

Hypothalamic Estrogen Signaling and Adipose Tissue Metabolism in Energy Homeostasis

Valeria C. Torres Irizarry^{1,2}, Yuwei Jiang^{1,2*}, Yanlin He^{3*} and Pingwen Xu^{1,2*}

¹ Division of Endocrinology, Diabetes, and Metabolism, Department of Medicine, The University of Illinois at Chicago, Chicago, IL, United States, ² Department of Physiology and Biophysics, The University of Illinois at Chicago, Chicago, IL, United States, ³ Pennington Biomedical Research Center, Louisiana State University System, Baton Rouge, LA, United States

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*Correspondence:

Yuwei Jiang yuweij@uic.edu Yanlin He Yanlin.He@pbrc.edu Pingwen Xu pingwenx@uic.edu

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Torres Irizarry VC, Jiang Y, He Y and Xu P (2022) Hypothalamic Estrogen Signaling and Adipose Tissue Metabolism in Energy Homeostasis. Front. Endocrinol. 13:898139. doi: 10.3389/fendo.2022.898139 Obesity has become a global epidemic, and it is a major risk factor for other metabolic disorders such as type 2 diabetes and cardiometabolic disease. Accumulating evidence indicates that there is sex-specific metabolic protection and disease susceptibility. For instance, in both clinical and experimental studies, males are more likely to develop obesity, insulin resistance, and diabetes. In line with this, males tend to have more visceral white adipose tissue (WAT) and less brown adipose tissue (BAT) thermogenic activity, both leading to an increased incidence of metabolic disorders. This female-specific fat distribution is partially mediated by sex hormone estrogens. Specifically, hypothalamic estrogen signaling plays a vital role in regulating WAT distribution, WAT beiging, and BAT thermogenesis. These regulatory effects on adipose tissue metabolism are primarily mediated by the activation of estrogen receptor alpha (ER α) in neurons, which interacts with hormones and adipokines such as leptin, ghrelin, and insulin. This review discusses the contribution of adipose tissue dysfunction to obesity and the role of hypothalamic estrogen signaling in preventing metabolic diseases with a particular focus on the VMH, the central regulator of energy expenditure and glucose homeostasis.

Keywords: estrogen, ER α , ER β , VMH, energy homeostasis, WAT, BAT

INTRODUCTION

Obesity is a health condition characterized by excessive accumulation of body fat and has become a global epidemic (1). Obesity is associated with numerous metabolic complications such as insulin resistance, glucose intolerance, and hyperlipidemia (2–4). Up to date, therapeutic strategies often include changes in diet and exercise. However, these interventions exhibit little long-term impact. Therefore, there is a critical need to understand the influencing factors that regulate fat expansion and identify alternative long-lasting solutions.

Notably, sex plays a key role in adipose tissue distribution and energy balance regulation. Abundant clinical and epidemiological studies show that men and women differ in body fat accumulation and distribution (5–8). Women have more fat accumulated in subcutaneous white adipose tissue (WAT) (8) and have greater metabolic activity in brown adipose tissue (BAT) when compared to men (9). In contrast, males tend to accumulate visceral fat, which promotes metabolic disorders. This sex-specific fat storage is presumed to implicate evolutionary pressures. Women are

adapted to store fat with a low lipolytic rate to respond to chronic energetic challenges such as gestation and lactation (10), protecting them against metabolic diseases (11). In contrast, men have evolved to store metabolically active visceral fat depots as energy fuel under short-term energetic challenges such as hunting (10). However, the sex-specific mechanisms that regulate adipose tissue metabolism are not fully understood.

One of the sex steroids accounting for these sex differences is estrogen. Accumulating evidence suggests that estrogen plays a vital role in sex dimorphic control of metabolic diseases and adipose tissue metabolism (12-14). More specifically, brain estrogen signaling has been shown to enhance sympathetic nervous system (SNS) output, promote thermogenesis in BAT, prevent hyperplasia and hypertrophy in WAT, and regulate WAT distribution (15-20). However, the underlying mechanisms by which central estrogen modulates adipose tissue metabolism are not well characterized. Previous studies were mainly focused on the local effects of estrogen in the brain or adipose tissue (21, 22), but the roles of estrogen-initiated crosstalk between brain and fat in determining sex- and depot-specific adipose tissue function have not been well investigated. This review will summarize the existing evidence regarding brain estrogen-initiated inputs regulating adipose tissue metabolism.

OBESITY AND METABOLIC SYNDROME

Individuals are considered obese when their Body mass index (BMI) is higher than 30kg/m^2 (23). Obesity occurs as the result of energy imbalance when the caloric intake surpasses energy expenditure, which is influenced by diverse behavioral, socioeconomic, and genetic factors (24). Examples of these factors are sedentary lifestyles or limited physical activity, and excessive caloric intake. Improvement of lifestyle by engaging in activities that require physical exercise and reducing the consumption of high fat/carbohydrate foods can positively impact obesity and metabolic syndrome (25-27). Genetic factors also play a crucial role in energy imbalance and excessive adiposity. Genome-wide associated studies (GWAS) had successfully identified 445 single nucleotide polymorphisms (SNPs) and 389 genes associated with obesity (28). Moreover, several SNPs were found to target miRNAs linked with adipogenesis and lipid metabolism (28). Clinical studies in patients with obesity have revealed that mutations in leptin, pro-opiomelanocortin (POMC), and melanocortin 4 receptor (MC4R) are positively associated with hyperphagia, hyperinsulinemia, and excessive adiposity (29-34). Accordingly, several research efforts have focused on developing therapies against obesity by targeting these genes, mainly to enhance satiety and reduce energy intake.

ADIPOSE TISSUE FUNCTION

To better understand the pathogenesis of obesity, it is crucial to define adipose tissue function. Adipose tissue is no longer considered only a storage depot for excess energy. This tissue also acts as a regulator of numerous metabolic and endocrine responses. For instance, adipose tissue regulates appetite, thermogenesis, lipid metabolism, sexual reproduction, immunological responses, insulin signaling, and glucose homeostasis. Moreover, adipose tissue is a highly dynamic organ that consists of numerous cell types, including adipocytes, fibroblasts, endothelial cells, immune cells, and adipocyte progenitor cells. These progenitor cells can undergo adipogenesis to create new adipocytes (35). There are two major types of adipose tissue, white adipose tissue (WAT) and brown adipose tissue (BAT). As detailed below, the excessive adiposity observed in obesity results from the dysregulation of both WAT and BAT metabolic function.

White Adipose Tissue

In living organisms, energy consumption and storage are essential to ensure survival in times of energetic challenges and limited caloric availability. WAT functions as an energy reservoir to store lipids in the form of triglycerides. WAT is mainly composed of white adipocytes, unilocular cells with one large lipid droplet where free fatty acids (FFA) are stored in the form of triglycerides (Figure 1). When there is an energy demand for example, after physical activity or under food scarcity, white adipocytes release energy by undergoing lipolysis, a process orchestrated by hormone-sensitive lipase (HSL) and adipose triglyceride lipase (ATGL), where triglycerides are catabolized into FFAs that enter circulation and are delivered to energydemanding organs such as muscle and liver (35, 37). In addition to its energy storage function, WAT is also an endocrine organ that secrets adipokines such as leptin, adiponectin, and resistin, all of which contribute to food intake and energy balance regulation (38-41)

WAT depots can be further categorized based on their anatomical distribution. Subcutaneous WAT (sWAT) is mainly located in gluteal and femoral regions, whereas visceral/omental adipose tissue (vWAT) is primarily accumulated in the abdomen and internal organs (42, 43). sWAT is associated with optimal metabolic health, whereas vWAT contributes to metabolic dysregulation. For instance, in humans, sWAT expansion is linked to improvement in insulin sensitivity, diminished lipolysis rate, decreased circulation of cytokines, and augmented levels of adipokines (44). In contrast, vWAT expansion is associated with increased insulin resistance, systemic inflammation, and other metabolic syndrome features (45-47). However, it is not clear why vWAT is more metabolically detrimental. It is likely attributed to the anatomical location of vWAT, which is surrounding internal organs within the body cavity. Moreover, vWAT appears to be a highly metabolically active depot, which constantly mobilizes lipids and increases circulating FFA levels. In addition to gender difference, adipose tissue expansion is dependent on genetic components and diet (48). White adipocytes can increase in number (hyperplasia) or size (hypertrophy) in response to excessive energy intake (48, 49). While WAT is highly plastic that can expand from 4% to 50-70% of an individual's body composition, the total capacity is limited. When WAT expansion goes beyond its capacity, excess fat begins to accumulate in tissues, such as skeletal muscle, liver, heart, and pancreas, ultimately



leading to lipotoxicity, systemic inflammation, insulin resistance, and metabolic disorders (50-52)

