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Divergent Roles of $\alpha 5$ and $\beta 4$ Nicotinic Receptor Subunits in Food Reward and Nicotine-Induced Weight Loss in Male Mice

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Conflict of interest

The authors declare that they have no conflict of interest.

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Abstract

A major obstacle to successful smoking cessation is the prospect of weight gain. Despite a clear relationship between cigarette smoking and body weight, surprisingly little is known about the physiological and molecular mechanism by which nicotine affects energy homeostasis and food motivated behaviors. Here we use loss-of-function mouse models to demonstrate that two nicotinic acetylcholine receptor (nAChR) subunits encoded by the CHRNA5-CHRNA3-CHRNA4 gene cluster, $\alpha 5$ and $\beta 4$, exhibit divergent roles in food reward. We also reveal that $\beta 4$ -containing nAChRs are essential for the weight-lowering effects of nicotine in diet-induced obese mice. Finally, our data support the notion of cross-talk between incretin biology and nAChR signaling, as we demonstrate that the glycemic benefits of glucagon-like peptide-1 receptor (GLP-1R) activation partially relies on $\beta 4$ -containing nAChRs. Together, these data encourage further research into the role of cholinergic neurotransmission in regulating food reward and the translational pursuit of site-directed targeting of $\beta 4$ -containing nAChRs for treatment of metabolic disease.

Key Words

Nicotinic Receptor, nAChR, Body Weight, Metabolism, Reward, Nicotine

1. Introduction

Successful management of obesity and its metabolic comorbidities necessitates a deeper understanding of biological mechanisms that regulate body weight and the pathological mechanisms underlying excessive gain of body fat. Epidemiologic studies imply a strong relationship between cigarette smoking and body weight (1). Smokers weigh less than non-smokers of the same age and sex (2,3), and smoking cessation has been linked to increased food intake, decreased metabolic rate, and weight gain (4–6). Nicotine is regarded as the primary molecular effector of smoking on energy balance, and animal studies have demonstrated that nicotine modulates body weight through combined actions on energy expenditure and hedonic and homeostatic feeding (7–10).

Nicotine signals via nicotinic acetylcholine receptors (nAChRs), a family of ligand-gated ion channels that are expressed in both the central and peripheral nervous systems and also in non-neuronal cells (11–13). In 2011, a landmark discovery revealed that the $\beta 4$ nAChR subunit in arcuate nucleus (ARC) pro-opiomelanocortin (POMC) neurons is important for the effects of nicotine on short-term food intake (14). $\beta 4$ is located on chromosome 15 in a gene cluster with $\alpha 3$ and $\alpha 5$, and the three subunits co-assemble to form functional receptors (15,16). Interestingly, the $\alpha 5$ - $\alpha 3$ - $\beta 4$ (*CHRNA5-CHRNA3-CHRNB4*) gene cluster has been linked to smoking heaviness, nicotine dependence (17–21), and higher body mass index (BMI) (22,23). Animal studies have implicated all individual subunits of the $\alpha 5$ - $\alpha 3$ - $\beta 4$ gene cluster in regulating nicotine consumption (titration of reward and aversion) (17,18,24). However, despite a clear implication of $\alpha 5$ and $\beta 4$ in nicotine sensitivity, the role of these subunits in food reward has received very little attention.

Here, we employ loss-of-function mouse models to investigate the role of nAChR subunits $\alpha 5$ and $\beta 4$ in body weight homeostasis and food reward. Moreover, we set out to clarify if nAChRs containing $\alpha 5$ or $\beta 4$ subunits are essential for the weight-lowering effects of nicotine. Finally, the fact that signaling cross-talk between the glucagon-like peptide-1 receptor (GLP-1R) and nAChRs is important for nicotine sensitivity (25) prompted us to scrutinize if $\alpha 5$ or $\beta 4$ is important for the benefits of GLP-1R agonists on weight loss and glucose metabolism.

2. Methods

2.1 Animal experiments

2.1.1 Animals

Global $\alpha 5$ knockout (KO) and wild type (WT) mice on C57BL6 background were generated as previously described (26). Global $\beta 4$ KO and WT mice on a C57BL6J background were generated as previously described (27). Male mice were double or single housed at all times in a temperature and humidity-controlled environment (22°C, 33-35% humidity) and on a 12-hour light-dark cycle. Mice had *ad libitum* access to standard chow diet (Altromin #1310, Lage, Germany) and tap water. For diet induced obesity (DIO), mice were given *ad libitum* access to a high fat, high sucrose (HFHS) diet (D12331; Research Diets, New Brunswick, USA). All animal experiments were approved by the Danish Animal Experimentation Inspectorate (license: 2018-15-0201-01457) and performed according to institutional guidelines.

2.1.2 Food reward

Male $\alpha 5$ KO/WT (n=7) and $\beta 4$ KO/WT (n=9) mice were kept on chow diet and single or double housed. To assess food reward, they were given a HFHS diet pellet/animal for 1 hour daily during the pm time of the light cycle. The pellet was placed in the cage bottom and was weighed before and after to determine the amount of HFHS diet consumed by each mouse. The experiment was conducted over 8 days: 3 days on, 2 days off, and 3 days on.

