

Impairment of the G β y-SNAP25 brake on exocytosis enhances insulin action, protects against diet-induced obesity, and promotes adipocyte browning

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The Gβγ complex inhibits vesicle exocytosis by two mechanisms: inhibiting calcium entry by binding to voltage gated calcium channels, and binding to SNAP25 in the SNAP Receptor (SNARE) complex. To de-convolute the role of each of these mechanisms *in vivo*, we have made a mouse with the second mechanism disabled. The SNAP25 $\Delta 3$ mutation renders the SNARE complex deficient in binding to Gβγ and was used to investigate the importance of the Gβγ-SNAP25 interaction in glucose stimulated insulin secretion (GSIS) and global metabolic homeostasis. GSIS and α_{2A} adrenergic receptor-mediated inhibition of GSIS were not altered in SNAP25 $\Delta 3/\Delta 3$ mice. Nevertheless, SNAP25 $\Delta 3/\Delta 3$ mice exhibited a marked improvement in insulin sensitivity and were resistant to weight gain when challenged with a high fat diet (HFD). Reduced food consumption in the early stages of HFD feeding were partly responsible for the inability of SNAP25 $\Delta 3/\Delta 3$ mice to gain weight on HFD. Additionally, improved insulin-mediated glucose uptake into white adipose tissue and increased ‘browning’ were observed in SNAP25 $\Delta 3/\Delta 3$ mice, which is consistent with an impaired ability to retain energy stores. These phenotypic changes in SNAP25 $\Delta 3/\Delta 3$ mice are all metabolically protective, indicating that pharmacological targeting of the Gβγ-SNAP25 interaction may have a metabolic benefit.

Introduction:

G-protein coupled receptors (GPCRs) are canonically known to mediate downstream signaling events through the activation of heterotrimeric G-proteins. While the classical $G\alpha$ -mediated signaling events are well known, and many represent important pharmacological targets (1), signaling via the $G\beta\gamma$ -subunits have received less attention and the therapeutic targeting of these signaling pathways has not been as widely exploited (2). One function of $G\beta\gamma$ is to inhibit exocytosis by two mechanisms, modulation of calcium entry, and direct binding to the exocytotic fusion complex (3). We have shown that $G\beta\gamma$ binding to the SNARE complex is mainly through the last three amino acids of SNAP25 (Soluble *N*-ethylmaleimide-sensitive factor Attachment Protein), in the SNAP Receptor (SNARE) complex (4). In order to specifically study the contribution of exocytosis at the exocytotic fusion step *in vivo*, we developed an allele of SNAP25 which lacks these last three amino acids, which we have called SNAP25 $\Delta 3$ (5). The ability of SNAP25 $\Delta 3$ to form SNARE complexes that undergo calcium-synaptotagmin mediated zipping and regulate exocytosis is identical to that of SNAP25, but its ability to interact with $G\beta\gamma$, and GPCR-mediated inhibitory effects on exocytosis, is ablated. The SNAP25 $\Delta 3/\Delta 3$ mouse has demonstrated the importance of the $G\beta\gamma$ -SNARE pathway in a number of neurological processes, including stress and pain processing, as well as long-term potentiation (6) and spatial memory (5). We hypothesized that the SNAP25 $\Delta 3/\Delta 3$ mouse may have an altered metabolic phenotype because neurological processes use SNARE-dependent signals to regulate a number of metabolically important processes, such as feeding behavior, energy balance, and thermoregulation (7-12), and the secretion of important metabolic hormones, such as insulin (13-15).

Metabolic diseases, such as diabetes and obesity, are estimated to have an annual global economic impact in the trillions of (US) dollars (16-20). The therapeutic value of the recently developed glitides and gliptins highlights the benefits of pharmacological agents that augment glucose stimulated insulin secretion (GSIS) (21). We hypothesized that the SNAP25 $\Delta 3$ mutation would similarly enhance GSIS because previous studies have indicated that norepinephrine inhibits insulin exocytosis via the α_{2A} adrenergic receptor (AR) through $G\beta\gamma$ inhibition of SNARE in the pancreatic β -cell (22). While the recent advances that have led to these anti-diabetic agents, a lack of effective therapeutics for obesity persists. Obesity is co-morbid with insulin resistance and type II diabetes in addition to many other diseases such as cancer, asthma, mental disorders, nonalcoholic fatty liver disease, cardiovascular diseases, and heart failure (23, 24). This lack of obesity therapeutics is a serious

deficiency in the currently available pharmacological cornucopia, as recent studies have reported that about 40% of Americans are clinically obese and the prevalence of obesity worldwide is rising (25).

We took advantage of the $\beta\gamma$ -SNAP25 interaction and its ability to serve as a brake on secretion to determine if limiting this physiologic inhibitory pathway could affect insulin secretion and/or weight gain in response to a high fat diet. Though SNAP25 is a well-known component of the insulin exocytosis machinery (13-15), we observed no alteration in GSIS from SNAP25 $\Delta 3/\Delta 3$ mice. However, adiposity and glucose homeostasis were markedly improved. These studies went on to probe the basis for this metabolic protection by examining energy intake and energy expenditure, and potential cellular targets that contribute to this protection. Though SNAP25 $\Delta 3/\Delta 3$ mice did not exhibit enhanced GSIS, their resistance to diet induced obesity and improved glucose homeostasis suggest that targeting of the $G\beta\gamma$ -SNAP25 interaction may have therapeutic benefits for metabolic diseases.

Results:

In order to test our hypothesis that removal of the $G\beta\gamma$ inhibition of SNAP25-mediated exocytosis would improve insulin secretion, we measured circulating glucose and insulin levels in SNAP25 $\Delta 3$ mice during a glucose tolerance test (GTT). In our first study, 14 week-old, chow fed male mice (weighing 29.3 \pm 0.710 g vs. 28.9 \pm 0.749 g, SNAP25 $^{+/+}$ vs. SNAP25 $\Delta 3/\Delta 3$ respectively, $P = 0.695$) received an IP injection of 2 mg/kg glucose. The SNAP25 $\Delta 3/\Delta 3$ mice had a normal glucose tolerance compared to SNAP25 $^{+/+}$ mice (**Fig. 1A**). Plasma insulin was measured during the IP-GTT. We found that insulin levels in SNAP25 $\Delta 3/\Delta 3$ mice showed a significantly reduced insulin response at 15 and 30 minutes compared to SNAP25 $^{+/+}$ mice (**Fig. 1B**). In a separate cohort of chow fed male mice (28.2 \pm 0.753 g vs. 27.3 \pm 0.596 g, SNAP25 $^{+/+}$ vs. SNAP25 $\Delta 3/\Delta 3$ respectively, $P = 0.375$) we performed an oral glucose tolerance test because we hypothesized that the incretin effect, which we bypassed with the IP injection of glucose, might be important for the augmentation of insulin secretion in the SNAP25 $\Delta 3$ mouse. Oral glucose tolerance was again not different between genotypes (**Fig. 1C**). However as was observed with the IP route of delivery plasma insulin levels were lower in SNAP25 $\Delta 3/\Delta 3$ mice after the glucose gavage (**Fig. 1D**). Given the lower levels of insulin in SNAP25 $\Delta 3/\Delta 3$ mice, but a normal glucose tolerance, it would appear that insulin sensitivity is improved in SNAP25 $\Delta 3/\Delta 3$.

In order to separate the insulin secretion phenotype from the insulin sensitivity phenotype, we measured α_2A AR dependent inhibition of GSIS from SNAP25 $\Delta 3/\Delta 3$ islets *ex vivo*. It has been previously reported in studies

using INS-1 832/13 rat insulinoma cells that inhibition of insulin exocytosis by norepinephrine- α_{2A} AR signaling is mediated by the G $\beta\gamma$ -SNAP25 interaction (26). Therefore, we also sought to determine whether an α_{2A} AR agonist would similarly inhibit insulin secretion in intact islets from SNAP25 $\Delta 3/\Delta 3$ mice due to the lack of G $\beta\gamma$ -SNAP25 interaction. Islets from SNAP25 $\Delta 3/\Delta 3$ mice did not have a statistically different alteration in GSIS (**Fig. 2A,B**). The α_{2A} AR selective agonist, brimonidine (Br), inhibited GSIS in a concentration-dependent manner in islets regardless of SNAP25 genotype (**Fig. 2C**). Insulin content of the islets was also not affected by SNAP25 genotype (**Fig. 2D**). This indicates that in intact mouse islets, α_{2A} AR-mediated signaling is able to inhibit GSIS via a mechanism that is not dependent upon the G $\beta\gamma$ -SNAP25 interaction.

