In vivo metabolic effects after acute activation of skeletal muscle Gs signaling

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ABSTRACT

Objective: The goal of this study was to determine the glucometabolic effects of acute activation of Gs signaling in skeletal muscle (SKM) in vivo and its contribution to whole-body glucose homeostasis.

Methods: To address this question, we studied mice that express a Gs-coupled designer G protein-coupled receptor (Gs-DREADD or GsD) selectively in skeletal muscle. We also identified two Gs-coupled GPCRs that are endogenously expressed by SKM at relatively high levels (β2-adrenergic receptor and CRF2 receptor) and studied the acute metabolic effects of activating these receptors in vivo by highly selective agonists (clenbuterol and urocortin 2 (UCN2), respectively).

Results: Acute stimulation of GsD signaling in SKM impaired glucose tolerance in lean and obese mice by decreasing glucose uptake selectively into SKM. The acute metabolic effects following agonist activation of β2-adrenergic and, potentially, CRF2 receptors appear primarily mediated by altered insulin release. Clenbuterol injection improved glucose tolerance by increasing insulin secretion in lean mice. In SKM, clenbuterol stimulated glycogen breakdown. UCN2 injection resulted in decreased glucose tolerance associated with lower plasma insulin levels. The acute metabolic effects of UCN2 were not mediated by SKM Gs signaling.

Conclusions: Selective activation of Gs signaling in SKM causes an acute increase in blood glucose levels. However, acute in vivo stimulation of endogenous Gs-coupled receptors enriched in SKM has only a limited impact on whole-body glucose homeostasis, most likely due to the fact that these receptors are also expressed by pancreatic islets where they modulate insulin release.

Keywords Skeletal muscle; GPCR; G protein; DREADD; Clenbuterol; Urocortin 2; Glucose homeostasis

1. INTRODUCTION

G protein-coupled receptors (GPCRs) represent the largest group of cell membrane receptors. Considering the large repertoire of ligands that act through GPCRs, it is not surprising that GPCRs are involved in nearly all physiological functions in the human body. This includes regulation of food intake, insulin secretion, lipid storage, and tissue glucose uptake [1], all processes that are highly relevant in the study of metabolic disorders. It is very difficult to assess the in vivo metabolic functions of a specific GPCR expressed by a particular tissue or cell type, primarily because each GPCR is expressed in many organs or tissues and may be able to activate more than one class of heterotrimeric G proteins. During the past decade, DREADD (Designer Receptor Exclusively Activated by Designer Drugs) technology [2,3] has emerged as a powerful chemogenetic tool to study the in vivo physiological relevance of GPCR-dependent pathways in a temporally and spatially controlled manner [1,4]. The most commonly used DREADDs are mutant muscarinic acetylcholine receptors that can be selectively activated by clozapine-N-oxide (CNO) [2,3] or certain CNO derivates such as deschloroclozapine (DCZ) [5] and compound 21 (C21) [6]. When used in the proper concentration or dose range, CNO is otherwise pharmacologically inert. During the past 15 years, DREADDs that are selectively linked to each of the four major G protein families (Gαi, Gαs, Gαq, and G12) have been developed [7]. Since SKM is responsible for the majority of insulin-mediated glucose disposal, SKM plays an important role in regulating glucose homeostasis [8]. Impaired SKM insulin sensitivity represents a key metabolic defect in the pathogenesis of type 2 diabetes [8]. An improved understanding of SKM physiology may lead to the development of novel classes of antidiabetic drugs specifically targeting SKM.

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Received August 8, 2021 • Revision received November 21, 2021 • Accepted December 2, 2021 • Available online 6 December 2021

https://doi.org/10.1016/j.molmet.2021.101415
SMK expresses dozens of GPCRs that differ in their G protein-coupling properties [9]. We have shown that activation of SMK G_α-coupled receptors has a beneficial effect on whole-body glucose homeostasis by promoting glucose uptake into SKM [10]. In this study, we explored the in vivo metabolic effects caused by acute activation of SMK G_α-coupled receptors. Analysis of mutant mice that expressed a G_α-coupled DREADD (Gs DREADD, hereafter GsD) selectively in SMK (SMK-GsD mice) showed that acute stimulation of G_α signaling in SMK in vivo impairs glucose tolerance by decreasing glucose uptake into SMK. In parallel, we used selective agonists to stimulate β_2-adrenoceptors (β_2-AR) and the corticotropin releasing hormone receptor 2 (CRF_2), two receptors that are expressed at relatively high levels in SMK. We found that acute in vivo stimulation of these receptors did not mimic the metabolic effects observed in CNO-treated SMK-GsD mice, most likely due to the ability of these receptors to regulate hormone secretion from pancreatic islets. Our study provides an excellent example of how tissue expression patterns of receptors can affect the acute effects of GPCR agonists on whole-body glucose homeostasis.

