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## Mechanisms for Sex Differences in Energy Homeostasis

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### Abstract

Sex differences exist in the regulation of energy homeostasis. Better understanding of the underlying mechanisms for sexual dimorphism in energy balance may facilitate development of gender-specific therapies for human diseases, e.g. obesity. Multiple organs, including the brain, liver, fat and muscle, play important roles in the regulations of feeding behavior, energy expenditure and physical activity, which therefore contribute to the maintenance of energy balance. It has been increasingly appreciated that this multi-organ system is under different regulations in male vs. female animals. Much of effort has been focused on roles of sex hormones (including androgens, estrogens and progesterone) and sex chromosomes in this sex-specific regulation of energy balance. Emerging evidence also indicates that other factors (not sex hormones/receptors and not encoded by the sex chromosomes) exist to regulate energy homeostasis differentially in males vs. females. In this review, we summarize factors and signals that have been shown to regulate energy homeostasis in a sexually dimorphic fashion and propose a framework where these factors and signals may be integrated to mediate sex differences in energy homeostasis.

### Introduction

Obesity, resulting from imbalance of energy homeostasis, is now recognized as a serious global health problem, due to its high prevalence and strong association with hypertension, coronary heart disease, stroke, and other metabolic disorders. A clear sexual dimorphism exists in the regulation of feeding behavior and energy homeostasis in rodents (Shi, et al. 2009; Sugiyama and Agellon 2012). For example, total daily energy intake in male rats is higher than that in females, even when corrected by their larger lean body mass and metabolic rate (Woodward and Emery 1989). In addition, high fat-diet (HFD) feeding leads to larger body weight gain in male rats/mice than in female counterparts (Benz, et al. 2012; Dorfman, et al. 2017; Grove, et al. 2010; Morselli, et al. 2016; Stubbins, et al. 2012; Yang, et al. 2014). However, the mechanisms for this sexual dimorphism remain elusive, and better

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understanding of this fundamental sex difference in energy homeostasis will no doubt benefit our current combat against obesity pandemics.

Males and females are obviously different in the circulating levels of sex hormones and the sex chromosomes their cells carry. Thus, these two categories of factors have been generally thought to be major contributors to the sexual dimorphism in obesity. In this review, we will discuss the functions of sex hormones and sex chromosomes in the context of body weight control. In addition, we will also discuss additional factors, which do not belong to these two categories, as additional contributors to the sex dimorphism of energy homeostasis.

## 1. Sex hormones

Androgens are described as male sex hormones and estrogens are described as female sex hormones (Chan and O'Malley 1976). The most important biologically relevant forms of estrogens and androgens are  $17\beta$ -estradiol (E2) and testosterone (T), respectively. Both males and females have these hormones to varying degrees. Androgens are produced in male testis and female ovary. Androgens, mainly **testosterone** and **androstenedione**, can be converted to estrogens by aromatase (Jarvie and Hentges 2012). Thus, even in males, despite the low circulating levels of estrogens, estrogens can be produced through local aromatization using circulating testosterone as a substrate. Tissues that express aromatase include not only the gonads, but also the breast, brain, muscle, bone, and adipose tissue (Nelson and Bulun 2001; Simpson 2003; Simpson, et al. 2005). Another major female sex hormone is progesterone, which can also be synthesized by the adrenal glands (Johnston, et al. 2015; Wittmann, et al. 2013) and **nervous tissue**, especially in the brain (Schumacher, et al. 2004). Classically, androgens, estrogens and progesterone can bind to their nuclear receptors androgen receptors (AR), estrogen receptors (ER) and progesterone receptors (PR), respectively, and these receptors function as transcription factors to regulate gene expression (Yang and Shah 2014). Accumulating evidence also indicates that these sex hormones can bind to membrane receptors and exert rapid signal transduction effects in target cells (Mamounis, et al. 2014). As outlined below, these sex hormones and their receptors contribute to the regulation of energy balance with complex mechanisms.

### 1.1. Estrogens

**1.1.1 Ligand:** It is well documented that estrogens play an essential role in preventing body weight gain in females. For example, the withdrawal of endogenous estrogens by ovariectomy (OVX) in female animals leads to body weight gain and hyperadiposity, and these obese phenotypes can be prevented by E2 treatment but not by progesterone (Drewett 1973; Geary, et al. 2001; Roesch 2006; Rogers, et al. 2009; Schwartz and Wade 1981; Wallen, et al. 2001). Since E2 can be produced via conversion from testosterone by the aromatase, aromatase knockout mice represent a good model to examine functions of endogenous E2 not only in female but also in male animals. Female aromatase knockout mice show increased body weight from 3 months of age, while male mutant mice show late onset obesity one year later (Jones, et al. 2000). Both male and female aromatase knockout mice show increased gonadal and infrarenal fat pad compared to control littermates (Jones et al. 2000). This increased adiposity is associated with reduced spontaneous physical activity

levels, reduced glucose oxidation, and a decrease in lean body mass (Jones et al. 2000). These findings indicate that endogenous E2 prevent obesity in both sexes.