Our prior studies suggested that SNAP25 $\Delta 3/\Delta 3$ had an altered behavioral response to acute physiological stressors (5, 6). We performed calorimetry studies where we assessed energy expenditure and feeding behavior in chow fed male SNAP25 $^{+/+}$ and SNAP25 $\Delta 3/\Delta 3$ mice at standard housing temperature (22 °C). We also exposed them to an acute cold challenge (6 °C) to assess whether they could mount a physiologic response to a cold stress and to see if they exhibit a normal increase in energy expenditure and food intake. At normal housing temperatures SNAP25 $\Delta 3/\Delta 3$ mice had similar rates of energy expenditure (**Fig. 3A**) and food intake (**Fig. 3B**), with similar circadian patterns for both variables peaking at the onset of the dark cycle. In response to a decrease in environmental temperature both SNAP25 $\Delta 3/\Delta 3$ mice and their SNAP25 $^{+/+}$ littermates mounted a robust increase in energy expenditure. This was also accompanied by an increased duration of feeding activity in SNAP25 $\Delta 3/\Delta 3$ mice (**Fig. 3E**). Cardiovascular parameters were also measured, and no differences were observed in the SNAP25 $\Delta 3/\Delta 3$ mice (**Supplementary Fig. S1**). Thus, on a chow diet SNAP25 $\Delta 3/\Delta 3$ mice display subtle changes in feeding behavior that overall does not impact energy balance, and the physiologic response to cold is intact.

Despite having no apparent difference in islet function, SNAP25 $\Delta 3/\Delta 3$ mice did appear to have improved insulin sensitivity in the absence of a change in body weight or energy expenditure on a chow diet. It was unclear if the improvement in insulin action would persist in the presence of a high fat diet. Therefore, we placed 8-week-old male and female SNAP25 $^{+/+}$ and SNAP25 $\Delta 3/\Delta 3$ mice on a HFD (Bio-serv diet, 60% kcal from fat) for 8 weeks. In both male and female mice, SNAP25 $^{+/+}$ mice rapidly gained weight on the HFD, whereas SNAP25 $\Delta 3/\Delta 3$ mice were resistant to HFD-induced weight gain (**Fig. 4A,E**). Body composition analyses over the course of the study revealed that the reduced weight gain in SNAP25 $\Delta 3/\Delta 3$ mice was due to reduced adiposity (**Fig. 4B,C,F,G**). This was confirmed in postmortem analyses of these mice; gonadal fat (epididymal WAT (eWAT) in male and

periovarian (poWAT) in female), inguinal WAT (iWAT), retroperitoneal WAT (rWAT), and interscapular BAT (iBAT) were all reduced in SNAP25 $\Delta 3/\Delta 3$ mice (**Fig. 4D,H**). Liver weight was reduced only in male SNAP25 $\Delta 3/\Delta 3$. We also found no significant difference in fecal triglyceride content, indicating that the reduction in body weight in SNAP25 $\Delta 3/\Delta 3$ mice was not due to a decrease in lipid absorption by the gut (**Supplementary Fig. S2**).

Having observed a profound difference in weight gain and adiposity in SNAP25 $\Delta 3/\Delta 3$ mice when challenged with a HFD, we hypothesized that glucose homeostasis would also be improved. Indeed, HFD fed SNAP25 $\Delta 3/\Delta 3$ male mice showed a significant improvement in both the IP-GTT (**Fig. 5A**) and ITT (**Fig. 5B**) compared with SNAP25 $^{+/+}$ mice. SNAP25 $\Delta 3/\Delta 3$ male mice had lower fasting glucose levels (**Fig. 5C**) congruent with their lower IP-GTT and ITT curves. Nevertheless, plasma insulin was also significantly lower (**Fig. 5D**). Together, these data indicate that there is a bona fide improvement in the insulin sensitivity of the SNAP25 $\Delta 3/\Delta 3$ male mice. Female mice did not gain as much weight in response to HFD as their male counterparts. This likely explains why female SNAP25 $\Delta 3/\Delta 3$ mice had a similar glucose (**Fig. 5E**) and insulin tolerance (**Fig. 5F**), as well as blood glucose (**Fig. 5G**) and plasma insulin (**Fig. 5H**), as their SNAP25 $^{+/+}$ littermates. We then decided to investigate feeding behavior in the SNAP25 $\Delta 3/\Delta 3$ mouse more thoroughly as the difference in weight gain occurred within the first few weeks on diet. We examined the weekly food intake in mice on HFD for 5 weeks (**Fig. 6**). During the first two weeks of HFD, SNAP25 $^{+/+}$ mice exhibit a robust increase in consumption (which is commonly observed in wild-type mice). However, this was not found in SNAP25 $\Delta 3/\Delta 3$ mice. By the third week, the hyperphagia in the SNAP25 $^{+/+}$ mice had diminished, and by week 4 the two genotypes began to consume similar amounts of HFD.

These data indicate that SNAP25 $\Delta 3/\Delta 3$ mice have a significantly altered food consumption phenotype, which is likely a major contributor towards the reduced weight gain and improved glucose homeostasis. However, the contribution of specific tissues to the improved glucose homeostasis of SNAP25 $\Delta 3/\Delta 3$ mice was unknown. To investigate this we performed hyperinsulinemic euglycemic clamps in chronically catheterized conscious male mice that had been consuming the HFD for 8 weeks (27 \pm 2 vs 39 \pm g body weight; SNAP25 $\Delta 3/\Delta 3$ vs. SNAP25 $^{+/+}$) to determine if insulin suppression of endogenous (i.e. hepatic) glucose production and/or stimulation of peripheral glucose uptake were altered. The glucose infusion rate required to maintain euglycemia was markedly increased in SNAP25 $\Delta 3/\Delta 3$ despite lower clamp insulin concentration (**Fig. 7A**), with arterial blood glucose maintained at euglycemia in both groups throughout the clamp procedure (**Fig. 7A'**). Arterial insulin levels were lower in the

basal period in SNAP25 $\Delta 3/\Delta 3$ (2.5 ± 0.2 vs. 5.8 ± 1.1 ng/ml) and increased during clamp period (5.1 ± 0.8 vs. 9.7 ± 1.7 ng/ml) during insulin infusion. Using [^3H]glucose we determined that the increase in glucose requirements was because of increase in whole body glucose uptake (**Fig. 7B**). Basal endogenous glucose production as well as insulin suppression of endogenous glucose production were comparable (**Fig. 7C**). To determine which tissues contributed to the increase in glucose disposal we assessed glucose uptake in multiple tissues during the clamp using [^{14}C]2-deoxy-D-glucose (2-DG). 2DG-determined glucose uptake was markedly increased in SNAP25 $\Delta 3/\Delta 3$ mice in multiple skeletal muscles (gastrocnemius, soleus, and vastus lateralis) and white adipose tissue depots (perigonadal and inguinal) (**Fig. 7D**) as well as in heart. Glucose uptake was unchanged in the interscapular brown adipose tissue (iBAT) and brain. Together, these data suggest that multiple tissues of SNAP25 $\Delta 3/\Delta 3$ mice take up glucose more readily in response to insulin than SNAP25 $^{+/+}$ mice.

The marked increase in glucose uptake in subcutaneous AT indicated that there may be additional changes to this tissue. We first examined this tissue histologically. The iWAT of SNAP25 $\Delta 3/\Delta 3$ mice had much smaller adipocytes (**Fig. 8A**). This was especially noticeable in iWAT of female mice. There were also significantly more small multilocular adipocytes that exhibited strong staining for uncoupling protein 1 (UCP1): the signature marker of brown and beige adipocytes. The morphology of the iBAT from SNAP25 $\Delta 3/\Delta 3$ mice also showed smaller lipid droplets (**Fig. 8B**). UCP1 staining appeared stronger in the SNAP25 $\Delta 3/\Delta 3$ iBAT, but this could be due to the higher density of the adipocytes per visual field. This indicates that SNAP25 $\Delta 3/\Delta 3$ mice have improved browning of their subcutaneous WAT, which is likely contributing to the increased glucose uptake in their subcutaneous AT.