2.1.3 Sucrose preference test

Sucrose preference was measured using a 2-bottle free-choice test. For habituation, $\alpha 5$ KO/WT (n=12) and $\beta 4$ KO/WT (n=8) mice were housed individually for 4 days in cages containing 2 water bottles. The bottles consisted of a 50 mL centrifugal tube with a metal drinking nipple. Following habituation, one of the water bottles was replaced with a bottle containing sucrose water. The volumes of water and sucrose solution along with body weight and food intake were measured daily. On consecutive days, the concentration of sucrose (Sigma Aldrich, St. Louis, USA) was increased from 0.5% to 3% to 10% to 30%. Mice had unlimited access to both bottles at all times and, to control for potential side-preferences, the positions of the bottles were changed daily.

2.1.4 In vivo pharmacology

To study the role of nAChR $\alpha 5$ and $\beta 4$ subunits in nicotine-induced weight loss, nicotine was subcutaneously injected daily for 14 days. Male DIO $\alpha 5$ KO/WT (n=7) and $\beta 4$ KO/WT (n=11) mice with an average body weight of ~50 g, received subcutaneous injections of 2 mg/kg body weight nicotine ((-)-nicotine ditartrate, Santa Cruz Biotechnology, Dallas, USA, injection volume: 5 μ L/g body weight) or vehicle (isotonic saline, Amgros I/S, Copenhagen, Denmark, injection volume: 5 μ L/g body

weight). Injections were performed 2 hours prior to the onset of the dark cycle. At time of injection, food intake and body weight were measured. On the final day of the study, an intraperitoneal glucose tolerance test (ipGTT) was performed (see 2.1.5).

For liraglutide pharmacology, male DIO $\alpha 5$ KO/WT (n=5) and $\beta 4$ KO/WT (n=6) mice with an average body weight of ~52g received subcutaneous injections of 10 nmol/kg body weight liraglutide (Novo Nordisk, Copenhagen, Denmark, injection volume: 5 μ L/g body weight) or vehicle (isotonic saline, Amgros I/S, Copenhagen, Denmark, injection volume: 5 μ L/g body weight) daily for 7 days. Compounds were administered 2 hours prior to the onset of the dark cycle. At time of injection, body weight and food intake were measured. On the final day of the study, an ipGTT was performed (see 2.1.5).

2.1.5 Intraperitoneal glucose tolerance test (ipGTT)

Shortly after entering the light phase, mice were moved to a quiet procedure room, where they were fasted for 6 hours. After 6 hours of fasting, the test commenced with an intraperitoneal injection of 1.5 g/kg body weight glucose (D-(+)-Glucose, Sigma Aldrich, St. Louis, USA) dissolved in isotonic saline, Amgros I/S, Copenhagen, Denmark). Blood glucose levels were determined at times 0 (prior to injection), 15, 30, 60, and 120 minutes after injection, using a handheld glucometer (Abbott GmbH & Co. KG, Wiesbaden, Germany).

2.2 Gene expression analysis (qPCR)

Hypothalamus and striatum were dissected, immediately frozen on dry ice, and stored at -80°C. Tissue was homogenized in a Trizolreagent (QIAzol Lysis Reagent, Qiagen, Hilden, Germany) using a 5mm stainless-steel bead (Qiagen, Hilden, Germany) and a TissueLyser (Qiagen, Hilden, Germany) at

50Hz for 3 minutes, followed by 5 minutes incubation at room temperature. 200 μ L chloroform (Sigma-Aldrich, St. Louis, USA) was added and tubes were shaken vigorously for 15 seconds and incubated for 2 minutes at room temperature. The samples were then centrifuged for 15 minutes at 12,000 \times g at 4°C. The aqueous phase was mixed 1:1 with 70% ethanol. The samples were further processed using RNeasy Lipid Tissue Mini Kit (Qiagen, Hilden, Germany) following the instructions provided by the manufacturer. RNA content was measured using a NanoDrop 2000 (Thermo Fisher, Waltham, USA). 500 ng of RNA was converted into cDNA by mixing FS buffer and DTT (Thermo Fisher, Waltham, USA) with random primers (Sigma-Aldrich, St. Louis, USA) and incubated for 3 min at 70 °C. Then, dNTPs, RNase out, and Superscript III (Thermo Fisher, Waltham, USA) were added, and samples were placed in a thermal cycler for 5 min at 25 °C, 60 min at 50 °C, and 15 min at 70 °C. cDNA was diluted 1:20 and kept at -20 °C until further processing. qPCR was performed using PrecisionPLUS qPCR Master Mix containing SYBR green (Primer Design, Camberly, UK). cDNA, primers (Table 1 and Supplementary Table 1) (28), and PrecisionPLUS qPCR Master Mix were mixed in a 384-well plate and incubated in a LightCycler (LightCycler 480 II, Roche, Basel, Switzerland) using 2 minutes preincubation at 95°C followed by 45 cycles of 60 sec at 60 °C. Melting curve analysis was performed by stepwise increasing the temperature from 60 to 95 °C (Table 1).

2.3 Statistical Analyses

The data were analyzed using a paired two-tailed t test, an unpaired two-tailed t test, or a two-way ANOVA with Bonferroni post-hoc test, as described in the figure legends. $P < 0.05$ was set as the criterion for significance. All data are expressed as mean \pm SEM. Analyses were performed using Prism version 9 (GraphPad, San Diego, USA).