Brown and Beige Adipose Tissue

In contrast to WAT, BAT is a highly vascularized organ mainly composed of brown adipocytes. Brown adipocytes have small lipid droplets and are rich in mitochondrial content, which contributes to their brown appearance (Figure 1) (53, 54). The primary function of BAT is thermogenesis, a process by which the uncoupling protein 1 (UCP1, expressed in the inner mitochondrial membrane) uncouples oxidative phosphorylation to dissipate energy as heat (54-56). In both humans and rodents, BAT thermogenesis is mainly dependent on the sympathetic nervous system (SNS) activation (55-59). Activation of SNS drives the release of norepinephrine (NE) which activates β 3-adrenoceptors (β 3-AR) expressed in brown adipocytes. Mitochondria typically generate energy by synthesizing ATP via oxidative phosphorylation. However, β3-AR activates the cyclic AMP/Protein kinase A pathway to release FFA, which will be used as fuel by UCP1 to uncouple ATP synthesis and release the produced energy in the form of heat (55, 58). BAT thermogenesis can be induced by multiple factors such as diet, cold exposure, and β -adrenoceptors agonists (53, 57, 58, 60-63). In addition to UCP1, other important thermogenic and BAT markers have been identified including peroxisome proliferator-activated receptor gamma coactivator 1alpha (PGC1a), PR domain containing 16 (PRDM16), cell deathinducing DFFA-like effector A (CIDEA), and type II iodothyronine deiodinase (DIO2) (20, 57, 59, 64). It was once thought that BAT was absent in adulthood. However, positron emission tomographycomputed tomography (PET/CT) scanning technologies with ¹⁸Ffluorodeoxyglucose (¹⁸FDG) have revealed that adult humans have metabolically active BAT depots (9, 56, 61, 62). BAT content in humans can be found in cervical supra-clavicular, axillary, and paraspinal regions, accounting for approximately 2% of total fat depots (Figure 1) (9, 61). As an endocrine organ, BAT secretes batokines as well as cytokines such as fibroblast growth factor 21 (FGF21) and interleukin 6 (IL-6) (65, 66). However, the endocrine

functions of BAT in physiological and pathological conditions still need to be fully explored.

In addition to its thermogenic and endocrine function, BAT also plays an essential role in glucose homeostasis. BAT glucose uptake can be stimulated by SNS-activated thermogenesis or insulin signaling (67). BAT expresses insulin receptors that allow translocation of GLUT4 into the plasma membrane, thus facilitating glucose clearance (35, 67). Studies have shown that human-to-mice BAT transplantation results in increased insulin sensitivity, decreased body weight gain, and attenuated high fat diet (HFD)-induced insulin resistance (66). In line with this, fasting-induced insulin resistance is accompanied by decreased BAT glucose uptake and thermogenic activity in humans (68). These findings provide additional potential connection between BAT dysfunction and metabolic syndrome.

Under specific physiological stressors, brown-like adipocytes can arise from white adipocytes or adipocyte progenitor cells. These cells are termed beige/brite adipocytes and the process by which they emerge is called WAT browning or beiging. Beige adipocytes share morphological features with both white and brown adipocytes. Beige adipocytes have larger lipid droplets for fat storage, and although they have less mitochondrial density, they are UCP1 positive cells and they possess thermogenic capacity (69, 70). Similar to BAT activation, WAT beiging can be induced by cold exposure, physical activity, and β -adrenergic receptors (Figure 1) (57, 63, 71). BAT thermogenesis and WAT browning are potential therapeutic targets to treat obesity and metabolic syndrome because they can consume glucose and fatty acid. Cold exposure in humans results in increases in BAT content, thermogenic activity, and energy expenditure, accompanied by lower BMI and body fat content (61, 72, 73). Conversely, BAT content was lower in individuals with higher visceral fat deposition and BMI (72). Additionally, individuals with detectable active BAT depots display lower circulating cholesterol and glucose levels (74). In rodents, UCP1 ablation leads to body weight gain, increased food intake, and decreased diet-induced thermogenesis, phenotypes that are exacerbated by

HFD (75). Activation of β 3-AR using CL316,243 agonist remodeled white adipocytes into more multilocular structures and smaller lipid droplets, associated with increased expression of brown adipocyte markers and UCP1 protein (57). Murano et al. reported an increase in noradrenergic fiber density in BAT and WAT following 10 day-cold exposure in adult female mice (63). The abundant evidence from the animal and clinical studies strongly support that increasing BAT activity and inducing WAT beiging represent viable strategies to prevent and treat obesity and its related comorbidities.

However, clinical evidence on BAT activation contribution to body weight loss is scarce. To this date, there are more than 40 ongoing clinical trials aiming to better understand chronic BAT activity in humans and test potential therapeutic approaches. In healthy males, long-term cold exposure (17°C, 2 hours/day for 6 weeks) stimulated BAT activity, increased energy expenditure, and significantly reduced body fat mass (76). Recently, mirabegron, a pharmacological agent and β3-adrenergic receptor agonist, has been tested in both men (NCT01783470) and women (NCT03049462). Healthy men receiving 200 mg mirabegron (3 interventions within 28 days) showed increased acute BAT activation and resting metabolic rate (RMR) (77). On the other hand, healthy women received daily 200 mg mirabegron for 28 days, resulting in enhanced BAT metabolic activity, increased resting energy expenditure, and improved insulin sensitivity. However, no changes in body weight mass were detected in these women (78). These findings position mirabegron as a promising therapeutic agent to stimulate endogenous BAT thermogenic activity in patients with metabolic syndrome. Further research is needed to identify the cellular, molecular, and transcriptional mechanisms governing the BAT/beige development and function.

SEXUAL DIMORPHISM IN ADIPOSE TISSUE

Sex-specific distribution of adipose tissue has been described in humans and animal models. Females are more abundant in sWAT, while males tend to have more abdominal-visceral depots (79, 80). Males have evolved to store highly metabolically active visceral fat depots that can be quickly mobilized and used as energy fuel under short-term energetic challenges such as hunting. On the other hand, females might have evolved to store fat subcutaneously with low lipolytic rates, which would store the energy for long-term challenges such as pregnancy and lactation (10). This evolutionary adaptation might confer protection against obesity and metabolic syndrome. Indeed, in studies using mice exposed to HFD for twelve weeks, males tend to gain more body weight and have higher expression of proinflammatory genes in the adipose tissue when compared to females (81). In contrast, women with vWAT accumulation are at higher risk of developing impaired glucose and lipid metabolism and cardiovascular complications when compared to men (7, 8). Clinical studies have shown that catecholamineinduced lipolysis is predominant in female but not male abdominal adipocytes (82). Additionally, during fasting, while FFA increases in the blood from both sexes, women have lower levels of circulating glucose when compared to men. This diminished glucose production is speculated to protect women from FFA-induced insulin resistance (83). Interestingly, a study addressing sex-specific responses to exercise revealed that women derived more energy expenditure from fat oxidation. In contrast, men utilized carbohydrates as the primary fuel during the exercise sessions (84). Taken together, these studies suggest that a higher lipolytic rate in women is possible because they are more dependent on fat as an energy source.

Importantly, the protective role of sWAT in females seems to be age-dependent, as postmenopausal women suffer fat redistribution. Fat depots from subcutaneous regions are transferred to visceral regions (85). This has led to the conclusion that sex hormones might play a critical role in gender-specific fat distribution and overall metabolic health. Particular attention has been given to estrogen as it has been shown that the decreased circulation of estrogen contributes to increased adiposity, insulin resistance, low metabolic rate, and adipose tissue inflammation (59, 86, 87).