**1.1.2. ER $\alpha$ :** Estrogen receptors include estrogen receptor- $\alpha$  (ER $\alpha$ ), estrogen receptor- $\beta$  (ER $\beta$ ) and G protein-coupled receptor 30 (GPR30). In particular, ER $\alpha$  is perhaps most studied in the context of energy balance. Humans or mice with mutations in the ER $\alpha$  (Esr1) gene are obese (Heine, et al. 2000; Okura, et al. 2003). Further, deletion of ER $\alpha$  in mice blocks the anorexigenic effects of E2 treatment (Geary et al. 2001). Early studies showed that microinjections of E2 into various brain regions change animal's feeding behavior (Butera and Beikirch 1989; Palmer and Gray 1986), suggesting that ER $\alpha$  expressed in the brain is important for the regulation of food intake. This notion was further supported by observations from various genetic mouse models. For example, we crossed mice carrying loxP-flanked ER $\alpha$  alleles (Esr1<sup>fl/fl</sup>) (Feng, et al. 2007) to the Nestin-Cre transgenic mice (Bruning, et al. 2000) to produce mice lacking ER $\alpha$  only in the brain (Xu, et al. 2011b). These female mutant mice develop obesity, characteristic of increased body weight and body fat. Obesity in these mice is associated with hyperphagia, decreased energy expenditure and decreased physical activity, which may all contribute to the development of obesity (Xu et al. 2011b). Notably, female mice lacking ER $\alpha$  in the brain display significantly elevated E2 in the circulation (Xu et al. 2011b), presumably due to the impaired negative feedback regulation by estrogens. Given that these mice develop robust obese phenotypes despite the higher E2 in the circulation, these observations argue that compared to ER $\alpha$  expressed in peripheral tissues, brain ER $\alpha$  plays predominant roles in the regulation of energy balance. Notably, male mice lacking ER $\alpha$  in the brain also develop modest obesity, indicating that brain ER $\alpha$  is required to prevent body weight gain in males (Xu et al. 2011b).

ER $\alpha$  is abundantly expressed in multiple brain regions that are implicated in the regulation of feeding behavior. These include the arcuate nucleus (ARH), the ventromedial hypothalamic nucleus (VMH), the nucleus of the solitary tract (NTS), the medial amygdala (MeA), the dorsal Raphe nuclei (DRN), and the medial preoptic area (MPOA) (Merchenthaler, et al. 2004). Interestingly, while many of these brain regions, e.g. ARH, VMH, and MPOA, in female brains express more abundant ER $\alpha$  than in male brains (Cao and Patisaul 2011; Kyi-Tha-Thu, et al. 2015), a few other regions in male brains, including the MeA, show higher ER $\alpha$  expression than in females (Xu, et al. 2015). In particular, about 20–30% pro-opiomelanocortin (POMC) neurons in the ARH co-express ER $\alpha$  (de Souza, et al. 2011; Miller, et al. 1995; Xu, et al. 2011a). E2 can increase excitatory synaptic inputs onto ARH POMC neurons and activate these neurons (Gao, et al. 2007; Malyala, et al. 2008; Saito, et al. 2015). Female mice lacking ER $\alpha$  only in POMC neurons develop hyperphagia and modest body weight gain (Xu et al. 2011b). In addition, E2-induced anorexigenic effects are blunted in these mutant female mice lacking ER $\alpha$  only in POMC neurons (Zhu, et al. 2015). Notably, male mice lacking ER $\alpha$  in POMC neurons do not have any feeding phenotypes (Xu et al. 2011b). Another hypothalamic ER $\alpha$  population with sexually dimorphic functions is that in the VMH. Martinez de Morentin et al. reported that injections of E2 into the VMH promote BAT-mediated thermogenesis in female rats (Martinez de Morentin, et al. 2014). Correa et al. developed a mouse model with loss of 26% of ER $\alpha$ -positive neurons in the VMH; this loss of ER $\alpha$ -positive neurons cause profound obesity only

in females but not in males (Correa, et al. 2015). Consistently, genetic deletion of ER $\alpha$  specifically in the VMH leads to female obesity, but does not affect male energy balance in rodents (Musatov, et al. 2007; Xu et al. 2011b). Collectively, these observations indicate that ER $\alpha$  in POMC neurons and VMH neurons contributes to the sex differences in body weight balance.

Other hypothalamic ER $\alpha$  populations may also be involved in body weight control. For example, Santollo et al. reported that microinjections of E2 into the MPOA decreases food intake in female rats (Santollo, et al. 2011). Further, earlier studies showed that E2 implanted in the paraventricular nucleus of the hypothalamus (PVH) decreases food intake and body weight in OVX female rats (Butera and Beikirch 1989). Additionally, the anorexigenic effects of subcutaneous E2 were blunted in female rats with PVH lesions (Butera, et al. 1992). However, subsequent studies failed to reproduce these phenotypes in female rats with PVH implants (Hrupka, et al. 2002). In addition, it needs to be pointed out that the PVH expresses low levels of ER $\alpha$  but high levels of ER $\beta$  (Merchenthaler et al. 2004). Thus, the functions of ER $\alpha$  in the MPOA and the PVH warrant further validation with genetic models.

ER $\alpha$  is also present in the hindbrain, including the NTS (Merchenthaler et al. 2004; Osterlund, et al. 1998; Schlenker and Hansen 2006). Treatment of E2 in wild type female mice suppresses food intake and potentiates CCK-induced satiation, which are accompanied by increased activity in NTS neurons (Asarian and Geary 2007; Geary et al. 2001). Interestingly, these responses are all abolished in female mice lacking ER $\alpha$  (Asarian and Geary 2007; Geary et al. 2001). ER $\alpha$  is abundantly expressed in the DRN (Merchenthaler et al. 2004), and the majority of these ER $\alpha$ -positive neurons in the DRN are 5-HT neurons (Cao, et al. 2014). E2 increases neural activities of DRN 5-HT neurons via an ER $\alpha$ -dependent mechanism (Cao et al. 2014; Dalmasso, et al. 2011; Robichaud and Debonnel 2005). Santollo et al. reported that microinjections of E2 into the DRN decreases food intake in female rats (Santollo et al. 2011). Female mice lacking ER $\alpha$  only in 5-HT neurons are resistant to E2's effects to suppress binge-like eating (Cao et al. 2014), a type of eating behavior that is not driven by hunger but rather than by rewards or hedonic cues. Collectively, ER $\alpha$  in the NTS and DRN mediates the anorexigenic effects of E2 in females, while functions of these ER $\alpha$  populations in male animals remain unknown.