3. Results

3.1 nAChR subunit distribution in hypothalamus and striatum

To delineate the physiological and pharmacological properties of $\beta 4$ and $\alpha 5$ subunits in homeostatic and hedonic feeding, we profiled $\beta 4$ and $\alpha 5$ mRNA expression in the hypothalamus and striatum and compared the relative expression levels to other major central nervous system (CNS) nAChR subunits (Figure 1A). We found that the $\beta 4$, $\alpha 4$, $\alpha 7$, and $\beta 2$ subunits are expressed at similar levels in the hypothalamus and striatum (Figure 1B), but that $\alpha 5$ mRNA is significantly higher in the striatum relative to the hypothalamus, whereas the opposite is the case for $\alpha 3$ mRNA. Next, we wanted to compare the relative distribution of the same subunits within each region. In agreement with previous work (29), we found that $\beta 2$ is the most abundant subunit in both regions (Figure 1C). Of note, the $\alpha 3$ subunit is abundantly expressed in the hypothalamus. Finally, we wanted to evaluate the relative mRNA levels of the $\alpha 5$ - $\alpha 3$ - $\beta 4$ gene cluster subunits in the two global KO mouse models (Figure 1D-1G). As expected, these analyses confirmed complete ablation $\alpha 5$ mRNA in the $\alpha 5$ KO model and complete ablation of $\beta 4$ mRNA in the $\beta 4$ KO model. Notably, we demonstrate that KO of either $\alpha 5$ or $\beta 4$ subunits results in the partial KO of the $\alpha 3$ subunit in both the hypothalamus and striatum (Figure 1D-G), and KO of the $\beta 4$ subunit suppresses $\alpha 5$ mRNA levels in the hypothalamus (Figure 1F). Despite the substantially reduced levels of $\alpha 3$ mRNA in both $\alpha 5$ and $\beta 4$ KO models, this may be sufficient for $\alpha 3$ -dependent nAChR signaling (30).

3.2 Divergent roles of $\alpha 5$ and $\beta 4$ subunits in food reward

To understand the physiological implication of $\alpha 5$ and $\beta 4$ subunits on energy homeostasis, we measured body weight of $\alpha 5$ KO and WT and $\beta 4$ KO and their respective WT control mice, maintained on a chow diet or following 25 weeks on a HFHS diet. We found that ablation of either the $\alpha 5$ or $\beta 4$ subunit did not elicit perturbations in long-term weight homeostasis (Figure 2A and 2B), which might relate to compensatory mechanisms in the KO strains. Additionally, gene expression analyses of inguinal white adipose tissue (iWAT), brown adipose tissue (BAT) and hypothalamus from both KO strains reveal similar expression of genes related to thermogenesis (iWAT and BAT) and appetite regulation (hypothalamus) in both KO mouse strains relative to their respective WT controls (Supplementary Figure 1) (31). To gauge if $\alpha 5$ and $\beta 4$ subunits are involved in food reward and palatable food intake, we established a simple experimental paradigm. Animals maintained on a chow diet were given 1-hour access to a single pellet of HFHS diet for 3 consecutive days. Subsequently, mice had 2 days of wash-out with no HFHS diet exposure, before being re-introduced to the HFHS pellet paradigm for 3 additional days (Figure 2C1). Whereas the initial intake might relate to palatability and novelty, intake at later stages of the experiment is likely governed by the motivational value of the HFHS pellet. For the first 2 days, $\alpha 5$ KO and WT mice had a comparable intake of the HFHS pellet, but, from day 3 and onwards, the $\alpha 5$ KO mice had a greater intake of HFHS diet, leading to a significantly higher cumulative HFHS diet intake by the end of the experiment ($p = 0.0208$, Figure 2D and 2E). In contrast, $\beta 4$ KO mice displayed a lower daily intake of the HFHS reward and plateaued at ~ 0.3 g consumed during the 1-hour exposure as opposed to ~ 0.6 g for the WT mice (and as opposed to >1 g for $\alpha 5$ KO mice). Thus, $\beta 4$ KO mice had a significantly lower cumulative intake relative to WT mice in this food reward paradigm ($p = 0.0084$, Figure 2F and 2G).

Given the divergent roles of the $\alpha 5$ and $\beta 4$ subunits in mixed high fat, high sugar reward, we employed 2-bottle free choice paradigm to directly evaluate the sweet preference component of

this behavior. For this, an up-titrated sucrose preference test was employed (Figure 2C2). These experiments revealed similar preference for sucrose between $\alpha 5$ KO, $\beta 4$ KO, and WT mice (Figure 2H-2K). Together, these data reveal that both $\alpha 5$ and $\beta 4$ subunits are implicated in food reward, but whereas loss of $\beta 4$ decreases food reward, loss of $\alpha 5$ has the opposite effect. Further, the body weight data imply that any motivational aspects consequential to genetic ablation of either $\alpha 5$ or $\beta 4$ subunits are accounted for homeostatically over time.

