FGF19 action in the brain induces insulin-independent glucose lowering

Gregory J. Morton,1 Miles E. Matsen,1 Deanna P. Bracy,2 Thomas H. Meek,1 Hong T. Nguyen,1 Darko Stefanovski,3 Richard N. Bergman,3 David H. Wasserman,2 and Michael W. Schwartz1

1Diabetes and Obesity Center of Excellence, Department of Medicine, University of Washington, Seattle, Washington, USA.
2Department of Molecular Physiology and Biophysics, Vanderbilt School of Medicine, Nashville, Tennessee, USA.
3Diabetes and Obesity Research Institute, Cedars-Sinai Medical Center, Los Angeles, California, USA.

Insulin-independent glucose disposal (referred to as glucose effectiveness [GE]) is crucial for glucose homeostasis and, until recently, was thought to be invariable. However, GE is reduced in type 2 diabetes and markedly decreased in leptin-deficient ob/ob mice. Strategies aimed at increasing GE should therefore be capable of improving glucose tolerance in these animals. The gut-derived hormone FGF19 has previously been shown to exert potent antidiabetic effects in ob/ob mice. In ob/ob mice, we found that systemic FGF19 administration improved glucose tolerance through its action in the brain and that a single, low-dose i.c.v. injection of FGF19 dramatically improved glucose intolerance within 2 hours. Minimal model analysis of glucose and insulin data obtained during a frequently sampled i.v. glucose tolerance test showed that the antidiabetic effect of i.c.v. FGF19 was solely due to increased GE and not to changes of either insulin secretion or insulin sensitivity. The mechanism underlying this effect appears to involve increased metabolism of glucose to lactate. Together, these findings implicate the brain in the antidiabetic action of systemic FGF19 and establish the brain’s capacity to rapidly, potently, and selectively increase insulin-independent glucose disposal.

Introduction

In addition to insulin secretion and insulin sensitivity (Si), insulin-independent mechanisms are critical to normal glucose homeostasis (1). Given that such mechanisms contribute at least as much to normal glucose tolerance as does insulin itself (1), it is surprising how little is known about them. This lack of research interest can be traced to the widespread perception of insulin-independent glucose disposal, which has been termed glucose effectiveness (GE; the ability of glucose to promote its own disposal, independently of insulin), as the fixed, obligate, and unregulated mechanism whereby insulin-insensitive tissues meet ongoing fuel needs. Yet glucose intolerance and type 2 diabetes (T2D) are characterized by decreases of GE as well as of insulin secretion and action (1). In leptin-deficient ob/ob mice, for example, GE was shown recently to be reduced by approximately 70%, based on minimal model analysis of data obtained from a frequently sampled i.v. glucose tolerance test (FSIGT) (2). Based on its importance to glucose homeostasis, we hypothesized that any intervention capable of normalizing glucose tolerance in these mice should do so, at least in part, by increasing GE.

FGF19 and its rodent homolog, FGF15, is one of three members of the family of hormonal FGFs. FGF19 is secreted by enterocytes located in the distal small intestine following activation of the nuclear bile acid receptor, FXR, by bile acid binding (3). In addition to its well-established role in the negative feedback control of hepatic bile acid synthesis (4, 5), FGF19 exerts potent antidiabetic effects in rodent models, including ob/ob mice (6). Similarly, transgenic overexpression of FGF19 improves glucose tolerance in diet-induced obese mice (6, 7), whereas FGF15-deficient mice display impaired glucose tolerance that is corrected by FGF19 administration (8).

Although initially thought to act primarily via FGFR4 receptors in the liver, the antidiabetic effects of FGF19 appear to involve a different FGF receptor subtype, because they are preserved in FGR4-deficient mice (9). Instead, several findings implicate FGFR1 in this effect. First, systemic administration of a FGF19 variant that activates FGFR1 but not FGFR4 ameliorates diabetes in ob/ob mice, while this effect is absent for a FGF19 variant that activates FGFR4 but not FGFR1 (10). Second, the potent glucose-lowering effect of FGF19 is mimicked by monoclonal antibodies that selectively activate a specific FGFR1 isoform (FGFR1c) (11). Although activation of FGFR1 in peripheral tissues, such as brown and white adipose tissue, may contribute to glucose lowering, this receptor is also expressed in mediobasal hypothalamic areas involved in glucose homeostasis (12, 13), and administration of FGF19 directly into the brain improves glucose tolerance in both ob/ob mice (6) and diet-induced obese rats (14).