One proposed mechanism for gender-specific fat distribution is the estrogens-mediated modulation of sympathetic inputs and lipolytic/lipogenic rates in adipose tissues. In human WAT, adrenergic activation of β -adrenergic receptor (AR) promotes lipolysis, whereas adrenergic stimulation of a2A-AR inhibits lipolysis (88, 89). As discussed earlier, sWAT in women is characterized by a lower lipolytic rate to ensure energy preservation. In line with this, healthy women under long-term estradiol (E2) treatment have increased α 2A-AR mRNA levels and protein binding capacity in subcutaneous depots compared to placebo-treated individuals. E2 did not affect the α 2A-AR mRNA profile in vWAT (88). In vitro studies with cultured adipose tissue fragments consistently show that E2 treatment exerts anti-lipolytic effects in subcutaneous adipocytes (88). This suggests that estrogens modulate female WAT distribution in part by upregulating α 2A-AR signaling in sWAT while stimulating β -AR-mediated high lipolytic rate in vWAT, thus promoting subcutaneous energy storage.

Moreover, these previous findings were associated with estrogen receptor alpha (ER α), which has been evidenced to be the main target for estrogen's anti-obesity effects (further discussed below) (88). In line with this, mutations in ER α are linked with higher BMI, waist circumference, and increased fat mass in middle-aged women (90). In rodents, central E2/ER α signaling has been shown to contribute to changes in fat distribution. Deletion of ER α in the VMH SF1 neurons leads to increased body weight, massive gonadal WAT (gWAT, rodent visceral adipose depot) expansion, increased lipogenesis, and decreased sympathetic tone (20). The role of estrogen in the regulation of energy balance will be discussed in more detail in the following section.

Similar to WAT, sex differences in BAT have also been described. The volume of BAT depots between the sexes is similar. However, women have higher BAT mass in the

cervical-supraclavicular region when compared to men, and female BAT is metabolically more active than males (9, 61, 91). Cold exposure results in significantly higher energy expenditure in women (61). Interestingly, the higher prevalence of metabolically active BAT in women is positively correlated with plasma estradiol levels (92, 93). Treatment of β 3adrenergic receptor agonist CL316,243 (CL) upregulates thermogenic markers and mitochondrial respiratory chain proteins in gWAT of female mice but not in males, indicating sex-specific WAT browning phenotypes (57). Female rats have higher BAT activity and oxygen consumption than males under *ad libitum* conditions. However, under caloric restriction, these phenotypes are reversed, as indicated by diminished mitochondrial density and downregulation of thermogenic genes in females (94).

A recent report by MacCannell et al. used high-resolution respirometry to assess sex-specific mitochondrial function and metabolic flexibility in mice challenged with HFD. HFD-fed females show increases in electron transport chain (ETC) components, specifically Complex I and II respiration, in BAT mitochondria compared to females fed a standard diet, suggesting adaptive mitochondrial BAT respiration. Notably, HFD-fed males did not exhibit this phenotype. In addition, improved mitochondrial respiration is accompanied by upregulation of BAT-associated genes. These findings show that under HFD condition, female mice have greater metabolic flexibility and adapt to nutritional challenges by maximizing the activity of ETC complexes to modulate energy expenditure and restore energy balance (95). Altogether, these findings provide evidence for sex-specific regulation of BAT metabolism and suggest mitochondrial metabolic flexibility as a potential mechanism underlying the physiological relevance of BAT function in women.

ESTROGEN

Estrogen is a sex hormone mainly produced in the ovaries, corpus luteum, placenta, and other non-gonad regions like the heart, liver, and brain (96). In females, the three types of estrogen present are estrone (E1), estradiol (E2/17β-estradiol), and estriol (E3). Estrogen biosynthesis initiates with cholesterol. A series of chemical reactions result in the synthesis of testosterone. The action of aromatase mediates the conversion from testosterone to estradiol (96). E2 or 17β -estradiol is the most active form of estrogen in premenopausal women. The levels of estrogen are decreased post-menopause, which is associated with a higher risk of metabolic complications and body weight gain. The antiobesity effects of estrogen and its role in energy homeostasis are well evidenced. For instance, Hong et al. demonstrated that compared to females, male mice were more susceptible to increased body fat. However, women begin to experience body weight gain after menopause (97). Importantly, ovarian hormone depletion by OVX removed female mice protection against body weight gain (86). In line with this, hormone replacement therapy in postmenopausal women or E2 replenishment in OVX mice reverts the obesogenic phenotypes (97, 98). It has also been

reported that food intake and body weight gain in rats varies depending on the estrous cycle stage, pregnancy, or lactation (59). In female mice, disrupted estrogen signaling leads to obesity, decreased locomotion, and reduced heat production and energy expenditure (19), demonstrating estrogen's impact on thermogenesis. Estrogen signaling is mediated by its receptors, Estrogen receptor alpha (ER α), Estrogen receptor beta (ER β), G protein-coupled receptor 30 (GPR30) and Gq protein-coupled receptor (Gq-mER), all of which are expressed in the brain and adipose tissue (96).

ERα

From all the estrogen receptors, $ER\alpha$ is the most extensively studied in the context of energy homeostasis, given the abundant evidence suggesting that ERa mediates estrogen's anti-obesity effects. ERa is expressed in gonadal organs, liver, adipose tissue, and brain. In the brain, $ER\alpha$ expression predominates in the bed nucleus of the stria terminalis, the nucleus tractus solitarius (NTS), amygdala, hypothalamus, periaqueductal gray and dorsal raphe nucleus (DRN) (99). Specific hypothalamic regions that express ERa include POMC neurons and neuropeptide Y (NPY)/agouti-related protein (AgRP) neurons, which are found in the arcuate hypothalamic nucleus (ARH), ventromedial hypothalamic nucleus (VMH) and medial preoptic area (MPOA) (99). All these brain regions are implicated in food intake and thermoregulation. Mutations in ESR1, the gene that encodes $ER\alpha$, have been associated with higher BMI and increased fat mass in women (90). In rodents, ERa congenital global KO results in increased adiposity and elevated serum glucose and insulin in both male and female mice (100). Consistently, a point mutation of the palmitoylation site of ERa (C451A-ERa), which results in membrane-specific loss of function, impaired glucose sensing in female mice (101), and disturbed sexual differentiation in the perinatal programming of the male brain (102). Depletion of ER α in the mouse brain abolishes the beneficial metabolic effects of estrogen, resulting in hyperphagia, body weight gain, increased visceral adiposity, and impaired energy expenditure (17, 20, 59, 103). In human adipose tissue, expression of ERa is positively associated with UCP1 protein density (80). Similarly, adipose tissue ER α expression in mouse is correlated with lower adiposity, higher UCP1 content, optimal BAT thermogenic function, and insulin sensitivity (104). Activation of ERa with synthetic agonist propylpyrazoletriol (PPT) upregulates UCP1 and causes WAT beiging both in vitro and in vivo (103).

ERβ

The role of ER β in adipose tissue function is more controversial. This receptor is primarily expressed in bone marrow, endothelium, lungs, adipose tissue, and the brain. In the brain, ER β is most abundant in the hippocampus, bed nucleus of the stria terminalis, amygdala, DRN, and very limited in the hypothalamus (99). ER β has not been paid as much interest as ER α , not only because of the evidence discussed earlier favoring ER α as the principal mediator of estrogen signaling, but also because most studies targeting ER β have concluded that it does not affect energy homeostasis. For example, depletion of ER β in animals has no effect on total adiposity in mice (105). In OVX rats, activation of ER α by its agonist propylpyrazole triol (PPT) causes body weight loss and hypophagia associated with a reduction in meal size. However, using diarylpropionitrile (DPN) to activate ER β does not influence body weight and food intake (98, 106). In contrast, recent reports have shown that ER β might have a beneficial impact on metabolic health and potentially a protective function. González-Granillo et al. recently demonstrated that subcutaneous injection of ER β agonist 4-(2-(3,5-dimethylisoxazol-4-yl)-1H-indol-3-yl)phenol (DIP) induces fat redistribution from visceral to subcutaneous depots, elevates heat production, and stimulates WAT browning in male mice (107). Similarly, activation of ER β using a synthetic agonist (b-LGND2) decreases body weight and fat mass in WT mice. Deletion of ER β reversed these phenotypes (108).

It has also been reported that $ER\beta$ is more abundant in healthy females when compared to men. Recently Porter and colleagues studied the transcriptional and protein content of estrogen receptors in obese individuals, including men, premenopausal and postmenopausal women. They report that ER α and ER β protein expression in adipose tissue was positively correlated with UCP1 content, further evidencing the role of ER α in BAT thermogenesis and suggesting that ER β may also play an important part. Importantly, they showed that ERB gene expression was higher in adipose tissue of women when compared to men (80). It has been suggested that in the absence of ERa, the available estrogen (produced in adipose tissue, the second primary source of estrogen after ovaries) is forced to activate ERB as compensatory mechanism, indicating that ER β may have a protective metabolic role in the absence of estrogen or ER α (109). However, further studies are required to clarify the discrepancies in the field and elucidate the potential role of ERβ in metabolic health.