2. METHODS

2.1. Mouse maintenance and diet
All mice were fed ad libitum and kept on a 12-h light/12-h dark cycle at room temperature (23 °C). Mice were maintained either on a standard mouse chow (7022 NIH-07 diet, 15% kcal fat, energy density 3.1 kcal/g, Envigo Inc.) or a high-fat diet (HFD, 60% kcal fat; #F3282, Bioserv). Transgenic mouse models received HFD after tamoxifen induction at the age of 10 weeks for a duration of at least 8 weeks. Lean (B6-M, C57BL/6N-Tac) and obese (DBA-B6-M, C57BL/6N-Tac, >8 weeks of HFD starting at the age of 6 weeks) WT mice were purchased from Taconic. Metabolic tests with mice fed standard chow were performed starting at the age of 10 weeks. Mice consuming HFD underwent metabolic tests starting at the age of 14 weeks (C57BL/6N-Tac WT mice) or 18 weeks (transgenic mice), respectively. All experiments were approved by the NIDDK Institutional Animal Care and Use Committee.

2.2. Generation of SMK-specific knock-out and knock-in mice
To generate SMK-specific knock-out (KO) mice, floxed mice (Adbrb2^fl/fl or Gnas^fl/fl mice) [11,12] were crossed with HSA-Cre(ER^{T2}) mice [13]. Mice used for these matings had been backcrossed onto a C57BL/6 background. To avoid potential developmental changes, Cre recombinase activity was induced in 8–12 week-old HSA-Cre(ER^{T2})-positive homozygous floxed mice by intraperitoneal injection of tamoxifen (2 mg per day dissolved in corn oil) for 5 consecutive days. Tamoxifen-injected Cre-negative homozygous floxed littermates served as control mice. To generate SMK-specific Gα-DREADD knock-in mice (SMK-GsD mice), heterozygous ROSA26-LSL-Gs-DREADD-CRE-luc mice [14] were crossed with HSA-Cre(ER^{T2}) mice [13]. To induce GsD expression in SMK, Cre-positive hemizygous ROSA26-LSL-Gs-DREADD-CRE-luc mice were injected with tamoxifen as described above. Tamoxifen-injected Cre-negative hemizygous littermates served as control mice in all experiments.

2.3. In vivo metabolic studies
All metabolic tests were performed on male littermates (age range 10–30 weeks) using standard protocols. For i.p. glucose tolerance tests (ipGTT), mice were fasted overnight for 15 h and blood glucose levels were measured before (0 min) and at defined time points after i.p. injection of normal saline containing glucose (2 g/kg for mice consuming normal chow; 1 g/kg for mice on HFD). For insulin tolerance tests (ITT), mice were fasted for 4 h and blood glucose levels were measured before (0 min) and at specific time points after i.p. injection of human insulin (0.75 or 1 IU/kg, as indicated; Humulin R, Eli Lilly). Blood glucose levels were determined using a blood glucose meter (Contour; Bayer). To study glucose-stimulated insulin secretion (GSIS), mice were fasted overnight for 15 h, and blood samples were collected in heparinized capillary tubes (Fisher Scientific) before (0 min) and 5, 15, and 30 min following the i.p. injection of glucose (2 or 1 g/kg, as indicated). Samples were centrifuged (5,000 g, 10 min, 4 °C) for plasma collection, and plasma insulin levels were determined using an ultra-sensitive mouse insulin ELISA kit (Crystal Chem, Inc). Where indicated, mice were co-injected or pre-injected i.p. with GPCR agonists (clenbuterol (1 mg/kg) or urocortin 2 (0.1–0.3 mg/kg)) or vehicle for acute treatment experiments.

2.4. Streptozotocin (STZ) model
We used a STZ treatment protocol that destroys about 70–80% of mouse β-cells [15]. Lean WT mice were fasted for 5 h and injected with a relatively low dose (50 mg/kg i.p.) of STZ for 5 consecutive days. Metabolic studies were initiated seven days after the last STZ injection.

2.5. Clozapine-N-Oxide (CNO) administration
To acutely stimulate G_α signaling in SMK, control and SMK-GsD mice received an i.p. injection of CNO (10 mg/kg), either in combination with glucose (ipGTT) or insulin (ITT), respectively.