The current studies were undertaken to determine whether the antidiabetic effect of systemically administered FGF19 involves a central site of action and, if so, to determine the contribution(s) made by insulin-dependent and -independent mechanisms to this effect. We report that, in ob/ob mice, the glucose-lowering effect of systemic FGF19 is reduced by approximately 50% when its action in the brain is blocked by i.c.v. administration of an FGFR antagonsist. Thus, the brain plays a key role in the antidiabetic effects of this hormone. To investigate how FGF19 action in the brain affects systemic glucose metabolism, we performed a FSIGT 2 hours after i.c.v. injection of either FGF19 or its vehicle in ob/ob mice as well as in vehicle-injected C57BL/6 controls. Glucose and insulin data from the FSIGT were analyzed using the minimal model method, which has been widely used in humans (1) and was recently validated in mice (including in ob/ob mice) (2), as a tool with which to measure insulin secretion, Si, and GE. Surprisingly, we found that the potent antidiabetic effect of i.c.v. FGF19 in ob/ob mice did not involve increases of either insulin secretion or Si. Instead, the
Effect was mediated entirely by a 3-fold increase of GE. Furthermore, our data suggest that reduced GE in ob/ob mice is linked to reduced glycolysis of glucose to lactate and that FGF19 action in the CNS of these animals reverses this defect. Thus, whereas insulin-independent glucose lowering has heretofore been considered invariant, obligate, and unregulated, our findings reveal the brain’s inherent capacity to rapidly, potently, and selectively increase glucose disposal via this mechanism.

Results

Effect of i.c.v. FGF19 on glucose tolerance in ob/ob mice. Since i.c.v. FGF19 can reduce food intake and body weight as well as blood glucose levels (6, 14), we initially sought to identify a protocol in which effects on glucose metabolism are not confounded by changes of energy balance. After verifying an earlier report (6) showing that chronic (i.e., over 6 days) i.c.v. administration of FGF19 exerts potent glucose lowering in ob/ob mice (data not shown), we asked whether this effect can be detected following just a single i.c.v. injection of FGF19, with food removed during the interval (1.5 hours) between i.c.v. injection and glucose tolerance testing. We found that, in ob/ob mice (mean body weight: 49.8 ± 1.2 g), i.c.v. injection of FGF19 lowered fasting blood glucose levels (P < 0.05) and dose-dependently improved glucose tolerance relative to that of vehicle-treated controls (P < 0.05) (Figure 1, A and B). Since the mice did not have access to food following i.c.v. injection, and, as expected, this beneficial effect was fully blocked by i.c.v. pretreatment with PD173074 at a dose that had no effect when given alone (Figure 2, A and B).

Prior to undertaking this study, we verified that, when given i.c.v., the inhibitor blocks the response to i.c.v. FGF19. Consistent with our earlier observations, a single i.c.v. injection of FGF19 lowered fasting blood glucose levels and significantly improved i.p. glucose tolerance in ob/ob mice that received an i.c.v. vehicle pretreatment injection, and, as expected, this beneficial effect was fully blocked by i.c.v. pretreatment with PD173074 (Figure 2, C and D), the glucose-lowering effect of systemic FGF19 appears to be mediated at least in part via a central mechanism.

Effect on glucose tolerance of a single injection FGF19 given either centrally or systemically to ob/ob mice. (A and C) Glucose tolerance (0.5 g/kg) and (B and D) the integrated area under the glucose curve (AUCglucose) in leptin-deficient ob/ob animals that received either a single, acute i.c.v. or i.p. injection of FGF19 at either 1 or 3 μg or its vehicle. Data represent mean ± SEM. *P < 0.05 vs. i.c.v. vehicle. #P < 0.05 vs. i.c.v. FGF19 (1 μg).
levels were similar in i.c.v. vehicle-treated ob/ob and WT mice, basal plasma insulin levels were markedly elevated in the former group, consistent with previous evidence of severe insulin resistance in these animals (2). Similarly, glucose tolerance was markedly impaired in i.c.v. vehicle-treated ob/ob mice relative to that in WT controls following an i.v. glucose bolus (Figure 3, A and B), despite a markedly increased acute insulin response to glucose (a measure of first-phase insulin secretion) and sustained elevation of plasma insulin levels throughout the study (Figure 3, C and D).

As expected, this marked impairment of glucose tolerance was significantly improved in ob/ob mice that received FGF19 (P < 0.05) (Figure 3, A and B). Based on the area under the glucose curve, i.c.v. FGF19 treatment corrected the abnormality of glucose tolerance of ob/ob mice by 65% (P < 0.05). Therefore, within 2 hours of a single i.c.v. injection FGF19 dramatically improved glucose tolerance in ob/ob mice. As this improvement occurred despite no difference in plasma insulin levels at any time during the glucose tolerance test (Figure 3, C and D), it cannot be attributed to increased insulin secretion. We therefore turned our attention to its effects on the two other determinants of glucose tolerance, Si and insulin-independent glucose disposal.