GPR30

Although ER α and ER β are the most studied ERs in the context of energy homeostasis, there are other novel receptors that represent a promising therapeutic target for metabolic syndrome. The GPR30 or G protein-coupled estrogen receptor 1 (GPER1) is expressed in reproductive organs, adipose tissue, pancreas, bone tissue, and brain, including the hypothalamus (110–114). The binding of estrogen to GPR30 initiates a signaling cascade that activates the protein kinase A (PKA)/ERK pathway (113).

Although GPR30 has a lower binding affinity to estradiol when compared to ERα (113), various reports employing a whole-body deletion model (GPR30 KO) suggest that this receptor is essential for the maintenance of glucose and energy balance. Deletion of GPR30 in mice leads to increased body weight attributed to increased fat mass, augmented circulating levels of cholesterol and triglycerides, glucose intolerance, impaired insulin sensitivity, and elevation of proinflammatory cytokines (115). Consistent with these findings, Davis et al. reported that whole-body GPR30 KO mice show elevated body weight, increased adipocyte size, BAT remodeling, and decreased UCP1 gene expression and energy expenditure (112).

Interestingly, male mice exhibit more susceptibility to GPR30 deletion when compared to females, as male GPR30 KO mice

show significant changes in body weight as early as 8 weeks of age, but GPR30 KO females begin to significantly gain weight at 13 weeks of age. In addition, adipocyte area and UCP1 levels are not significantly changed in female GPR30 KO mice as opposed to male mice. In the same study, GPR30 KO females were ovariectomized and treated with E2. Estradiol supplementation did not affect body weight changes, glucose tolerance, and adipocyte morphology in these GPR30 KO OVX mice. These findings imply that GPR30 in males is required to maintain optimal body weight and energy expenditure, and in females, it confers protection against adipose tissue remodeling and glucose imbalance (112).

It is important to note that in female mice, deletion of GPR30 decreased hypothalamic ER α and increased ER β gene expression, suggesting that GPR30 may regulate the expression of classical ERs in the hypothalamus (112). Female mice lacking GPR30 also show hyperglycemia and decreased insulin expression and release from the pancreas (114). In contrast to these findings, another report indicates that GPR30 KO female mice fed an HFD exhibit decreased body fat mass, improved glucose balance and insulin sensitivity, reduced adipocyte size and adipogenesis, indicating that in HFD-fed females, GPR30 contributes to adipose tissue expansion and energy imbalance (110). Recently, Sharma et al. reported that GPR30 agonism with GPR30-selective small-molecule agonist G-1 reverses OVXinduced obesity by decreasing body weight and fat mass, increasing energy expenditure, promoting BAT remodeling, improving glucose homeostasis, and lowering proinflammatory responses. These phenotypes persisted in OVX mice challenged with HFD, demonstrating that GPR30 activation exerts anti-obesity effects in female mice (116).

Gq-mER

The Gq protein-coupled receptor (Gq-mER) has gained attention over the past years, but limited evidence links this membrane ER with energy balance. Gq-mER is mainly expressed in the hypothalamus, including both NPY/AgRP and POMC neurons in ARH, the paraventricular nucleus (PVN), and the MPOA (117-122). Early studies demonstrate the expression and activation of hypothalamic Gq-mER, as administration of STX (Gq-mER selective ligand) induces POMC neural responses in ERαKO, ERβKO, and GPR30KO mice (117, 122). In addition, STX mimics the effect of E2 replacement in OVX guinea pigs by decreasing body weight and uterine weight (122). The suggested mechanism by which Gq-mER regulates neural responses in ARH is through GABA_B receptors expressed in NPY/AgRP and POMC neurons. Gq-mER activation desensitizes GABA_B receptors, increasing membrane excitability in POMC neurons, while in NPY/AgRP neurons, it enhances GABA_B inhibitory effects, indicating that Gq-mER agonism triggers anorexigenic signals (118, 123). In line with this, daily subcutaneous injection of STX results in decreased food intake in gonadectomized male and female guinea pigs (118). Gq-mER signaling has also been associated with the regulation of body temperature. Treating OVX guinea pigs with STX results in reduced core body temperature accompanied by decreased fat pad weight and body weight and reductions in food intake and meal

frequency (121). In summary, Gq-mER has been shown to play an essential role in estrogenic regulation of energy balance. Further research is required to identify the potential therapeutic applications of these ERs.

ESTROGEN IN THE BRAIN

In 2011, we provided genetic evidence that brain ER α signaling is vital for body weight control. In this study, we crossed $ER\alpha^{flox/flox}$ with Nestin-Cre mice to selectively delete ERa from CNS (ER α^{CNS} -KO). We found that compared to their littermate controls, both male and female $ER\alpha^{CNS}$ -KO mice have increased body weight, which is solely reflected by an increase in fat mass, but not lean mass. In addition, we found these mice eat more and have decreased energy expenditure accompanied by reduced physical activity. Importantly, we showed that $ER\alpha^{CNS}$ -KO mice have a larger visceral fat depot than the controls, which in other words, they have abdominal obesity. Finally, we found that $ER\alpha^{CNS}$ -KO females have increased E2 in the blood (20). This is important when put in a broader perspective, as it indicates that increased peripheral estrogen signaling does not rescue obesity induced by brain-specific KO of ERa, suggesting an essential role of brain ERa in body fat distribution and energy expenditure.

ARH

Fluctuations in feeding behavior are estrous cycle-dependent in female rodents. Specifically, food intake elevates during the metestrous-diestrous phase, when estrogen levels are at the lowest. Conversely, food intake decreases to the lowest point during the proestrus-estrous stage, in which estrogen levels are high (124–126). Notably, the reduced food intake and body weight during the proestrus phase is associated with decreased NPY/AgRP and increased POMC mRNA expression, suggesting a potential role of ARH appetite-regulating neuropeptides in estrogenic regulation of food intake (125). Consistently, POMC excitatory synapses are enhanced during the proestrus phase (126). E2 supplementation in OVX mice decreases appetite and adiposity, associated with enhanced POMC neuronal activity, as indicated by increases in POMC excitatory synapses and c-fos expression (126). ERa congenital globe KO lowers POMC immunoreactivity, which was not reversed by E2 replenishment, suggesting that POMC synaptic plasticity is mediated by E2/ER signaling. These findings support a model that estrogens modulate food intake by inhibiting AgRP neurons and activating POMC neurons, resulting in an anorexigenic response (126).

In supporting this point, administration of E2 directly into the ARH fails to decrease food intake in mice lacking AgRP/NPY neurons (121), suggesting a vital role of AgRP/NPY neurons. However, although ER α is the primary mediating receptor for anorexigenic effects of estrogens, ER α does not express in AgRP/ NPY neurons (121). These findings indicate that estrogens may act through ER α expressed by presynaptic neurons to regulate AgRP/NPY neurons. Consistently, the following study demonstrated that the anorexigenic effects of E2 are blocked in female ER α ^{flox/flox}/POMC-Cre mice with ER α selectively deleted from POMC progenitor neurons (120), implying a mediating role of ER α expressed by POMC progenitor neurons. Mechanistically, phosphatidylinositol 3-kinase (PI3K) has been suggested to mediate the metabolic functions of ER α signals. Specifically, estrous cycle-dependent fluctuations in food intake are blunted in female mice with PI3K genetically inhibited in the POMC progenitor neurons (124). These results indicate that an ER α -PI3K cascade in POMC progenitor neurons mediates estrogenic actions to suppress food intake.

In line with this, ablation of ER α , specifically in POMC progenitor neurons, leads to chronic hyperphagia, decreased leptin sensitivity, and body weight gain in female mice (20). Interestingly, a recent study showed that region-specific deletion of ER α in the ARH of ER $\alpha^{flox/flox}$ mice by stereotaxic delivery of AAV virus has no effect on food intake, suggesting that estrogen may exert its anorexic effects by targeting other brain sites rather than the ARH (16). This discrepancy could be due to the non-ARH deletion of ER α in the ER $\alpha^{flox/flox}/POMC$ -Cre model. The recombination in off-target sites could contribute to physiological phenotypes of POMC-Cre transgenics (127). It is possible that ER α deleted in NTS POMC neurons led to changes in food intake.