However, it is clear that actions of ER $\alpha$  also prevent obesity in males. For example, ER $\alpha$  gene deficiency results in obesity in male mice (Callewaert, et al. 2009; Heine et al. 2000) and in men (Grumbach and Auchus 1999; Smith, et al. 1994). In addition, administration of E2 or its analogs reduces body weight in male mice (Finan, et al. 2012; Gao et al. 2007). Both male and female aromatase knockout mice develop obesity (Jones et al. 2000). Notably, abundant aromatase is expressed in a few brain regions, including the MeA (Wu, et al. 2009), which makes it possible that ER $\alpha$  in these male brain regions could be exposed to high levels of E2 despite the lack of circulating estrogens. Indeed, loss of ER $\alpha$  in the MeA not only causes obesity in female mice, but also in male mice (Xu et al. 2015). Thus, we speculate that ER $\alpha$  populations in brain regions with aromatase activity may play sexually monomorphic roles in the regulation of energy balance, while ER $\alpha$  in brain regions devoid of aromatase activity contribute to the sex differences in body weight balance.

The role of ER $\alpha$  in the peripheral tissues has been extensively reviewed before (Mauvais-Jarvis, et al. 2013). The deletion of ER $\alpha$  from the liver does not affect body weight fed with chow or HFD in either male or female mice (Matic, et al. 2013). Loss of ER $\alpha$  in the muscle causes obesity in female mice, associated with decreased energy expenditure and physical activity without changes in food intake (Ribas, et al. 2010), while effects of such deletion in male mice were not reported. Deletion of ER $\alpha$  in adipocytes causes obesity in female mice but not in male mice (Davis, et al. 2014). Together, the current literature indicate that ER $\alpha$  in some regions of the brain and in fat contributes to the maintenance of energy homeostasis in a sexually dimorphic fashion, although its effects in males still warrant further investigations. However, ER $\alpha$  in brain regions with enriched aromatase activity (e.g. the MeA) regulates energy balance in both sexes.

**1.1.3. Other estrogen receptors:** Compared to ER $\alpha$ , ER $\beta$ , another classic ER has received less attention at least in the context of body weight balance. An earlier study by Ohlsson et al reported that chow-fed mice (both male and female) with global deficiency in ER $\beta$  show normal body weight and fat mass compared to wild type mice (Ohlsson, et al. 2000). Consistent with this, both Santollo et al (Santollo, et al. 2007) and Roesch (Roesch 2006) found that an ER $\beta$  agonist, diarylpropionitrile (DPN), has no effect on food intake and body weight in chow-fed OVX female rats, while PPT (the ER $\alpha$  agonist) at similar doses can significantly reduce food intake and body weight. While these earlier studies suggest a minor role of ER $\beta$  in body weight control in chow-fed animals, Foryst-Ludwig et al demonstrated that female ER $\beta$  knockout mice, when fed on a HFD, developed obesity compared to HFD-fed wild type mice, which is associated with normal food intake, but increased energy expenditure and decreased fat oxidation (Foryst-Ludwig, et al. 2008). However, it has not been reported whether male mice lacking ER $\beta$  also show similar obese phenotypes. Certainly, the ER $\beta$ -mediated control of energy homeostasis in difference sexes warrants further investigations.

GPR30 (also known as GPER) is a G protein-coupled estrogen receptor, bound to the cell membrane (Thomas, et al. 2005). Body weight phenotypes among several independent GPR30 knockout mouse lines are controversial. For example, both Haas et al (Haas, et al. 2009) and Sharma et al (Sharma, et al. 2013) observed obese phenotypes in male and female GPR30 knockout mice, which were generated by Wang et al (Wang, et al. 2008); however, Liu et al reported no difference in body weight in the same GPR30 knockout strain (Liu, et al. 2009). Otto et al constructed an independent GPR30 knockout line, and found no obese phenotypes in female mutants (Otto, et al. 2009). Interestingly, another GPR30 knockout line generated by Martensson et al show reduced body weight only in females, but not in males (Martensson, et al. 2009). More recently, Davis et al re-characterized Wang's GPR30 knockout mice and reported that both male and female mutants are significantly heavier than wild type littermates, which appears to depend on reduced energy expenditure independent of physical activity, but not on food intake (Davis et al. 2014). The discrepancy from these studies may be attributed to different strategies to construct the GPR30 knockout alleles, different genetic background that mice were maintained on, and/or different facility environment, etc. Further investigations are warranted to confirm THE roles of GPR30 in estrogenic regulation on body weight homeostasis in both sexes.

Collectively, high levels of circulating estrogens prevent obesity in female animals. Many of these estrogenic actions are mediated by multiple ER $\alpha$  populations, which therefore at least partly account for sex differences in body weight. However, ER $\alpha$  in certain brain regions with enriched aromatase activity, e.g. the MeA, also contributes to the maintenance of male energy balance. ER $\beta$  and GPR30 also regulate body weight balance, but their sex-specific functions remain to be further investigated.