To determine whether the effect of FGF19 to increase GE involves reductions of either plasma glucagon or nonesterified fatty acid (FFA) levels, we perfused: 340 ± 51 mg/dl for vehicle vs. 241 ± 34 mg/dl for FGF19; P < 0.05). However, the glucose-lowering effect of FGF19 was not associated with significant changes in the plasma level of either glucagon or FFAs (Figure 4, C and D).

Liver parameters. As a first step to investigate whether changes of liver glucose metabolism contributed to the glucose-lowering effect of i.c.v. FGF19, we measured baseline liver glycogen and triglyceride content as well as hepatic expression of gluconeogenic
genes in the absence of a glucose challenge. Our data show that while hepatic levels of G6Pase were significantly reduced in ob/ob mice treated with i.c.v. FGF19 relative to vehicle-treated ob/ob and WT controls, there were no differences in hepatic Pepck mRNA levels or in either liver glycogen or triglyceride content (Figure 5).

To determine whether the effect of i.c.v. FGF19 to enhance disposal of a glucose load is associated with increased liver glycogen content, as expected if the mechanism involves increased liver glucose uptake and glycogen synthesis, we measured liver glycogen content 30 minutes after an i.p. glucose bolus (0.5 g/kg) in ob/ob animals that had received i.c.v. injection of either vehicle or FGF19 (3 μg) 2 hours earlier. Consistent with our previous observations, central FGF19 administration reduced blood glucose levels relative to those of vehicle-treated controls both at baseline and 30 minutes after the glucose bolus (baseline: 179 ± 31 mg/dl for vehicle vs. 106 ± 26 mg/dl for FGF19; P < 0.05; 30 minutes: 330 ± 46 mg/dl for vehicle vs. 243 ± 5 mg/dl for FGF19; P < 0.05). However, the glucose-lowering effect of FGF19 was not associated with a significant increase in hepatic glycogen levels (9.66 ± 0.93 mg/g tissue for vehicle vs. 9.37 ± 0.37 mg/g tissue for FGF19; P = NS). These results indicate that, in ob/ob mice, the effect of FGF19 to improve glucose tolerance, selectively increase GE, and raise plasma lactate levels occurs despite no change in liver glycogen content.

Role of melanocortin signaling in glucoregulatory effects of i.c.v. FGF19. As the melanocortin pathway is implicated in central regulation of glucose metabolism by leptin (18, 19), insulin (20, 21), and glucose, we asked whether (a) FGF19 activates proopiomelanocortin (POMC) neurons in the arcuate nucleus (as judged by induction of c-Fos staining) and (b) whether intact melanocortin signaling is required for the glucose-lowering effect of i.c.v. FGF19. To facilitate detection of c-Fos immunoreactivity in POMC neurons, we studied the effect of i.c.v. FGF19 in C57BL/6 mice in which POMC cells are fluorescently labeled (Pomc-Tau-GFP mice). As depicted in Figure 6, there was no effect of i.c.v. FGF19 to induce c-Fos in labeled POMC cells relative to i.c.v. vehicle-treated controls.

To determine whether melanocortin signaling is required for the glucose-lowering effects of central FGF19, we asked whether it lowers fasting blood glucose levels or improves glucose tolerance in melanocortin-4 receptor–deficient (Mc4r-deficient) mice. Indeed, i.c.v. FGF19 clearly exerts a glucose-lowering action in these animals relative to that in i.c.v. vehicle-treated animals (P < 0.05) (Figure 6, C and D). Therefore,
FGF19-induced glucose lowering is mediated at least in part via a melanocortin-independent mechanism.