2.6. In vivo [14C]2-deoxyglucose uptake
To measure glucose uptake into individual tissues in vivo, mice were fasted overnight for 14–16 h and then injected i.p. with saline containing 2 g/kg glucose and 10 µCi of [14C]2-deoxyglucose ([14C]2-DG) (PerkinElmer). Where indicated, 10 mg/kg CNO was added to acutely stimulate SMK G_α signaling in SMK-GsD mice. Mice were euthanized 2 h later, and SMK and other tissues were harvested. The tissue content of [14C]2-DG-6-phosphate was determined as a measure of SMK glucose uptake, as described previously [16].

2.7. Glycogen measurements
Glycogen content in SMK was determined using a glycogen assay kit (Cayman Chemical) according to the manufacturer’s instructions.

2.8. Plasma somatostatin and free fatty acids measurements
Blood samples were collected in EDTA tubes, centrifuged (5,000 g, 10 min, 4 °C) for plasma collection, and somatostatin or free fatty acid levels were determined using a somatostatin EIA kit (Phoenix Pharmaceuticals, Inc.) or the Wako NEFA HR (2) assay (Fujifilm Healthcare Solutions), respectively.

2.9. Static insulin release experiments with isolated mouse islets
Pancreatic islets were isolated from wild-type male mice (age: ~20 weeks; strain: C57BL/6N-Tac) by collagenase digestion and Histopaque (Sigma Aldrich, Cat # 1077–1) gradient separation and cultured overnight in RPMI 1640 medium (Thermo Fisher Scientific, Cat # 11,879,020) containing 5.5 mM glucose [17]. On the day of the experiments, islets were preincubated in Krebs–Ringer bicarbonate/HEPES buffer (KRB; 119 mM NaCl, 5 mM KCl, 2.5 mM CaCl_2, 1.2 mM MgCl_2, 1.2 mM KH_2PO_4, 25 mM NaHCO_3, 10 mM HEPES, 0.5% BSA, pH 7.4) containing 3 mM or 16 mM glucose for 30 min. Batches of ten islets were transferred to a 24-well plate (10 islets/well) and incubated in 400 µl of KRB medium with 3 or 16 mM glucose for 1 h at 37 °C, followed by
Figure 1: Acute CNO treatment of SKM-Gs-DREADD mice causes impaired glucose tolerance by reducing SKM glucose uptake. (A) Glucose tolerance test (GTT), lean control and SKM-GsD mice (overnight fasted) co-injected i.p. with glucose (2 g/kg) and CNO (10 mg/kg) (n = 6 or 7/group). (B) Area under the curve (AUC) of GTT shown in (A). (C) Insulin tolerance test (ITT), lean control and SKM-GsD mice (fasted for 4 h) co-injected i.p. with insulin (0.75 IU/kg) and CNO (10 mg/kg) (n = 7/group). (D) AUC of ITT shown in (C). (E) Glucose-stimulated insulin secretion (GSIS), lean control and SKM-GsD mice (overnight fasted) co-injected i.p. with glucose (2 g/kg) and CNO (10 mg/kg) (n = 10/group). (F) AUC of GSIS shown in (E). (G) GTT, HFD-induced obese control and SKM-GsD mice (overnight fasted) co-injected i.p. with glucose (1 g/kg) and CNO (10 mg/kg) (n = 7/group). (H) AUC of GTT shown in (G). (I) ITT, HFD-induced obese control and SKM-GsD mice (fasted for 4 h) co-injected i.p. with insulin (0.75 IU/kg) and CNO (10 mg/kg) (n = 6 or 7/group). Data were normalized to initial blood glucose values (30 mg/dl vs. 30 mg/dl for control and SKM-GsD mice, respectively). Two hours after injections, tissues were collected, and the [14C]2-DG-6-phosphate content was determined in the indicated tissues (n = 5–7/group). Quad, quadriceps muscle; WAT, white adipose tissue; BAT, brown adipose tissue. All studies were carried out on male mice that were at least 8 weeks old. Where indicated mice were fed a HFD for at least eight weeks. Data are presented as means ± SEM. Differences were tested for statistical significance using an unpaired two-tailed Student’s t-test (*p < 0.05, Figure 1B, D, H, M) or two-way ANOVA (*G, p < 0.05 for genotype effect; *T, p < 0.05 for time effect; #I, p < 0.05 for genotype × time interaction effect; Figure 1A, C, G, I) followed by Sidak’s post hoc test (adjusted *p < 0.05).

