weight gain, adiposity, reduced energy expenditure and glucose intolerance, in female NHP, OVX-mediated E<sub>2</sub> depletion inconsistently increases body weight<sup>3,4</sup>. Despite the lack of a robust E<sub>2</sub>-associated metabolic phenotype following OVX in NHPs, selective estrogen receptor modifiers (SERMs) can induce modest weight loss in OVX rhesus monkeys<sup>5</sup>. Interestingly, in OVX cynomolgus macaques, E<sub>2</sub> replacement therapy has no effect on body weight gain and ovarian estrogen depletion fails to diminish insulin sensitivity<sup>4</sup>. In addition to ovarian E<sub>2</sub>, E<sub>2</sub>

synthesized in the brain (neuro-E<sub>2</sub>) can stimulate top-down neuronal signaling cascades. Such neuro-E<sub>2</sub> is proposed to act as a neurotransmitter and neuromodulator<sup>6,7</sup>. Aromatase, the enzyme responsible for converting testosterone to E<sub>2</sub> as well as the conversion of androstenedione to estrone, before its onward enzymatic conversion to E<sub>2</sub>, has been documented as present in rhesus macaque brain regions, including the hypothalamus<sup>8</sup>. Therefore, it is a plausible hypothesis that the failure for OVX to increase female NHP body weight is due to the residual actions of neuro-E<sub>2</sub> orchestrating energy balance in the hypothalamus of OVX NHPs, and likely women.

### *5.2 Neural Regulation of Female Metabolic Function in Adult Marmoset and Rhesus Monkeys*

The studies included in this thesis identify ER $\alpha$  as the predominate hypothalamic estrogen receptor governing E<sub>2</sub>-regulated energy homeostasis in adult female marmoset and female rhesus monkeys. Adult female marmosets with reduced ER $\alpha$  protein expression in the MBH exhibited reduced locomotor activity and trended towards glucose intolerance, while ER $\alpha$ KD adult female rhesus monkeys exhibited body weight percent increase and reduced morning locomotor activity. Although these metabolic findings are less severe and less timely than those exhibited by ER $\alpha$ KD rodents<sup>9,10</sup>, they highlight the importance of hypothalamic ER $\alpha$  in regulating certain aspects of female metabolism. In both of our ER $\alpha$ KD NHP studies, we identified disruption of locomotor activity with reduced hypothalamic ER $\alpha$  protein expression, but in neither study was knockdown complete or uniformly achieved within and between individual female monkeys. It has been well established in rodent studies that reduced hypothalamic ER $\alpha$  contributes to reduced locomotor activity, specifically, estrogens acting via ER $\alpha$  in the

mPOA<sup>11,12</sup>. Therefore, it is possible that the less robust locomotor activity phenotype we observed in female marmosets and female rhesus monkeys is due to (1) less than complete ER $\alpha$  knockdown, especially since the extent of ER $\alpha$  knockdown negatively correlated with aspects of induced metabolic phenotype, and (2) contributions of modest off-target knockdown of ER $\alpha$  protein expression, such as within mPOA incurred the inability of our current MRI-guided viral vector neuro-infusion inducing gene knockdown within the brain to be localized within a single hypothalamic nucleus, alone. Further, ER $\alpha$  expressing neurons within specific hypothalamic nuclei and sub-regions therein such as the ventral VMN (VMNv) may mediate specific metabolic processes.

### *5.3 Estrogen Regulation of Female Bone Density in Adult Female Marmoset and Rhesus Monkeys*

In adult female marmoset monkeys, we demonstrated that OVX + aromatase inhibition did not exaggerate body weight gain beyond that demonstrated by ovarian E<sub>2</sub> depletion, alone. Additionally, neither OVX nor aromatase inhibition contributed to discernable skeletal bone loss. In almost all female mammals with regular, frequent estrous, ovarian, or menstrual cycles, a reduction in circulating E<sub>2</sub> concentrations, either spontaneous or experimentally induced, leads to a reduction in bone mass<sup>13-17</sup>. This has been demonstrated in NHP species and occurs in as little as 3 months in rhesus monkeys<sup>13-17</sup>. It is therefore surprising that we found no deficit in bone mass or bone density of either the total body or lumbar spine associated with systemic or systemic and hypothalamic E<sub>2</sub> deficiency in female marmosets after 7 months on study, though, importantly, estrogen and aromatase activity within the bone microenvironment<sup>18</sup> were not assessed. It is possible that marmoset monkeys evolved metabolic control systems