We then began to investigate the molecular changes that occurred in the adipose tissue of the SNAP25 $\Delta 3/\Delta 3$ mice. Norepinephrine secreted from sympathetic neurons in response to cold is the classical mechanism by which brown adipose tissue (BAT) thermogenesis is triggered and white adipocytes are pushed towards a more brown-like phenotype (7, 27). Norepinephrine- β AR signaling drives the activation of PKA and thereby the BAT signature protein, UCP1. PKA signaling appeared to be increased as indicated by the greater intensity of bands detected in Western blots using a PKA substrate antibody (**Fig. 9**) in both the iWAT and gWAT of SNAP25 $\Delta 3/\Delta 3$ mice. Note that in SNAP25 $^{+/+}$ mice the HFD tended to suppress many of these phosphoproteins, but, importantly, they were largely preserved in the SNAP25 $\Delta 3/\Delta 3$ mice, particularly in the iWAT. This was associated with an increase in mitochondrial proteins and markers of lipolytic including adipose triglyceride lipase

(ATGL). HFD feeding also reduced these markers, most prominently in SNAP25^{+/+} mice, while in HFD fed SNAP25^{Δ3/Δ3} mice these markers were similar to those of mice fed a chow diet. These changes are consistent with a scenario of increased norepinephrine release from the neurons that innervate the iWAT and gWAT depots. In the iBAT there were few changes observed in response to either genotype or diet.

Discussion

Hormones that signal through GPCRs have long been known to be important modulators of signals and cell types that impact type 2 diabetes (28), since they are regulators of GSIS (29-31), feeding behavior (32), gastrointestinal biology (33), and adipocyte lipolysis and thermogenesis (34). This signaling has most often been attributed to receptor-driven activation of Gα subunits engaging their intracellular effectors. Here, we demonstrate that signaling through the Gβγ subunits that regulate exocytosis by interacting with the SNARE complex have important physiological consequences on feeding behavior and adipocyte thermogenesis. By utilizing a mouse model that lacks the last three amino acids of SNAP25, we have been able to investigate the role of Gβγ-inhibition of SNAP25-mediated vesicle release on glucose handling and energy balance. While the mechanism in the various cell types that may be responsible for this will require further in-depth investigation, these finding may have therapeutic importance for metabolic diseases such as obesity and type 2 diabetes, since targeting of Gβγ-mediated signaling has yet to be exploited for these diseases.

We predicted that the SNAP25^{Δ3/Δ3} mice would exhibit improved GSIS because Gβγ is a regulator of insulin exocytosis (22, 35). Instead, we found that there is no difference in GSIS in islets from SNAP25^{Δ3/Δ3} mice. Furthermore, α₂AR-mediated inhibition of GSIS was unaltered. In addition to these *ex vivo* studies, we examined GSIS *in vivo*. Contrary to our expectations, *in vivo* GSIS actually was reduced. This led us to hypothesize that SNAP25^{Δ3/Δ3} mice have improved insulin-dependent glucose uptake so we conducted a series of studies to determine if these mice had alterations in their energy homeostasis.

In our prior work we observed that SNAP25^{Δ3/Δ3} have behavioral deficits that impact learning, memory, and the response to stressors (5, 6). We asked whether similar deficits could impact energy homeostasis. Using the Promethion comprehensive calorimetry system we monitored energy expenditure and feeding behavior in chow fed SNAP25^{Δ3/Δ3} and their SNAP25^{+/+} litter mates. We found that overall energy expenditure and food intake was unremarkable. We then metabolically challenged these mice with cold exposure. They were able to mount a robust increase in exergy expenditure and food intake in response to this metabolic stressor. Thus,

while SNAP25^{Δ3/Δ3} have some behavioral deficits it does not seem to negatively impact maintenance of metabolism.

In the present studies, we found that SNAP25^{Δ3/Δ3} mice were remarkably resistant to weight gain when challenged with a HFD. We first asked if this phenotype was driven by alterations in energy intake. In SNAP25^{Δ3/Δ3} mice, the initial hyperphagia seen in wild type mice to a HFD was absent. After 2 weeks the hyperphagia is less robust in the SNAP25^{+/+} mice and the food of intake gradually matches of SNAP25^{Δ3/Δ3} mice, The lack of the hyperphagia may reflect a neophobia of the novel food, disliking it only because it is different than what they are used to eating. This delay in augmenting caloric intake sets these mice up on a different weight gain trajectory that persists throughout the study. There was no alteration in gut absorption of dietary triglycerides indicating that feeding behavior was the sole alteration in energy intake.

SNAP25^{Δ3/Δ3} mice have marked improvement in glucose homeostasis. Our initial hypothesis was that this would be present because an improvement in insulin secretion as SNAP25 has been show to transduce signals that inhibit insulin secretion (13-15). In fact, we found that SNAP25^{Δ3/Δ3} mice have normal β-cell function and the improvement in glucose homeostasis is because of an improvement in insulin action. This improvement was observed on a chow diet, but this benefit was most evident when animals were presented with a HFD. The protection from weight gain on a HFD synergized with the underlying metabolic protection to lead to robust metabolic protection. The sites of this improvement was evident from clamp studies. We saw marked improvement in glucose uptake in skeletal muscle and white adipose tissue. The improvement in inguinal white adipose tissue was profound (a 4-fold increase).

If this temporary reduction in caloric intake was the sole driver of the lean phenotype, then we would expect the body weights to remain parallel. Instead, they continue to diverge suggesting that the lean phenotype of SNAP25^{Δ3/Δ3} mice may be more complicated. Studies utilizing a radiolabeled glucose tracer showed that SNAP25^{Δ3/Δ3} mice have a marked increased glucose uptake in adipose tissue. Upon closer evaluation of the adipose tissue, it was found that SNAP25^{Δ3/Δ3} mice have increased browning of their white adipose tissue. This browning corresponded with increased PKA signaling in SNAP25^{Δ3/Δ3} adipose tissue. We hypothesize that increased sympathetic tone from the SNAP25^{Δ3/Δ3} neurons innervating the WAT is driving the enhanced PKA signaling and thermogenic marker expression, though we cannot rule out the possibility of other hormones contributing to this effect. It has recently been appreciated that adult humans possess significant amounts of

brown adipocytes that are rich in mitochondria and UCP1 (36-39), and the activation of adipocyte thermogenesis is now seen as an attractive therapeutic target for obesity and metabolic disease. In humans adipose tissue, the β_1 AR is more important than the β_3 AR for the regulation of metabolism (40). Inhibiting the G $\beta\gamma$ -SNAP25 interaction to stimulate endogenous catecholamine release in the adipose tissue, instead of pharmacologically targeting the β ARs directly, may be a way to circumvent the side effects of β_1 AR agonists.

These findings highlight the importance of the G $\beta\gamma$ -SNAP25 interaction in the regulation of body weight and glycemic control. Though we did not observe the augmentation of GSIS that we predicted, we did find that SNAP25 $\Delta 3/\Delta 3$ mice have beneficial metabolic effects in adipose tissue, skeletal muscle, and feeding behavior. Importantly, there are also no apparent cardiovascular effects in these mice which could have limited the therapeutic relevance of targeting this pathway. Nevertheless, these studies are limited in their ability to determine how these beneficial metabolic effects occur. Tissue specific gene knock-in of SNAP25 $\Delta 3$ will be invaluable to further understand the processes behind these metabolic changes. While there are many pathways that may be targeted to increase adipose tissue metabolism and energy expenditure, these signaling pathways are also heavily involved in various physiological processes in other tissues which constrains their clinical utility (34). Pharmacological targeting of the G $\beta\gamma$ -SNAP25 interaction may be a therapeutic alternative to directly targeting the GPCRs because SNAP25 $\Delta 3/\Delta 3$ mice have improvements in food intake, activity, and adipose browning, all without significant side effects.

Materials and Methods

Animal procedures

Mice used for these experiments were generated from heterozygous breeding of SNAP25 $^{+/\Delta 3}$ mice on a C57BL/6 background. The strategy for the generation of these mice was previously described (5). Mice were maintained ad libitum on chow, Laboratory Rodent Diet 5001 (LabDiet), unless otherwise stated. Mice were maintained on a 12-hour light, 12-hour dark cycle and housed with 3–5 animals per cage. All procedures were approved by the Institutional Animal Care and Use Committee at Vanderbilt University

HFD feeding

SNAP25 $^{+/+}$ and SNAP25 $\Delta 3/\Delta 3$ mice were placed on a HFD (60% calories from fat and 36% fat by weight, 3282; Bio-Serv). Throughout the course of the study, body composition of live mice was measured by pulsed

NMR with Minispec Model mq7.5 (Bruker Instruments). Mice were euthanized by isoflurane overdose at the end of the study for collection of tissues.

Glucose tolerance tests (GTTs) and insulin tolerance tests (ITTs)

For all tolerance tests, mice were fasted for 5 hours, and fasting blood glucose was measured from a drop of tail vein blood with a Bayer CONTOUR glucometer and glucose test strips at the indicated time points. For GTTs in figure 1, separate cohorts of mice were given either an intraperitoneal (IP) injection or oral gavage of 2.0 g/kg glucose. Insulin content was analyzed in duplicate by RIA by Vanderbilt University Hormone Assay and Analytical Services Core.

