

Gut-brain communication and obesity: understanding functions of the vagus nerve

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Given the crucial role of the gastrointestinal tract and associated organs in handling nutrient assimilation and metabolism, it has long been known that its communication with the brain is important for the control of ingestive behavior and body weight regulation. It is also clear that gut-brain communication is bidirectional and utilizes both rapid neural and slower humoral mechanisms and pathways. However, progress in understanding these mechanisms and leveraging them for the treatment of obesity and metabolic disease has been hindered by the enormous dimension of the gut mucosa, the complexity of the signaling systems, and lack of specific tools. With the ascent of modern neurobiological technology, our understanding of the role of vagal afferents in gut-brain communication has begun to change. The first function-specific populations of vagal afferents providing nutritional feedback as well as feed-forward signals have been identified with genetics-guided methodology, and it is hoped that extension of the methodology to other neural communication pathways will follow soon. Currently, efficient clinical leveraging of gut-brain communication to treat obesity and metabolic disease is limited to a few gut hormones, but a more complete understanding of function-specific and projection-specific neuronal populations should make it possible to develop selective and more effective neuromodulation approaches.

Introduction

Given that the prevalence of obesity is approaching 50% in some countries (1), it is hard to argue that body weight/adiposity is regulated at healthy levels through biological mechanisms (2–7). It has been speculated that the best explanation for this conundrum is evolutionary genetic drift (7) and the dependency of such regulation on the environment and lifestyle (8, 9). In genetically susceptible individuals, pressures from food, physical, and social environments (10) seem to be too much for a regulatory system that evolved mainly to defend against starvation (7, 11). Being unable to change the root cause, namely an environment that facilitates overnutrition and discourages physical activity, the fight against obesity is currently limited to using known physiological systems controlling energy intake, assimilation, and expenditure for behavioral, pharmacological, and surgical therapies. As such, the gut-brain axis is a key element for the control of ingestive behavior with important implications for the development of obesity and metabolic diseases. Here we discuss recent advances in understanding how nutrients are perceived by the gastrointestinal tract and how relevant signals are mediated to the brain, with emphasis on the vagus nerve.

The bases of bidirectional gut-brain crosstalk

Although use of the term “gut-brain axis” on PubMed has increased more than tenfold only in the last decade, it has long been known that connections between these organs exist and

that they have important functional implications in health and disease. For example, Galenus of Pergamum, a surgeon and philosopher in the Roman Empire (129–199 CE), noted: “A large portion of nerves is emitted from the brain to the entrance of the stomach, because nature has made this an instrument of appetite for food, which is at the door ... of all instruments nature has prepared for management of nutrients” (translated from ref. 12). The recent fascination with the term is almost solely due to literature on the gut microbiome (e.g., ref. 13), covered in other articles in this Review series. However, for students of ingestive behavior and metabolic physiology, understanding the gut-brain axis has been a crucial element for half a century (e.g., ref. 14). It was this interest in ingestive behavior and the rising obesity epidemic that spurred some of us to pursue intensive research on the functional anatomy of the vagus nerve. As schematically depicted in Figure 1, the vagus nerve with its sensory and motor fibers is one of the key players in reciprocal gut-brain communication.

The sympathetic nervous system and dorsal root spinal afferents are no doubt also important for reciprocal fast signal transmission between the gut and the brain, though we know much less of their potential implications in ingestive behavior and metabolic regulation. Spinal primary afferent neurons with cell bodies in dorsal root ganglia (DRGs) innervate the entire gastrointestinal tract and associated glands (pancreas, liver, gallbladder) as well as immune organs (lymph nodes, gut-associated lymphoid tissue, spleen).

Humoral factors secreted by the gut, and pituitary hormones discharged by the brain, additionally serve for slower communication with each other. However, it would be shortsighted to consider only direct connections between the gut and the brain, as

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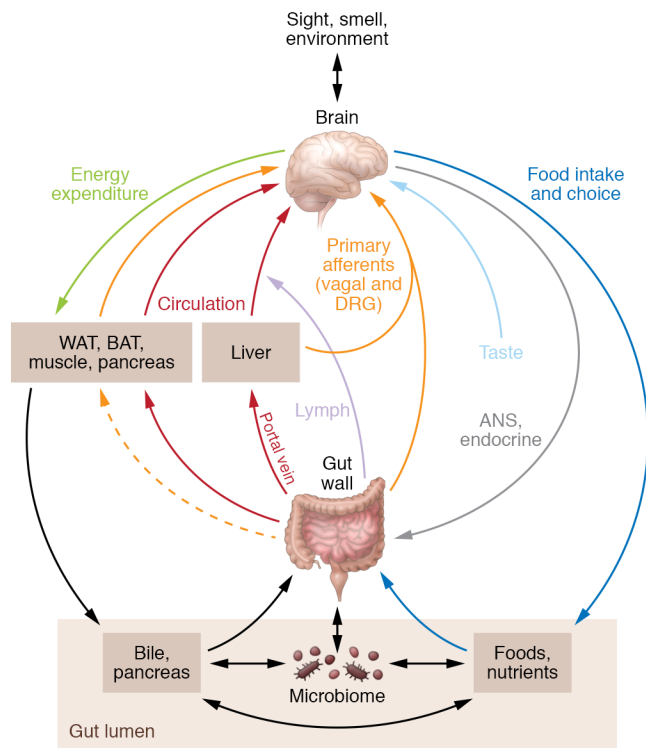