2.10 Western blotting experiments
Tissues were isolated quickly, frozen in liquid nitrogen, and stored at −80 °C until use. For western blotting studies, frozen tissues were homogenized in ice-cold RIPA buffer (Sigma Aldrich), and protein concentrations were determined using a BCA protein assay (Thermo Fisher Scientific). Protein extracts were separated on NuPAGE 4–12% Bis-Tris or, for high molecular mass proteins, on 3–8% Tris-Acetate gels (Thermo Fisher Scientific) and blotted onto nitrocellulose membranes (Bio-Rad). Membranes were blocked for 1 h at room temperature in TBS-T (0.1%) containing 5% BSA. Membranes were then incubated overnight with primary antibodies at 4 °C. Following three washing steps with TBS-T (0.1%), membranes were incubated with HRP-conjugated secondary antibodies for 1 h at room temperature. After thorough washing, proteins were visualized with SuperSignal West Dura Extended Duration Substrate (Thermo Fisher Scientific) on the c600 Imaging System (Azure Biosystems). Immunoreactive bands were quantified using Image J Software (NIH).

2.11 Statistics
Data are expressed as mean ± SEM for the indicated number of observations. Data were assessed for statistical significance by 2-way ANOVA tests, followed by the indicated post hoc tests, or by using a two-tailed unpaired Student’s t-test, as appropriate. A p-value of less than 0.05 was considered statistically significant. The specific statistical tests that were used are indicated in the figure legends.

3. RESULTS AND DISCUSSION
3.1. Acute stimulation of a Gs-coupled designer GPCR expressed in SKM decreases glucose uptake into SKM
Since individual GPCRs are widely expressed and systemic administration of agonists in vivo usually affects multiple tissues, we took advantage of DREADD technology to clearly define the role of acute SKM Gs stimulation in regulating glucose homeostasis. To explore the in vivo metabolic effects caused by acute stimulation of a Gs-linked
We generated mice that expressed a Gs-coupled DREADD (GsD) [18] under the transcriptional control of the SKM-specific human a-skeletal actin (HSA) promoter (SKM-GsD mice). Control and SKM-GsD mice did not show any metabolic differences in the absence of CNO (Meister et al. Nature Communications, in press). Following CNO (10 mg/kg i.p.) treatment, SKM-GsD mice maintained on regular chow or a HFD displayed a significant impairment in glucose tolerance (Figure 1A, B, G, H). In contrast to lean SKM-GsD mice (Figure 1C, D), HFD SKM-GsD mice displayed a slight deficit in insulin tolerance (Figure 1I, J) as compared to their control littermates. Acute administration of CNO to SKM-GsD and control mice had no significant effect on glucose-stimulated insulin secretion (GSIS, Figure 1E, F, K, L). To investigate whether glucose uptake by SKM was altered in CNO-treated SKM-GsD mice, we co-injected (i.p.) SKM-GsD and control mice with CNO (10 mg/kg) and glucose (2 g/kg) containing 10 μCi of [14C]-2-deoxyglucose ([14C]-2-DG). Accumulation of the [14C]-2-DG...
metabolite, \(^{14}\text{C}-2\text{-DG}-6\text{-phosphate}\), was determined as a measure of glucose uptake. We found that CNO-treated SKM-GsD mice showed reduced glucose uptake into SKM (quadriceps muscle), as compared to CNO-injected control littermates (Figure 1M). Glucose uptake into adipose tissues, heart, and brain remained unaffected following CNO injection of SKM-GsD mice (Figure 1M). Taken together, these data strongly suggest that acute stimulation of SKM Gs signaling impairs blood glucose uptake into SKM. In contrast, we previously demonstrated that chronic activation of SKM Gs signaling leads to improved glucose tolerance without affecting insulin sensitivity or secretion (Meister et al. Nature Communications, in press).

We next mined RNA-Seq data to identify GPCRs expressed by mouse quadriceps muscle. This analysis showed that the \(\beta_2\)-adrenergic receptor (\(\beta_2\)-AR) and the corticotropin-releasing hormone receptor 2 (CRF2), two Gs-coupled receptors, were highly expressed in SKM (Supplemental Fig. 1). These receptors are also expressed in other metabolically relevant tissues, although at lower levels [9]. Previous studies have led to contradictory results regarding the roles of these receptors in regulating glucose homeostasis [19–25].

