

Androgen excess in pancreatic β cells and neurons predisposes female mice to type 2 diabetes

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Androgen excess predisposes women to type 2 diabetes (T2D), but the mechanism of this is poorly understood. We report that female mice fed a Western diet and exposed to chronic androgen excess using dihydrotestosterone (DHT) exhibit hyperinsulinemia and insulin resistance associated with secondary pancreatic β cell failure, leading to hyperglycemia. These abnormalities are not observed in mice lacking the androgen receptor (AR) in β cells and partially in neurons of the mediobasal hypothalamus (MBH) as well as in mice lacking AR selectively in neurons. Accordingly, i.c.v. infusion of DHT produces hyperinsulinemia and insulin resistance in female WT mice. We observe that acute DHT produces insulin hypersecretion in response to glucose in cultured female mouse and human pancreatic islets in an AR-dependent manner via a cAMP- and mTOR-dependent pathway. Acute DHT exposure increases mitochondrial respiration and oxygen consumption in female cultured islets. As a result, chronic DHT exposure in vivo promotes islet oxidative damage and susceptibility to additional stress induced by streptozotocin via AR in β cells. This study suggests that excess androgen predisposes female mice to T2D following AR activation in neurons, producing peripheral insulin resistance, and in pancreatic β cells, promoting insulin hypersecretion, oxidative injury, and secondary β cell failure.

Conflict of interest: The authors have declared that no conflict of interest exists

Submitted: February 26, 2018 **Accepted:** May 10, 2018 **Published:** June 21, 2018

Reference information:

JCI Insight. 2018;3(12):e98607. https://doi.org/10.1172/jci. insight.98607.

Introduction

The association between testosterone (androgen) excess and dysglycemia in women has been known since the original description of diabetes in bearded women by Achard and Thiers (1). In most prospective and cross-sectional studies, increased testosterone concentrations are associated with increased risk of type 2 diabetes (T2D) in women (2), and polycystic ovarian syndrome (PCOS), the main cause of testosterone excess, predisposes women to T2D (3–5). Despite this background, the exact mechanism of hyperglycemia in women with androgen excess is unknown. In extreme syndromes of insulin resistance and in obese women with PCOS, insulin resistance and the resulting hyperinsulinemia drive the ovarian production of androgens (6). However, extensive evidence suggests that in the most common form of PCOS androgen excess is instrumental in altering glucose homeostasis. First, PCOS is characterized by primary functional



ovarian hyperandrogenism that is independent from insulin resistance (7). Second, in hyperandrogenic women, about half of the intervention studies to reduce androgen levels or block androgen action improved insulin sensitivity and glucose homeostasis (8–12). Finally, testosterone administration to healthy women or female-to-male transsexuals also promotes insulin resistance (13, 14). Therefore, it is critical to dissect the mechanisms by which androgen excess alters glucose homeostasis in women using preclinical models because it opens new avenues for therapy.

Male mice with neuronal androgen receptor (AR) deletion exhibit hypothalamic insulin resistance, demonstrating the importance of AR action in the brain in promoting insulin sensitivity (15). In female mice, testosterone alters kisspeptin and gonadotropin-releasing hormone neuronal function in the arcuate nucleus (ARC) of the hypothalamus (16), an area that controls peripheral insulin sensitivity (17–21). We reported that developmental or adult androgen excess in female mice alters the function of ARC neurons in an AR-dependent manner (22, 23). Thus, testosterone action in hypothalamic neurons could contribute to insulin resistance in hyperandrogenic women (24, 25).

It is widely recognized that pancreatic β cell failure is essential to the pathogenesis of diabetes (26, 27), suggesting that hyperglycemic women with androgen excess exhibit β cell dysfunction. In fact, women with androgen excess exhibit either higher basal insulin secretion and decreased postprandial insulin secretion (28) or an exaggerated acute insulin response to glucose (29). These abnormalities are not explained by insulin resistance and are closely associated with androgen excess (30). In these women, there is a robust relationship between β cell function and testosterone, raising the possibility that excess testosterone produces insulin hypersecretion and β cell dysfunction (31). The AR is expressed in β cells in both sexes, and AR activation in β cells enhances glucose-stimulated insulin secretion (GSIS) from male rodent and human islets (32). Thus, in women, testosterone action on the β cell AR may produce insulin hypersecretion and β cell dysfunction.

To explore the hypothesis that androgen excess produces T2D via action on AR in β cells and/or neurons, we generated female mice with androgen excess and conditional AR deletion in β cells and/or neurons. We show that androgen excess predisposes these mice to hyperglycemia via the combined activation of the AR in pancreatic β cells (producing dysfunction) and hypothalamic neurons (producing peripheral insulin resistance).

Results

Generation of female mice with AR deletion in β cells and hypothalamus. We previously reported that male β cell–specific AR-knockout (β ARKO^{RIP}) mice, generated using a RIP2-Cre transgene, exhibit total β cell AR elimination associated with a 50% elimination of AR in the ARC and ventromedial hypothalamus (VMH) (32). Therefore, we used this model to assess the effect of androgen excess in β cells and hypothalamic neurons of female mice. Since mice expressing the RIP2-Cre transgene develop glucose intolerance and impaired insulin secretion even in the absence of genes targeted by loxP sites (33), we have used the RIP-Cre mice as a control group in female as well as in male mice (32). Elimination of the AR protein in islets (Figure 1A) and decreased AR expression in the ARC (Figure 1, B and C) and VMH (Figure 1, B and D) was confirmed in female β ARKO^{RIP} mice.

Androgen excess predisposes female controls, but not $\beta ARKO^{RIP}$ mice, to diabetes. To study the metabolic consequences of androgen excess in vivo, female control and $\beta ARKO^{RIP}$ mice were exposed to dihydrotestosterone (DHT) — the nonaromatizable AR agonist — or vehicle for 4 weeks (vehicle, 121.5 ± 12 pg/ml, DHT 489 ± 23.2 pg/ml, mean \pm SEM). On a normal chow diet, female $\beta ARKO^{RIP}$ mice showed no alteration in glucose homeostasis compared with controls (Supplemental Figure 1, A–G; supplemental material available online with this article; https://doi.org/10.1172/jci.insight.98607DS1). Following exposure to DHT, both control and $\beta ARKO^{RIP}$ mice retained normal glucose tolerance (Supplemental Figure 1A) and normal fasting and fed blood glucose levels (Supplemental Figure 1, B and C) and fasting insulin levels (Supplemental Figure 1D). However, DHT exposure produced a marked fed hyperinsulinemia in control mice, which was not observed in $\beta ARKO^{RIP}$ mice (Supplemental Figure 1E). Control and $\beta ARKO^{RIP}$ mice treated with vehicle and DHT retained similar islet architecture and pancreatic insulin concentration (Supplemental Figure 1, F and G). Exposure to DHT slightly increased β cell mass in both control and $\beta ARKO^{RIP}$ mice (Supplemental Figure 1F).