For tolerance tests in figure 5, mice were given IP injections of 1.0 g/kg dextrose (Agri Laboratories, Ltd.) in 0.9% saline (Hospira, Inc.) at 22 weeks of age and an IP injection of 0.5 U/kg insulin (Humulin R, Lilly USA) in 0.9% saline (Hospira, Inc.) at 23 weeks of age. Mice were fasted for five hours and blood glucose was collected from the tail vein after which the mice were subsequently euthanized by CO₂. Blood was collected into EDTA coated tubes via cardiac puncture and plasma insulin was measured by ELISA (Mercodia).

Mouse islet perfusion

Pancreatic islets were isolated and perfusion assays were performed on fresh islets at the Vanderbilt Islet Procurement and Analysis Core as previously described (PMID: 22167125). Islet preparations were equilibrated and stable baseline response established at 5.6 mmol/L glucose, and insulin secretion was stimulated with 16.7 mmol/L glucose.

Energy balance

Energy balance studies were performed on chow-fed, male mice in a Promethion system (Sable Systems International) by the Vanderbilt Mouse Metabolic Phenotyping Center.

Cardiovascular imaging

Cardiac parameters, as measured by parasternal M-mode echocardiography¹⁰³ were collected. Mice had their chest fur shaved and a topical depilatory agent was used to remove any remaining fur in the area. Ultrasound coupling gel heated to 34°C was applied to the chest area and a linear array transducer (18-23 MHz) was positioned to obtain two-dimensional B-mode parasternal long and short axis views at the mid-ventricular level (Vevo 2100, VisualSonics). One-dimensional M-mode images were obtained for measurement in the short axis view to obtain cardiac wall thickness and chamber dimensions. Left ventricular (LV) chamber size and wall

thickness were measured off-line in the M-mode from at least three consecutive beats and averaged. LV wall thickness: intraventricular septum (IVS) and posterior wall (PW) at systole and diastole; and LV internal dimensions (LVID) during systole and diastole were measured.

Hyperinsulinemic euglycemic clamps

Clamp studies were done chronically catheterized (carotid artery and jugular vein) conscious mice. Catheters were inserted 4-5 days prior to a study by the Metabolic Pathophysiology Core (41-43). In chronically catheterized mice [$3\text{-}^3\text{H}$] glucose was used to measure whole body basal and clamp glucose flux. A $4\text{ mU}\cdot\text{kg}^{-1}\cdot\text{min}^{-1}$ insulin infusion will be initiated to increase insulin to a physiologic range. Red blood cells from a donor animal are infused at a constant rate to replace blood taken during study. Basal and clamp period blood samples for glucose, insulin and tracer. At the end of the clamp period, multiple tissues were collected to measure the accumulation of $^{14}\text{C}2\text{DG}$. Using tracer methods [$3\text{-}^3\text{H}$]glucose and ^{14}CDG during the clamp, we assessed tissue glucose uptake (44) and whole body (and hepatic) glucose flux (45, 46).

Histology

Adipose tissues were fixed with were fixed in 10% formaldehyde overnight and subsequently stored in 70% ethanol then processed routinely, embedded, sectioned and stained with hematoxylin and eosin (H&E) or immunohistochemical stained for UCP1 (ab10983, Abcam). Histology was performed by the Vanderbilt Translational Pathology Shared Resource. Slides were imaged at $20\times$ with a Leica SCN400 Slide Scanner by the Vanderbilt Digital Histology Shared Resource.

Western blotting

Adipose tissues were lysed and sonicated in a buffer containing 25 mM HEPES (pH 7.4), 150 mM NaCl, 5 mM EDTA, 5 mM EGTA, 5 mM glycerophosphate, 0.9% Triton X-100, 0.1% IGEPAL, 5 mM sodium pyrophosphate, 10% glycerol, plus 1 tablet each of cOmplete™ protease inhibitor cocktail (04693124001, Roche) and PhoSTOP phosphatase inhibitors (04906845001, Roche) per 10 ml of lysis buffer. The lysates of 40 μg total protein were resolved in 10% Tris-glycine gels, transferred to nitrocellulose membranes, which were incubated overnight at 4°C with specific primary antibodies, followed by secondary antibody incubations for 1 h at room temperature. Image acquisition was performed on Bio-Rad digital ChemiDoc MP with IR.

The following Antibodies were purchased from Cell Signaling Technology (Danvers, MA): anti-ACLY (dilution 1:1000, #4332), anti-aP2 (1:1000, #2120), Anti-ATGL (1:1000, #2439), anti-CoxIV (1:1000, #4844), anti-

HSL (1:1000, #4107), PKA substrate (1:1000, #9624), and β -actin (1:2000, #4967). Total OXPHOS Rodent WB antibody cocktail (1:1000, MS604-300) was from Abcam (Cambridge, MA). Secondary antibodies anti-rabbit (1:20000, A3687) and anti-mouse (1:20000, A3562) were obtained from MilliporeSigma (Burlington, MA).

Statistics

Data are mean \pm SEM, using GraphPad Prism version 8.3.1 for Windows 64-bit (GraphPad Software). Analysis comparing genotype only were performed with an unpaired t-test. Analysis comparing genotype along with time, light/dark cycle, or +/- clamp, were performed with a two-way ANOVA or a mixed-effects model if data were missing. Multiple comparison tests were performed with the Bonferroni correction for SNAP25 genotype only and are indicated on figures by asterisks corresponding to *, $P < .05$; **, $P < .01$; ***, $P < .001$.

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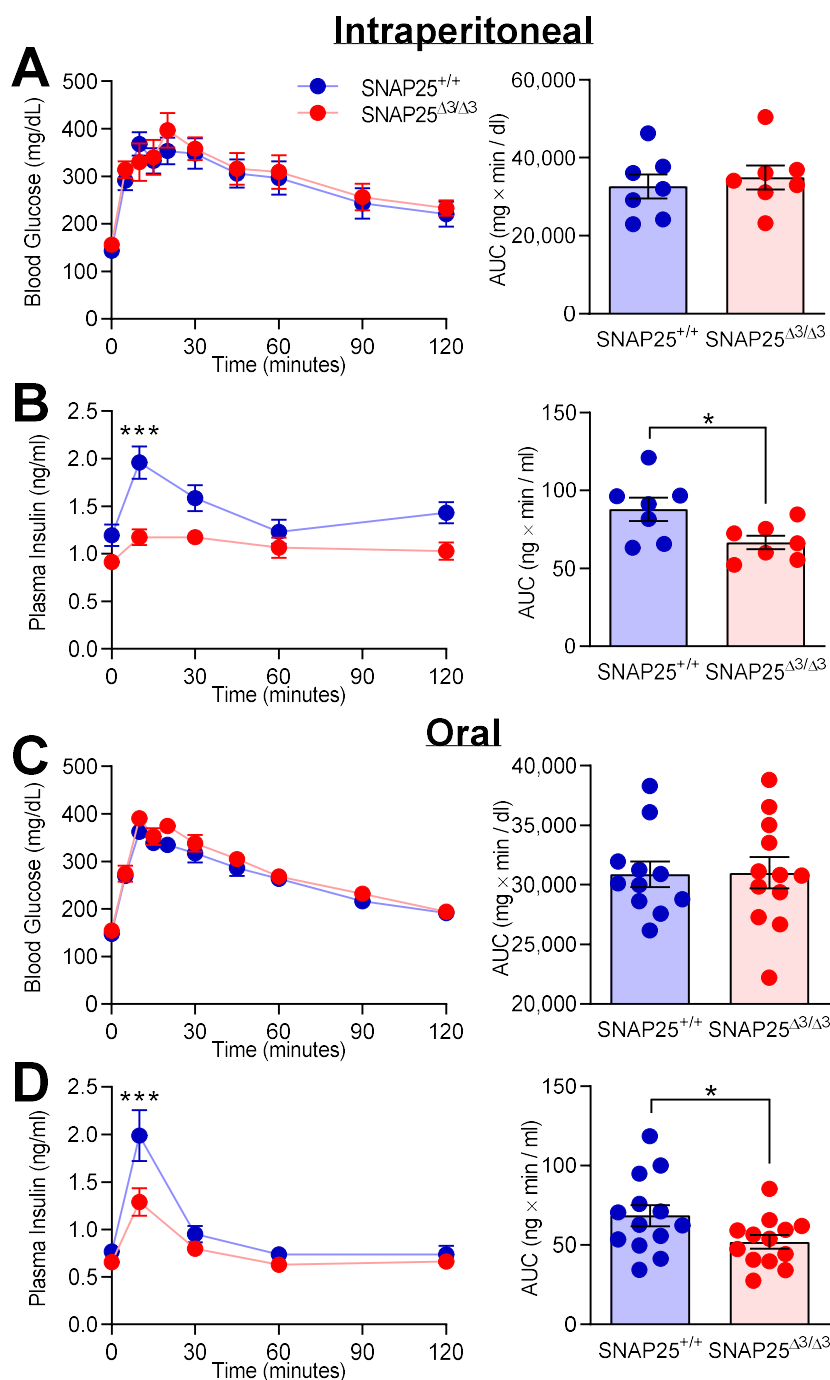