3.2. Acute \(\beta_2\)-AR activation improves glucose tolerance only in healthy lean mice

To study the acute in vivo metabolic effects of \(\beta_2\)-AR stimulation, we initially administered clenbuterol, a \(\beta_2\)-AR-selective agonist, to lean wild type (WT) mice. After a single clenbuterol injection (1 mg/kg i.p.), lean WT mice (either freely fed or fasted for 15 h) showed only slight differences in blood glucose excursions after a single clenbuterol injection (1 mg/kg i.p.), as compared to control mice injected with vehicle alone (Figure 2A, B). The clenbuterol bolus caused a significant increase in plasma insulin levels (Figure 2C, D). During a glucose tolerance test (GTT), lean WT mice co-injected i.p. with glucose (2 g/kg) and clenbuterol showed significantly reduced blood glucose levels, as compared to control mice injected with glucose alone (Figure 2E, F). Acute clenbuterol injection led to a dramatic increase in GSIS in lean WT mice (Figure 2G, H). An insulin tolerance test (ITT) demonstrated that co-injection of insulin (0.75 IU/kg i.p.) with clenbuterol decreased insulin sensitivity (Figure 2I, J). Furthermore, clenbuterol injection during the ITT caused an increase in plasma free fatty acid (FFA) levels (Figure 2K), probably due to the activation of \(\beta_2\)-ARs expressed by adipose tissue [26].

We tested whether acute clenbuterol administration was able to improve glucose homeostasis in a model of diet-induced obesity. WT mice that were fed a high fat diet (HFD) for at least 8 weeks were characterized in several metabolic tests, either in the absence or presence of co-injected clenbuterol. When obese HFD mice were subjected to a GTT, i.p. co-injection of glucose (1 g/kg) and clenbuterol showed no effect on glucose tolerance (Figure 2L, M). As observed with lean WT mice, acute clenbuterol injection of obese WT mice led to a severe decrease in insulin tolerance (Figure 2N, O). In contrast to the outcome of GSIS studies with lean WT mice (Figure 2G, H), acute clenbuterol treatment did not potentiate GSIS in obese WT mice (Figure 2P, Q).

To mimic the loss of \(\beta\)-cell mass characteristic for advanced T2D, we used a streptozotocin (STZ) treatment protocol known to destroy about 70–80% of mouse \(\beta\)-cells [15]. When STZ-treated mice were co-injected i.p. with glucose (1 g/kg) and clenbuterol, they displayed even more pronounced glucose intolerance than STZ-treated mice treated with glucose alone (Figure 2R, S). This phenotype is opposite to the one observed with lean healthy mice (Figure 2E, F), mimicking the effects of SKM-GsD stimulation in vivo (Figure 1A, B). These results strongly suggest that the clenbuterol-induced improvements in glucose tolerance observed in lean healthy mice (Figure 2E, F) are primarily driven by enhanced insulin release (Figure 2G, H). To test the ability of clenbuterol to directly stimulate pancreatic insulin secretion, we incubated isolated WT mouse islets in low (3 mM) and high glucose (16 mM) medium in the presence or absence of clenbuterol (1 \(\mu\)M). Clenbuterol treatment significantly enhanced GSIS (Figure 2T), indicating that activation of islet \(\beta_2\)-ARs promotes insulin secretion [27,28]. Because of the high expression of the \(\beta_2\)-AR in SKM and the striking phenotypes displayed by the SKM-GsD mice, we next explored the potential contribution of SKM \(\beta_2\)-ARs to the acute in vivo metabolic effects of clenbuterol. To address this question, we used mice that lacked \(\beta_2\)-ARs selectively in SKM (SKM-\(\beta_2\)-AR-KO mice).

Acute clenbuterol injection retained the ability to trigger improved glucose tolerance in SKM-\(\beta_2\)-AR-KO (Figure 3A, B), consistent with the concept that this beneficial metabolic effect was mainly driven by increased insulin release (Figure 2G, H, T). However, we observed a slight reversal of the clenbuterol-mediated impairment in insulin sensitivity in SKM-\(\beta_2\)-AR-KO (Figure 3C, D), suggesting that the clenbuterol effect during the ITT is at least partially dependent on the activation of SKM \(\beta_2\)-ARs.