Discussion

Observations made in a rat model of uncontrolled, insulin-deficient diabetes suggest that the brain has the capacity to lower blood glucose levels via insulin-independent means. Specifically, continuous or repeated daily i.c.v. leptin administration can normalize elevated blood glucose levels in these animals despite persistent, severe insulin deficiency (22–25). Building upon this initial finding, we hypothesized that, since the insulin-independent component of glucose tolerance, GE, is markedly reduced in ob/ob mice, this defect will be remedied by FGF19 (2), because it has potent antidiabetic effects in these animals. We further hypothesized that since FGF19 is effective following either peripheral or low-dose central administration in these animals (6), its effect to increase GE is mediated centrally. Our finding that glucose lowering induced by systemic FGF19 administration in ob/ob mice is attenuated by central administration of an inhibitor of FGF receptors implicates the brain in this effect. Moreover, centrally administered FGF19 improves glucose tolerance in these animals via a mechanism that cannot be explained by changes of insulin secretion or insulin action but arises instead from a 3-fold increase of GE. Although a detailed understanding of mechanisms mediating this effect awaits further study, our data support a model in which central FGF19 increases GE by stimulating glycolysis of glucose to lactate and in which the neuronal systems involved are at least partially independent of melanocortin signaling. Taken together, these data implicate the CNS in the glucose-lowering effect of FGF19 and, to our knowledge, are the first to establish the ability of the brain to improve glucose tolerance by rapidly, potently, and selectively increasing insulin-independent glucose disposal. These findings therefore establish the existence of a novel brain mechanism for insulin-independent control of blood glucose levels.

Growing evidence suggests that the CNS plays a key role in the regulation of glucose homeostasis. In obese rodents, increased hypothalamic input from hormonal (i.e., insulin and leptin) or nutrient-related (i.e., FFA) signals improves hepatic Si via a mechanism involving vagal input to the liver, whereas disruption of this hypothalamic input causes insulin resistance (26–30). In addition to these effects, the recent finding that continuous i.c.v. infusion of leptin completely normalizes glycemia in rats with streptozotocin-induced diabetes (22–25) suggests that the brain has the capacity to potently lower glucose levels via mechanisms that are insulin independent. To investigate the hypothesis that FGF19 has the capacity to lower blood glucose by mechanisms that are insulin independent, we used minimal model analysis of glucose and insulin data from a FSIGT, an approach developed in humans and recently validated for use in mice (2). We found that, in ob/ob mice, a single i.c.v. injection of a low dose of FGF19 improved glucose tolerance markedly, despite having no effect on either insulin secretion or action. The dramatic effect of centrally administered FGF19 to increase GE is remarkable, in that with the exception of vigorous exercise (31), there is no precedent for rapid upregulation of insulin-independent glucose disposal, much less by the brain.

The brain’s inherent capacity to promote glucose lowering through insulin-independent mechanisms is novel in the sense that previous work in this area has focused largely on hypothalamic control of hepatic Si (32, 33). That insulin-independent effects were not identified in such studies could reflect the fact that the methods used are not optimized to detect them. The euglycemic, hyperinsulinemic clamp method, widely accepted as the “gold standard” for quantitative assessment of glucose metabolism and Si, has been used extensively in human as well as rodent studies. By comparison, although minimal model analysis of FSIGT data has seen wide usage in human research, it was only validated for use in mice in the last year (2). This method measures not only Si and GE, but also first-phase insulin secretion and the disposition index (the product of Si and insulin secretion, a validated predictor of progression to T2D in humans). By comparison, the clamp (but not the minimal model analysis) quantifies rates of glucose production and utilization and, with some modification, can assess rates of glycogenolysis, gluconeogenesis, and glucose uptake into specific (nonhepatic) tissues.
Si is defined in clamp studies as the increase of glucose disposal (or decrease of glucose appearance) observed when insulin levels are raised while glucose levels are held constant. Stated differently, changes in rates of glucose production or disposal measured in the face of high insulin are attributed solely to the action of insulin. However, in the event that insulin-independent glucose disposal (or insulin-independent suppression of hepatic glucose production) is also increased by the intervention under study, the extent to which observed changes are attributable to insulin action cannot be known with certainty, since some of the glucose infused during the clamp (to maintain euglycemia in the presence of high insulin) would have been disposed of via insulin-independent mechanisms. Thus, whether insulin-independent mechanisms contributed to some of the centrally mediated effects attributed previously to changes of Si (32, 33) awaits additional study.

The insulin-independent component of glucose disposal in sedentary subjects has historically been considered to be a fixed, obligate, and invariant means whereby insulin-independent tissues obtain glucose primarily via the process of glucose mass action, by which increasing concentrations of glucose passively drive glucose into cells in a largely unregulated manner (1). It is for this reason that research into mechanisms underlying insulin-independent glucose disposal has languished in comparison with the substantial effort and progress made toward understanding insulin secretion and its action in peripheral tissues. Nevertheless, data from clinical studies performed years ago offer evidence suggestive of regulation. For one, i.v. infusion of glucagon-like peptide-1 (GLP-1) in humans improves i.v. glucose tolerance by increasing GE as well as insulin secretion (34), while, conversely, GE was reduced following i.v. administration of a somatostatin analog, an effect that impairs glucose tolerance beyond what can be explained by the associated inhibition of insulin secretion (35). Although the mechanism underlying these effects on GE is unknown, our data with FGF19 raise the possibility of an action mediated in the brain.