Figure 1: SNAP25^{Δ3/Δ3} mice have normal glucose tolerance but secrete less insulin

A. Glucose tolerance was assessed by an IP-GTT in chow-fed male SNAP25^{+/+} and SNAP25^{Δ3/Δ3} mice at 14 weeks of age. **B.** Insulin secretion during the IP-GTT revealed that SNAP25^{Δ3/Δ3} mice have reduced plasma insulin in response to glucose challenge ($P = 0.0176$ for the effect of genotype, $P = 0.0031$ for the effect of interaction). For A and B, $n = 8$ SNAP25^{+/+} and $n = 8$ SNAP25^{Δ3/Δ3}. **C.** Glucose tolerance was assessed by an oral GTT in chow-fed male SNAP25^{+/+} and SNAP25^{Δ3/Δ3} mice at 15 weeks of age. **D.** Insulin secretion during the oral GTT revealed that SNAP25^{Δ3/Δ3} mice have reduced plasma insulin in response to glucose challenge ($P = 0.073$ for the effect of genotype, $P = 0.0009$ for the effect of interaction). For C and D, $n = 13$ SNAP25^{+/+} and $n = 13$ SNAP25^{Δ3/Δ3}. Values are expressed as mean \pm SEM.

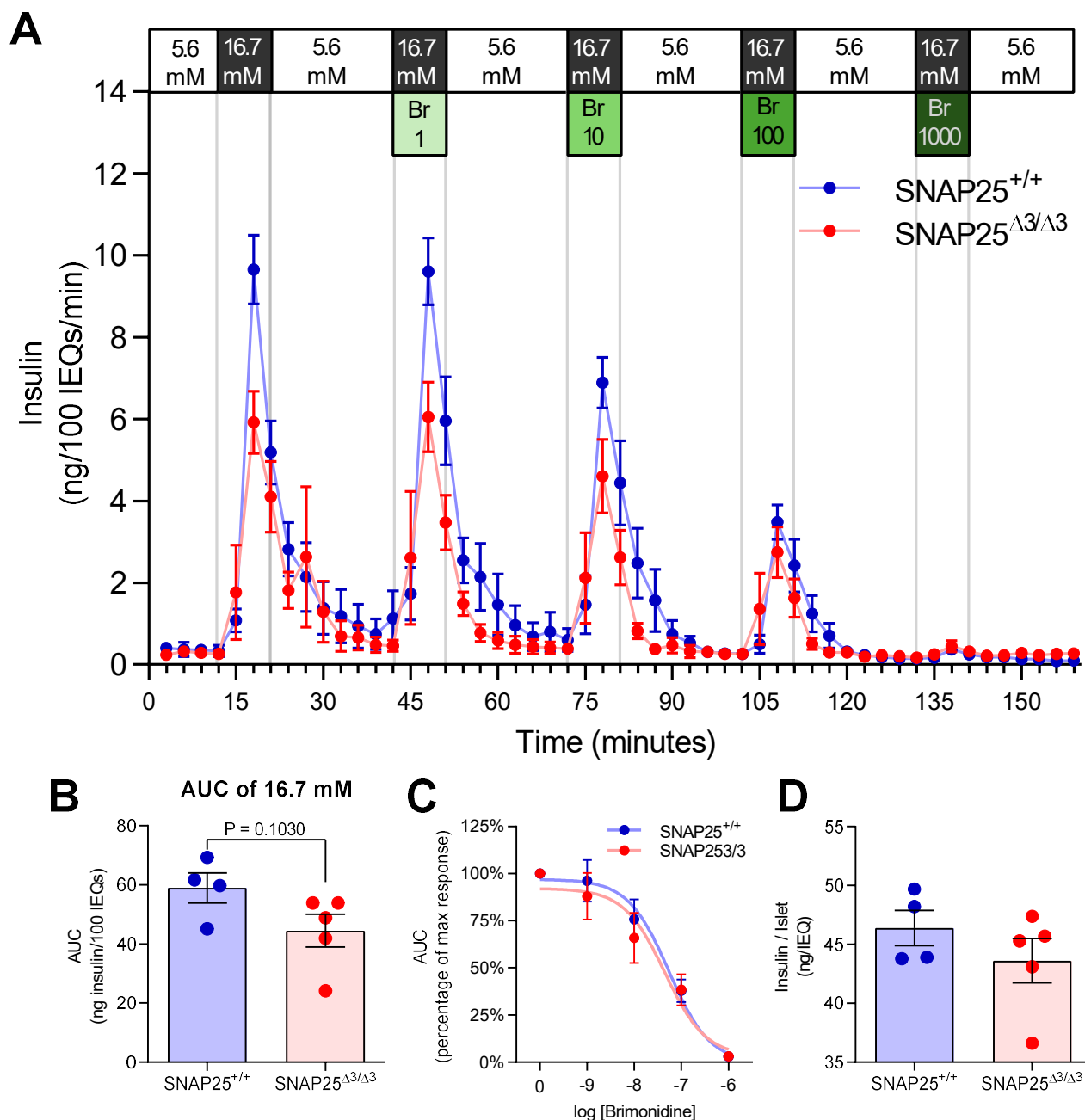


Figure 2. Ablation of the Gβγ-SNAP25 interaction does not alter glucose stimulated insulin secretion from islets *ex vivo*

A. Perfusion of islets from 12-week-old male SNAP25^{+/+} and SNAP25^{Δ3/Δ3} mice. **B.** Comparison of GSIS did not reveal a statistically significant difference between SNAP25^{+/+} and SNAP25^{Δ3/Δ3} islets. **C.** AUC values were normalized to the individual's maximal GSIS and a dose-response curve for the inhibition of GSIS by the α₂AR selective agonist, brimonidine (Br), was generated. The log IC₅₀'s were similar, being -7.262 for SNAP25^{+/+} and -7.348 for SNAP25^{Δ3/Δ3}. **D.** Islet insulin content was not different between the two genotypes. n = 4 SNAP25^{+/+}, n = 5 SNAP25^{Δ3/Δ3}. Values are expressed as mean ± SEM.