To mimic metabolic stress seen in humans, mice were exposed to a high-sucrose, high-fat Western diet (WD). Under these conditions, exposure to DHT produced abnormalities characteristic of early T2D in female control mice. They displayed fasting and fed hyperinsulinemia (Figure 2, A and B) associated with exaggerated GSIS (Figure 2C) and insulin resistance during an insulin tolerance test (ITT) (Figure 2D). Accordingly, these mice developed compensatory β cell hyperplasia, with increased islet size and β cell mass as well as pancreas insulin content (Figure 2, E–G). However, DHT exposure produced a failure of



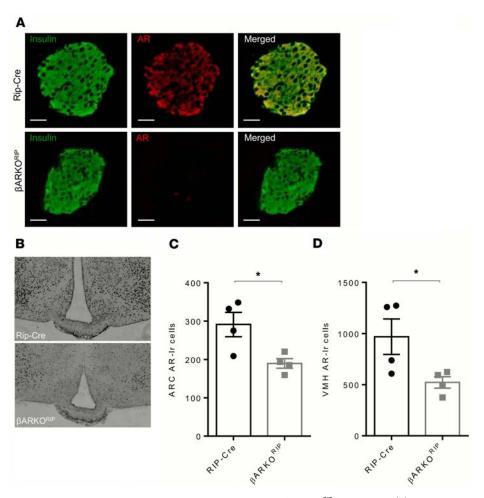


Figure 1. AR expression in pancreatic β cells and the hypothalamus of βARKO^{RIP} female mice. (A) Pancreas sections showing AR immunofluorescent staining (red) colocalizing with insulin (green) in β cells of control RIP-Cre female mice and confirming successful AR deletion in representative βARKO^{RIP} islets at 12 weeks. Original magnification, \times 40. (B) AR expression in the arcuate nucleus (ARC) and ventromedial hypothalamus (VMH) of RIP-Cre and βARKO^{RIP} mice, as assessed by immunostaining. (C) AR-immunoreactive cell number in the ARC. (D) AR-immunoreactive cell number in the VMH. Values represent the mean \pm SEM with dot plots (n = 4). *P < 0.05, Student's t test.

β cells to compensate for insulin resistance, as DHT-exposed control mice exhibited hyperglycemia in the fasting and fed state despite hyperinsulinemia (Figure 2, H and I) and following a glucose challenge (Figure 2J). In control mice, DHT exposure was also associated with increased expression of PEPCK (pckI) mRNA in liver, suggesting increased gluconeogenesis (Figure 2K). In contrast, in female $βARKO^{RIP}$ mice, exposure to DHT did not produce fasting hyperinsulinemia and did not induce insulin resistance with associated β cell hyperplasia (Figure 2, A–G). These mice remained normoglycemic in the fasting and fed states and during a glucose challenge (Figure 2, H–J), and DHT produced no increase in liver PEPCK mRNA (Figure 2K). Thus, under the conditions of a WD, chronic DHT exposure has produced a phenotype of insulin resistance with islet hyperplasia and hyperinsulinemia, culminating in hyperglycemia in control mice. This is not observed in $βARKO^{RIP}$ mice on WD exposed to chronic DHT.

Androgen excess predisposes female controls, but not NARKO-/- mice, to diabetes. In female mice, AR protein was expressed in the ARC and the VMH (Figure 3, A and B). Expression of AR protein in these nuclei was similar in males and females (Figure 3, C and D). Likewise, in microdissected ARC, AR mRNA expression was similarly expressed in female and male mice (Figure 3E). Consistent with the functionality of AR in female neurons of the mediobasal hypothalamus (MBH), whole-cell patch-clamp recordings were conducted to determine the effect of DHT application on the excitability of neurons in the ARC and VMH. In the ARC, application of DHT altered the firing rate in a subset of recorded neurons (Figure 3, F–H and L, and Supplemental Table 1). Following DHT application, hyperpolarization was observed in 7 of 19 recorded



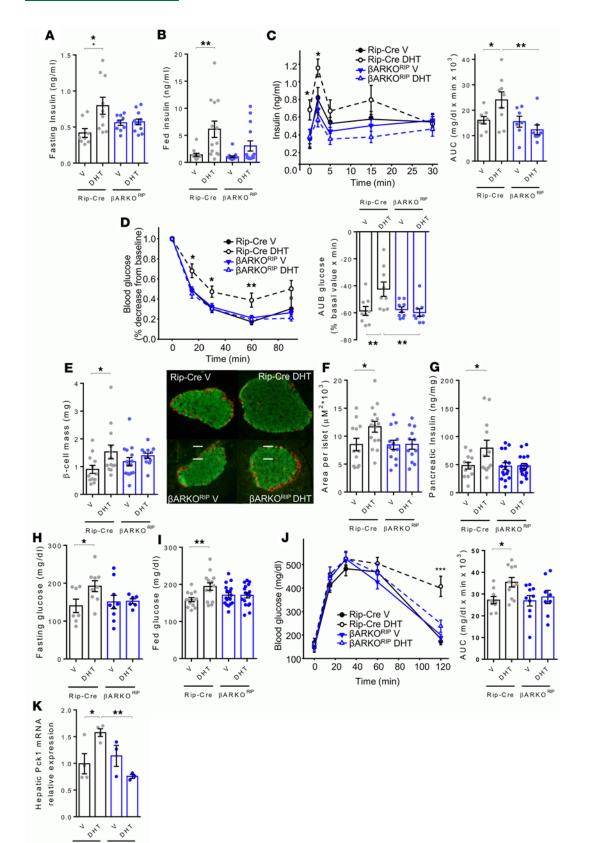


Figure 2. Chronic DHT excess predisposes female controls, but not BARKORIP mice, to diabetes. Female mice were fed WD for 9 weeks followed by s.c. treatment with vehicle (V) or DHT for 4 weeks. (A) Fasting serum insulin levels. (B) Random fed serum insulin levels. (C) Insulin secretion during an i.p. glucose-stimulated insulin secretion (GSIS) test (3 g/kg) and corresponding AUC. (D) i.p. insulin tolerance test (ITT) with blood glucose represented as percentage decrease from baseline, with corresponding area under baseline (AUB). (E) β Cell mass and representative islets with insulin staining (green). Scale bar: 10 μm. (**F**) Islet area. (G) Pancreas insulin concentration. (H) Fasting blood glucose. (I) Random fed blood glucose. (J) Blood glucose during an i.p. glucose tolerance test (GTT) (2 g/ kg) and AUC. (K) Hepatic pck1 relative expression quantified by qPCR. Values represent the mean ± SEM with dot plots (n = 6-19). *P < 0.05, **P < 0.01, Student's t test or 2-way ANOVA.

Rip-Cre βARKO^{RE}



neurons, accompanied by a decrease in action potential frequency (Figure 3, F–H and L, and Supplemental Table 1). In contrast, another subset of neurons responded with depolarization and an increase in firing rate, which did not reach statistical significance (Figure 3, I–K, and Supplemental Table 1). The remainder of the recorded neurons was unresponsive to DHT (-42.93 ± 3.56 mV vs. -41.98 ± 3.65 mV, n = 8). In the VMH, bath application of DHT showed similar results. A decrease in action potential frequency was determined in 6 of 10 recorded VMH neurons (2.44 ± 0.65 Hz vs. 1.05 ± 0.38 Hz) (P < 0.05), while two neurons responded with an increasing trend (1.35 ± 0.62 Hz vs. 4.27 ± 0.38 Hz) and two did not respond to DHT (5.30 ± 1.61 Hz to 5.23 ± 2.05 Hz) (Figure 3L and Supplemental Table 1).