## 1.2. Progesterone

Unlike estrogens, the administration of progesterone alone does not affect food intake, body weight, or adiposity in OVX female mice (Toth, et al. 2001). However, the anorectic effect of estrogens can be attenuated by concurrent administration of progesterone (Schwartz and Wade 1981; Wade 1975). In gonadal intact rodents, exogenous progesterone causes increase in body weight and inguinal white adipose tissue mass in female animals but not in male animals, by regulating the endocrine properties of adipocytes (Komatsu, et al. 2011; Kudo, et al. 2014; Sarton, et al. 2001). However, female mice with the global deletion of progesterone receptor do not show any difference in body weight (Rickard, et al. 2008). Notably, Klump et al reported that in cycling women, the tendencies of binge eating are negatively associated with circulating estrogen level but positively associated with progesterone level (Klump, et al. 2013; Klump, et al. 2014), suggesting that progesterone may promote binge eating. However, administration of progesterone in OVX female rats does not affect binge-like eating behavior (Yu, et al. 2011), although it remains debatable whether the binge-like eating behaviors in rodents really simulates human binge eating. Collectively, some evidence exists to support an orexigenic effect of progesterone, but further investigations are needed to confirm these effects and to explore the underlying mechanisms.

## 1.3. Androgens

**1.3.1. Ligand—**Androgens are also implicated to regulate male and female energy balance. Most women diagnosed with polycystic ovary syndrome (PCOS) are hyperandrogenemia, and higher level of serum androgen is correlated with higher BMI (Alexiou, et al. 2017; Dumesic, et al. 2016; Yuan, et al. 2016). In female animals, chronic treatment of dihydrotestosterone (DHT), a non-aromatizable androgen, promotes visceral adiposity with reduced energy expenditure but normal food intake (Nohara, et al. 2014). These effects are associated with reduced POMC neuronal innervations in the hypothalamus (Nohara et al. 2014). Effects of testosterone in males appear to be more complex. Serum testosterone levels are inversely correlated with obesity in men (Jorgensen, et al. 1996; Katznelson, et al. 1998). Man with lower androgen level showed increased fat mass which can be reversed by the testosterone administration (Rolf, et al. 2002). However, in male castrated adult mice, DHT treatment increases adiposity (Moverare-Skrtic, et al. 2006). Consistently, treatment of these male mice with testosterone in the presence of an aromatase inhibitor, but not testosterone alone, induces retroperitoneal fat accumulation (Moverare-Skrtic et al. 2006). Together, these observations demonstrate that in both sexes, DHT promote adiposity, probably via actions on the androgen receptor, while effects of testosterone may be complex due to its direct actions on the androgen receptor combined with indirect effects on estrogen receptors.

**1.3.2. Androgen receptor**—The role of androgen receptor in body weight control was suggested by an abnormal body weight profile in male mice with global androgen receptor knockout (Fan, et al. 2005; Lin, et al. 2005; Sato, et al. 2003). Interestingly, while young mutant male mice show decreased body weight, aged mutant male mice have increased body weight and adiposity compared to age-matched wild type males (Fan et al. 2005; Lin et al. 2005; Sato et al. 2003). Although selective deletion of androgen receptor in the brain (resulting from crosses between synapsin 1-Cre mice and *AR<sup>fl/y</sup>*) does not affect body weight in young male mice, such deletion leads to a late onset obesity (Yu, et al. 2013). These observations indicate that brain androgen receptor is required to prevent aging-associated obesity in males. Further characterization of the brain-specific androgen receptor deletion mouse model revealed increased expression of an anorexigenic neuropeptide, agouti-related peptide, and impaired insulin sensitivity in the hypothalamus, which presumably contribute to development of obesity in these mice (Yu et al. 2013). Interestingly, HFD feeding was found to accelerate obesity development in mice with brain-specific androgen receptor deletion, and this phenomenon is associated with a heightened activity of hypothalamic NF- $\kappa$ B and increased expression of protein-tyrosine phosphatase 1B (PTP1B) (Yu et al. 2013).

Androgen signaling in the liver also contributes to energy homeostasis, as male mice with deletion of androgen receptor from the liver develop obesity when fed with HFD, but female mutants do not (Lin, et al. 2008). Deletion of the androgen receptor in adipocyte (using the AP2-Cre as a putative fat-specific Cre allele) does not affect adiposity in male mice fed with chow diet (Yu, et al. 2008), but this deletion results in increased visceral fat in mice fed with HFD (McInnes, et al. 2012). These results suggest that androgen signaling in the adipose tissue protect against HFD-induced visceral obesity in males. However, the AP2-Cre used for the deletion of androgen receptor may also cause deletion of the receptor in the brain (Mullican, et al. 2013). Thus, the effects of the androgen receptor in the fat need to be further validated. The deletion of the androgen receptor in the muscle decreases the weight of both fat mass, lean mass, and body weight in male mice, despite normal food intake (Ophoff, et al. 2009). These indicate that the androgen signaling in the muscle is essential for the masculinization of energy homeostasis in males. Thus, effects of the androgen receptor in male animals are tissue-specific. While its actions in the brain, liver and adiposity function to prevent male mice from obesity, its actions in the muscle promote growth of both lean mass and adiposity. Notably, except for liver-specific deletion, female phenotypes in mice lacking androgen receptor globally or in specific tissues were not reported in these studies. However, Fagman et al reported that global deletion of androgen receptor in female mice on an apolipoprotein E (apoE)-deficient background are prone to diet-induced obesity compared to apoE-deficient female mice (Fagman, et al. 2015). While these results suggest potential anti-obesity actions of androgen receptor in females, these effects need to be further confirmed in animals with normal apoE functions.

#### 1.4. Developmental Programming

The structure and functions of many organs that regulate energy balance, e.g. the brain, are immature at birth (Markakis 2002). In rodent brains, the majority of neural circuits do not fully develop till about postnatal day 21 (Grove and Smith 2003; Rinaman 2006, 2007).