regulated by extra-ovarian E<sub>2</sub> or that are generally less subject to E<sub>2</sub> regulation. A conclusion arising from Chapter 2 suggested that brain derived E<sub>2</sub> may be enough to maintain bone mass even in the context of circulating E<sub>2</sub> deficiency. Bone in marmosets may therefore have evolved to be less subject to estrogen regulation when, in the wild and captivity, adult females can spend two years or more exhibiting intermittent or absent ovarian cycles, and thus low E<sub>2</sub>, while they help raise the offspring their social group's dominant female<sup>18</sup>.

In contrast to our marmoset study (Chapter 2), we found in adult rhesus monkeys that OVX + aromatase inhibition induced greater skeletal bone loss compared to E<sub>2</sub> females than OVX alone. These macaque findings are consistent with human studies and the association of bone loss and osteoporosis with the menopausal transition<sup>19</sup>. Osteoblasts are cells of the bone responsible for osteogenesis or "bone growth"<sup>27</sup>. Human osteoblasts have all necessary biological components to synthesize estrogens<sup>26</sup>. Although the mechanisms by which estrogens influence bone metabolism have not been fully elucidated, it is evident that diminished circulating E<sub>2</sub> during menopause contributes to bone loss. In addition, post-menopausal cancer patient aromatase inhibitor use exaggerates bone loss<sup>25-28</sup>. Since menopause typically begins in the later half of life, it can be difficult to tease apart menopause associated metabolic effects from the effects of aging. Studies investigating bone integrity in young women (~20-40 years old) with amenorrhea have identified that estrogen deficiency at these ages is sufficient to significantly reduce bone mineral density and the age correlated with demineralization<sup>32</sup>. This highlights that although age is an important variable to consider when investigating menopause associated phenotypes, bone integrity seems

to be strongly reliant on E<sub>2</sub>. It is therefore possible that E<sub>2</sub> synthesized within osteoclasts, and thus a component of extra-ovarian E<sub>2</sub>, may contribute independently towards maintenance of bone density, independent of ovarian produced E<sub>2</sub>. Our evidence suggests that when bone synthesized E<sub>2</sub> is reduced via aromatase inhibition, bone loss accelerates compared to OVX, alone. Molecular and hormonal assessments within bone and homeostatic endocrine regulation of bone density were not performed. Such assessments may provide insight into how letrozole alters bone steroidogenesis and turnover. Additionally, our findings confirm the previously published promptness during which bone loss occurs following systemic E<sub>2</sub> depletion in rhesus monkeys<sup>13-17</sup> and humans<sup>1</sup>. Further, our identification of E<sub>2</sub>-mediated bone loss in adult female rhesus monkeys suggests that bone is regulated by E<sub>2</sub> in female Old World monkeys, as found in women, and unlike marmoset monkeys, and possibly other New World primates.

#### *5.4 Potential importance of high calorie diets or DIO to experimental designs exploring estrogen depletion altering female weight gain in NHPs, and likely women*

In regard to body weight and energy expenditure, we did not observe OVX or OVX + aromatase inhibition altered metabolic function in rhesus monkeys. In marmosets, however, we identified OVX and OVX + aromatase inhibition induced body weight gain. Importantly, the marmoset study was performed in context of DIO. It is possible that we did not observe body weight differences in rhesus monkeys because their metabolic processes were not challenged by an experimentally induced positive energy balance and thus did not have to accommodate a greater potential for positive dysfunctional onset. Previous research has shown rhesus monkeys fed a high fat diet demonstrated body weight increase<sup>20</sup>, hyperinsulinemia and hyperglycemia<sup>21</sup> within 6-