3.3 The $\beta 4$ nAChR subunit is indispensable for nicotine-induced weight loss

Studies with viral-mediated ablation of the hypothalamic $\beta 4$ subunit demonstrated that this subunit is essential for the acute effects of nicotine on food intake (14,32). However, whether $\beta 4$ is exclusively responsible for this pharmacological effect, whether other nAChR subunits are implicated, or whether there is subunit redundancy, is less clear. Similarly, it is unknown if the $\beta 4$ subunit is important for the long-term effect of nicotine on energy balance. To address these questions, we exposed $\alpha 5$ KO, $\beta 4$ KO, and WT mice to a 2-week nicotine pharmacology treatment study. In agreement with previous studies, we used 2mg/kg nicotine administered daily to DIO mice maintained on a HFHS diet (33). For the $\alpha 5$ KO strain, we observed a similar weight loss in response to nicotine for both KO and WT mice (Figure 3B and 3C, main effect of nicotine; $p=0.0005$). The nicotine-induced weight loss was not accompanied by significant changes in cumulative food intake, and nicotine treatment did not influence glucose tolerance (Figure 3D-3G). In contrast, nicotine treatment of $\beta 4$ KO mice did not lower body weight in contrast to the ~6% weight loss observed in the WT mice (Figure 3H-3I). For the $\beta 4$ subunit study, nicotine, at least in part, mediated the weight loss in the WT mice via reduced food intake (Figure 3J-3K, $p<0.0001$).

Finally, following the 2-week treatment study, a glucose tolerance test revealed that the $\beta 4$ subunit KO had a slightly impaired glucose tolerance, but this was independent of nicotine treatment (Figure 3L-3KM).

These findings clearly demonstrate that the $\beta 4$ subunit, but not the $\alpha 5$ subunit, is required for nicotine-induced weight loss, and loss of the $\beta 4$ subunit leads to perturbed glucose metabolism in the context of a HFHS diet. Thus, our data further contribute to the specific role of $\beta 4$ -nAChRs as a relevant pharmacological target for weight loss.

3.4 Glycemic benefits of liraglutide implicates $\beta 4$ -nAChRs

GLP-1R agonists have emerged as potent agents for treatment of type 2 diabetes and obesity. Interestingly, central GLP-1 signaling modulates nicotine intake via the habenular-interpeduncular pathway (25). Given that the habenular-interpeduncular pathway is particularly enriched with $\alpha 5$ - and $\beta 4$ - nAChRs (34,35) and site-specific loss- and gain-of-function models have demonstrated key roles for both receptor subunits in governing nicotine consumption (17,18), neural cross-talk between GLP-1R and nAChR signaling might be relevant to this neurobiology. To evaluate this, we exposed $\alpha 5$ KO, $\beta 4$ KO, and WT mice to a pharmacological treatment study with the long-acting GLP-1R agonist liraglutide (36). The experiment was conducted over 7 days with daily injections of either liraglutide or vehicle followed by an ipGTT on day 7 (Figure 4A). We demonstrate that liraglutide decreases body weight and food intake to a similar degree in both WT mice and $\alpha 5$ KO mice (Figure 4B-G), and in WT mice and $\beta 4$ KO mice (Figure 4H-K), but the anti-diabetic effect of liraglutide is diminished in $\beta 4$ KO mice (Figure 4L-M).

4. Discussion

Despite ~1 billion tobacco users globally (37), remarkably little is known about the physiological and molecular mechanism by which nicotine affects energy homeostasis. In the brain, nicotine signals through pentameric nAChRs comprised of 7 different α subunits and 3 different β subunits. Here, we evaluated the role of the $\alpha 5$ and $\beta 4$ subunits from the $\alpha 5$ - $\alpha 3$ - $\beta 4$ (*CHRNA5-CHRNA3-CHRNB4*) gene cluster in the physiological regulation of energy homeostasis and food reward and in response to nicotine and GLP-1 weight-lowering pharmacology. We show that both subunits are abundantly expressed in the hypothalamus and striatum and that they have opposing roles in food reward. Moreover, we demonstrate that nAChRs containing $\beta 4$ subunits are indispensable for the effects of nicotine on food intake and weight loss and that they are important for the benefits of GLP-1R agonism on glucose metabolism. In agreement with previous work (Jall et al., 2020), these findings suggest that ligands selective for $\beta 4$ -nAChRs hold promise for targeting the inherent benefits of nicotinic receptor activity on energy metabolism.

The rewarding effects of nicotine are often ascribed to nAChRs containing $\beta 2$ subunits in the ventral tegmental area (38–40). High doses of nicotine also result in co-activation of nAChRs containing $\alpha 5$, $\alpha 3$, and $\beta 4$ subunits in the medial habenula that, through projections to the interpeduncular nucleus, promote nicotine avoidance (17,19,41,42). Loss-of-function models have confirmed key roles for both $\alpha 5$ and $\beta 4$ subunits in governing nicotine sensitivity (43). At low doses, $\beta 4$ KO mice administer less nicotine compared to WT mice (18), whereas $\alpha 5$ KO mice are reported to have reduced nicotine sensitivity and thus self-administer more nicotine than WT mice at high concentrations of nicotine (17,44,45). Here, we find that $\beta 4$ KO mice consume less HFHS diet than WT mice in a food reward paradigm. This observation agrees with recent work showing diminished food reward in $\beta 4$ KO mice using an operant conditioning task for palatable food reward (18). Speaking against a robust role for $\beta 4$ in food reward and palatable food intake, we failed to

demonstrate a sucrose drinking phenotype. Conversely, and in agreement with a role for the $\alpha 5$ subunit in providing an inhibitory reward signal, we find that $\alpha 5$ KO mice overconsume the rewarding HFHS diet relative to WT mice. Further, our data agree with recent work showing increased food seeking relapse in rats with a *CHRNA5* gene variant (46). Together, this points to shared neurobiological pathways of nicotine- and food reward titration. Importantly, the observed genotype-specific food motivated behavior is abrogated in the context of chronic *ad libitum* access to HFHS diet as both $\alpha 5$ KO mice and $\beta 4$ KO mice develop a diet-induced obese phenotype similar to WT mice.