After weaning, the brain still undergoes substantial re-modeling (McNay, et al. 2011). Events (e.g. nutrient and hormone milieu) during the perinatal window have profound effects on the development of neural circuits controlling body weight and program energy balance later in life (Remmers and Delemarre-van de Waal 2011). For example, the placenta undergoes adaptations in response to maternal diet and metabolic status to alter fetal nutrient supply, which contributes to the developmental programming of offspring (Gabory, et al. 2013; Tarrade et al. 2015). This placental programming effect shows sexual dimorphism (Tarrade, et al. 2015). Male placenta decreases size in response to maternal high salt or high fat intake, which is associated with increased inflammation in the placenta and increased fetal exposure to free fatty acids and glucose. In contrast, female placenta is only affected by maternal high fat intake but is resistant to high salt stress (Reynolds, et al. 2015). In addition, it has been reported in mice that maternal high fat feeding during lactation period predisposes the male offspring for obesity and impaired glucose homeostasis, which is associated with an impairment of POMC neuron-originated neural circuitry. Strikingly, these effects are not seen in female offspring (Vogt, et al. 2014). While mechanisms for this sex difference is not clear, sex hormones have been implicated to regulate development of the brain to program energy balance. For example, immediately after birth, male rodents experience a transient testosterone surge in the circulation associated with a concurrent testosterone surge detected in the hypothalamus (Rhoda, et al. 1984). Interestingly, hypothalamic estrogen levels also increase at birth in males (Rhoda et al. 1984), presumably due to aromatization of testosterone. Estrogens are also detectable in female hypothalamus at birth although their levels are lower than those in males during the neonatal period (Amateau, et al. 2004). This neonatal surge of sex hormones may program the long-term regulations of energy balance in both sexes. For example, administration of testosterone in male and female rat pups lead to changes in structures and excitability of hypothalamic neurons (Matsumoto and Arai 1980). Further, neonatal exposure to excess testosterone renders a long-term obese phenotype in female mice, characteristic of increased energy intake and increased visceral adiposity during the adulthood (Nohara, et al. 2013b; Nohara, et al. 2011). These are associated with reduced leptin sensitivity, reduced expression of POMC mRNAs and POMC neuron innervations in the hypothalamus, which may contribute to obesity (Nohara et al. 2011). In males, neonatal exposure to testosterone leads to an early reduction in lean mass, decreases in locomotor activity, energy expenditure and food intake, but a late onset of hyperadiposity (Nohara, et al. 2013a). It is important to note that the reduced leptin sensitivity in females is not recapitulated by neonatal DHT exposure (Nohara et al. 2011), suggesting that this effect of testosterone may be mediated by estrogen receptors. Collectively, the current literature suggests robust effects of sex hormones on programming of energy balance during early development. Notably, most of genetic mouse studies so far use animal models with genes (e.g. ER $\alpha$ , ER $\beta$ , and AR) deleted at embryonic stage. Thus data obtained from these models could not fully distinguish effects of the sex hormones during early development vs. adulthood; further investigations using inducible genetic deletion models are warranted to tackle this issue.

## 2. Sex chromosomes

The sex dimorphism in gonadal hormones is ultimately determined by the sex chromosomes. The Sry gene located in the Y chromosome initiates the differentiation of the testes, which



produce androgens and cause the masculinization of male mice (Goodfellow and Lovell-Badge 1993). Without the Sry gene, ovaries will be developed in the female mice and produce ovary hormones including estrogens and progesterone (Goodfellow and Lovell-Badge 1993). To evaluate the pure effect of sex chromosomes, the Sry gene was deleted from the Y chromosome to generate XY mice with ovaries, and translocated to a non-sex chromosome to generate XX mice with testes, together with XX ovary mice and XY testis mice. These “four core genotypes” mouse model were used to distinguish the effects between gonadal sex (testes or ovaries) and sex chromosomes (XX or XY) (De Vries, et al. 2002). Karen Reue’s group found that gonadal males (XX testis and XY testis) are heavier than gonadal females (XX ovary and XY ovary) in the gonad-intact adult mice, indicating that physiologically sex hormones are dominant in the regulation of sex dimorphism of energy homeostasis. However, the body weight difference among the four groups of mice disappeared one month after the gonadectomy, and XX mice (XX ovary and XX testis) gradually become heavier than XY mice (XY ovary and XY testis) 2–10 months after the gonadectomy, associated with increased fat mass, increased daylight time feeding, and decreased lipid oxidation (Chen, et al. 2012; Reue 2017). They also found that XX and XXY mice are much heavier than XO and XY mice, further supporting an important role of the X chromosome dosage in the regulation of energy balance (Chen et al. 2012; Reue 2017).

Thus, the X chromosome dosage is a risk factor for obesity, with one extra X chromosome instead of Y chromosome causing body weight gain (Reue 2017). Notably, XX mice are reported to have lower aromatase expression in the brain than XY mice, independent of gonadal sex (Cisternas, et al. 2015), raising a possibility that the obesity seen in XX mice may be due to decreased brain estrogen bioavailability. In addition, the inactivation of one of the two X chromosomes in XX cells during early development silences most genes on the X chromosome, but some X-linked genes escape this process and cause a higher dose in females than males (Chen et al. 2012; Pessia, et al. 2012; Reue 2017). Some of the X-linked “escape” genes, e.g. *Eif2s3x*, *Kdm6a*, *Ddx3x*, *Kdm5c*, *Usp9x* and *Uba1*, have been validated to be expressed at higher levels in XX fat mass than XY gonadal fat in gonadectomized mice (Chen et al. 2012). However, whether these genes are responsible for the X chromosome’s regulation on energy homeostasis needs to be further studied.