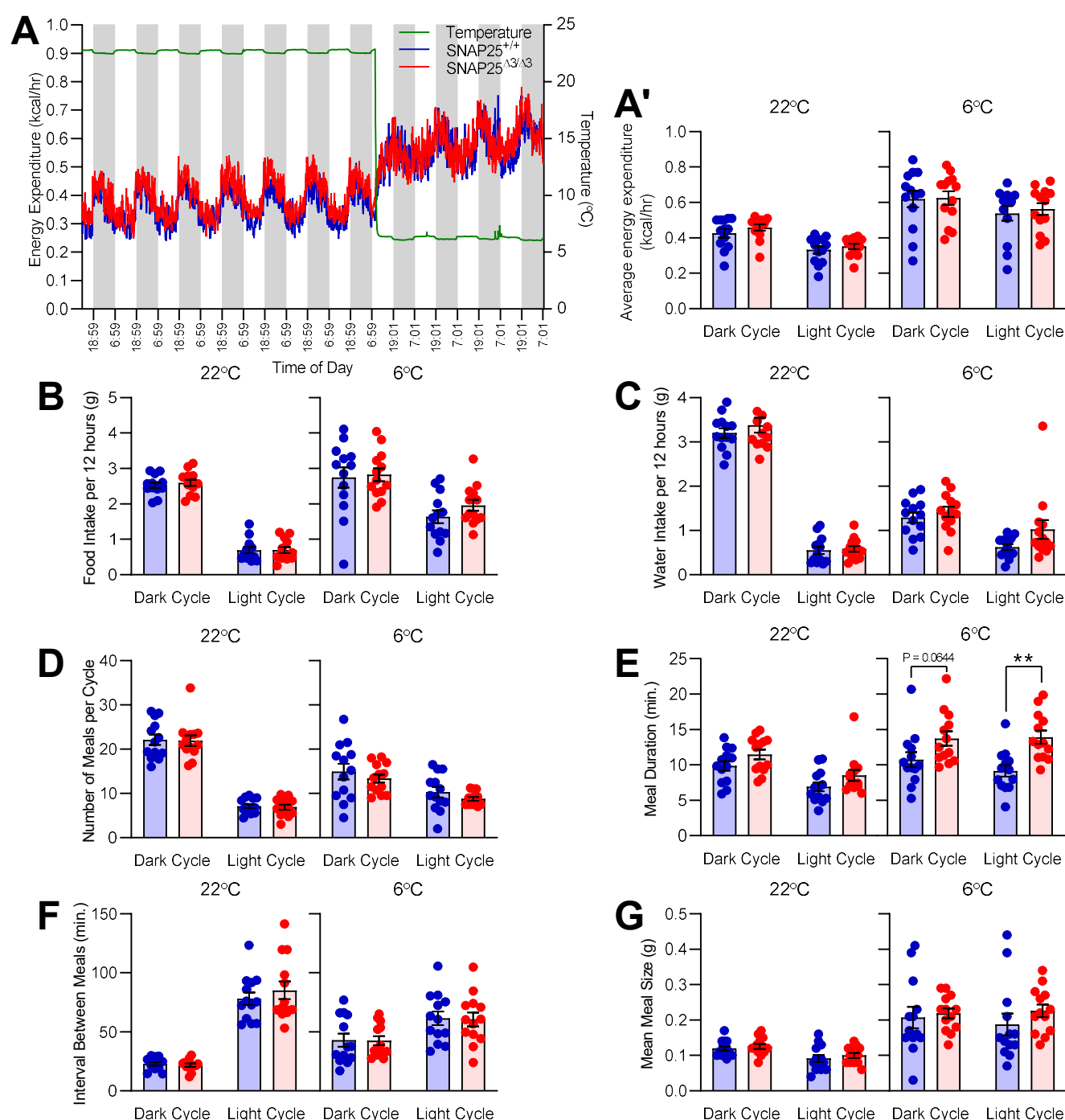


Figure 3. SNAP25^{Δ3/Δ3} mice have differences in feeding behavior but no difference in energy expenditure

A. Energy expenditure was measured by metabolic cages at 22°C or 6°C (as indicated by green line) in young, lean, male SNAP25^{Δ3/Δ3} mice or SNAP25^{+/+} littermates (A n=8 each, A' n=13 each). **B.** Food consumption and **C.** water consumption were not different between genotypes. **D.** Differences in the number of meals was not statistically significant. **E.** However, meals consumed by SNAP25^{Δ3/Δ3} mice lasted for a longer period which was especially noticeable when housed at 6°C (P = 0.0050 for the effect of genotype at 6°C). **F.** There was no noticeable difference for the interval between the meals. **G.** The size of the meal was also not changed. For B-G, n=13 for each genotype. * p<0.05, **p<0.01, Bonferroni post-hoc comparison. Values are expressed as mean ± SEM.

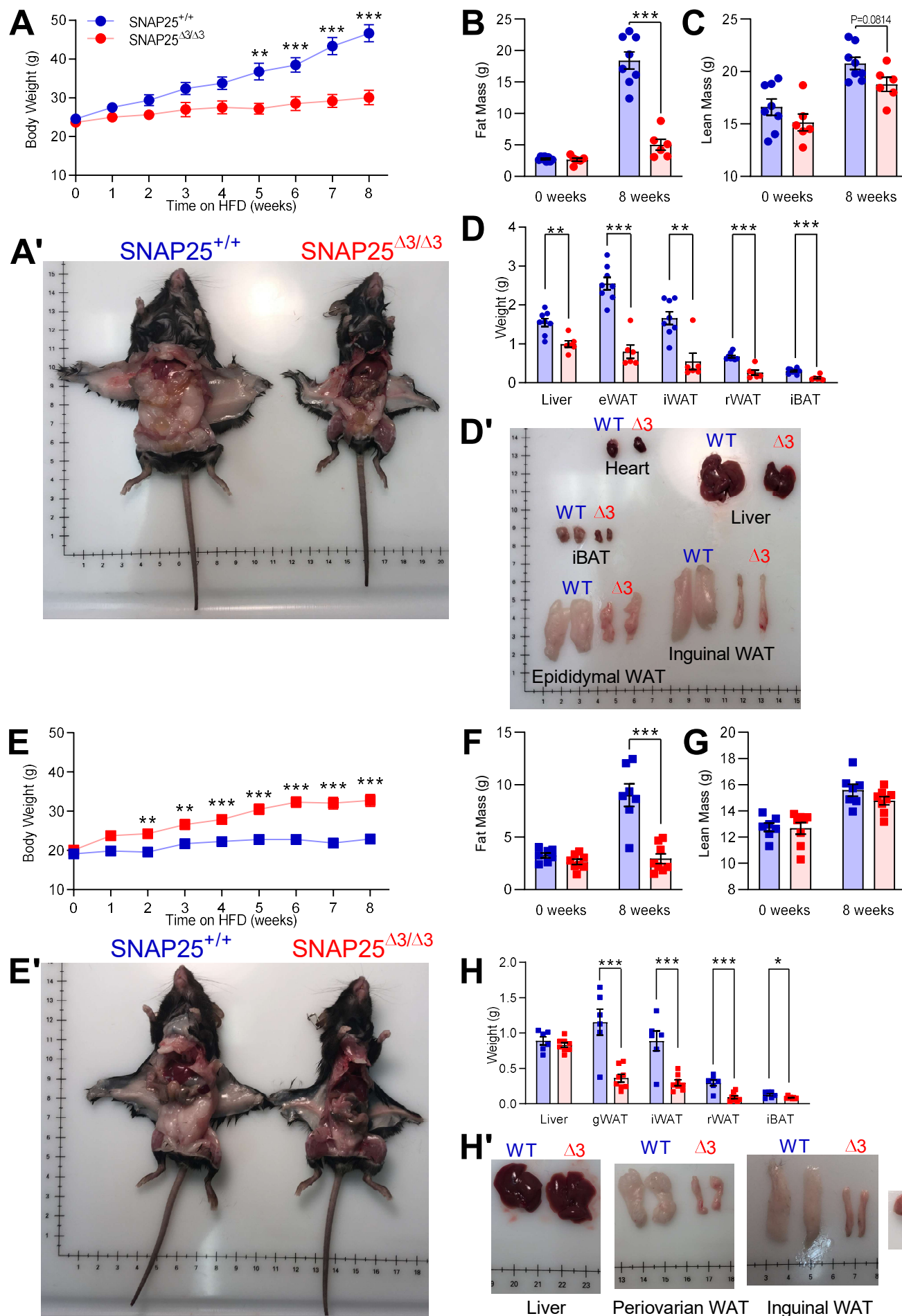


Figure 4. SNAP25^{Δ3/Δ3} male and female mice are resistant to diet-induced obesity

Male SNAP25^{+/+} and SNAP25^{Δ3/Δ3} mice were fed HFD beginning at 8 weeks of age. **A.** Male SNAP25^{+/+} mice became significantly heavier than SNAP25^{Δ3/Δ3} animals during the period of HFD feeding. Body composition was assessed at 4-week intervals by pulsed NMR. **B.** After 8 weeks of HFD feeding male SNAP25^{+/+} gained significantly more fat mass than SNAP25^{Δ3/Δ3}. **C.** The male SNAP25^{Δ3/Δ3} mice tended to have less lean mass throughout the study ($P = 0.0891$ for the effect of genotype). **D.** Liver and fat pads were weighed postmortem at the conclusion of the study. All adipose and liver tissues weighed were heavier in male SNAP25^{+/+} than in SNAP25^{Δ3/Δ3} mice. **E.** Female SNAP25^{+/+} mice became significantly heavier than SNAP25^{Δ3/Δ3} animals during the period of HFD feeding. Body composition was assessed at 4-week intervals by pulsed NMR. **F.** After 8 weeks of HFD feeding female SNAP25^{+/+} gained significantly more fat mass than SNAP25^{Δ3/Δ3}. **G.** Female SNAP25^{Δ3/Δ3} mice had similar less lean mass throughout the study. **H.** Liver and fat pads were weighed postmortem at the conclusion of the study. All adipose tissues weighed were heavier in female SNAP25^{+/+} than in SNAP25^{Δ3/Δ3} mice, but liver weight were similar. For all figures, $n = 8$ male SNAP25^{+/+}, $n = 6$ male SNAP25^{Δ3/Δ3}, $n = 6-8$ female SNAP25^{+/+}, $n = 8$ female SNAP25^{Δ3/Δ3}. Values are expressed as mean \pm SEM.