To assess the contribution of AR in the hypothalamus to DHT-induced insulin resistance and hyperglycemia, we generated female neuronal AR-knockout (NARKO^{-/-}) mice by crossing AR^{lox/lox} mice with the synapsin-Cre-transgenic mice. NARKO^{-/-} male mice have previously been shown to display decreased AR expression in the brain but normal expression in other tissues (32). In female NARKO^{-/-} mice, body weight, fasting and fed blood glucose and insulin, glucose tolerance, insulin sensitivity, and GSIS were similar to those of control AR^{lox/lox} mice on a normal chow diet (Supplemental Figure 2). Consistent with results obtained previously (Figure 2), DHT exposure in WD-fed female control mice produced fasting and fed hyperinsulinemia with insulin resistance (Figure 4, A–C) associated with hyperglycemia in the fasting and fed states (Figure 4, D and E) and during glucose challenge (Figure 4F). These effects were not observed in female NARKO^{-/-} mice (Figure 4). This demonstrates that excess DHT action on neuronal AR is instrumental to the diabetogenic effect of DHT in female mice. Note that DHT exposure produced an increase in body weight and adipose tissue mass in female control mice fed a WD, which was not observed in female NARKO^{-/-} mice (Supplemental Figure 3).

To further characterize the contribution of excess AR activation in neurons to hyperglycemia in female mice exposed to a WD, we infused DHT directly into the lateral ventricle of the brain in order to induce central DHT excess without peripheral DHT elevation. We confirmed that serum DHT concentrations were not increased in DHT-infused mice (vehicle, 162.6 ± 14.3 ; DHT, 171.5 ± 24.3 pg/ml). After 4 weeks of infusion, compared with vehicle-infused female mice, DHT-infused mice developed isolated fasting hyperinsulinemia (Figure 5, A and B) associated with insulin resistance during ITT (Figure 5C). Unlike those after peripheral exposure to DHT, central DHT had no effect on fed and fasted blood glucose and glucose tolerance (Figure 5, D–F). In addition, central DHT infusion had no effect on body weight and adipose tissue mass (Supplemental Figure 4). These findings confirm that excess AR activation selectively in neurons contributes to hyperinsulinemia and peripheral insulin resistance. However, it also suggests that, unlike in the case of general DHT exposure, i.c.v. DHT exposure does not lead to hyperglycemia.

AR activation in female β cells produces insulin hypersecretion in a cAMP- and mTOR-dependent manner. To explore the contribution of the AR in β cells in vivo without the partial AR hypothalamic deletion observed in β ARKO^{RIP} mice, we generated a second β cell–specific ARKO model (β ARKO^{MIP}) by crossing AR^{lox} mice with MIP-Cre-transgenic mice (also known as Ins1-Cre/ERT mice) that lacks Cre activity in the hypothalamus (34). Unfortunately, the MIP-Cre-transgenic mouse has documented limitations due to transgene-driven expression of human growth hormone in islets, decreasing glucagon secretion, and improving peripheral insulin sensitivity (35). Therefore, the ARlox MIP-Cre mice must be used as controls of βARKO^{MIP} mice. Although this model proved reliable to demonstrate the importance of the AR in male β cells (32), the model could not be used in females. Indeed, unlike in female AR^{lox} or RIP-Cre controls used previously (Figures 2 and 4), DHT failed to induce fasting hyperinsulinemia and hyperglycemia, glucose-induced insulin hypersecretion, and insulin resistance in female ARlox MIP-Cre mice (Supplemental Figure 5). Thus, the MIP-Cre transgene improves insulin sensitivity and blunts the effect of DHT. To circumvent this problem and to determine to what extent the DHT-induced hyperinsulinemia was islet cell autonomous, due to DHT action on AR in β cells, we studied the effect of DHT on insulin secretion in static incubation of cultured islets from WD diet-fed female control and βARKO^{RIP} mice. Because of the deleterious effect of the RIP-Cre transgene on islet function (33), which is exacerbated by WD (32), we assessed insulin secretion selectively at high glucose concentrations (16.7 mM glucose). DHT exposure produced insulin hypersecretion in control islets, but this was not observed in βARKO^{RIP} islets (Figure 6A). We translated these findings to cultured islets from adult female human donors. Consistent with results obtained in the mouse, exposure of human islets to DHT produced insulin hypersecretion at high glucose levels compared with vehicle, and this effect was reversed in the presence of the AR antagonist flutamide (Figure 6C). Neither DHT, nor AR genetic elimination, nor pharmacological inhibition affected islet insulin content in mouse or human islets (Figure 6, B and D). To examine the effect of DHT on the responsiveness of isolated



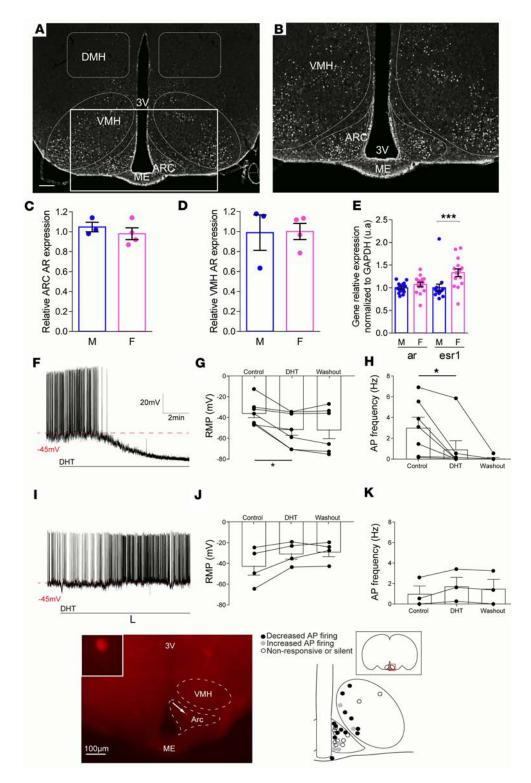


Figure 3. AR expression and activity in hypothalamic neurons of female mice. (A) AR staining in the MBH. Original magnification, ×40 (scale bar: 100 μm). **(B)** High-magnification image of the inset in **A. (C)** AR protein levels in the ARC and **(D)** the VMH. Data are expressed relative to the number of AR-positive cells in females (n = 3-4). **(E)** Estrogen receptor α (esr1) and AR relative mRNA expression in the MBH quantified by qPCR following microdissection. **(F)** Electrophysiological recordings of action potentials of a representative DHT-inhibited neuron or **(I)** a DHT-activated neuron in the ARC of wild-type females following treatment with DHT. **(G** and **J)** Resting membrane potential (RMP) and **(H** and **K)** action potential frequency (AP) before and after application of DHT and following washout. **(L)** Biocytin staining of one representative recorded neuron in the ARC (scale bar: 100 μm) and an illustration of the localization of the recorded neurons. Values represent the mean ± SEM with dot plots (n = 15 mice or n = 10-17 neurons). *P < 0.05, ***P < 0.001, Student's P < 0.001, and the plots of the recorded neurons in the test.