In parallel, a few other X-linked genes have been implicated in the regulation of energy balance. One such example is the gene encoding the androgen receptor, as we have discussed. Another well-studied X-linked gene is O-GlcNAc transferase (OGT), which catalyzes the posttranslational modification of proteins by O-GlcNAc and is regulated by nutrient access. Loss of OGT in orexigenic AgRP neurons inhibits neural activity of these neurons and leads to lean phenotypes in both male and female mice (Ruan, et al. 2014). On the other hand, loss of OGT in glutamatergic neurons causes robust obesity in male mice associated with hyperphagia, and these phenotypes can be rescued by restoration of OGT only in PVH neurons which are largely anorexigenic (Lagerlof, et al. 2016). Female mice carrying the same mutation were not characterized in this study (Lagerlof et al. 2016). Nevertheless, functions of OGT appear to be site-specific, promoting weight gain in AgRP neurons but preventing obesity in PVH neurons, but its sex-specific role needs to be further characterized.

5-HT<sub>2C</sub> receptor (5-TH<sub>2C</sub>R), encoded by an X-linked gene (*Ht2cr*), also plays essential roles in preventing obesity (Tecott, et al. 1995). Male mice with global deficiency of 5-TH<sub>2C</sub>R develop a late onset hyperphagic obesity, which can be exacerbated by HFD feeding (Nonogaki, et al. 1998; Tecott et al. 1995). 5-HT and its analogs, e.g. d-fenfluramine, have been shown to suppress food intake in animals, and these effects are largely mediated via the 5-TH<sub>2C</sub>R expressed by POMC neurons in male animals (Berglund, et al. 2013; Xu, et al. 2008). In addition, the 5-TH<sub>2C</sub>R expressed by dopamine neurons mediates 5-HT actions to inhibit binge-like eating in male mice (Xu, et al. 2017). Indeed, a selective agonist of the 5-TH<sub>2C</sub>R, lorcaserin, was approved by FDA as an anti-obesity medicine. Mechanisms for 5-TH<sub>2C</sub>R's effects on appetite control involve its actions to activate the transient receptor potential channel 5 (TrpC5) which leads to depolarization of POMC neurons (Gao, et al. 2017). Interestingly, TrpC5 is also encoded by the X chromosome, and it has been shown to mediate actions of multiple hormones on POMC neurons, including estrogens (Qiu, et al. 2018), leptin (Gao et al. 2017; Qiu, et al. 2010) and insulin (Qiu, et al. 2014). Importantly, deletion of TrpC5 in POMC neurons leads to obesity in male mice, which is associated with increased daylight feeding and decreased energy expenditure (Gao et al. 2017). However, effects of both 5-TH<sub>2C</sub>R and TrpC5 deletion on female energy balance have not been reported.

In summary, the Sry gene on the Y chromosome determines the development of male or female gonads, which influence the circulating sex hormones and therefore energy homeostasis. The X chromosome also influences the energy balance in a dose-dependent manner. A few X-linked genes have been implicated in energy balance, but their contributions to the sex difference in body weight balance remain to be further investigated. In addition, more X-linked genes may be related to the sexual dimorphism in energy balance, which remain to be identified.

### 3. Other factors

While the majority of the field focus on the functions of sex hormones and sex chromosomal genes, other factors (not sex hormones/receptors and not encoded by the sex chromosomes) may exist to regulate the sex differences in energy homeostasis.

For example, POMC mRNA and POMC neuron projections have been shown to be more abundant in female hypothalamus than in males, corresponding to lower food intake in female mice (Nohara et al. 2011). Further, we recently found that POMC neurons in female mice display higher neural activities compared to male counterparts (Wang, et al. 2018). In addition, enhanced POMC mRNAs and neural activity are both attributed to a transcription factor, namely TAp63. Strikingly, deletion of TAp63 in POMC neurons confers “male-like” diet-induced obesity to female mice, but does not affect body weight in male mice (Wang et al. 2018). Importantly, anorexigenic effects of estrogens are not affected by loss of TAp63 in POMC neurons (Wang et al. 2018), suggesting that TAp63 actions are largely independent of the female sex hormone. Notably, the gene encoding TAp63 is located on an autosome. Thus, these results indicate that TAp63 in POMC neurons is one such example that factors, not related to sex hormones/receptors and not on sex chromosomes, could contribute to the sex differences in energy balance. Interestingly, Sirt1, a nicotinamide adenine dinucleotide

(NAD<sup>+</sup>) dependent deacetylase (Imai, et al. 2000; Michan and Sinclair 2007), is a known transcriptional target of TAp63 (Su, et al. 2012). Deletion of Sirt1 in POMC neurons increases susceptibility to diet-induced obesity specifically in female mice but not in male mice due to reduced energy expenditure, indicating that Sirt1 in POMC neurons plays a sexually dimorphic role in energy homeostasis (Ramadori, et al. 2010). Similarly, loss of signal transducer and activator of transcription 3 (STAT3) in POMC neurons leads to modest obesity in female mice but does not affect male mice (Xu, et al. 2007). Notably, STAT3 is shown to mediate anorexigenic effects of estrogens (Gao et al. 2007). Thus, the sexually dimorphic functions of STAT3 in POMC neurons may result from different levels of circulating estrogens in males vs. females. In contrast, deletion of GABA<sub>B</sub> receptors in POMC neurons results in obesity in the male mice but not female mice fed a HFD, but it is unclear why GABA<sub>B</sub> receptors affect male and female differently (Ito, et al. 2013).