NTS

The NTS is a brainstem region that regulates satiety (128). E2 treatment has been shown to increase neuronal activity (as measured by c-fos expression) in the NTS of OVX rats. Moreover, c-fos expression colocalized with ER α in the NTS of E2-treated rats (129). Expression of ERs in the NTS of OVX rats is increased following 48-hour fasting (130). It has also been demonstrated that ER α mRNA expression in the NTS of OVX rats fluctuates with the estrous cycle, showing higher expression during estrous and lower expression during proestrus. OVX rats exhibit reduced ER α NTS mRNA expression compared to shamoperated rats (131).

DRN

The DRN is the primary site for 5-hydroxytryptamine (5-HT, serotonin) synthesis in the brain. Interestingly, 90% of the serotonergic neurons in the DRN (5-HT^{DRN}) coexpress ER α (132). Ovarian hormone depletion in female rats results in decreased c-fos expression in 5-HT^{DRN} neurons. Conversely, E2 replenishment increases serotonergic neural activity in the DRN (133). Specific deletion of ER α in the DRN of OVX mice results in binge-like eating behavior, and estrogen replacement fails to reverse this phenotype. Moreover, activation of 5-HT by either E2 or PPT suppresses binge-like eating (132). A study in OVX rats by Santollo et al. reported that food intake was significantly decreased within 24 hours after specific E2 microinfusion in ARH, DRN, and the MPOA (134).

MPOA

The MPOA has been identified as a major thermoregulatory hypothalamic region. MPOA subregions show either coldsensing or warm-sensing properties in sham female rats. However, after estrogen depletion by OVX, c-fos signaling is significantly decreased, and thermosensitivity of MPOA subregions is impaired (135). The MPOA has also been linked with torpor regulation in rodents. A recent study showed that activation of $ER\alpha^+$ neurons in MPOA decreases core body temperature and lowers energy expenditure and metabolic rate. Furthermore, ablation of ER α in MPOA reverses the torpor-like state in females but not in males (136). These findings suggest an integrating role of MPOA estrogen signaling in the regulation of energy and temperature homeostasis. Besides estrogens, tamoxifen, a selective estrogen receptor modulator, has also been shown to modulate the metabolic functions of MPOA neurons. For example, subcutaneous injection of tamoxifen alters the transcriptional profile in all MPOA cell types by repressing metabolic genes. Hypothalamic ablation of ERa using ERaflox/flox/Nkx2-1-Cre mice, resulting in ERa deletion in MPOA, vlVMH, and ARH, eliminates the tamoxifenmediated gene repression and upregulates metabolic genes in the MPOA. In line with this, hypothalamic ablation of ERa blunts tamoxifen-induced inhibition of BAT thermogenesis and locomotion. These findings suggest that tamoxifen acts through $ER\alpha$ expressed by the MPOA neurons to impair energy expenditure and suppress metabolism-related genes (137).

Sagittalis Nucleus of the Hypothalamus (SGN)

SGN, a unique hypothalamic nucleus discovered in 2008, is located between the ARH and VMH. Interestingly, this novel hypothalamic region was rich in ER α -immunoreactive cells (138). Studies focusing on SGN's role in energy balance are very limited. C-fos immunoreactivity was higher in the SGN of OVX rats treated with estradiol when compared to control OVX rats. In line with this, neural activity is enhanced when rats of both sexes engage in sexual behavior, supporting the notion that ER α^{SGN} might play a role in sex-specific sexual arousal (139, 140). Further research might provide a better understanding of the physiological relevance of ER α^{SGN} in sexual behaviors.

VMH

ER α signaling in the VMH has also been shown to modulate thermogenesis and glucose homeostasis. Silencing ER α in VMH leads to hyperphagia, hyperglycemia, and decreased energy expenditure (20). The role of estrogen signaling in VMH will be discussed in more depth later in this article.

CENTRAL ESTROGEN INTERACTION WITH HORMONES, ADIPOKINES, AND KINASES

Leptin

Leptin is an adipokine synthesized in adipocytes and encoded by the obesity (ob) gene with anorexigenic properties. Leptin is considered a biomarker for obesity as the levels of leptin are directly proportional to body fat mass accumulation (141). Leptin signals through leptin receptors (LepR, encoded by db gene) in the brain to communicate the status of body fat stores. If fat depots are not metabolized, leptin will inhibit NPY/AgRP neurons and activate POMC neurons to exert an anorexigenic response and increase satiety (141). During LepR activation, JAK2 tyrosine kinase phosphorylates LepR, which initiates the recruitment and activation of the STAT3 signaling cascade. STAT3 is essential for leptin signaling, as deletion of STAT3 leads to hyperphagia and impairs energy expenditure (142). Mutations in the LepR-B gene results in obesity in humans and rodents (128). Leptin-deficient ob/ob mice receiving intraperitoneal injections of leptin experience decreases in body mass and food intake compared to ob/ob mice receiving PBS treatment (143). Estrogen has been reported to be a key determinant of serum leptin levels and central leptin sensitivity. In diabetic Akita female mice -mice carrying Ins2 mutation-, ER α ablation exacerbates hyperphagia by further decreasing central leptin signals and downregulating POMC gene expression (144). OVX rats display dramatic increases in serum leptin levels, associated with significant changes in body weight gain. After E2 replacement, serum leptin levels are decreased (145, 146). Both ob/ob and db/db mice treated with E2 for 4 weeks showed body weight loss, diminished fat mass, hypophagia, and energy expenditure. This was accompanied by elevated hypothalamic pSTAT3 and increased POMC immunoreactivity (126). In the hypothalamus, LepR has been shown to coexpress with ER α and ER β , particularly in the MPOA, PVN, dorsomedial hypothalamus (DMH), VMH and ARH (142). Interestingly, estrogen depletion by OVX decreases leptin gene expression in these hypothalamic regions (147). Similarly, E2 replenishment in OVX rats significantly increased hypothalamic leptin content, accompanied by decreases in energy intake and body weight (145). Deletion of LepR in vagal afferent neurons was shown to increase body weight gain, fat accumulation, adipocyte size, and meal numbers. These obesogenic phenotypes were associated with lower circulating estrogen and decreased expression of ERa in vagal afferent neurons (148). Altogether this evidence shows that central estrogen signaling interacts significantly with leptin-induced anorexigenic signals.

As a hormone that directly interacts with hypothalamic neurons to modulate WAT distribution and energy intake, it is not surprising that leptin has also been reported to act on thermoregulating brain regions to influence body temperature and BAT thermogenesis. In addition to ARH, leptin-sensitive neurons can be found in the MPOA, PVN, DMH, VMH, and lateral hypothalamus (LH) (149). These brain regions are well known for their thermoregulatory and thermogenic properties. Ob/ob mice display decreased body temperature at thermoneutrality, which is brought back to basal levels following leptin administration (150). Leptin deficiency affects body temperature during thermoneutral conditions and when being challenged by cold temperatures. Mice lacking both leptin and UCP1 (ob/ob.Ucp1^{-/-}) are unable to survive temperatures below 12°C. Interestingly, leptin treatment allowed *ob/ob*.Ucp1^{-/-} mice to adapt to cold exposure by increasing their core temperature and oxygen consumption (151). Leptin has been shown to upregulate

Ucp1 mRNA levels as well as thermogenic and sympathetic markers in rodents with intact BAT sympathetic innervation (152, 153).

A clinical study performed in pre- and post-menopausal women evaluated the relationship between ER0/ERB ratio and obesity. They showed that a lower $ER\alpha/ER\beta$ ratio in omental adipose tissue (vWAT) is positively correlated with higher BMI and waist/ hip/thigh circumference. Moreover, serum leptin levels significantly correlated with these obesity markers, suggesting that circulating leptin may mediate the regulatory effects of estrogen signaling on adipose tissue homeostasis (154). Similar to leptin, estrogen's effect on BAT thermogenesis, thermoregulation, cold adaptation, and energy expenditure has been evidenced (9, 20, 57, 59, 61, 87, 136, 155). As discussed later in this article, estrogen and leptin were identified as energy sensors that signal through the AMPK(VMH)-SNS-BAT axis to activate BAT thermogenesis (59, 156). Additional studies might be valuable to elucidate a potential estrogen-leptin interaction and its effects on energy expenditure and overall metabolic health.