months. Moreover, a sedentary lifestyle is another load on regulatory mechanisms and in humans, has been shown to be associated with severe symptoms of menopause and obesity<sup>22</sup>, together with an increased likelihood of positive energy balance. In this regard, the subtle difference in reduced glucose effectiveness observed between E<sub>2</sub> depleted and E<sub>2</sub> replete female rhesus monkeys may be highlighting early indications of disruption to glucoregulation, and ultimately energy balance, unless compensated for effectively by increased pancreatic islet beta cell insulin production and release during meal-typical immediately post-prandial hyperglycemia. This hint of a trend towards hyperglycemia was nevertheless achieved without requiring DIO in the experimental design and challenging the E<sub>2</sub> depleted monkeys with a prevailing positive energy balance environment. Therefore, it is plausible that reduced E<sub>2</sub> is accommodated metabolically by alternate energy balance mechanisms for a limited duration, or until caloric intake increases, such as from DIO or higher calorie diet, or diminished locomotory activity level occurs. Figure 1 provides a diagrammatic representation of this newly acquired understanding of mechanism in female NHPs that may have application to clinical management of women experiencing E<sub>2</sub>-depleted environments.

#### *5.5 Potential hypothalamic metabolically regulatory mechanism in parallel to E<sub>2</sub> regulation*

One potential mechanism for maintaining energy balance in a diminished E<sub>2</sub> environment is the leptin-melanocortin pathway. Like leptin, E<sub>2</sub> is a strong anorexigenic signal regulating energy balance by stimulating hypothalamic preopiomelanocortin (POMC) and cocaine-and amphetamine-regulated transcript (CART) neurons and inhibiting neuropeptide Y (NPY) and agouti-related peptide (AgRP) neurons. It had been

assumed that E<sub>2</sub>'s effect on energy balance acts in tandem with leptin's role via co-expression of leptin receptor (LepRb) and ER $\alpha$  in the mediobasal hypothalamus<sup>23</sup>, but recent research in female rodents has shown co-expression is more pronounced in the preoptic area (POA) and is limited in the mediobasal hypothalamus, including the VMN and ARC. Quantification of LepRb and ER $\alpha$  colocalization throughout the murine hypothalamus has shown the absence of reliable colocalization in all areas aside from the POA. In addition, one month after OVX, co-expression of LepRb and ER $\alpha$  tend to decline<sup>24</sup>. E<sub>2</sub> mediation of energy homeostasis may therefore not be dependent on activation of LepRb on co-expressing cells and circulating levels of E<sub>2</sub> may act indirectly on the leptin-melanocortin pathway. It might therefore be speculated that in the absence of circulating levels of E<sub>2</sub>, circulating levels of leptin maintain the hypothalamic leptin-melanocortin pathway and energy balance. In this latter regard, our adult female rhesus monkey study (Chapter 4) indicated no between female group differences in circulating levels of leptin, but hypothalamic leptin expression was not assessed, as illustrated diagrammatically in Figure 1. Analysis of gene expression in the hypothalamus of these monkeys could provide insight into upregulated and thus, compensatory mechanisms in an E<sub>2</sub> depleted environment.

Another potential mechanism for maintaining female energy balance may involve a ligand-independent mechanism. We found reduced hypothalamic ER $\alpha$  expression in adult female rhesus monkeys exhibiting body weight percent increase and decreased morning activity. OVX and OVX + extra-ovarian E<sub>2</sub> depletion in adult female rhesus monkeys, however, failed to induce alterations to body weight or energy expenditure. Mouse and in vitro studies have demonstrated that estrogen receptors, in particular,



ER $\alpha$ , can be activated in the absence of E<sub>2</sub><sup>29-31</sup>. For example, in the absence of E<sub>2</sub>, IGF-1 and activators of the protein kinase A can activate ER $\alpha$  and induce otherwise E<sub>2</sub>-dependent gene transcription<sup>29-31</sup>. Therefore, it is plausible that ER $\alpha$  activation is necessary for female energy balance even when its ligand, E<sub>2</sub>, is diminished in concentration or absent, as in a post-menopausal environment or due to aromatase inhibitor use. In such E<sub>2</sub> depleted instances, ER $\alpha$  is activated by other factors. Further investigation into the specific ligand-dependent and ligand independent mechanisms of energy balance may elucidate how female metabolism is maintained in an E<sub>2</sub> depleted environment with hypothalamic ER $\alpha$  expression intact.