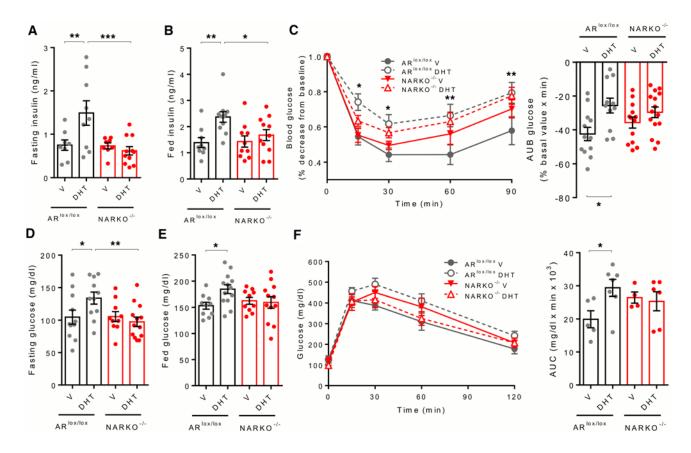


Figure 4. Chronic DHT excess predisposes female controls, but not NARKO^{-/-} mice, to diabetes. AR^{lox/lox} or NARKO^{-/-} mice were fed WD for 9 weeks followed by s.c. treatment with vehicle (V) or DHT for 4 weeks. (A) Fasting serum insulin levels. (B) Fed serum insulin levels. (C) Blood glucose represented as percentage decrease from baseline during an i.p. insulin tolerance test (ITT), with corresponding area under baseline (AUB). (D) Fasting blood glucose. (E) Random fed blood glucose. (F) Blood glucose during an i.p. glucose tolerance test (GTT) and corresponding AUC. Values represent the mean ± SEM with dot plots (n = 4–15). *P < 0.05, **P < 0.01, ***P < 0.001, ***P < 0.001,

islets with regard to first- and second-phase insulin secretion, we studied a batch of wild-type female mouse islets during a perifusion. Exposure to high glucose produced a classical rapid first-phase insulin release at 2 minutes followed by a second-phase starting at 5 minutes (36) (Figure 6E). Exposure to DHT produced an overall increased insulin secretion after both low and high glucose exposures (Figure 6E). Thus, in female islets, excess DHT produces insulin hypersecretion in response to glucose via AR activation in β cells. GSIS is triggered by glucose metabolism, ultimately leading to the opening of Ca2+ channels and influx of intracellular calcium [Ca²⁺], (37). We explored whether DHT could increase [Ca²⁺], influx in female mouse islets. Raising glucose concentration increased [Ca2+], influx to a similar extent in both vehicle- and DHT-exposed islets (Figure 6F). The finding that DHT acting on AR enhanced GSIS without increasing cellular [Ca²⁺], influx suggests that AR activation amplifies GSIS by acting downstream of the Ca²⁺ signal, as it does in male islets, by increasing cAMP signaling (32). We studied DHT-induced cAMP production in female islets dynamically using the FRET probe Epac2 camps. Exendin 4 induced a rapid and sustained 2-fold increase in cAMP levels compared with vehicle controls. DHT exerted a similar stimulatory effect on cAMP levels, although the onset of the cAMP increase was slightly delayed compared with that after exendin 4 induction (Figure 6G). Since DHT activation of AR stimulates insulin secretion via cAMP production but independently from increase in β cell mass, we reasoned that AR activation enhances GSIS via the protein kinase mTOR. Indeed, activation of mTOR in β cells stimulates insulin secretion downstream of cAMP and without affecting β cell mass (38). In addition, AR is known to activate mTOR in the prostate (39–41). In cultured islets from female mice fed a normal chow or a WD, inhibition of the TORC1 complex using rapamycin blocked the effect of DHT in enhancing GSIS after high glucose exposure (Figure 6H). However, the lack of specificity of rapamycin is a limitation of this agent. Therefore, we used a mouse overexpressing 1 copy of a kinase-dead mTOR mutant (KD-mTOR) transgene exclusively in β cells. In cultured



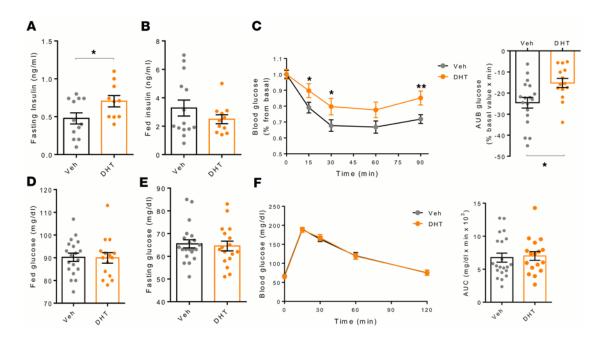


Figure 5. Chronic DHT perfusion in the brain produces insulin resistance in female mice. Female mice were fed WD for 9 weeks followed by i.c.v. infusion of vehicle (V) or DHT for 4 weeks. (A) Fasting serum insulin levels. (B) Fed serum insulin levels. (C) Blood glucose represented as an initial percentage decrease during an i.p. insulin tolerance test (ITT), with corresponding area under baseline (AUB). (D) Random fed blood glucose. (E) Fasting blood glucose. (F) Blood glucose during an i.p. glucose tolerance test (GTT) and corresponding AUC. Values represent the mean ± SEM with dot plots (n = 10–21). *P < 0.05, **P < 0.01, Student's t test or 2-way ANOVA.

islets from female KD-mTOR mice fed a normal chow, the effect of DHT in enhancing GSIS was abolished compared with that in WT islets (Figure 6I). Finally, in female mice fed a WD, the effect of chronic DHT exposure in promoting fasting and fed hyperinsulinemia and hyperglycemia, as described in Figures 2 and 4, was blunted following inhibition of the TORC1 complex with rapamycin treatment (Figure 6J). Thus, in female mice, excess DHT produces insulin hypersecretion in response to glucose via AR activation in β cells in a cAMP- and mTOR-dependent manner.

Excess AR activation in β cells produces mitochondrial hyperfunction and predisposes islets to oxidative injury. Since DHT excess produces insulin hypersecretion from female β cells, we reasoned that chronic DHT exposure could enhance β cell mitochondrial function. We studied mitochondrial oxygen consumption in female cultured islets exposed to DHT. Exposure of female islets to DHT increased the basal and total oxygen consumption rate (Figure 7A). Acute DHT exposure increases β cell mitochondrial respiration and produces insulin hypersecretion in response to glucose, suggesting that chronic DHT exposure might cause β cell exhaustion and injury. Consistent with this possibility, we previously reported that, following chronic exposure to testosterone, female mice were more susceptible to a second β cell stress induced by streptozotocin (STZ) and were prone to insulin-deficient diabetes (42). Since STZ augments the generation of islet reactive oxygen species, resulting in islet oxidative stress (43, 44), we hypothesized that chronic AR activation in female β cells promotes islet oxidative injury in vivo. To address this issue, we quantified lipid peroxidation (a marker of oxidative injury), using thiobarbituric acid reactive substances (TBARS) in female islets of WD-fed control and βARKO^{RIP} mice from experiments described in Figure 2. Chronic DHT exposure increased TBARS accumulation in control islets, consistent with the development of oxidative injury (Figure 7B). Although TBARS were increased at basal level in islets from βARKO^{RIP} mice compared with controls, DHT produced no further increase in TBARS accumulation in β ARKO^{RIP} islets, consistent with a role for AR activation in β cells in producing oxidative damage (Figure 7B). Having observed that chronic DHT exposure produces islet oxidative damage in female mice via AR in β cells, we sought to determine to what extent chronic DHT action on the AR in β cells exacerbates β cell stress induced by the ROS inducer STZ in vivo. Following STZ challenge, as previously described (45, 46), female control mice were relatively protected. However, control mice exposed to DHT became more vulnerable to STZ and showed a greater predisposition to insulin-deficient diabetes with