Ghrelin

Ghrelin is an orexigenic hormone produced in the stomach that stimulates food intake *via* its growth hormone secretagogue receptors (GHSRs), which are expressed in the ARH and VMH (157). In contrast to leptin, ghrelin is secreted under fasting conditions and signals through GHSRs to activate NPY/AgRP neurons and inhibit POMC neurons, leading to an orexigenic response and an increase in food intake (155).

In estrogen-depleted rats, ghrelin mRNA in the stomach is increased, suggesting that the absence of ovarian hormones facilitates ghrelin synthesis. In line with this, ERa was coexpressed with ghrelin in an intact female rat stomach. E2 replenishment in OVX rats decreases plasma ghrelin levels and mRNA content in the stomach (158). Ghrelin-induced increases in food intake are tightly associated with the estrous cycle in female rats. Intracerebroventricular (ICV) administration of ghrelin results in increased energy intake principally during the diestrus cycle when estrogen levels are lower. Consistently, after ghrelin ICV infusion, neural activity is greater in ARH during the diestrus phase rather than the proestrus phase (159). Yokota-Nakagi et al. explored the effects of estrogen on ghrelin effect in HFD-fed OVX rats. As expected, estrogen depletion accompanied by HFD dramatically increased body weight, food intake, and vWAT. Implantation of E2 pellets reversed the obesogenic phenotypes. Administration of GHRP-6, a GHSR agonist, increased HFD intake in OVX mice treated with placebo pellets, but those mice with E2 replenishment were protected from the orexigenic effects of GHRP-6. Similarly, E2 decreased c-fos expression in ARH following GHRP-6 administration (155). These findings suggest that estrogen influences ghrelin signaling in ARH neurons.

Although ghrelin is widely known for its effects on energy intake, multiple reports have evidenced the role of ghrelin in energy expenditure. Lin et al. worked with old Ghsr^{-/-} mice to show that ablation of ghrelin receptor decreases body weight and improves insulin sensitivity. These phenotypes were induced by energy expenditure rather than changes in food intake or

physical activity. Specifically, the increase of energy expenditure in $Ghsr^{-/-}$ mice is accompanied by increases in O_2 consumption, CO₂ production, RMR, and respiratory quotient (RQ). In vitro and in vivo studies demonstrated that GHSR ablation results in increased UCP1 content and upregulated thermogenic genes in BAT (160). We showed that neuronspecific deletion of GHSR partially prevents DIO in mice by improving insulin sensitivity and metabolic flexibility, increasing energy expenditure and physical activity. DIO-neuron-deficient GHSR mice demonstrated higher body core temperature and resistance to cold exposure when compared to control mice. Interestingly, expression of GHSR was upregulated in the ARH and VMH of wild-type mice fed an HFD when compared to mice fed a regular diet, suggesting that ghrelin signals through these hypothalamic neurons to counteract the effects of HFD (157). The potential role of ovarian hormones (including estrogen) in ghrelin-regulated energy expenditure remains to be elucidated.

Insulin

Adipose tissue dysfunction and obesity are tightly linked to insulin resistance and diabetes (161). Insulin is a hormone synthetized and secreted by β -cells in the pancreas in response to elevated circulating glucose levels. In homeostatic conditions, insulin induces lipogenesis to decrease plasma FFA. It also stimulates tissue glucose uptake by binding to the insulin receptors in adipose tissue and skeletal muscle, which activates glucose transporter GLUT4, thus lowering blood glucose levels. Under excessive caloric intake, the insulin demand increases and insulin receptor-expressing tissues become unresponsive to insulin as a protective mechanism to avoid excessive glucose uptake (162). As a result, pancreatic β -cells uncontrollably release insulin to overcome the desensitization, which eventually impairs β -cell function (35, 162). Insulin resistance is accompanied by adipose tissue inflammation, impaired adipokine release, downregulation of GLUT4, and ectopic fat accumulation resulting from increased FFA (52, 161, 163). Rodent studies have shown that high-fat diets contribute to insulin resistance by impairing peripheral glucose uptake and increasing pro-inflammatory cytokines gene expression in adipose tissue (52, 164). In humans, oral administration of palm oil decreased adipose tissue insulin sensitivity and increased hepatic fat deposition (165). Other clinical studies demonstrate that increased visceral fat deposition is significantly associated with insulin resistance, dyslipidemia, and hyperinsulinemia in healthy individuals (166).

Insulin is a pancreatic hormone that is crucial for glucose metabolism and energy balance. Insulin receptors are expressed in the hypothalamus, including the ARH and VMH (167, 168). In response to changes in circulating glucose, insulin is released from pancreatic β -cells. It enters the brain to directly act on ARH, blocking orexigenic effects of NPY/AgRP neurons and enhancing anorexigenic effects of POMC neurons to increase satiety and decrease blood glucose (167). Multiple studies have shown that OVX in rodents increases circulating insulin and glucose, decreases insulin sensitivity, and impairs β -cell function and insulin secretion. These phenotypes are exacerbated by HFD

and accompanied by increased body weight and total body fat. Estrogen replacement in these animals lowers plasma insulin and glucose levels, increases glucokinase and GLUT2 expression, improves glucose tolerance and insulin sensitivity (169–172). Central estrogen signaling is suggested to be involved in insulin sensitivity, as intact obese female mice are protected from DIO-induced insulin resistance in NPY/AgRP neurons. Conversely, ovarian hormone depletion abolishes NPY/AgRP insulin sensitivity (117), and deletion of ER α in steroidogenic factor 1 (SF1) - positive neurons in the VMH impairs glucose tolerance in female mice (17, 20).

Insulin resistance may result not only from β -cell dysfunction or glucose disposal impairment but can also be caused by a reduction in insulin-mediated thermogenesis (173). BAT is highly sensitive to insulin given the high expression of insulin receptors in this tissue (35, 174). Insulin-mediated BAT activity improves lipid/glucose clearance, contributing to decreased glucolipotoxicity, ultimately improving β-cell function and insulin sensitivity (174). In humans, insulin stimulation increases BAT glucose uptake, energy expenditure, and expression of UCP1 and glucose transport genes in BAT (175). Similarly, cold exposure for eight hours increases energy expenditure, glucose disposal, and insulin sensitivity in individuals with high BAT activity. Administration of insulin following cold exposure further enhances glucose disposal and reduces glucose production in these patients (176). A study by Benedict et al. directly targeted central insulin signaling by administrating intranasal insulin to healthy men following an overnight fast. Individuals treated with insulin showed a dramatic increase in energy expenditure (177).

In mice, it has been shown that insulin acts through SNS to upregulate UCP1 in BAT and thermogenic function (178). In line with this, adrenergic stimulation in mice increases WAT lipolysis, BAT insulin signaling, and energy expenditure. These phenotypes are reversed by insulin receptor antagonism (179). Deletion of insulin receptors specifically in BAT results in decreases in glucose uptake, mass, and thermogenic capacity of BAT (179, 180). A recent report targeting diosmetin, a natural flavonoid commonly found in citrus fruits, showed that diosmetin treatment reverses DIO in mice by decreasing fat mass, reducing plasma glucose and insulin, and increasing insulin sensitivity and glucose disposal. In addition, diosmetintreated mice show increased O2 consumption, enhanced heat production, and increased body temperature and WAT browning. Interestingly, mRNA levels of ERa and ERB were dramatically increased in WAT and BAT, suggesting that diosmetin-induced thermogenesis was mediated by estrogen signaling. Administration of estrogen antagonist fulvestrant abolished the beneficial metabolic effects of diosmetin in DIO mice. These findings link insulin sensitivity and thermogenesis with estrogenic signals in adipose tissue.

Mammalian Target of Rapamycin (mTOR)

Emerging evidence highlights the relevance of hypothalamic mTOR signaling in body weight balance and energy intake. mTOR is a serine-threonine kinase that senses fluctuations in nutrient availability (181, 182). In its phosphorylated (active,

pmTOR) form, mTOR phosphorylates its major substrate, the serine/threonine ribosomal protein S6 kinase B1 (S6K1), modulating multiple cellular processes, including cell growth, cell survival, proliferation, and glucose metabolism (181, 182). Hypothalamic mTOR expression and activity have been reported in ARH, VMH, PVN, and the mediobasal hypothalamic area (MBH) (181–183).