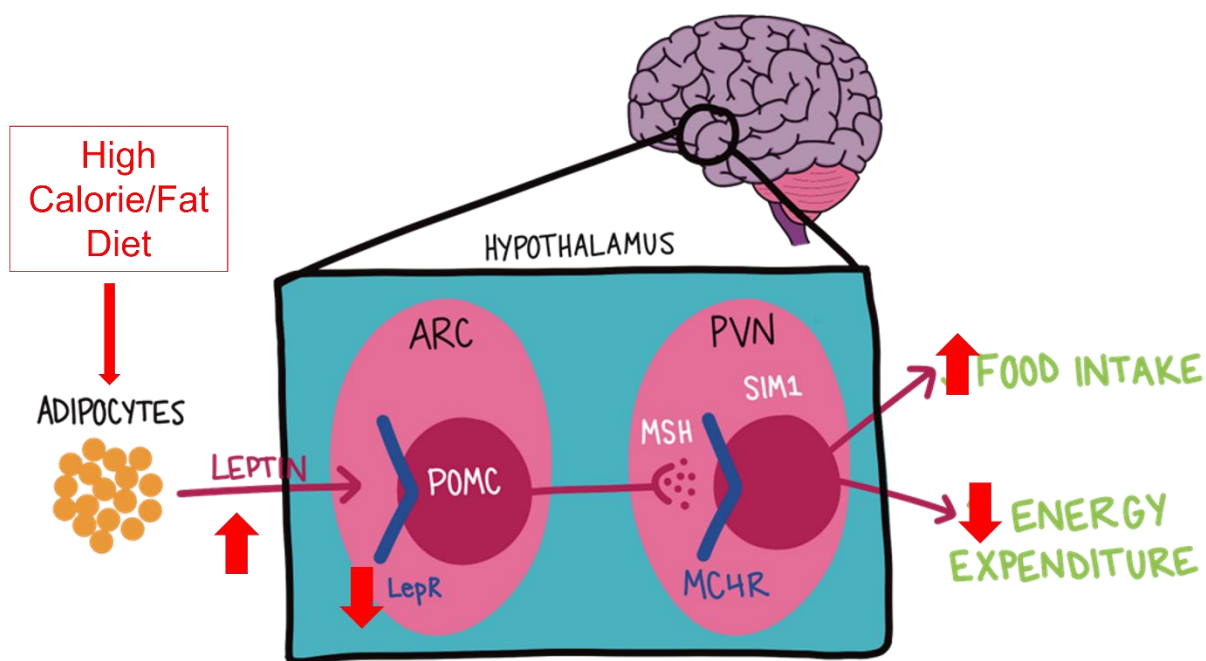
#### *5.6 Applications of this study to Understanding and Treating Metabolic Dysfunction in Post-menopausal Women and Female Cancer Survivors*

Our recently completed marmoset and rhesus macaque studies sought to investigate whether or not failure of ovarian E<sub>2</sub> depletion (OVX or VEH) to reliably increase female NHP body weight is due to residual actions of extra-ovarian E<sub>2</sub>, including neuro-E<sub>2</sub> via estrogen receptor alpha (ER $\alpha$ ) orchestrating energy balance in the hypothalamus. We demonstrated that NHP ER $\alpha$  expression may indeed be the predominate estrogen receptor governing female NHP metabolism and that there is a component of extra-ovarian E<sub>2</sub> involved in metabolic regulation, but the latter can be highly NHP species specific.

The studies in this thesis identified how organs or sub-regions therein, including the brain and its hypothalamus, may sense and adjust to changes in systemic hormone levels to maintain energy homeostasis. It is of utmost importance that future research aims to identify tissue-specific mechanisms of E<sub>2</sub>-mediated and E<sub>2</sub>-independent

metabolic regulation. Specifically, how organ specific hormonal mechanisms may change during systemic reduction of E<sub>2</sub>. Understanding the roles of endocrine organs such as adipose, liver and brain, to maintain energy homeostasis during the menopausal transition or following cancer survival will enable discovery of novel therapeutic targets to improve negative symptoms of menopause and cancer survival while reducing harmful off target effects of current HRTs or the inability to administer HRT because of the increased risk of recurring E<sub>2</sub>-associated cancer.

## 5.7 Figures



**Figure 1:** A high calorie/fat diet or diet induced obesity (DIO), increases adipose tissue via fat accumulation, resulting in increased levels of circulating leptin. The heightened levels of circulating leptin reduce the sensitivity of leptin receptors and reduce downstream connections, resulting in increased food intake and decreased energy expenditure.

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