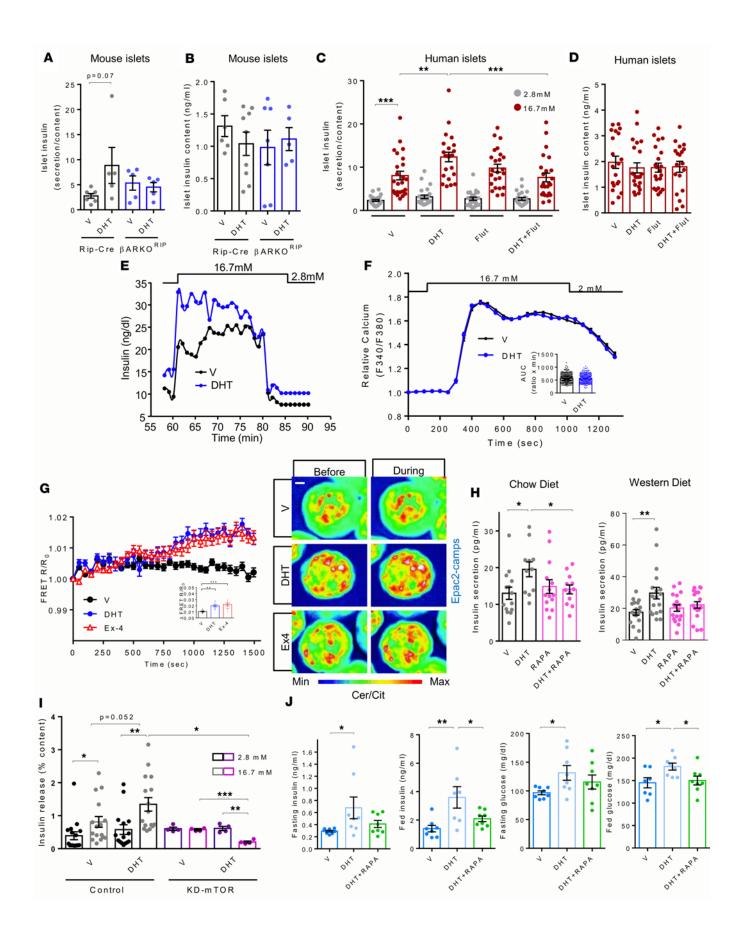




Figure 6. DHT enhances GSIS via an AR/cAMP/mTORC1 pathway. (A) GSIS measured in static incubation in islets from mice fed a WD for 8 weeks and treated with vehicle (V) or DHT (10 nM) for 48 hours ex vivo. (B) Islet insulin content from **A**. Results represent 10 islets/condition in triplicate from $n \ge 3$ independent experiments. (C) Glucose-stimulated insulin secretion (GSIS) measured after static incubation in islets from female human donors. (D) Islet insulin content from **C**. Islets were treated V, DHT (10 nM), and or flutamide (Flut, 10 μM) in vitro for 48 hours. Results represent 5 islets per condition of n = 6 independent experiments. (E) Basal and GSIS at the indicated glucose concentrations during a perifusion in WT female islets (vehicle, ethanol; DHT, 10 nM). Islets were isolated from mice at 14 weeks and perifused in batches of 60 per group. (F) Intracellular Ca²⁺ influx and corresponding AUC from the indicated glucose concentrations in isolated female mouse islets (n = 397 cells for V, n = 360 cells for DHT). (G) 10 nM DHT- and 10 nM Ex-stimulated cAMP concentrations monitored in female mouse islets infected with adenovirus harboring the FRET probe Epac2 camps, with corresponding AUC and representative live cell imaging. Scale bar: 20 μm. (H) GSIS measured in static incubation in islets from WT female mice fed chow or WD, treated with V, DHT (10 nM), rapamycin (RAPA, 10 nM), or DHT+RAPA for 48 hours (n = 12-14 batches of 10 islets/condition, 4 independent experiments). (I) GSIS measured in static incubation in islets from female WT and KD-mTOR mice, treated as in **A**. Results represent 10 islets per condition of n = 4 independent experiments. (J) WT female mice were exposed to WD for 4 weeks followed by 4 weeks of treatment with V, DHT, or DHT+RAPA (0.5 mg/kg/d; n = 7-8/group). Results represent the mean ± SEM, with dot plots of fasting and fed serum insulin as well as fasting and fed blood glucose. **P < 0.01, ***P < 0.001, 1-way ANOVA.

exacerbated hyperglycemia (Figure 7, C and D) and hypoinsulinemia (Figure 7, E and F) as well as a more pronounced reduction in β cell mass and pancreas insulin content (Figure 7, G and H). In contrast, in β ARKO^{RIP} mice, DHT produced no decrease in β cell mass or pancreas insulin content and did not exacerbate STZ-induced insulin-deficient diabetes (Figure 7, C–H). Thus, in female mice, chronic DHT activation of AR in β cells increases β cell respiration and vulnerability to other oxidative or proapoptotic injuries and predisposes to β cell failure.

Discussion

In this study, we have explored the hypothesis that androgen excess in female mice predisposes to T2D via chronic AR activation in the brain (producing peripheral insulin resistance) and in pancreatic islet β cells (producing insulin hypersecretion and secondary β cell failure). We have used a combination of mouse models in which (a) AR was activated globally (controls), (b) AR was activated globally except in β cells and MBH hypothalamic neurons (β ARKO^{RIP}), (c) AR was activated globally except in CNS neurons (NARKO^{-/-}), and (d) AR was activated selectively in the CNS. In addition, we used cultured islets from female mice and human donors. The combination of these models, when analyzed together, allows us to propose a paradigm in which excess and chronic AR activation in β cells and neurons is instrumental in promoting hyperglycemia in hyperandrogenic female mice (summarized in Figure 8).

On a WD, female control mice exposed to androgen excess exhibit exaggerated insulin secretory response, hyperinsulinemia, and insulin resistance, but they fail to adequately compensate for insulin resistance and develop β cell failure with hyperglycemia.

The first finding of this study is that excess AR activation in neurons is instrumental in the development of insulin resistance in these mice. The androgen-induced hyperinsulinemia and insulin resistance of female control mice was not observed in littermate NARKO-/- mice lacking AR selectively in neurons. Further, i.c.v. androgen infusion in wild-type female mice also produced fasting hyperinsulinemia and insulin resistance. Together, these results argue for the importance of neuronal ARs in androgen-induced insulin resistance. We reported that developmental or adult androgen excess in female mice alters the function of proopiomelanocortin (POMC) neurons in the ARC (22, 23). POMC neurons of the ARC are involved in peripheral insulin sensitivity. Leptin action in POMC neurons suppresses hepatic glucose production (18), and ablation of leptin and insulin receptors in POMC neurons produces insulin resistance and hyperandrogenism (19). Accordingly, enhancement of phosphoinositide 3-kinase signaling selectively in POMC neurons increases liver insulin sensitivity (20), and reexpression of leptin receptors exclusively in POMC neurons of leptin receptor-deficient mice normalizes blood glucose independently of obesity (21). Thus, excess AR activation in neurons of the ARC, VMH, or in other areas that influence MBH neurons may alter insulin or leptin action in POMC neurons and produce peripheral insulin resistance. This does not exclude the possibility that androgen excess can produce insulin resistance via AR activation in peripheral tissues. In fact, excess AR activation in hepatocytes of adult female mice causes hepatic insulin resistance (47). Studies are needed to identify the exact AR-expressing hypothalamic neurons that are targeted by androgens to promote peripheral insulin resistance.