Although our conclusions regarding the role of GE in the antidiabetic effect of FGF19 are based primarily on studies conducted in ob/ob mice, we believe that these findings have broader implications. First, our findings are compatible with several studies showing that, in rodent models of severe insulin-deficient diabetes, leptin administration can fully normalize blood glucose levels, despite persistent insulin deficiency (22–25). The effect is therefore insulin independent, and, moreover, it is mediated centrally, since the effect is observed when leptin is given i.c.v. at a much lower dose than is required during systemic administration. Combined with the current studies, these observations establish brain’s inherent capacity to induce dramatic, insulin-independent glucose lowering and provide a compelling rationale for investigation into underlying mechanisms and their clinical relevance. The question of whether centrally active mediators other than leptin and FGF19 can increase GE is also relevant. For example, FGF21 exerts antidiabetic effects analogous to those induced by FGF19, and both hormones appear to signal via the same receptor (FGFR1), but whether FGF21 increases GE is unstudied. In addition to candidate peptides, an increase of circulating glucose levels is itself well known to increase GE, and the possibility that this involves an effect of glucose in the brain is consistent with available data (36).

The contribution made by insulin-independent mechanisms to overall glucose homeostasis is surprisingly large. Available evidence suggests that GE accounts for approximately 50% of glucose disposal in humans following an i.v. glucose challenge (in rodents, the contribution may be greater), with the remaining 50% attributable to the combined effects of insulin action and insulin secretion (1). It is noteworthy in this context that, like insulin resistance, reduced GE makes an important contribution — perhaps comparable to that of insulin resistance — to impaired glucose tolerance and T2D in humans (1, 37). Consequently, reduced GE is a strong risk factor for the development of T2D (38). Combined with our current findings, the possibility that a central defect contributes to impaired GE in obesity-associated metabolic impairment can be considered. Related questions of strong translational interest include whether defects in CNS sensing or responsiveness to FGF19 or other inputs contribute to common clinical disorders of glucose homeostasis and whether interventions that increase GE have therapeutic potential in individuals with obesity and diabetes.

As a first step to investigate how FGF19 action in the brain increases GE, we measured the increment of plasma lactate levels following i.v. injections of either FGF19 (3 μg) or its vehicle. Data represent mean ± SEM. *P < 0.05 vs. i.c.v. vehicle.

Figure 5
Hepatic effects of i.c.v. FGF19 in ob/ob mice. Hepatic expression of (A) G6Pase and (B) Pepck using real-time PCR and (C) liver triglyceride and (D) liver glycogen content in leptin-deficient ob/ob animals that received daily i.c.v. injections of either FGF19 (3 μg) or its vehicle. Data represent mean ± SEM. *P < 0.05 vs. i.c.v. vehicle.
levels that follows a glucose challenge. The rationale for this approach begins with evidence published some 40 years ago that plasma lactate levels increase following consumption of a mixed meal in humans (39). Subsequent studies showed that this response results not from lactate absorption from the GI tract, but from the uptake of glucose into tissues, followed by its metabolism to lactate that is subsequently released into the circulation (40, 41). Since the same glucose→lactate metabolic response occurs after an i.v. glucose challenge, calories do not need to enter the body from the GI tract to elicit this response—a increased circulating glucose level is sufficient (15). Interestingly, the effect of either oral or i.v. glucose to increase lactate levels is reduced in obese humans (15), as is GE (1). We therefore asked (a) whether reduced GE in ob/ob mice is associated with blunting of the glucose→lactate response and (b) whether FGF19 action in the brain increases GE in part by stimulating this process. Our data are consistent with both predictions. We found that the plasma lactate level increased rapidly following an i.v. glucose challenge in WT mice, whereas lactate levels did not increase at all in vehicle-treated ob/ob mice, consistent with their markedly reduced GE (by ~70%). In contrast, both GE and the plasma lactate response to i.v. glucose increased markedly in ob/ob mice receiving i.c.v. FGF19. Together, these findings support a model in which (a) reduced GE in ob/ob mice results in part from a decreased glucose→lactate metabolic response, (b) the effect of FGF19 to increase GE involves an increase in this response, and (c) the brain mediates this effect of FGF19.