Figure 1. Schematic diagram showing potential gut-brain communication pathways relevant to obesity and metabolic diseases. See article text for details. BAT, brown adipose tissue; WAT, white adipose tissue; ANS, autonomic nervous system.

indirect neural and humoral signaling pathways via other organs are potentially also important (Figure 1). For eating behavior and metabolic regulation, the liver, adipose tissue, muscle, and pancreas, with their secretion of hormones and cytokines as well as their innervation by the autonomic nervous system and primary afferent neurons, are of particular interest. Besides hormones and cytokines, the recently discovered exosomes and microRNAs should also be considered for organ-to-organ communication (15, 16). Finally, bidirectional communication is further highlighted by the fact that the brain dictates how much and what kind of food is ingested (ingestive behavior), which subsequently determines how the gut and associated organs (e.g., microbiome, bile system, and pancreas) react and ultimately determines the type of feedback sent back to the brain and other organs (Figure 1).

The role of gut-brain communication in eating behavior

While earlier research focused mainly on mechanisms for signaling satiation, there is now much appreciation for signaling of appetite and food reward from the gut to the brain (17–23). The term “appetition” was coined by Sclafani and his colleagues based on the observation that the interaction of certain nutrients with the intestinal mucosa elicits more appetite, rather than less, as in satiation (22). We have long known the French proverb, “*L'appétit vient en mangeant*” (appetite comes with eating), referring to the fact that external cues such as sight, smell, and taste signaling the availability of delicious beneficial food stimulate appetite for and consumption of this food (24, 25). It turns out that such a feed-

forward mechanism extends to the small intestine. By using non-nutritive saccharin-sweetened and differentially flavored solutions combined with intragastric or intraduodenal nutrient infusions, Sclafani and colleagues comprehensively demonstrated that animals learn to associate taste and flavor of ingested foods with its postingestive consequences (25, 26). This learning mechanism has two important consequences. First, the increased intake of the flavored solution when paired with intragastric glucose or lipids as compared with intragastric water indicates that intragastric nutrients act as positive reinforcers that can activate the brain reward system (20, 21). Second, this mechanism is fundamental for subsequent food choice and survival, as it allows acceptance of beneficial foods or rejection of harmful foods before any large amount is sampled and ingested. Identification of the signal transduction pathway from arrival in the gut to changing behavior in the brain is thus of great interest and has recently led to exciting new discoveries.

The role of the vagus nerve

Historical background on functional vagal anatomy. Based on the early premise that their activation leads to satiation and reduced energy intake, vagal afferents have attracted considerable attention. Before the availability of current molecular genetics-based neurobiology techniques, the vagal afferent system was thoroughly mapped using retrograde and anterograde labeling methods and electrical nerve recording. At least three fundamentally different types of terminal architectures have been demonstrated in both rats and mice. Intramuscular arrays are contained in the external muscular layers, particularly in the stomach wall, and are thought to represent slowly adapting stretch receptors (27, 28). Intraganglionic laminar endings (IGLEs) cover parts of myenteric ganglia that lie between the longitudinal and circular external muscle layers throughout the esophagus and gastrointestinal tract (29). Based on indirect evidence obtained from guinea pig stomach (30), they are thought to sense the level of tension, whether generated by passive tissue stretch or generated by active muscle contraction. Ultrastructural observations also support a mechanosensory function (31). Combined with recent immunohistochemical data, ultrastructural observations suggest additional chemosensory functions and complex interactions of IGLEs with myenteric neurons and calcitonin gene-related peptide-positive spinal afferents passing through myenteric ganglia (31–33), suggesting additional chemosensory functions. Finally, mucosal endings have been traced to the entire gastrointestinal tract, with the highest density in the villi and crypts of the proximal small intestine (34, 35). Originally thought to mediate nutrient-induced satiation signals, they now are implicated in nutrient-induced appetite signaling (36). In addition to the gut itself, the associated hepatic portal vein (37, 38) and pancreatic β cells (39, 40) are also innervated by vagal afferents.