In addition to POMC neurons, microglial cells in the hypothalamus also contribute to the sex differences in body weight control. Dorfman et al recently reported that male mice fed a HFD reduce hypothalamic levels of CX3CL1 (a neuron-released chemokine) and its receptor expressed by microglia, CX3CR1, while the CX3CL1-CX3CR1 levels remain normal in the hypothalamus of HFD-fed female mice (Dorfman et al. 2017). Interestingly, female Cx3cr1 knockout mice develop “male-like” hypothalamic microglial accumulation and activation, associated with increased susceptibility to diet-induced obesity (Dorfman et al. 2017). Conversely, increasing brain CX3CL1 levels in the hypothalamus of male mice converts them to a “female-like” metabolic phenotype with reduced microglial activation and reduced body weight gain (Dorfman et al. 2017). These data identify the CX3CL1-CX3CR1 signaling in hypothalamic microglia as an important mediator for sex differences in energy balance.

Kisspeptin (Kiss1) and its receptor, GPR54, are key regulators of reproduction (Murphy 2005). Female mice lacking GPR54 develop massive obesity fed with either chow or HFD, associated with increased fat mass and serum leptin level, and decreased energy expenditure but normal food intake (Tolson, et al. 2016; Tolson, et al. 2014). On the other hand, male mice lacking GPR54 have none to minimal phenotypes in energy balance (Tolson et al. 2016; Tolson et al. 2014). Thus, the Kiss1-GPR54 signaling regulates energy homeostasis in a sexually dimorphic fashion. Notably, since loss of Kiss1-GPR54 signaling substantially reduces the circulating levels of testosterone in males and reduces estrogens in females (d'Anglemon de Tassigny, et al. 2007; Seminara, et al. 2003), the obesity phenotypes observed in GPR54 knockout mice could be at least partially attributed to the altered levels of sex hormones in these mice. Similarly, global deletion of angiotensin II receptor (AT2R) renders female mice more prone to diet-induced obesity with impaired lipid metabolism, but the same mutation does not affect male mice (Samuel, et al. 2013). This female-specific obesity is associated with reduced estrogen level in female mutants, which may account for the sexually dimorphic obesity seen in mice lacking AT2R (Samuel et al. 2013).

While many of the aforementioned factors appear to prevent obesity in females, lecithin cholesterol acyltransferase (LCAT) plays an exactly opposite role. LCAT is an enzyme in the regulation of high density lipoprotein (HDL) metabolism. Deletion of LCAT results in a profound HDL deficiency in both male and female mice (Li, et al. 2011). However, LCAT

knockout female mice are protected from high fat high sucrose diet-induced obesity, but male mutant mice are not (Li et al. 2011). The most striking sexually dimorphic pattern was found in mice lacking the cytokine IL-6 in the muscle. On one hand, deletion of IL-6 from the muscle results in decreased body weight in male mice, which is associated with lower core body temperature during the light phase and increased respiratory exchange ratio (Ferrer, et al. 2014; Molinero, et al. 2017). On the other hand, loss of IL-6 in female muscles leads to increased body weight (Ferrer et al. 2014; Molinero et al. 2017). Thus, IL-6 in the muscle regulates energy balance in exactly opposite directions in males vs. females.

Together, these findings indicate that many autosome genes contribute to the sex differences in energy balance through multiple mechanisms. Conceivably, some factors/genes may be transcriptional targets of sex hormones, since estrogens, progesterone and androgens are all known to regulate gene expression via their nuclear receptors. Alternatively, some factors/molecules may mediate body weight-regulatory effects of the sex hormones, e.g. STAT3 being a mediator of anorexigenic effects of estrogens. Further, some factors could regulate levels of sex hormones or receptors, e.g. Kiss1-GPR54 and AT2R. Of course, some genes may be independent of sex hormone actions and function as parallel mechanisms to mediate sex differences in energy balance, e.g. TAp63. Of course, how these factors functionally interact with the sex hormones/chromosomes remain largely unknown and future studies are certainly warranted. It also needs to be pointed out that a lot more factors/molecules in this category may exist but are missed simply because female animals are purposely avoided in most studies to reduce efforts.

## Conclusions

The existence of sex difference in biology and physiology, including the energy homeostasis, has been increasingly appreciated (Arnold 2014), and the underlying mechanisms started to emerge (Figure 1). The sex chromosomes are the fundamental difference between males and females and genes in the X or Y chromosome are conceivably major contributors to the sexual dimorphism. The Sry gene in the Y chromosome determines the fate of gonadal development, and therefore influence the levels of sex hormones. The X chromosome dosage adds the risk of obesity in females with mechanisms that are yet to unfold; in parallel, some X-linked genes, including those encoding the androgen receptor, OGT, 5-HT<sub>2C</sub>R, trpC5, and likely more, are involved in the regulation of energy balance, and some of these molecules are therapeutic targets for obesity treatment. All sex hormones, including estrogens, progesterone and androgens, are involved in the regulation of body weight balance in both sexes, but their actions and mechanisms are largely sex-specific. In addition, increasing number of autosome genes are found to contribute to the sex differences in body weight control. While some of these genes are either downstream targets or upstream factors of sex hormones, some of them appear to provide parallel mechanisms and function independent of the hormones.