The identification of tissues in which this glucose→lactate response occurs in mice awaits further study. In dogs and humans, the liver is implicated, and it is notable in this context both that liver glucose uptake is insulin independent and that the brain plays a fundamental role to regulate this process (38). However, our finding that the effect of FGF19 to increase GE is not associated with increased liver glycogen content in ob/ob mice raises questions regarding whether liver glucose uptake is increased, and both fat and skeletal muscle can also metabolize glucose to produce lactate that is subsequently released into the circulation (42, 43). Additional studies are therefore needed to definitively identify the tissue(s) involved.

Although our findings support the hypothesis that the antidiabetic effects of FGF19 involve a central site of action, we emphasize that they do not preclude a role for peripheral tissue effects as well (8, 44). FGFR1 activation exerts glucose-lowering effects in liver as well as both brown and white adipose tissue (11), and the beneficial effects of systemic FGF19 on glucose tolerance that we observed in ob/ob mice were only partially blocked by central FGF receptor blockade. Future studies using targeted disruption of FGF receptor signaling in a tissue-specific manner will help to clarify the extent to which glucoregulatory effects of FGF19 are mediated centrally versus peripherally.

In conclusion, little is known regarding the neurocircuitry underlying central FGF19 action, although our data implicate a mechanism independent of melanocortin signaling. This assertion is based on our findings that FGF19 failed to induce c-Fos (a marker of neuronal activation) in hypothalamic POMC neurons and that FGF19 improved glucose tolerance in Mc4r-deficient mice (albeit, not to the same extent as was observed in ob/ob mice). Whatever the underlying neurocircuitry, convergent lines of evidence that implicate activation of FGFR1 in the antidiabetic effects of FGF19 (and FGF21, as well) (9, 10) and the observation that FGFR1 is expressed in key hypothalamic areas implicated in metabolic regulation (12, 13) support a model in this brain area that mediates the increase of GE induced by FGF19. Additional studies are warranted to test this hypothesis.

Whether FGF19 plays a physiological role in glucose homeostasis or diabetes pathogenesis is still largely unstudied. Also of interest is the as yet untested hypothesis that beneficial metabolic effects of bariatric surgical procedures are tied to increased FGF19 secretion from gut enterocytes. Consistent with this possibility, circulating levels of bile acids (45, 46) and FGF19 increase following roux-en-Y gastric bypass (RYGB) surgery in humans (47, 48). As bile acid stimulation of FGF19 secretion is mediated via FXR, these data collectively raise the possibility that increased bile acid–induced stimulation of FGF19 secretion (and increased GE) contributes to the high rate of diabetes remission induced by RYGB (49), but a direct test of this hypothesis is still awaited.

In conclusion, our findings suggest that the glucose-lowering effect of systemic FGF19 involves a novel central mechanism whereby insulin-independent glucose disposal is rapidly, potently, and selectively increased. The finding that the brain has the inherent ability to lower blood glucose levels via such a mechanism opens new potential avenues for diabetes drug discovery.
Methods

Animals. All animals were housed individually under specific pathogen-free conditions in a temperature-controlled room with a 12-hour light/12-hour dark cycle. Animals were provided with ad libitum access to water and standard laboratory chow (PMI Nutrition International Inc.) unless otherwise stated. Adult, male ob/ob mice on the C57/Bl6 background and their WT littermates were obtained from The Jackson Laboratory. To better identify POMC cells, we used POMC-Tau-eGFP mice that were provided to us by Tomas Horvath at Yale University (New Haven, Connecticut, USA), in which T-topaz GFP is expressed under the transcriptional control of the POMC promoter and backcrossed onto the C57/Bl6 background for >10 generations (50). Mc4r−/− mice, originally derived as described in ref. 51, were obtained from breeding colonies maintained at the University of Washington.

Surgery. Animals underwent cannulation of the lateral ventricle (Plastics One) under isoﬂurane anesthesia at stereotactic coordinates based on the brain atlas of Paxinos and Watson (52): –0.7 mm posterior to bregma; 1.3 mm lateral, and 1.3 mm below the skull surface. The cannula was permanently ﬁxed to the skull using glass ionomer cement (Instech Solo- mon) and was closed with an obturator. Animals received buprenorphine hydrochloride (Reckitt Colman Pharmaceuticals) at the completion of the surgery and were allowed to recover for at least 5 days while food intake and body weight were recorded. i.c.v. injections were administered using a (33-gauge) needle that extended 1 mm beyond the tip of the cannula over a period of 60 seconds in a ﬁnal volume of up to 1.5 μl. The injector was then allowed to remain in place for an additional 60 seconds before removal. For studies involving a FSIGT, adult male C57/Bl6 and ob/ob mice underwent lateral ventricular cannulation as above as well as catheterization of both the carotid artery and jugular vein, as described in detail previously (53).