In recent years, several reports have revealed the association between estrogen's anorexic effects and mTOR signaling. An elegant study from González-García et al. provided evidence of mTOR-dependent weight loss and decreased food intake induced by E2. In addition to the typical obesogenic phenotypes associated with estradiol depletion, OVX rats showed reduced phosphorylation of mTOR and S6K1 in the ARH compared to sham rats. Conversely, E2 supplementation led to decreased body weight and food intake, associated with increased phosphorylation of mTOR and pS6K1 in the ARH. Consistently, ICV delivery of PPT reversed the OVX-induced body weight gain and strongly enhanced phosphorylation of mTOR and pS6K1 in the ARH (174). These findings support a model that E2/ER α activation increases mTOR signaling in the ARH to regulate food intake and body weight (182).

A recent report identified the role of the kappa-opioid receptor (k-OR) in the obesogenic effects of E2 withdrawal (175). They found that a k-OR/mTOR/p70S6K axis compensates for estrogens depletion in OVX mice by promoting body weight loss, increased energy expenditure, and WAT browning. Conversely, central antagonism of orexigenic k-OR increased phosphorylation of mTOR in the MBH of WT OVX mice, resulting in beneficial metabolic effects. Selective expression of constitutive activated S6K in the MBH decreased body weight and WAT mass and stimulated WAT browning, as indicated by increased UCP1 expression in gWAT. These findings suggest a model that inhibition of k-OR activates mTOR signaling in MBH to prevent body weight gain and adiposity induced by OVX (183).

VENTROMEDIAL HYPOTHALAMUS (VMH)

The VMH plays a critical role in modulating energy expenditure, appetite regulation, sexual aggressive behaviors, lipid mobilization, and thermoregulation. The VMH was the first hypothalamic region identified to stimulate BAT thermogenesis and energy expenditure (19, 168). It is well established that VMH lesions in animals induce increases in food intake, adiposity, and body weight gain (184). Electrical stimulation of the VMH increases BAT temperature and thermogenic activity. This was abolished by the β -adrenergic blockade, confirming that SNS is an essential intermediary between the VMH-BAT crosstalk (185). SF1 is the most abundant neural population in the VMH. SF1 neurons express LepR, vesicular glutamate transporter 2 (vGLUT2), and ERa (19, 168). SF1 neurons project to the medial amygdala (MeA), DMH, periaqueductal gray (PAG), and ARH, all of which are linked to feeding regulation (186). Interestingly, recent reports have reveal that glutamatergic VMH neurons project to POMC neurons and

that this VMH \rightarrow POMC neural circuit senses changes in nutrient availability and helps maintain energy homeostasis (187). In addition to regulation of energy expenditure, the VMH neurons have been recently shown to be glucose-sensitive and play a key role in glucose homeostasis (188).

Estrogen Signals in VMH Regulate Energy Balance

AMP-activated protein kinase (AMPK) is a critical element in the regulatory effects of VMH on BAT thermogenesis and glucose homeostasis. AMPK is activated by phosphorylation of Thr172 in the alpha subunit. Activation of AMPK signaling in the hypothalamus is associated with increased food intake and decreased energy expenditure (189). In the VMH, multiple factors have been found to repress AMPK activation and subsequently promote anti-obesogenic signals. These factors include bone morphogenetic protein 8B (BMP8B) (190), glucagon-like peptide 1 (GLP-1) (191), thyroid hormones (TH) (192), leptin (156), and estradiol (59). These AMPK-regulating factors have been extensively reviewed in a recent publication (193). The raphe pallidus (RPa) and inferior olive (IO) are two brainstem nuclei that are anatomically connected to the VMH and participate in the activation of SNS to activate BAT thermogenesis (Figure 2) (189).

Several studies have evidenced the requirement of VMH estrogen-ER α signaling for the modulation of BAT thermogenesis and fat distribution (16, 17, 19, 20, 57). Martínez de Morentin and colleagues unveiled the role of AMPK in the stimulatory effects of VMH estrogen signaling on BAT thermogenesis and energy expenditure (59). In their study, they showed that OVX rats treated with E2 directly into the VMH *via* ICV infusion show dramatic loss of body weight gain and food intake, increases mRNA levels and protein content of UCP1 in BAT and increases BAT temperature. All these effects were associated with decreased AMPK phosphorylation (active)

in the VMH, suggesting a protentional role of VMH AMPK signaling. In supporting this, specific injection of E2 into the VMH decreased VMH AMPK activity, increased c-fos numbers in RPa and IO, and upregulated expression of thermogenic markers in the BAT. Conversely, VMH-specific co-infusion of AMPK completely abolished these anti-obesity effects (59). These findings reveal a VMH-AMPK-SNS-BAT axis that is modulated by estrogenic signals to maintain energy balance (**Figure 2**). Other studies have shown that silencing $ER\alpha$ in the VMH of rodents induces body weight gain, increased visceral adiposity, decreased physical activity, downregulation of thermogenic genes in the BAT, and increased circulating glucose (16, 17). In a study performed by Xu and colleagues, female mice with ERa deleted specifically in the SF1 neurons demonstrate increased body weight, increased visceral and gonadal adiposity, impaired glucose tolerance, decreased heat production, as well as decreased SNS activity and BAT UCP1 transcripts. The group concluded that estrogens activate ERa expressed by the SF1 neurons of the VMH to increase sympathetic outflow and modulate fat distribution, which ultimately results in increased BAT thermogenesis and decreased lipid storage (20).

Besides the VMH-AMPK-SNS-BAT axis, E2 also modulates ceramides metabolism and endoplasmic reticulum (ER) stress in the brain to regulate thermogenesis and energy balance (187). Specifically, OVX increased levels of ER stress markers in MBH and VMH, while both E2 replenishment and ICV ceramide inhibition abolished these phenotypes in rats. This group further targeted serine palmitoyltransferase long chain base subunit 1 (SPTLC1), an enzyme that catalyzes ceramide synthesis, to explore ceramide action in the VMH. They found that VMH-specific injection of SPTLC1 shRNA decreased body weight, enhanced BAT and core temperature, and upregulated BAT UCP1 protein levels in OVX rats. These phenotypes were accompanied by the downregulation of ER stress markers in





the VMH. Consistently, VMH-specific overexpression of ER glucose chaperone GPR78, a protein that decreases ER stress by facilitating degradation pathways, also ameliorated ER stress and improved body weight and thermoregulation in OVX rats. These findings demonstrate that estrogens in the VMH contribute to energy balance by maintaining optimal levels of ceramides and decreasing ER stress (187).

ER α is highly expressed in the ventrolateral subregion of the VMH (vlVMH), where it has been shown to regulate fertility, social behaviors, and locomotor activity in a sex-specific manner (16, 18-20, 99). We and others have shown that DREADD activation of $ER\alpha^{vIVMH}$ neurons augmented BAT and core temperature as well as physical activity in female mice (18, 195). Moreover, this $ER\alpha^{vIVMH}$ chemogenetic activation ameliorates body weight gain caused by HFD, indicating beneficial effects of $ER\alpha^{vIVMH}$ activation in metabolic health (195). Correa et al. targeted Nkx2-1⁺ neurons in the vlVMH, which are associated with physical activity in mice. ERa and NKX2-1 are co-expressed in the vlVMH. Neural activation of Nkx2-1⁺ vlVMH using chemogenetic approaches increased ambulatory and vertical movement as well as heat production in Nkx2-1Cre female mice. Deletion of NKX2-1 in the vlVMH of female mice led to increased body weight, increased WAT depots, decreased locomotion, and lower heat production. More importantly, these phenotypes were associated with decreased Esr1 transcripts and ERa protein content (19). A single-cell transcriptomic study targeting SF1⁺ neurons in the VMH identified differentially expressed gene transcripts that included reprimo (Rprm), tachykinin Precursor 1 (Tac1), prodynorphin (Pdyn), and somatostatin (Sst). In females, Tac1 and Rprm were highly expressed in the vlVMH when compared to males, and these genes were found to coexpress with ER α in the vlVMH (ER α ^{vlVMH}), positioning these genes as femalespecific markers in this neuronal population (16, 18).