Despite the expanding list of factors found to be involved in sex differences in body weight control, and an emerging framework where these factors and signals may be integrated to regulate energy homeostasis in different sexes, a number of areas need further investigations. First, for any tissues or cells that can regulate energy balance, their functions in different

sexes would depend on the strength of estrogenic or androgenic tone locally in that region, which is ultimately determined by the availability of the sex hormones, the aromatase activity, and the relevant receptors. Thus, a few important questions need to be systemically addressed. Are these factors (sex hormones, aromatase, and the receptors) present in metabolic organs (including body weight-regulatory brain regions) in each sex? Are their levels dynamic throughout the life period? Most importantly, what are their functions in the context of energy balance? In addition, given the established role of the X chromosome dosage in body weight control, efforts are needed to identify what X-linked genes regulate energy balance and to determine whether these genes regulate body weight only in females, males or both. Further, we predict that a large number of other factors, not directly related to sex hormones/receptors and not encoded by sex chromosomes, also contribute to the sex differences in energy balance. Therefore we call attention to these “other factors” as potential regulators of sex differences in energy balance. Pharmacological and genetic investigations into energy balance in animal models are warranted and should contrast males with females, as required by the funding agencies, e.g. the NIH (Tannenbaum, et al. 2016). After these factors are identified, new studies can be designed to further delineate how they interact with sex hormones as well as sex-linked genes.

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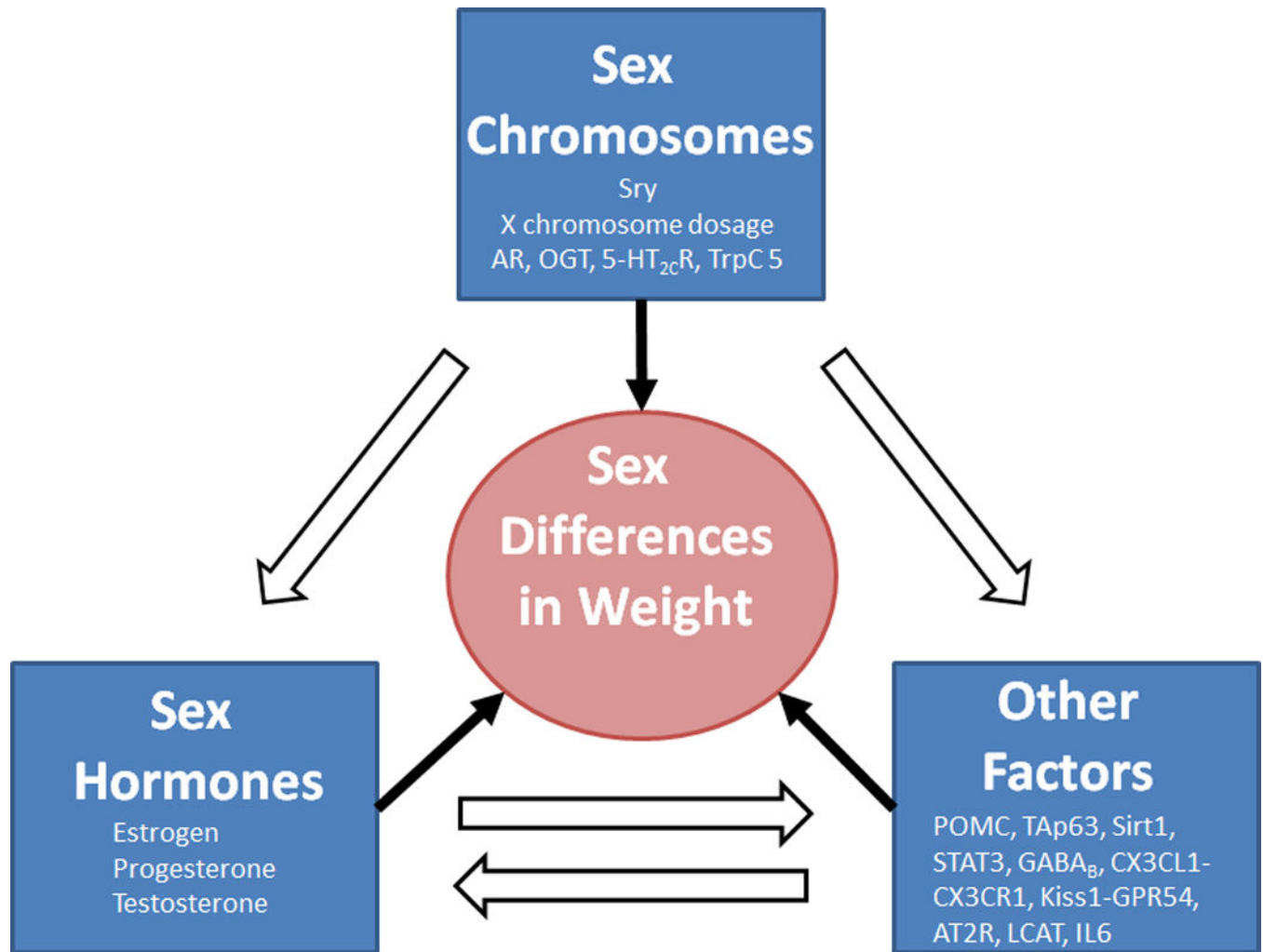
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**Figure 1. A proposed framework for molecular mechanisms underlying sex differences in body weight control.**

The Sry gene in the Y chromosome determines gonadal development and therefore the levels of sex hormones. The X chromosome dosage adds the risk of obesity in females; some X-linked genes have been implicated in the regulation of energy balance. Sex hormones are involved in the regulation of body weight balance in both sexes. In addition, other factors (autosomal-encoded factors that are not sex hormones/receptors) contribute to the sex differences in body weight control. These three categories of factors are integrated through their reciprocal interactions to ultimately determine the sex differences in body weight balance.