It is notable that ablation of both $\alpha 5$ and $\beta 4$ subunits leads to a comparable and concomitant reduction in $\alpha 3$ mRNA in the hypothalamus and striatum. As such, the respective KO models can be regarded as full KOs of the $\alpha 5$ and $\beta 4$ subunits and partial knock-down of the $\alpha 3$ subunit. The remnant $\alpha 3$ mRNA expression, however, does not necessarily result in reduced levels of $\alpha 3$ protein expression. Importantly, the similar reduction in $\alpha 3$ mRNA between $\alpha 5$ and $\beta 4$ KO mice, along with $\alpha 5$ mRNA reduction in $\beta 4$ KO mouse hypothalamus, implies that the phenotypes observed in this study are not confounded by differential changes in redundant subunit expression. This underscores the exclusive role for $\beta 4$ -nAChRs in driving nicotine induced weight loss. This finding substantiates previous work focusing on $\beta 4$ knock-down in the ARC (14,32). $\alpha 7$ - and $\beta 2$ -nAChRs have been implicated previously in homeostatic food intake regulation (47–49), and here we show a role for $\alpha 5$ and $\beta 4$ subunits in food reward. However, in terms of pharmacological nicotine, the effects on weight loss appear to be exclusively governed by $\beta 4$ -nAChRs.

Central GLP-1 receptors are reported to act on habenular avoidance circuits to control nicotine intake (25), emphasizing neural crosstalk between nAChRs and GLP-1Rs. This encouraged us to

evaluate if the full metabolic benefits of pharmacological GLP-1R agonism implicates functional $\alpha 5$ - or $\beta 4$ -nAChRs. Whereas the liraglutide-induced weight loss was intact in both $\alpha 5$ KO and $\beta 4$ KO mice, the effect of liraglutide on reversing diet-induced glucose intolerance was impaired in $\beta 4$ KO mice. The implication of $\beta 4$ -nAChRs in glucose metabolism is supported by previous work demonstrating that pharmacological activation of $\alpha 3\beta 4$ -nAChRs improves peripheral insulin sensitivity and reverses diet-induced glucose intolerance in mice (33,50). Together, these findings encourage further investigations into nAChRs, and in particular $\beta 4$ -nAChRs, in physiological and pharmacological regulation of glucose metabolism. It should be of particular interest to investigate metabolic benefits of coordinated pharmacological agonism at both $\beta 4$ -nAChRs and GLP-1R as well as to evaluate if variations in the *CHRNA5-CHRNA3-CHRNA4* gene region influence responsiveness to long-acting GLP-1R agonists in humans.

In summary, we report a divergent role of $\alpha 5$ - and $\beta 4$ -nAChRs in food reward. More work is needed to decipher the mechanistic underpinnings by which the cholinergic system governs food motivated behaviors and how this integrates with canonical energy homeostatic signals. Smoking cessation leads to weight gain and short-term risk of type 2 diabetes. Importantly, the negative influence on metabolic parameters is surpassed by the benefits of quitting smoking through reduction of cardiovascular and all-cause mortality (51). The question is whether or not the metabolic benefits of nAChR-targeting can be harnessed *without* compromising safety. The results presented here substantiate the importance of the $\beta 4$ subunit in energy metabolism and encourage further work evaluating selective targeting of $\beta 4$ -nAChRs for treatment of metabolic disease.

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Data availability

All data generated and analyzed are included in this article or in the data repositories; Figshare repository for supplementary Figure 1 and Supplementary Table 1, which are listed in References;

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Table 1:

Gene	Forward primer (5'-3')	Reverse primer (5'-3')
CHRNA5	CGTCCGCGAGGTTGTTGAAG	AGCTGCTTGACTGCTCACTAAG
CHRNA3	GCCAAAGAGATTCAAGATGATTGG	TCTGGGGCTATTGAGAAAGTGC
CHRNB4	ATCAGAGTGTTCATCGAGGACTG	CACTAGGCTGCTCATATCATCC
CHRNB2	TGACCAGAGTGTGAGGGAGG	AGCTGCAAATGAGAGACCTCAC
CHRNA4	GACTTCTCGGTGAAGGAGGAC	GGAAGATGTGGGTGACTGACG
CHRNA7	CCTAAGTGGACCAGGATCATTC	ATGTAGAGCAGGTTGCCATTGC
Rpl13a	GGA GGG GCA GGT TCT GGT AT	TGT TGA TGC CTT CAC AGC GT

Table 1: Primers for qPCR

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Legends for figures:

Figure 1: nAChR subunit distribution in the hypothalamus and striatum. **A.** Illustration of the two brain regions in focus: hypothalamus and striatum **B.** mRNA levels of nAChR subunits $\alpha 5$, $\alpha 3$, $\beta 4$, $\alpha 4$, $\alpha 7$, $\beta 2$ relative to *Rpl13a* in hypothalamus and striatum. **C.** mRNA levels of nAChR subunits $\alpha 5$, $\alpha 3$, $\beta 4$, and $\beta 2$ relative to *Rpl13a* normalized to $\beta 4$ mRNA in hypothalamus and striatum. **D.** Levels of $\alpha 5$, $\alpha 3$, and $\beta 4$ mRNA relative to *Rpl13a* in WT and $\alpha 5$ KO mice in hypothalamus. **E.** Levels of $\alpha 5$, $\alpha 3$, and $\beta 4$ mRNA relative to *Rpl13a* in WT and $\alpha 5$ KO mice in striatum. **F.** Levels of $\alpha 5$, $\alpha 3$, and $\beta 4$ mRNA relative to *Rpl13a* in WT and $\beta 4$ KO mice in hypothalamus. **G.** Levels of $\alpha 5$, $\alpha 3$, and $\beta 4$ mRNA relative to *Rpl13a* in WT and $\beta 4$ KO mice in striatum. Data analyzed by paired t-test (**B**) and unpaired t-test (**D-G**). Data presented as mean \pm SEM. * $p < 0.05$, ** $p < 0.01$, **** $p < 0.0001$.