Unlike others (47), we did not observe androgen-induced insulin resistance with normal rodent chow. The most likely explanation for this discrepancy is the impact of different vivarium environments. Indeed, differences in exogenous sources of soy phytoestrogens coming from normal rodent chow and the influence of different metagenomes (the interaction between host and microbiome genes) influence the experimental reproducibility of in vivo studies on glucose metabolism (48).



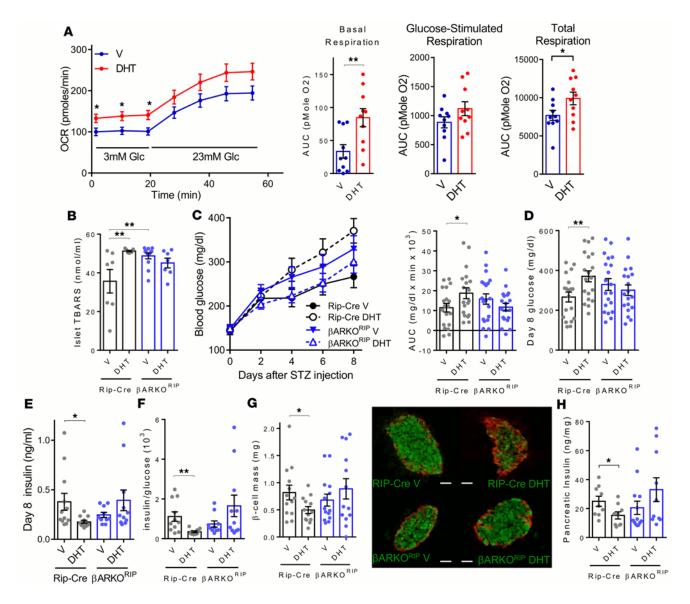


Figure 7. Chronic AR activation in β cells predisposes to oxidative stress. (A) Oxygen consumption rate (OCR) measured from islets isolated from 10 female mice treated with vehicle (V) or DHT (10 nM) (n = 10 wells/condition). (B) Islet TBARS from 13-week-old mice from Figure 2 (n = 7-12). (C-H) Mice were treated s.c. with V or DHT for 1 week followed by induction of diabetes with a single high dose of streptozotocin (STZ) (n = 9-20). (C) Blood glucose measured for 8 days following STZ-induced diabetes and corresponding AUC. (D) Random-fed blood glucose (day 8). (E) Random fed insulin levels (day 8). (F) Insulin/glucose ratio. (G) β Cell mass quantification and IHC images of representative islets. Scale bar: 10 μm. (H) Pancreatic insulin concentration. Values represent the mean ± SEM with dot plots. *P < 0.05, *P < 0.05, *P < 0.01, 1-way ANOVA.

The second finding of this study is that during WD feeding, excess AR activation in β cells is necessary for the development of hyperglycemia. Several lines of evidence support this possibility. First, acute androgen exposure produces insulin hypersecretion in an AR-dependent manner in cultured female mouse and human islets, with an exaggerated first- and second-phase of insulin secretion during perifusion. Second, chronic androgen excess in vivo produces islet oxidative injury via AR in β cells. Finally, and importantly, in wild-type female mice, peripheral androgen infusion (which targets neurons and β cells) produces hyperinsulinemia, insulin resistance, and hyperglycemia, consistent with β cell failure. In contrast, i.c.v. androgen infusion (which selectively targets central neurons but not β cells) produces fasting hyperinsulinemia and insulin resistance without hyperglycemia, consistent with retained β cell function. Thus, during WD feeding, excess AR activation in β cells and neurons causes insulin resistance and hyperglycemia, but excess AR activation selectively in neurons produces insulin resistance without hyperglycemia. This argues for a role of chronic β cell AR activation in the development of hyperglycemia, independent of insulin resistance. How does



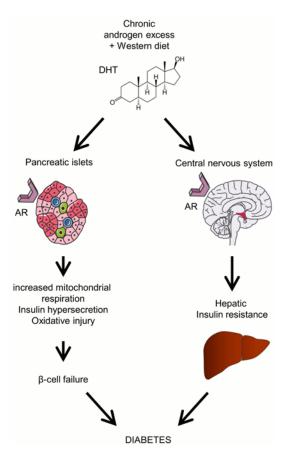


Figure 8. Proposed mechanism for the development of diabetes in female mice with chronic androgen excess. In female mice exposed to a Western diet, chronic androgen excess predisposes to T2D through the combined activation of AR in the hypothalamus, producing hepatic insulin resistance, and activation of AR in β cells, increasing oxygen consumption and producing insulin hypersecretion, oxidative injury, and secondary failure.

excess AR activation in β cells predispose them to failure? Testosterone increases mitochondrial respiration and accordingly promotes islet oxidative damage in vivo. Female mice exposed to testosterone become more susceptible to additional islet oxidative stress induced by either a WD or STZ, which is not observed in β ARKO^{RIP} mice, arguing for the importance of the β cell AR in β cell injury. This suggests that, in the presence of a prior islet metabolic stress, the deleterious effect of chronic AR activation causes β cell hyperfunction, mitochondrial activation, and oxidative stress, which predisposes to β cell demise in female mice.

Interestingly, there is a bidirectional modulation of metabolic homeostasis by testosterone in males and females. In males, physiological concentrations of testosterone are necessary to

maintain insulin sensitivity and β cell function, and testosterone deficiency predisposes to diabetes (24, 25, 32). In contrast, in females, increasing circulating testosterone concentrations to levels observed in males produces insulin resistance and β cell dysfunction and predisposes to diabetes. This sex dimorphism underscores the fundamental differences between male and female biological systems that are rooted in genetics, development, and puberty (48).

There are important clinical ramifications to this study. Women with PCOS exhibit insulin resistance, with testosterone levels positively correlated to the degree of insulin resistance (49). In addition, these women exhibit exaggerated insulin secretion unrelated to insulin resistance and closely associated with androgen excess, suggesting that testosterone produces insulin hypersecretion and β cell dysfunction (28–31). Thus, the predisposition to T2D in women with PCOS and androgen excess could result, as in the case of our mouse model, from the effect of testosterone on brain and β cells.

In summary, chronic androgen excess in female mice fed a WD predisposes to T2D. These effects require the combined effect of a metabolic stress and AR activation in the hypothalamus (producing insulin resistance) and in β cells (producing insulin hypersecretion and injury), which together contribute to secondary β cell failure and hyperglycemia (Figure 8).