A recent report has identified another sexually dimorphic subset of ER α^{vIVMH} neurons expressing MC4R, a receptor of anorexigenic ligand melanocyte-stimulating hormones (MSH) (16, 196). Loss-of-function mutations in *MC4R* are the most frequent genetic cause of obesity in humans (196). In this study, ER α /MC4R^{vIVMH} neurons were identified as a subset population of ER α^{vIVMH} neurons. They showed colocalization of MC4R and ER α in the vIVMH during proestrus in mice supplemented with E2. Although OVX mice display reduced locomotion and hypometabolism, chemogenetic manipulation of ER α /MC4R^{vIVMH} neurons significantly stimulated physical activity in these estrogen-depleted mice. The restoration of physical activity by ER α /MC4R^{vIVMH} stimulation was accompanied by reduced body weight and white adipocyte size. These findings suggest that ER α /MC4R^{vIVMH} plays an essential role in sexdimorphic physical activity in mice (16).

To gain mechanistic insight into the physiological functions of $\text{ER}\alpha^{\text{vIVMH}}$ neural subsets, the anatomical distribution of the downstream neural circuitries and their functionality need to be studied. Excitingly, MC4R^{vIVMH} neurons were shown to send projections to distinct brain regions including dorsal CA1, periaqueductal grey (PAG), and hindbrain pontine region (16). Similarly, we have recently identified 5-HT⁺ neurons in the DRN as a downstream neuronal population receiving monosynaptic innervation from $ER\alpha^{v\rm IVMH}.$ Using electrophysiology and optogenetic tools, we showed the glutamatergic nature of these $ER\alpha^{vlVMH} \rightarrow DRN$ monosynaptic inputs. Interestingly, we demonstrate that $ER\alpha^{vIVMH} \rightarrow DRN$ neural circuit has thermo-sensing and nutrient-sensing properties, which are abolished in OVX mice, suggesting an important role of ovarian hormones in $ER\alpha^{vIVMH} \rightarrow DRN$ responsiveness to ambient temperature and nutritional state. Optogenetic activation of the $ER\alpha^{vIVMH} \rightarrow DRN$ circuit stimulates BAT thermogenesis and physical activity, while its inhibition has the opposite effect, demonstrating an in vivo physiological role $ER\alpha^{vlVMH} \rightarrow DRN$ neural circuit. We show that activation of $5\text{-}\text{HT}^{\text{DRN}}$ neurons is required for these stimulatory effects induced by $ER\alpha^{vIVMH}$ activity. Importantly, deletion of $ER\alpha$ from vlVMH neurons projecting to DRN reduced physical activity and BAT thermogenesis in female mice, demonstrating that $ER\alpha^{vlVMH} \rightarrow DRN$ neural circuit is relevant for regulation of physical activity and energy expenditure in female mice (195). Altogether, these findings demonstrate the physiological relevance of ER α^{vlVMH} neurons in the maintenance of physical activity and thermogenic balance.

As discussed previously, the VMH communicates with both BAT and WAT through SNS to modulate thermogenesis and lipolysis. The VMH regulation of BAT thermogenesis has been discussed. However, VMH modulation of WAT lipolysis is less clear. Like BAT's thermogenesis mechanism, SNS induces the release of NE, which activates \$\beta3-adrenoceptors in white adipocytes. The subsequent downstream signaling cascade stimulates PKA, which phosphorylates perilipin A and HSL, two key factors that initiate lipolysis (197). One of the possible contributors of impaired SNS activation during obesity is insulin. It is suggested that insulin resistance may increase sympathetic firing and therefore increase lipolytic rate in WAT, which results in elevated free fatty acids in circulation (197). The VMH-AMPK-SNS-BAT axis has been discussed before. However, despite the evidence indicating the role of estrogen VMH signaling in adiposity and fat distribution, the neuroanatomical projections that mediate the estrogen VMH-SNS-WAT communication, as well as the underlying molecular mechanisms, remain to be elucidated.

Estrogen Signals in the VMH Regulate Glucose Homeostasis

In the brain, two different types of neurons can sense changes in circulating glucose levels: glucose-excited (GE) neurons and glucose-inhibited (GI) neurons (198). The VMH is highly enriched with both GE and GI neurons. Within the vlVMH, GE neurons are more abundant than GI. Additionally, the majority of ER α^{vlVMH} neurons are glucose-sensing neurons (188). The enzyme glucokinase is expressed in GE and GI neurons. It is required for the glucose-sensing property, as ablation of this enzyme in the VMH leads to increased fat mass, decreased glucagon secretion, and reduced sympathetic activity. These phenotypes were predominant in female mice, highlighting the already mentioned sexual-biased functions in

the vlVMH (199). In GE neurons, when glucose levels are elevated, the ATP/ADP ratio is increased. This, in turn, causes the closure of K_{ATP} channels, resulting in the depolarization and excitation of these neurons. In GI neurons, the increased glucose also increases the ATP/ADP ratio, but this results in the inhibition of AMPK. As a result, the K⁺ and Cl⁻ channels are opened, allowing the hyperpolarization and inhibition of GI neurons (198, 200). These mechanisms ultimately decrease the levels of glucose in circulation. Under low glucose conditions, K_{ATP} channels are opened, and GE neurons are hyperpolarized. GI neurons are depolarized by activation of AMPK, which induces closure of K⁺ and Cl⁻ channels. Inhibition of GE neurons and activation of GI neurons leads to an increase in circulating glucagon, which ultimately results in increased glucose levels (188, 200).

Since estrogen is a key factor in the VMH-AMPK-SNS-BAT axis, it is not surprising that under hypoglycemic conditions, estradiol impedes the activation of GI neurons in female mice by inhibiting AMPK signaling, thus preventing the decrease in glucose concentrations in a sex-specific manner (201). Yu et al. demonstrated the requirement of membrane-bound ERa for fasting- and glucopenia-induced refeeding. Mutations in membrane-bound ER α decreased the firing rate of ER α^{vIVMH} even after administration of PPT, and importantly, impaired the ability of ERavivMH neurons to respond to hypoglycemia (101). Our previous study reveals the mechanisms underlying the glucosesensing properties in $ER\alpha^{vIVMH}$ neurons (188). During hypoglycemia, the activation of GI neurons in $ER\alpha^{vlVMH}$ neurons is mediated by the calcium-activated chloride channel protein channel encoded by anoctamin 4 (Ano4), which contributes to their depolarization. Conversely, GE neurons are hyperpolarized by the activation of Abcc8, which encodes KATP subunit Sur1. Additionally, we reported that GI-ER α^{VVMH} neurons project to the medioposterior part of the ARH (mpARH) and GE-ER α^{vIVMH} neurons project to the DRN. Activation of $ER\alpha^{vIVMH} \rightarrow mpARH$ and inhibition of ER α^{vIVMH} \rightarrow DRN complementarily contribute to the prevention of hypoglycemia (188). Altogether, the discussed evidence supports estrogen as a critical element in the VMH for the modulation of glucose and energy homeostasis.

CONCLUSION

Obesity is considered a global epidemic affecting 650 million people worldwide (128, 202). This disease is often exacerbated by other metabolic disorders that have a significant impact on patient quality of life. In humans and animal models, sexual dimorphism has been described not only in adipose tissue

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deposition and distribution but also in BAT thermogenic capacity. This is partially mediated by the female sex hormone estrogen, which confers protection against metabolic diseases in premenopausal women. Hypothalamic estrogen signaling has been shown to have positive effects on energy homeostasis by decreasing food consumption, diminishing excessive adiposity, improving insulin sensitivity, and promoting BAT thermogenesis and WAT beiging. Although it is well established that SNS mediates estrogen brain-to-adipose tissue signals (57, 59), less is known about the downstream neural circuits mediating the regulatory effects of brain estrogen on adipose metabolism. As discussed earlier, multiple hypothalamic projections have been described, including the VMH-POMC neural circuit that detects fluctuations in nutritional status and the $\text{ER}\alpha^{vlVMH} \rightarrow$ DRN that modulates physical activity and energy expenditure (187, 195). It would be interesting to test the role of estrogen in VMH-POMC projection and explore in more depth the estrogen-modulated hypothalamic circuitry that regulates adipose tissue function and remodeling. Moreover, while $ER\alpha^+$ neurons in the VMH have been shown to be essential for heat production and energy expenditure (20), a recent report also revealed that $ER\alpha^+$ neurons in the MPOA are thermosensitive and decrease energy expenditure when activated (136). These findings support a model where $ER\alpha^+$ neurons in distinct hypothalamic regions act synergistically to respond to changes in nutrient availability and environmental temperature. Collectively, further studies are needed to unravel the estrogengoverned neural networks modulating adipose tissue function, which is critical for a better understanding of the pathophysiology of obesity.

AUTHOR CONTRIBUTIONS

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