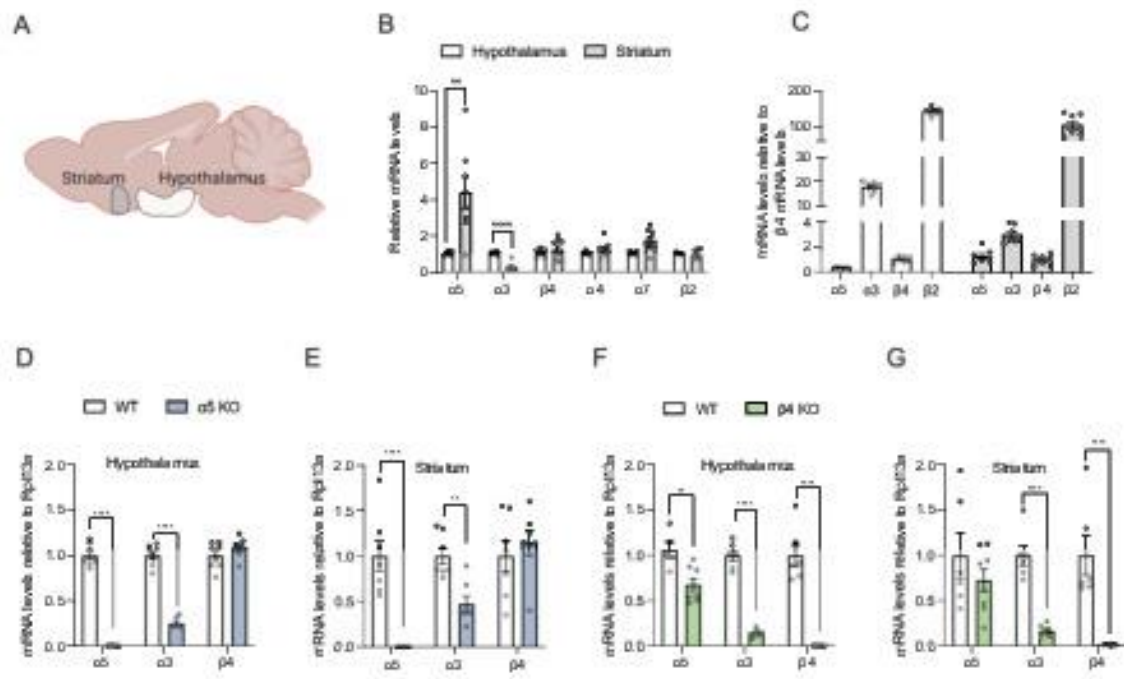
Figure 2: Divergent roles of $\alpha 5$ and $\beta 4$ in food reward. **A.** Body weight of WT and $\alpha 5$ KO on chow and HFHS diet. **B.** Body weight of WT and $\beta 4$ KO on chow and HFHS diet. **C.** Schematic illustration of experimental setups for food reward and sucrose preference. **D.** Daily intake of HFHS diet for WT and $\alpha 5$ KO mice. **E.** Cumulative HFHS diet intake for WT and $\alpha 5$ KO mice. **F.** Daily intake of HFHS diet for WT and $\beta 4$ KO mice. **G.** Cumulative HFHS diet intake for WT and $\beta 4$ KO mice. **H.** Daily intake of sucrose water for WT and $\alpha 5$ KO mice. **I.** Sucrose preference in %. Sucrose intake divided by total intake of sucrose and water for WT and $\alpha 5$ KO. **J.** Daily intake of sucrose water for WT and $\beta 4$ KO mice. **K.** Sucrose preference in %. Sucrose intake divided by total intake of sucrose and water for WT and $\beta 4$ KO. Data presented as mean \pm SEM. Data analyzed by unpaired t-test. * $p < 0.05$, ** $p < 0.01$.

Figure 3: The $\beta 4$ nAChR subunit is responsible for nicotine-induced weight loss. **A.** Schematic illustration of experimental setup for nicotine pharmacology. **B-G.** Data from $\alpha 5$ KO and WT mice. **H-M.** Data from $\beta 4$ KO and WT mice. **B.** Percentage change in body weight in response to 2 mg/kg nicotine or vehicle (saline) for $\alpha 5$ KO and WT mice. **C.** Percentage change in body weight at day 14 for $\alpha 5$ KO and WT mice. **D.** Cumulative HFHS diet