Methods

Experimental animals. The βARKO^{RIP} and neuronal specific knockout (NARKO^{-/-}) mice were generated by crossing AR^{lox/+} mice with RIP-Cre^{+/-} and Syn-Cre^{+/-} transgenic mice (Jackson Laboratory), respectively, as described previously (32). AR^{lox/lox} MIP-CreERT mice were generated by crossing AR^{lox/lox} mice with MIP-CreERT1^{Lphi} (MIP-CreERT) transgenic mice (Jackson Laboratory). The inducible βARKO^{MIP-ERT} mice were produced by treating AR^{lox/lox} MIP-CreERT mice with Tam after puberty as described previously (15). βARKO^{MIP} mice were generated by crossing AR^{lox/+} mice with Ins1^{tm1.1(cre)Thor/} J-Cre^{+/-} transgenic (MIP-Cre-transgenic) mice (Jackson Laboratory). Androgen excess was induced via implantation of a 20 mm DHT pellet prepared as described previously (47). KD-mTOR mice were generated as described previously (38). All transgenic mice described above were on a C57BL/6 mouse



background. Brain infusion studies were performed using female C57BL/6 mice from Charles River, while electrophysiological studies and gene expression studies used C57BL/6J mice from Jackson Laboratories. All studies were performed on littermate mice aged 10–16 weeks.

WD and in vivo induction of androgen excess. Mice were weaned on a WD high in saturated fat and simple sugars (sucrose and fructose) (30% AMF diet; 14.9% Kcal protein, 33.2% Kcal carbohydrates, 51.9% Kcal fat; Harlan Teklab). DHT and vehicle were administered to mice after 4 weeks of WD by s.c. pellets (10 mg/pellet, 60-day release; Innovative Research of America) or via i.c.v. injection for an additional 4–5 weeks. To infuse DHT i.c.v., a cannula was implanted into the lateral ventricle (coordinates compared with the bregma L +1 mm, AP –0.2 mm, DV –2 mm) using a stereotaxic frame (Kopf), linked to a DHT-filled osmotic minipump (Alzet) inserted subcutaneously.

Metabolic studies. Metabolic studies were performed 1 week apart following 4 weeks of s.c. or i.c.v. DHT exposure. Blood glucose and insulin concentrations were measured from blood drawn from tail veins using a TRUEResults glucose meter (NitroDiagnostics) and insulin ELISA kit (Linco Research and Millipore), respectively. For i.p. glucose tolerance test (2 g/kg) or i.p. GSIS (3 g/kg), mice were fasted overnight before glucose injection. For i.p. ITT (0.25–0.5 U/kg), mice were fasted for 6 hours and given insulin. Blood glucose and plasma insulin were assessed at the indicated times as described. Pancreas insulin concentration was quantified by acid ethanol extraction as described previously (32).

Inhibition of mTORC-1 in vivo. Mice were weaned on a WD as described above. After 4 weeks of WD, mice were exposed to vehicle, DHT, and DHT plus rapamycin (Santa Cruz Biotechnology) for an additional 4 weeks. Rapamycin was injected i.p. at 5 mg/kg every day.

Induction of experimental diabetes. One week following DHT pellet insertion, 9-week-old mice were exposed to a single i.p. injection of 150 mg/kg STZ (MilliporeSigma) to induce diabetes as described previously (42). Blood glucose was measured every 48 hours following STZ injection. At day 8 following STZ injection, insulin was assessed and pancreases were processed for measurement of pancreatic insulin concentration and β cell mass.

Immunohistochemistry and β cell mass quantification. Pancreases were fixed, paraffin embedded, and sectioned at 5 μ m. Brains were fixed via transcardial perfusion of 10% formalin and sectioned at 20 μ m. Antigen retrieval and section incubation with primary and secondary antibodies were performed as described previously (32). Images were captured at ×20 magnification using a fluorescent microscope (Nikon Eclipse E400 or TissueGnostics Imaging). The β cell mass was quantified by morphometric analysis as described previously (32). All islets were imaged from 1 section per animal (n = 3-4) per group.

Islet isolation and insulin secretion in static incubation. Female mouse islets were isolated following pancreatic duct catheterization, collagenase digestion, and Histopaque gradient separation as described previously (32). Following overnight recovery, islets were hand-picked under a dissection microscope, transferred to a 24-well plate (10 islets/well), and cultured in phenol red–free RPMI (11 mM glucose, 10% charcoal-stripped FBS). Islets from female human donors (provided by the Integrated Islet Distribution Program) were cultured in phenol red–free CMRL 1066 with 10% charcoal-stripped FBS. Islets were treated with DHT (10^{-8} M; Steraloids), Flutamide (10μ M; MilliporeSigma), or vehicle (95% ethanol) for 48 hours. Insulin release from islets was measured by static incubation using 10 islets/well as described previously (32). Insulin was measured by ELISA (Millipore).

Islet perifusion. Batches of 60 mouse islets (C57BL/6 females) were perifused at 37°C, at a flow of 100 µl/min, as described previously (32). Islets were first equilibrated for 60 minutes with Krebs-Ringer buffer (KRB) solution containing 2.8 mM glucose and then stimulated for 30 minutes with KRB solution containing either DHT (10⁻⁸ M) or vehicle (95% ethanol) and 16.7 mM glucose. Samples were collected in a 96-well plate. A perifusion system (Biorep Technologies) was used to determine biphasic response and insulin concentration was determined by ELISA.

Cytoplasmic calcium imaging. Whole islet intracellular Ca²⁺ influx was measured as described previously (32). Briefly, islets were plated on glass-bottom MatTek plates in RPMI media without phenol red and 10% charcoal-stripped FBS (Invitrogen) with either DHT (10⁻⁸ M) or vehicle (95% ethanol). Islets were cultured in a TC incubator (37°C, 5% CO₂) for 48 hours, during which time the media were replaced every 12 hours. Islets were then loaded with fura-2 (2 μmol) for 20 minutes in modified KRB containing 2 mM/l glucose, 119 mM/l NaCl, 2.5 mM/l CaCl2·[(H2O)6], 4.7 mM/l KCl, 10 mM/l HEPES, 1.2 mM/l MgSO₄, 1.2 mM/l KH₂PO₄, pH 7.3. Ca²⁺ imaging was performed using a Nikon microscope (TE2000-U) with illumination from a 300-watt xenon arc lamp (Lamba LS, Sutter instruments) and imaging with a cooled CCD camera (CoolSNAP HQ2 camera, Photometrics). Islets were perifused (2 ml/min) with KRB media at



37°C containing either 2 or 16.7 mM glucose and DHT or vehicle. The relative $[Ca^{2+}]_i$ was determined from the ratio of emitted fura-2 fluorescence with 340 nm or 380 nm excitation (F340/F380).