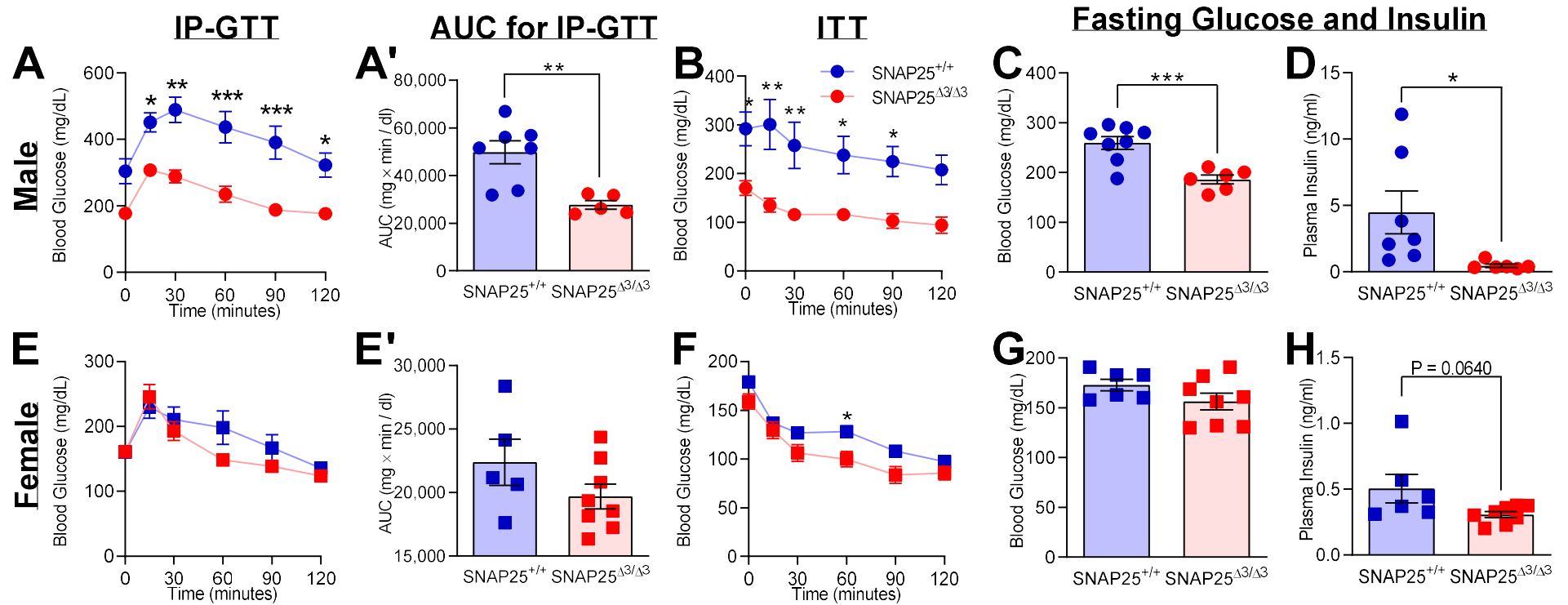


Figure 5. Male SNAP25^{Δ3/Δ3} mice have improved glucose and insulin tolerance when fed a HFD

A. Glucose tolerance was assessed by an IP-GTT in HFD-fed male SNAP25^{+/+} and SNAP25^{Δ3/Δ3} mice at 22 weeks of age after 14 weeks of HFD feeding and was found to be improved in the SNAP25^{Δ3/Δ3} mice. **B.** Insulin tolerance was assessed by an ITT in these mice at 23 weeks of age after 15 weeks of HFD feeding and was found to be lower in the SNAP25^{Δ3/Δ3} mice. **C.** 5-hour fasting blood glucose and **D.** plasma insulin were both reduced in these SNAP25^{Δ3/Δ3} mice. **E.** Glucose tolerance was assessed by an IP-GTT in HFD-fed female SNAP25^{+/+} and SNAP25^{Δ3/Δ3} mice at 22 weeks of age after 14 weeks of HFD feeding and was not found to be significantly altered. **F.** Insulin tolerance was assessed by an ITT in these mice at 23 weeks of age after 15 weeks of HFD feeding with only minimal changes being observed. **G.** 5-hour fasting blood glucose and **H.** plasma insulin were both not significantly changed. For GTT: n = 7 male SNAP25^{+/+}, n = 5 male SNAP25^{Δ3/Δ3}, 5 female SNAP25^{+/+}, n = 8 female SNAP25^{Δ3/Δ3}. For ITT: n = 7 male SNAP25^{+/+}, n = 7 male SNAP25^{Δ3/Δ3}, 5 female SNAP25^{+/+}, n = 8 female SNAP25^{Δ3/Δ3}. Values are expressed as mean ± SEM.

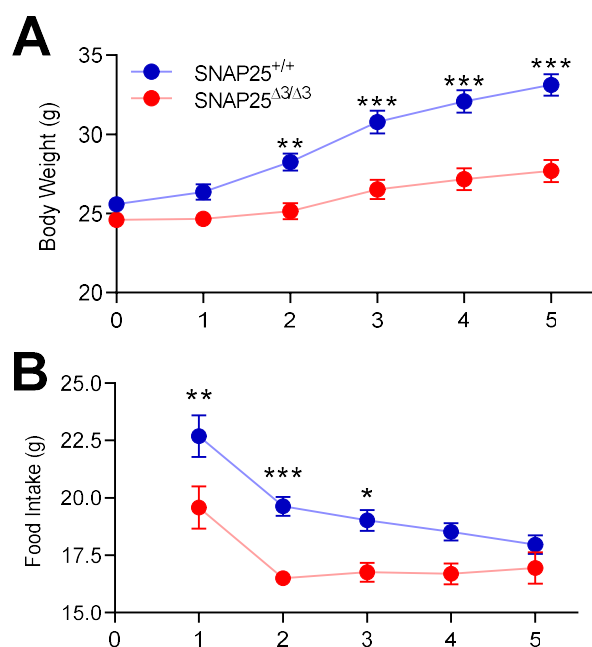


Figure 6. Male SNAP25^{Δ3/Δ3} mice do not consume as much food as SNAP25^{+/+} when presented with a HFD

Male SNAP25^{+/+} and SNAP25^{Δ3/Δ3} mice were fed HFD beginning at 8 weeks of age and food consumption was measured weekly. **A.** SNAP25^{+/+} mice became significantly heavier than SNAP25^{Δ3/Δ3} animals during the period of HFD feeding. **B.** Food intake was initially lower in SNAP25^{Δ3/Δ3} mice but became similar after four weeks. n = 23 SNAP25^{+/+}, n = 19 SNAP25^{Δ3/Δ3}. Values are expressed as mean ± SEM.