intake through experiment for $\alpha 5$ KO and WT mice. **E.** Cumulative HFHS diet intake at day 14 for $\alpha 5$ KO and WT mice. **F.** Blood glucose during ipGTT on day 14 of experiment for $\alpha 5$ KO and WT mice. **G.** Area under curve (AUC) calculated from individual blood glucose traces after the ipGTT for $\alpha 5$ KO and WT mice. **H.** Percentage change in body weight in response to 2 mg/kg nicotine or vehicle (saline) for $\beta 4$ KO and WT mice. **I.** Percentage change in body weight at day 14 for $\beta 4$ KO and WT mice. **J.** Cumulative HFHS diet intake through experiment for $\beta 4$ KO and WT mice. **K.** Cumulative HFHS diet intake at day 14 for $\beta 4$ KO and WT mice. **L.** ipGTT at day 14 of experiment for $\beta 4$ KO and WT mice. **M.** AUC calculated from individual blood glucose traces after the ipGTT for $\beta 4$ KO and WT mice. Data presented as mean \pm SEM. Data analyzed by 2-way ANOVA with Bonferroni post-hoc test. Main effect of treatment: $^{###}p < 0.001$. Main effect of genotype: $^{§§§§}p < 0.0001$. Post-hoc effect: $^{**}p < 0.01$, $^{***}p < 0.001$.

Figure 4: Glycemic benefits of GLP-1R agonism implicates $\beta 4$ -containing nAChRs. **A.** Schematic illustration of experimental setup for liraglutide pharmacology. **B-G.** Data from $\alpha 5$ KO and WT mice. **H-M.** Data from $\beta 4$ KO and WT mice. **B.** Percentage change in body weight in response to 10 nmol/kg liraglutide or vehicle (saline) for $\alpha 5$ KO and WT mice. **C.** Percentage change in body weight at day 7 for $\alpha 5$ KO and WT mice. **D.** Cumulative HFHS diet intake through experiment for $\alpha 5$ KO and WT mice. **E.** Cumulative HFHS diet intake at day 14 for $\alpha 5$ KO and WT mice. **F.** Blood glucose during ipGTT at day 14 of experiment for $\alpha 5$ KO and WT mice. **G.** AUC calculated from individual blood glucose traces after the ipGTT for $\alpha 5$ KO and WT mice. **H.** Percentage change in body weight in response to 10 nmol/kg liraglutide or vehicle (saline) for $\beta 4$ KO and WT mice. **I.** Percentage change in body weight at day 7 for $\beta 4$ KO and WT mice. **J.** Cumulative HFHS diet intake through experiment for $\beta 4$ KO and WT mice. **K.** Cumulative HFHS diet intake at day 14 for $\beta 4$ KO and WT mice. **L.** Blood glucose during ipGTT at day 14 of experiment for $\beta 4$ KO and WT mice. **M.** AUC calculated from individual blood glucose traces after the ipGTT for $\beta 4$ KO and WT mice. Data presented as mean \pm SEM. Data analyzed by 2-way ANOVA with Bonferroni post-hoc test. Main effect of treatment: $^{\#}p < 0.05$, $^{##}p < 0.01$, $^{####}p < 0.0001$. Main effect of genotype: $^{\$}p < 0.05$.