Dynamic measurement of islet cAMP. Islets were isolated from 8-week-old female Ins1-Cre^{+/-} knockin mice on a C57BL6/J background. These animals have previously been shown to be indistinguishable from Ins1-Cre^{-/-} littermates in terms of growth, glucose tolerance, insulin tolerance, in vivo and in vitro insulin release, and β cell mass (50, 51). The levels of cAMP were monitored in islets infected with adenovirus harboring the FRET probe Epac2 camps, as previously described (52). Live-cell imaging was performed using a Crest X-Light spinning disk head, coupled to a Lumencor SPECTRA X light engine and Nikon $10\times/0.45$ NA objective, controlled using custom scripts in MetaMorph software (Molecular Devices). Excitation was delivered at $\lambda = 430-450$ nm and emission was collected at $\lambda = 460-500$ nm and 520-550 nm for cerulean (Cer) and citrine (Cit), respectively, using a highly sensitive Photometrics Delta Evolve EMCCD camera. FRET response was calculated as the fluorescence ratio of Cer/Cit and normalized as R/R_0 , where R denotes fluorescence at any given time point and R_0 denotes fluorescence at time 0. Islets were maintained in HEPES-bicarbonate buffer, containing 120 mM NaCl, 4.8 mM KCl, 24 mM NaHCO₃, 0.5 mM Na₂HPO₄, 5 mM HEPES, 2.5 mM CaCl₂, 1.2 mM MgCl₂, 3–17 mM D-glucose and Ca²⁺. Vehicle (0.01% EtOH), DHT (10 nM), and exendin 4 (10 nM) were applied at the indicated time points.

TBARS assay. Islet oxidative stress was assessed in islets isolated from mice fed a WD and treated with DHT in vivo by measuring the concentration of TBARS (Zeptometrix) as described previously (42).

Seahorse assay. Islets isolated from 10 female C57/BL6N mice were washed once with assay buffer (made from Agilent Seahorse XF Base Medium supplemented with 3 mM glucose and 1% charcoal-striped FBS). Around 70 islets were transferred to each well of a Seahorse XF24 Islet Capture Microplate (Agilent) and were incubated in assay buffer at 37°C for 60 minutes before being transferred to Seahorse XF Analyzer (Agilent). Islets were maintained in the assay medium throughout the experiment, and the oxygen consumption rate was measured at the basal level (3 mM) as well as at a glucose-stimulated level (20 mM). The results were normalized to the results of the first time point of the vehicle-treated group.

Microdissection of MBH. Male and female (n = 15) mice were sacrificed with CO₂ and decapitated, and brains were dissected into 1-mm coronal sections with the help of an ice-cold mouse brain matrix (ASI-Instruments). From the brain slice covering approximately Bregma -1.0 mm to -2.0 mm, the hypothalamus was detached by a horizontal cut through the mammillothalamic tracts. The lateral hypothalamus was removed by a bilateral cut through the fornix, and the remaining hypothalamic piece was horizontally divided with a cut across the fornix to generate a ventral portion with VMH/ARC enrichment and a dorsal portion with DMH enrichment. The dissected brain tissue was snap frozen and stored at -80°C until RNA extraction.

RNA extraction, cDNA synthesis, and qPCR. RNA from ARC/VMH tissue was extracted by first homogenizing each sample in Trizol (Life Technologies) with a hand-held homogenizer (PELLET PESTLE Cordless Motor, Kimble Chase) and sterile pellet pestles (Kimble Chase), according to the manufacturer's recommendations. To minimize DNA and protein contamination, 100 μl phase separation reagent was added (1-bromo-3-chloropropane, Molecular Research Center Inc.) during the extraction protocol. cDNA was synthesized by reverse transcription PCR (SuperScript VILO, Invitrogen) according to the manufacturer's recommendations.

For quantitative PCR, $3 \mu (1.4 \text{ ng/}\mu)$ template cDNA was used in triplicate to determine gene expression with TaqMan assays (TaqMan Universal Master Mix II, with UNG, Applied Biosystems by Thermo Fisher Scientific) for Gapdh (Mm99999915_g1), Ar (Mm00442688_m1), and Esr1 (Mm00433149_m1), with Gapdh serving as a housekeeping gene. Reactions were pipetted by a robotic liquid handling system (Perkin Elmer Multiprobe II EX) and run on the 7500 HT Fast Real-Time PCR System (Applied Biosystems). Fold induction of gene expression was analyzed with the $\Delta\Delta$ CT method, as recommended for TaqMan assays.

Whole-cell patch-clamp recordings and identification of recorded neurons. Whole-cell patch-clamp recordings were performed at 34°C–37°C from neurons of the VMH or the ARC of 10-week-old female C57BL6J mice as previously described (53). DHT (10 nM) was used to determine the effects of AR activation on neuronal activity. After recordings, slices were fixed in paraformaldehyde (4%) and stored at 4°C. Following a rinse in PBS, slices were incubated in a PBS (10 mM), Texas red Avidin D (1:400) and Triton X-100 (1%) solution at room temperature for 4 hours, mounted, and coverslipped with Vectashield (Vector Laboratories).

Statistics. Results are presented as dot plots and mean ± SEM, as specified in figures. Statistical analyses



were performed using the unpaired 2-tailed Student's *t* test, 1-way ANOVA or Mann-Whitney tests, and Kruskal-Wallis test, when necessary. Cumulative incidence of diabetes was determined by Kaplan-Meier estimates, and statistical analysis of differences was determined by a log-rank test. A *P* value of less than 0.05 was considered statistically significant.

Study approval. Studies were performed with the approval of the Northwestern University, Tulane University (New Orleans, Louisiana, USA), Pennington Biomedical Research Center, and University of Miami Animal Care and Use Committees and in accordance with NIH guidelines.

Author contributions

GN designed and conducted the experiments (Figure 1; Figure 2; Figure 4; Figure 6, A–E, H, and J; Figure 7, B–H; and Supplemental Figures 1–3), analyzed the data, and wrote/reviewed the manuscript. CA (Figure 5 and Supplemental Figures 4 and 5), JJM (Figure 3, A–D; Figure 5; and Supplemental Figures 4 and 5), and WX (Figure 7A) designed experiments, analyzed data, contributed to discussion, and reviewed the manuscript. SL (Figure 7, B–H), AJRM and SMB (Figure 3, F–L), SY and RZ (Figure 3E), NHFF (Figure 6G), and MBR (Figure 6I) conducted experiments and analyzed data. DAJ conducted intracellular Ca²⁺ influx experiments (Figure 6F). VNS and PVK designed experiments, analyzed data, contributed to discussion, and reviewed the manuscript (Figure 7A). DJH, HM, EBM, and AZ designed experiments, contributed to discussion, and reviewed the manuscript. FMJ conceived the idea, designed experiments, analyzed the data, and wrote and reviewed the manuscript. FMJ is the guarantor of this work and, as such, had full access to all the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis.

Acknowledgments

This work was supported by grants from the NIH (R01 DK074970, DK107444, and P50 HD044405) and a Department of Veterans Affairs Merit Review Award (BX003725) to FMJ. CA was supported by an American Diabetes Association Postdoctoral Fellowship (1-16-PDF-004). AZ was supported by NIH grant R01 DK099598, EBM was supported by NIH grants R01 DK073716 and DK084236 and a Veterans Affairs Merit Review Award (IBX002728A), HM was supported by NIH grant R01 DK092587, and PVK was supported by NIH grant R01 NS094834. DJH was supported by a Diabetes UK RD Lawrence Fellowship (12/0004431), a Wellcome Trust Institutional Support Award, and MRC (MR/N00275X/1) and Diabetes UK (17/0005681) project grants as well as by the European Research Council under the European Union's Horizon 2020 research and innovation programme (starting grant 715884). Part of this work was presented during the American Diabetes Association meeting in New Orleans, Louisiana, USA, on June 10–14, 2016.

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insight.jci.org https://doi.org/10.1172/jci.insight.98607