Body composition analysis. Liver triglyceride content and measures of total fat and lean body mass were determined using quantitative magnetic resonance spectroscopy (EchoMRI 3-in-1 Animal Tissue Composition Analyzer; Echo Medical Systems) using the Energy Balance and Glucose Metabolism Core of the Nutrition Obesity Research Center at the University of Washington. Liver glycogen levels were determined using a colori- metric assay (Biovision) and were standardized to grams wet weight.

i.p. glucose tolerance testing. i.p. glucose tolerance tests (ipgtt; 30% d-glucose; 0.5 or 1.5 g/kg) were conducted in 5-hour fasted animals. Blood glucose levels were measured at t = 0, 15, 30, 60, 90, and 120 minutes using a hand-held glucometer (Accu-Chek) to test blood obtained from tail capillary samples.

CNS FGF receptor signaling in the glucose-lowering effects of FGF19 in ob/ob mice. To determine whether increased FGF19 signaling in the brain is sufﬁcient to improve glucose tolerance, adult male ob/ob mice were matched for body weight, food intake, and blood glucose levels and fasted at 09:00. At 11:30, animals received a single, acute i.c.v. injection of either vehicle (0.9% saline) or FGF19 (Phoenix Pharmaceuticals) at a dose of 1 μg or 3 μg. At 13:00, animals were subjected to an ipgtt (0.5 g/kg). At 15:00, blood (20 μl) was sampled for measurement of glucose and subsequent assay of plasma insulin at time points 1, 2, 4, 8, 12, 16, 20, 30, and 60 minutes. Additional samples were obtained for glucose measurement at 3, 5, 6, 10, 14, 18, 25, 40, and 50 minutes. Throughout the entire procedure, beginning at 10 minutes, mice received i.v. infusion of saline-washed erythrocytes (5 μl/min) to prevent the significant fall in hematocrit that would otherwise occur. Blood glucose levels were measured on a hand-held glucometer (Accu-Chek).

Minimal model analysis of FSIGT data. Based on glucose and insulin curves, glucose disposal was divided into 3 phases: mixing phase (0–2 minutes), GE-dominated phase (2–5 minutes), and insulin-domi- nated phase (5–60 minutes) (2). The mixing phase was not used for Si or GE modeling. Acute insulin response to glucose was based on insulin infusion of saline-washed erythrocytes (5 μl/min) to prevent the significant fall in hematocrit that would otherwise occur. Blood glucose levels were measured on a hand-held glucometer (Accu-Chek).

To determine whether the glucose-lowering effect of FGF19 involves activation of FGF receptors in the CNS, a separate cohort of ob/ob mice was matched for body weight, food intake, and blood glucose levels and fasted at 09:00. Ninety minutes later, animals received either an i.c.v. pre-injection of the FGFR inhibitor, PD173074 (Sigma-Aldrich; 25 μg), or its vehicle (100% DMSO). Relative to vehicle-treated controls, this dose had no effect by itself on either fasting blood glucose levels or glucose tolerance (data not shown). To confirm that the glucose-lowering effect of i.c.v. FGF19 is receptor mediated, animals received an i.c.v. injection of FGF19 (3 μg) or its vehicle 1 hour after pretreatment injection. At 13:00, animals were then subjected to an ipgtt (0.5 g/kg). Once the efﬁcacy of the inhibitor was validated against i.c.v. FGF19, we asked whether pretreatment blunts the antidiabetic effect of FGF19 following systemic administration at a higher dose. Adult male ob/ob mice followed the same protocol as described above, except that they received a systemic (rather than i.c.v.) injection of FGF19 (1 mg/kg).

To determine whether the glucose-lowering effects of CNS FGF19 could be explained by leakage into the periphery, a similar paradigm was used, except that ob/ob mice received daily peripheral injections of FGF19 (3 μg) or its vehicle at the same dose that was administered i.c.v.

FSIGT. A FSIGT was performed in 5-hour fasted ob/ob mice or their littermate controls at least 5 days following LV cannulation and arterial and venous catheterization. Blood sampling was performed via the arterial catheter in unrestrained, conscious animals. A baseline fasted blood sample was taken at –10 and 0 minutes. Based on a previous protocol (2), a 1 g/kg bolus of 50% dextrose was injected i.v. over a period of 15 seconds at t = 0 minutes. Blood (20 μl) was sampled for measurement of glucose and sub- sequent assay of plasma insulin at time points 1, 2, 4, 8, 12, 16, 20, 30, and 60 minutes. Additional samples were obtained for glucose measurement at 3, 5, 6, 10, 14, 18, 25, 40, and 50 minutes. Throughout the entire procedure, beginning at –10 minutes, mice received i.v. infusion of saline-washed erythrocytes (5 μl/min) to prevent the significant fall in hematocrit that would otherwise occur. Blood glucose levels were measured on a hand-held glucometer (Accu-Chek).