Figure 1



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Figure 2

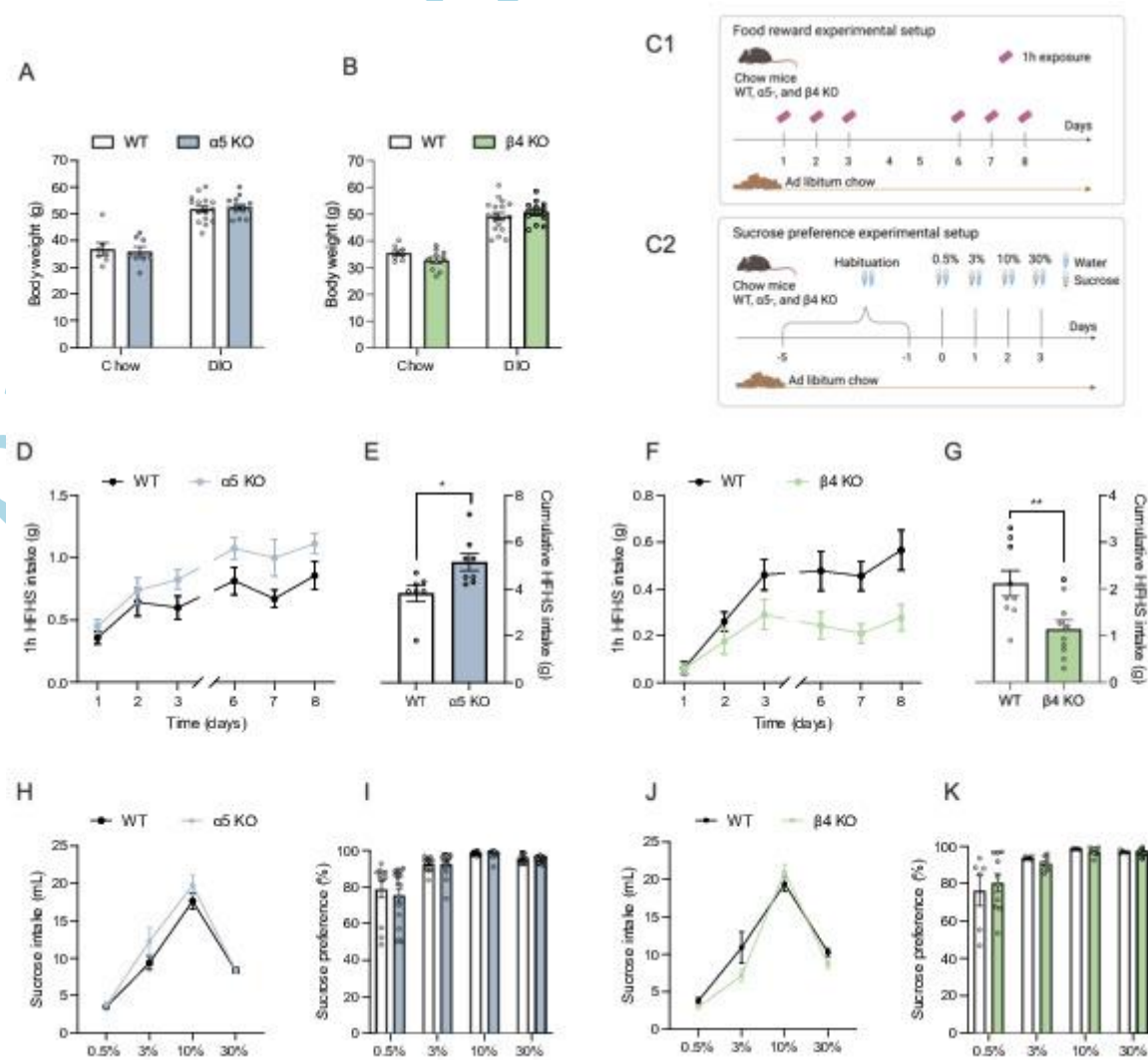
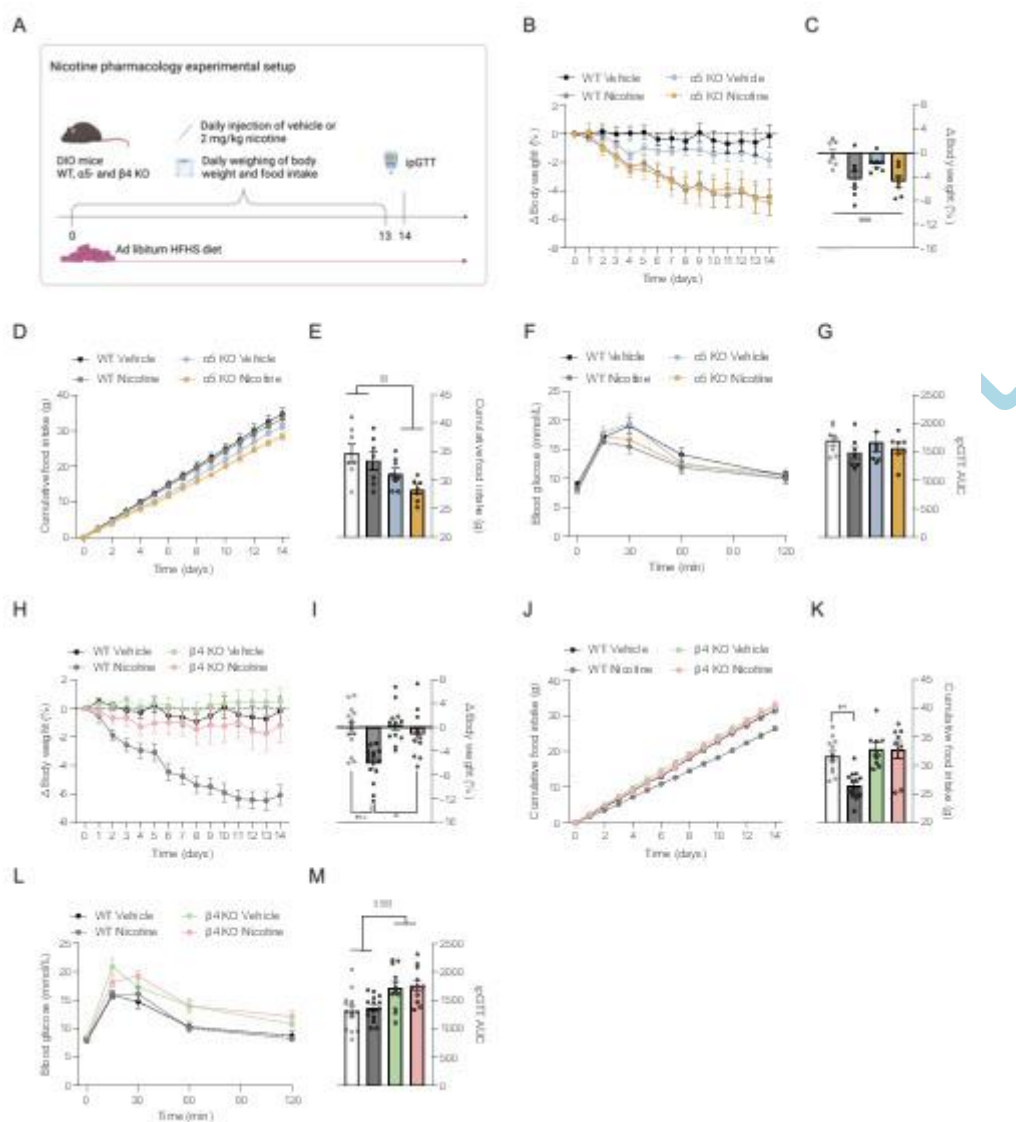


Figure 3



ACCEPT

Figure 4