To explore the potential roles of SKM \(\beta_2\)-ARs in regulating the responsiveness of SKM to insulin, we co-injected lean WT mice with insulin in the presence or absence of clenbuterol and isolated quadriceps muscles 15 min later. While clenbuterol had no significant effect on insulin-induced Akt or GSK3\(\beta\) phosphorylation (Supplemental Figs. 2A and 8), co-injection of insulin with clenbuterol led to a significant increase in the inhibitory phosphorylation of glycogen synthase at S641 in WT mice (Figure 3E, F). In agreement with this observation, glycogen levels were significantly decreased in SKM of WT mice that were co-injected with insulin and clenbuterol, as compared to mice injected with insulin alone (Figure 3G). This clenbuterol effect was not observed in SKM derived from SKM-\(\beta_2\)-AR-KO mice, indicating that the clenbuterol-induced decrease in SKM glycogen levels is mediated by SKM \(\beta_2\)-ARs (Figure 3H).

Our data strongly suggest that acute injection of clenbuterol improves glucose tolerance primarily by increasing plasma insulin levels through direct stimulation of islet \(\beta_2\)-ARs (Figure 2G–H, T). In agreement with this observation, it has been reported that activation of \(\beta_2\)-ARs expressed by mouse or human pancreatic \(\beta\)-cells promotes insulin secretion [27,28]. In contrast to a previous proposal [25], acute SKM \(\beta_2\)-AR signaling did not contribute to this beneficial metabolic effect, since acute clenbuterol injection retained the ability to improve glucose tolerance in mice lacking \(\beta_2\)-ARs in SKM (Figure 3A, B). These new data clearly demonstrate the importance of using cell type-specific mutant mouse models in clarifying the physiological roles of widely expressed GPCRs like the \(\beta_2\)-AR.

Acute clenbuterol injection failed to improve whole-body glucose homeostasis in obese mice consuming a HFD (Figure 2L, M) or after STZ-induced loss of \(\beta\)-cells (Figure 2R, S). The most likely explanation for these findings is that the clenbuterol-induced decrease in insulin tolerance (Figure 2L, J, N, O) is due to elevated plasma FFA levels caused by increased lipolysis (Figure 2K). We have recently shown that selective activation of Gs-coupled receptors in adipose tissue induces lipolysis [29]. It is likely that clenbuterol-stimulated glycogen breakdown in SKM also contributes to impaired insulin tolerance (Figure 3E, G). Increased glycogenolysis is predicted to slow down glucose influx into SKM [30]. The negative metabolic effect of acute stimulation of SKM \(\beta_2\)-ARs on whole-body glucose homeostasis becomes evident in the mouse model of HFD-induced insulin resistance and STZ-induced \(\beta\)-cell loss, where acute clenbuterol was unable to increase insulin.
secretion (obese mice; Figure 2P) or caused further impaired glucose tolerance (STZ-treated mice; Figure 2R, S).

It is well known that activation of the sympathetic nervous system promotes glycogen breakdown in SKM cells via activation of Gβγ-coupled β2-ARs [31]. The resulting increase in intracellular glucose levels is most probably the cause of the impaired SKM glucose uptake observed after acute Gs stimulation in SKM cells. In comparison, chronic stimulation of SKM Gs signaling by clenbuterol led to a significant increase in glucose uptake, most likely resulting from metabolic reprogramming of SKM cells causing enhanced glucose utilization (Meister et al. Nature Communications, in press).

3.3. Acute CRF2 activation increases blood glucose levels

As mentioned earlier, the Gβγ-coupled CRF2 receptor is enriched in SKM tissue (Supplemental Fig. 1). An acute bolus of urocortin 2 (UCN2; 0.1 mg/kg i.p.), a selective CRF2 agonist [32], increased blood glucose levels in overnight fasted and ad libitum fed lean WT mice (Figure 4A, B, E, F). UCN2 injection reduced basal plasma insulin levels in these mice (Figure 4C, D, G, H). Co-injection of glucose (2 g/kg) with UCN2 resulted in impaired glucose tolerance (Figure 4I, J) and reduced GS synthesis (Figure 4K).

In vitro studies with isolated pancreatic islets have shown that stimulation of CRF2 receptors causes an increase in somatostatin release [33]. We did not observe differences in plasma somatostatin levels following the co-administration of glucose and UCN2, as compared to glucose alone (Supplemental Fig. 3A). Surprisingly, UCN2 (100 nM) did not affect insulin release in isolated pancreatic islets derived from WT mice during static incubation assays (Supplemental Fig. 3B). Plasma FFA levels also remained unaffected by the UCN2 bolus (Supplemental Fig. 3C), and UCN2 injection did not affect insulin sensitivity in lean WT mice (Figure 4M, N).

UCN2 decreased glucose tolerance in HFD-induced obese mice (Figure 4O, P). In contrast to lean mice, UCN2 reduced insulin sensitivity in a time-dependent manner in obese mice (Figures 4Q, R and Supplemental Fig. 3D). The effect was most prominent when UCN2 was administered 15 min prior to the insulin bolus (Figure 4S, T).