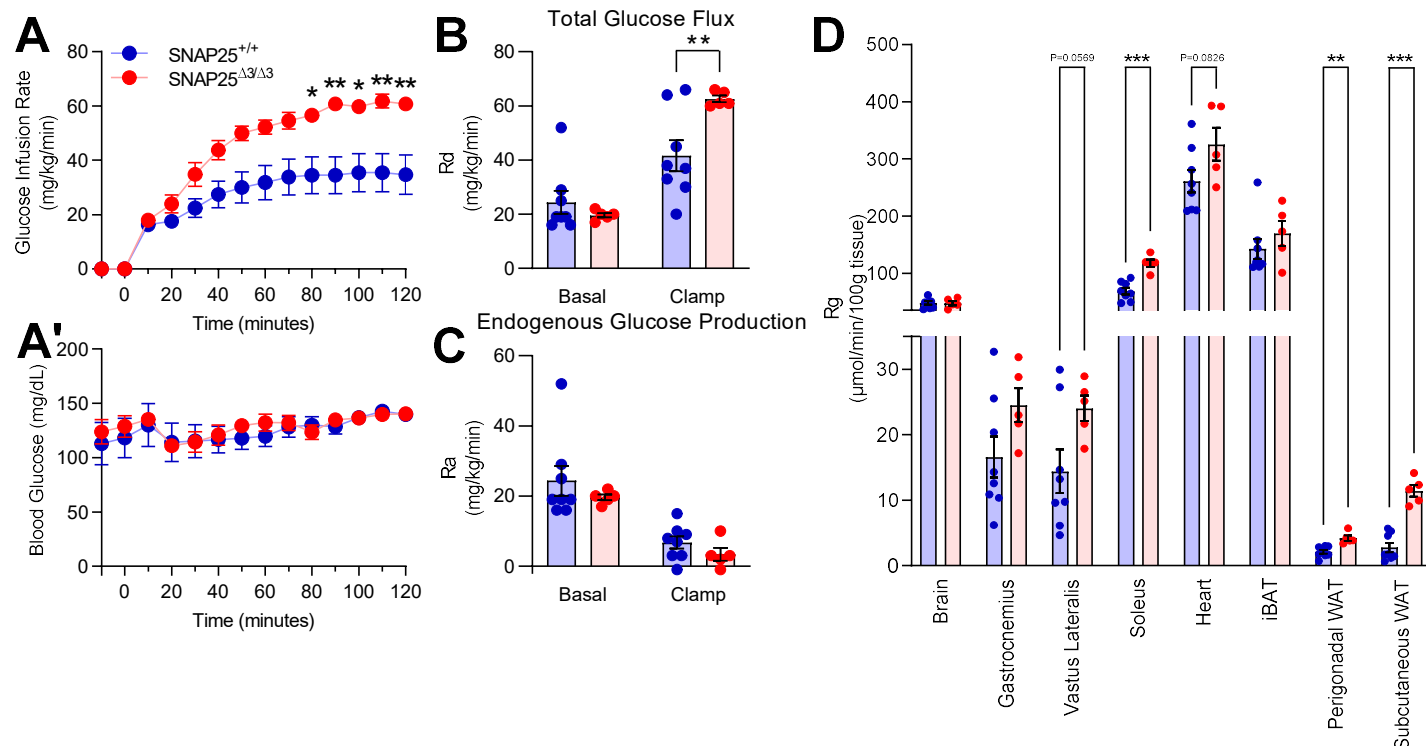
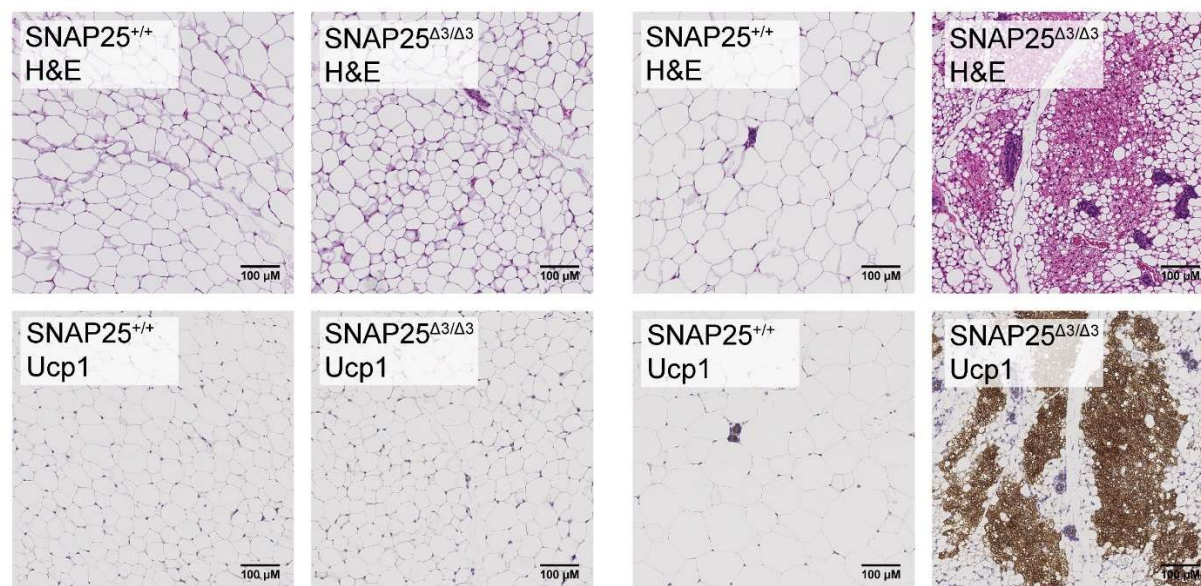


Figure 7. SNAP25^{Δ3/Δ3} mice have improved insulin sensitivity and increased glucose uptake in muscle and adipose tissues
A. Glucose infusion rate was improved in male SNAP25^{Δ3/Δ3} mice during a hyperinsulinemic euglycemic clamp study (P = 0.0224 for the effect of genotype). **A'.** During the clamp, the blood glucose was maintained at similar levels. **B.** The rate of glucose disposal was also increased in the SNAP25^{Δ3/Δ3} mice as compared to SNAP25^{+/+} (P = 0.0010 for the effect of interaction). **C.** The rate of endogenous glucose production was not altered in SNAP25^{Δ3/Δ3} mice. **D.** Uptake of radiolabeled 2-deoxyglucose into the tissues revealed that SNAP25^{Δ3/Δ3} mice have greater glucose uptake in soleus muscle and white adipose tissue depots. n = 7 SNAP25^{+/+}, n = 5 SNAP25^{Δ3/Δ3}. Values are expressed as mean ± SEM.

Male

Female

A. Inguinal White Adipose Tissue



B. Interscapular Brown Adipose Tissue



Figure 8. HFD fed $\text{SNAP25}^{\Delta3/\Delta3}$ mice have improved adipocyte morphology and female $\text{SNAP25}^{\Delta3/\Delta3}$ mice exhibit browning in inguinal white adipose depots

(A) Inguinal white adipose tissue (iWAT) and **(B)** interscapular brown adipose tissue (iBAT) were harvested from male and female $\text{SNAP25}^{+/+}$ and $\text{SNAP25}^{\Delta3/\Delta3}$ mice following 8 weeks on HFD. Representative H&E- and UCP1-stained sections show that $\text{SNAP25}^{\Delta3/\Delta3}$ mice had smaller adipocytes in both iWAT and iBAT. Furthermore, female $\text{SNAP25}^{\Delta3/\Delta3}$ mice exhibited a marked increase in UCP1-positively stained adipocytes in the iWAT. Images are a representative sample from 2 mice from each group.

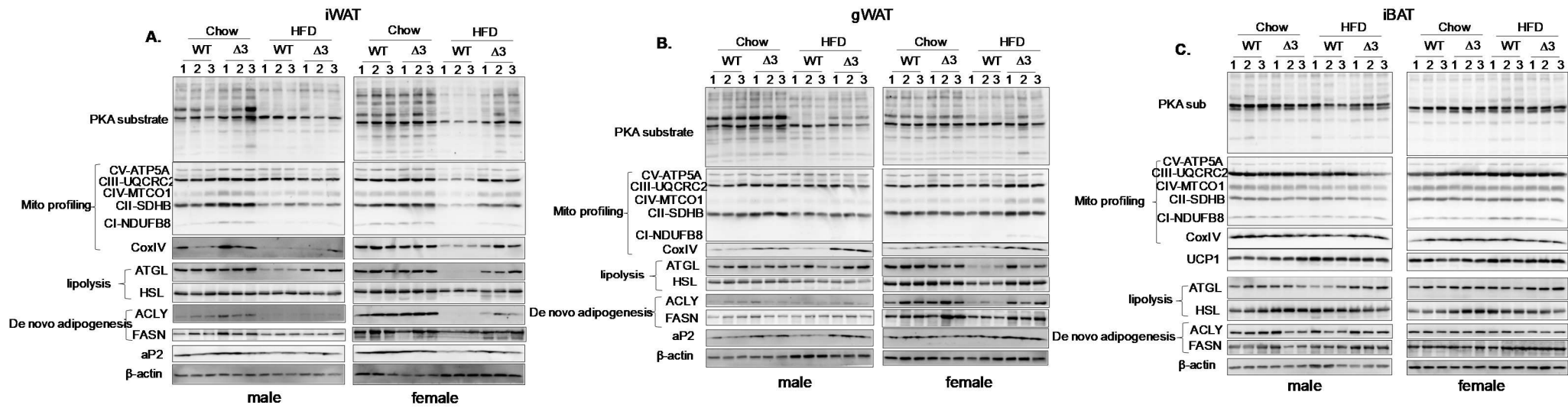


Figure 9. SNAP25^{Δ3/Δ3} mice have increased lipolysis, mitochondria markers, and PKA signaling in white adipose depots

(A) Inguinal white adipose tissue (iWAT), (B) gonadal white adipose tissue (gWAT), and (C) interscapular brown adipose tissue (iBAT) were harvested from male and female SNAP25^{+/+} and SNAP25^{Δ3/Δ3} mice following 8 weeks HFD or chow feeding. Both iWAT and gWAT from SNAP25^{Δ3/Δ3} mice exhibited increased markers of mitochondria and lipolytic activity. Western Blotting with anti-RRXS*/T* (the PKA consensus sequence) antibody, indicated that PKA signaling was increased in SNAP25^{Δ3/Δ3} mice.