Effect of i.c.v. FGF19 on c-fos induction in Pome-Tau-eGFP mice. To deter- mine whether FGF19 activates POMC neurons in the hypothalamus, female POMC-Tau-eGFP mice underwent lateral ventricular cannulation, as described previously. Following a 1-week recovery period, habituated animals were fasted for 3-hours and then received an i.c.v. injection of either saline or FGF19 (3 μg). Ninety minutes later, mice were perfused, with brains removed and processed for immunohistochemical out- comes, as described below.

Tissue collection and processing. For immunohistochemical studies, mice were anesthetized with ketamine/xylazine and perfused with PBS followed by 4% PFA in 0.1 M PBS. Brains were removed, post-fixed in 4% PFA overnight, embedded in 25% sucrose solution for 48 hours at 4°C, snap frozen in isopentane, and cooled with liquid nitrogen. The hypothalamus was sectioned at 14 μM in the coronal plane using a freezing cryostate, slide-
mounted, and stored at –80°C until selection for immunohistochemical staining or visualization of GFP by fluorescence microscopy.

Double-label immunofluorescence. For colocalization studies, anatomically matched sections throughout the hypothalamus were selected for staining. Briefly, slides were washed with 10 mM PBS at room temperature and were then blocked for 30 minutes with buffer containing 5% normal donkey serum, 0.1% BSA, and 0.01% Triton-X in 10 mM PBS. Following additional washes, slides were incubated overnight at 4°C in a “cocktail” of primary antibodies, including goat anti-fibroblast growth factor 15 ( Fitzgerald Industries) diluted 1:10,000 with rabbit polyclonal anti-c-Fos (Oncongene Research Products) diluted 1:100,000. Slides were washed with PBS and then incubated for 120 minutes at room temperature with the secondary antibodies (a) 1:1,000 donkey anti-goat IgG-Alexa Fluor 488 (Life Technologies) and (b) 1:1,000 donkey anti-rabbit Alexa Fluor 594 (Life Technologies), diluted in 5% normal donkey serum and 0.1% BSA in 10 mM PBS. After washing with PBS, slides were subsequently overlaid with an antifade aqueous mounting media. Control sections were incubated with normal serum and did not show staining.

Plasma analysis. Blood samples for measures of insulin, lactate, and FFAs were collected in EDTA-treated tubes, while blood for glucagon assay was collected on tubes containing 10 μl of 1 M benzamidine and 1 U of heparin. Blood was centrifuged and plasma was removed, aliquoted, and stored at –80°C. Plasma immunoreactive insulin levels were determined by ELISA (Crystal Chem), FFA levels were determined using a colorimetric assay (Wako Chemicals), plasma lactate levels were determined using a GM9D glucose direct analyzer (Analog Instruments), and plasma glucagon levels were determined using a glucagon RIA kit (Linco Research).

RT-PCR. Total RNA was extracted from liver using TRIzol B according to manufacturers’ instructions (MRC). RNA was quantitated by spectrophotometry at 260 nm (Nanodrop 1000, Thermo Scientific) and reverse-transcribed with AMV reverse transcriptase (1 μg) (Promega). Real-time PCR was performed on an ABI Prism 7900 HT (Applied Biosystems) using the commercially available PCR Master Mix (SYBR Green, Applied 2.0; Applied Biosystems) and analyzed as previously described (22, 26).

Statistics. All results are expressed as mean ± SEM. Statistical analyses were performed using Statistica (version 7.1). A 1-way analysis of variance with a LSD post-hoc test was used to compare mean values among multiple groups using log-transformed data when Bartlett’s test for unequal variance showed P < 0.05, and a 2-sample unpaired Student’s t test was used for 2-group comparisons. In all instances, probability values of < 0.05 were considered significant.

Study approval. All procedures were performed in accordance with NIH guidelines for the care and use of animals and were approved by the Institutional Animal Care and Use Committee at both the University of Washington and Vanderbilt University.

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Address correspondence to: Michael W. Schwartz, Department of Medicine, University of Washington at South Lake Union, 850 Republican St., N335, Box 358055, Seattle, Washington 98195, USA. Phone: 206.897.5288; Fax: 206.897.5293; E-mail: mschwart@u.washington.edu.


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