Previous studies examining the role of SKM CRF2 receptors in metabolic disorders have yielded conflicting results. Both increase [36] and decrease [19] in UCN2-stimulated SKM glucose uptake have been described using in vitro cell systems. UCN2- and CRF2-KO mice show improvements in whole-body glucose homeostasis [34,37], but mice overexpressing UCN2 showed the same phenotype [20,36]. Our study clearly demonstrates that acute UCN2 injection in vivo impairs glucose homeostasis by decreasing insulin secretion (Figure 4A–K).

CRF2 receptors are highly expressed in pancreatic β-cells [33], and previous in vitro studies suggest that UCN3, another CRF2 receptor agonist, can modulate somatostatin and glucagon secretion from pancreatic islets [33,38]. In this study, we did not observe an increase in plasma somatostatin levels following co-injection of glucose and UCN2 (Supplemental Fig. 3A), and UCN2 (100 nM) had no effect on

Figure 3: SKM β2-AR-mediated metabolic effects of acute clenbuterol injection. (A) Glucose tolerance test (GTT), lean control and SKM-β2-AR-KO mice (overnight fasted) co-injected i.p. with glucose (2 g/kg) and clenbuterol (1 mg/kg) or vehicle (n = 5/group). (B) Area under the curve (AUC) of GTT shown in (A). (C) Insulin tolerance test (ITT, 0.75 IU/kg insulin) performed with 4 h-fasted control and SKM-β2-AR-KO mice (n = 6 or 12/group). (D) AUC of ITT shown in (C). (E–G) Drug effects on SKM glycogen synthase (GS) phosphorylation (pGS) and SKM glycogen levels. Lean WT mice were fasted for 4 h and then co-injected i.p. with insulin (0.75 IU/kg) and clenbuterol (1 mg/kg) or vehicle. SKM tissues (quadriceps muscle) were isolated 15 min post-injection (n = 4 or 5/group), followed by the determination of pGS and total GS protein levels (E, F) and SKM glycogen content (G). (H) Glycogen levels in SKM (quadriceps muscle) of lean control and SKM-β2-AR-KO mice 15 min after i.p. co-injection of insulin (0.75 IU/kg) and clenbuterol (1 mg/kg) or vehicle (n = 3 or 4/group). All studies were carried out with male mice that were at least ten weeks old. Data presented as means ± SEM. Differences were tested for statistical significance using an unpaired two-tailed Student’s t-test (*p < 0.05) or two-way ANOVA (*C, p < 0.05 for clenbuterol effect, #I, p < 0.05 for clenbuterol × genotype interaction effect).
Figure 4: Metabolic effects of acute urocortin 2 injection. (A) Blood glucose levels in lean WT mice (overnight fasted) following an i.p. injection of urocortin 2 (UCN2, 0.1 mg/kg) or vehicle (n = 5/group). (B) Area under the curve (AUC) of (A). (C) Plasma insulin levels in lean WT mice (overnight fasted) following an i.p. injection of UCN2 (0.1 mg/kg) or vehicle (n = 5/group). (D) Area under the curve (AUC) of (C). (E) Blood glucose levels in lean, freely fed WT mice following an i.p. injection of UCN2 (0.1 mg/kg) or vehicle (n = 5/group). (F) AUC of (E). (G) Plasma insulin levels in lean, freely fed WT mice following an i.p. injection of UCN2 (0.1 mg/kg) or vehicle (n = 5/group). (H) AUC of (G). (I) Glucose tolerance test (GTT), lean WT mice (overnight fasted) co-injected i.p. with glucose (2 g/kg) and UCN2 (0.1 mg/kg) or vehicle (n = 6/group). (J) AUC of GTT shown in (I). (K) Glucose-stimulated insulin secretion (GSIS), lean WT mice (overnight fasted) co-injected i.p. with glucose (2 g/kg) and UCN2 (0.1 mg/kg) or vehicle (n = 6/group). (L) AUC of GSIS shown in (K). (M) Insulin tolerance test (ITT), lean WT mice (fasted for 4 h) co-injected i.p. with insulin (0.75 IU/kg) and UCN2 (0.1 mg/kg) or vehicle (n = 6/group). (N) AUC of ITT shown in (M). (O) GTT, HFD-induced obese WT mice (overnight fasted) co-injected i.p. with glucose (1 g/kg) and UCN2 (0.1 mg/kg) or vehicle (n = 4/group). (P) AUC of GTT shown in (O). (Q) ITT, HFD-induced obese WT mice (fasted for 4 h) co-injected i.p. with insulin (0.75 IU/kg) and UCN2 (0.3 mg/kg) or vehicle (n = 5 or 6/group). (R) AUC of ITT shown in (O). (S) ITT, HFD-induced obese WT mice (fasted for 4 h) co-injected i.p. with UCN2 (0.3 mg/kg) or vehicle, followed by an insulin bolus (0.75 IU/kg) (n = 11/group). (T) AUC of ITT shown in (S). All studies were carried out on male mice that were at least 8 weeks old. Where indicated, mice were fed a HFD for at least eight weeks. Data presented as means ± SEM. Differences were tested for statistical significance using an unpaired two-tailed Student’s t-test (*p < 0.05, Figure 4A, B, D, E, F, H, J, K, P, R, T) or two-way ANOVA (*U, p < 0.05 for UCN2 effect; *T, p < 0.05 for time effect; #I, p < 0.05 for UCN2 x time interaction effect; Figure 4I, M, O, Q, S) followed by Sidak’s post hoc test (*adjusted p < 0.05).
insulin release from isolated WT pancreatic islets (Supplemental Fig. 3B). One likely explanation for the discrepancy between the in vivo insulin data (Figure 4C,D, G, H, K) and the outcome of the in vitro studies with isolated islets (Supplemental Fig. 3B) is that specific circulating factors or neuronal inputs that contribute to UCN2-mediated insulin release in vivo are absent in the islet incubation assay. Studies with SKM-Gs-KO mice showed that the metabolic effects of acute UCN2 administration are not dependent on SKM Gs signaling (Supplemental Fig. 4). Further studies with conditional KO mice are necessary to determine whether the metabolic improvements seen after long-term treatment with UCN2 peptides [39] or UCN2 gene transfer [36] are indeed mediated by SKM CRF2 receptors.

Using a chemogenetic approach we showed that acute activation of SKM Gs signaling decreased glucose uptake into SKM in vivo, leading to significant impairments in whole-body glucose homeostasis. However, the acute in vivo stimulation of two SKM-enriched Gs-coupled receptors (B2-AR and CRF2) resulted in metabolic effects that were mostly SKM-independent and predominantly mediated by altered insulin levels, probably due to altered insulin secretion from pancreatic islets. Our study further highlights the need to distinguish between acute and long-term effects of GPCR agonists. Different drug exposure patterns in vivo can result in different tissue-specific effects, and in vitro and ex vivo data need to be carefully validated in whole-body settings in order to draw conclusions about the ultimate metabolic outcome. Our data provide an excellent example for the concept that acute administration of GPCR agonists needs to be complemented by chronic agonist treatment studies to properly assess translationally relevant metabolic outcomes.

In the present study, we demonstrated that acute activation of SKM Gs signaling caused an impairment in glucose tolerance. In contrast, a recent study (Meister et al. Nature Communications, in press) demonstrated that chronic activation of SKM Gs signaling leads to improved whole-body glucose homeostasis in both lean and obese mice. This latter finding raises the possibility that drugs able to selectively stimulate SKM Gs signaling may prove to be useful as novel antidiabetic drugs.

ACKNOWLEDGEMENTS

This research was funded by the Intramural Research Program of the National Institute of Diabetes and Digestive and Kidney Diseases (NIDDK, NIH). We thank Dr. Daniel Metzger (Université de Strasbourg, Illkirch, France) for providing the HSA-CreERT2 mice, Dr. Rebecca Berdeaux (Houston Medical School, Houston, USA) for providing the ROSA26-LSL-Gs-DREADD-CRE-luc mice, and Dr. Gerard Karsenty (Institute of Diabetes and Digestive and Kidney Diseases (NIDDK, NIH). We thank Dr. Erik A. Richter (University of Copenhagen, Denmark) for his advice and valuable experimental data. M.C. and L.S.W. provided mutant mouse models and helpful advice. J.R.K., M.K. and T.E.J. supervised part of the studies and provided critical guidance regarding the preparation of the manuscript. J.M., D.B.J.B and J.W. wrote and revised the manuscript. J.M. prepared the first draft. All authors approved the final version of the manuscript.

CONFLICT OF INTEREST

The authors have declared that no conflict of interest exists.

APPENDIX A. SUPPLEMENTARY DATA

Supplementary data to this article can be found online at https://doi.org/10.1016/j.molmet.2021.101415.

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