Assignment of specific functions to vagal afferent populations with different terminal architectures and different locations along the gastrointestinal tract has been hampered mostly by the inadequacy of tools limited to electrical stimulation and surgical or chemical interruption of the entire subdiaphragmatic vagal trunks or their main branches. Although more selective vagal branch manipulations have helped elucidate the viscerotopy of

vagal afferent and efferent innervation (41–44), these studies are still nonspecific with respect to sensory versus motor fibers and to sensory or motor function of subpopulations. Clearly, many scientists were attracted to the subdiaphragmatic vagus nerves and branches because of relatively easy access. The common hepatic branch, which separates from the left (anterior) subdiaphragmatic trunk just beneath the diaphragm, is particularly easy to locate and manipulate, and it has received by far the greatest attention. There has been a steady stream of publications using surgical transection of the common hepatic branch of the vagus nerve as a tool to study the role of the vagus nerve in liver function (for recent examples, see refs. 45–54). Unfortunately, most of these analyses seem unaware of the projection targets of vagal afferent and efferent fibers in the common hepatic branch, as identified in neuronal tracing and multiorgan functional analyses with electrical stimulation (37, 41, 55). These studies in rats have collectively demonstrated that a majority of vagal fibers passing through the (easily accessible) hepatic branch innervate targets in the proximal small intestine, pylorus, antrum, and pancreas, while a minority innervate the hepatic portal vein, bile ducts, and the liver hilum (37). That is, fibers in the common hepatic branch join the dense plexus surrounding the common hepatic artery, and the majority continue along the gastroduodenal artery. Only a minority of these fibers follow the hepatic artery proper toward the liver and portal vein (Figure 2). In recent studies addressing the benefit of preserving vagal pathways during cancer gastrectomy, similar findings were made in humans (56). Thus, cutting the common hepatic branch does a lot more than interrupt vagal motor input to the liver or vagal sensory output from the liver, and these potential collateral effects must be taken into consideration in interpreting such studies.

New generation of genetically guided vagal manipulations. Similar to the central nervous system, advances in genetics-based identification of molecularly distinct neurons have tremendously enriched our tool kit to study functional anatomy of the peripheral nervous system. First, with the use of reporter mice expressing brightly fluorescent proteins in all, or specific, populations of vagal sensory or motor neurons, the demanding and capricious anterograde tracer injections into the nodose ganglia or the dorsal motor nucleus could be circumvented, as well as often elaborate staining techniques (57–62). In general, these studies in mice confirmed the distribution and terminal architecture of both vagal afferents and efferents reported earlier in rats using anterograde labeling techniques.

Second, transcriptional profiling of neurons contained in spatially compact areas such as the jugular, nodose, dorsal root, or myenteric plexus ganglia with single-cell RNA sequencing (scRNA-Seq) has provided comprehensive atlases of molecularly distinct clusters of neurons (62–66), suggesting that they represent function-specific populations. Specifically, and most relevant to this discussion, applying scRNA-Seq to individual handpicked neurons in the nodose ganglia labeled by injections of a retrograde tracer into different segments of the gastrointestinal tract and the hepatic portal vein further revealed the organization of vagal afferent innervation of abdominal organs (62). These studies demonstrated that each site was innervated by vagal sensory neurons belonging to different clusters and thus different functionality, but with more or less enrichment from one or more clus-

ters. Given the specific organization of vagal motor outflow from the dorsal motor nucleus with respect to viscerotopy (41, 43, 55, 67–69) and function (70–72), similar scRNA-Seq analyses of functional and projection-specific vagal motor neurons in the dorsal motor nucleus are eagerly awaited.

Third, analysis of the effects of selective acute and chronic activation or inhibition of molecularly distinct subpopulations of vagal afferents on ingestive behavior and metabolic endpoints is well under way. Much of this research has focused on the mechanisms responsible for the post-oral detection of ingested nutrients, glucose and fat in particular, which provides the unconditioned stimulus for learning of food preferences as well as signals for appetite and satiation (21, 23, 25, 26, 73). Thus, optogenetic silencing of a specific population of enteroendocrine cells (EECs) that make synaptic contacts with vagal afferent neurons in the proximal small intestine, so-called neuropod cells, abolishes the vagal afferent signal to intestinal sucrose and prevents mice from distinguishing non-nutritive sweeteners from nutritive sucrose (74, 75). Elaborate *in vitro* and *in vivo* studies further identified a mechanism by which luminal glucose is transported into the EECs selectively through sodium-glucose transporter-1 (SGLT1) (76), which leads to cell depolarization, the release of glutamate from the neuropods, and rapid activation of glutamate receptors on vagal afferent terminals (ref. 36 and Figure 2). While the specific vagal afferent population was not identified and therefore was not accessible for manipulation, another recent study used the PhosphoTRAP methodology to identify the vagal afferent neurons in the nodose ganglia that were activated by intestinal glucose but not the artificial sweetener acesulfame potassium (Ace-K) (77). In addition, these researchers identified the activated downstream neurons expressing proenkephalin in the nuclei tractus solitarii (NTS) (Figure 2). Importantly, silencing either the specific neuron populations in the nodose ganglia or the NTS abolished the ability of mice to learn a preference for intestinal glucose (77). In fact, optogenetic activation of these neurons while the mice were receiving differentially preferred intestinal stimuli led to a reversal of preferences, with the previously avoided stimulus now preferred (77).

What is most surprising is that selectively silencing these vagal pathways (36, 77) results in complete abolition of discriminatory nutrient preference learning. Given that such learning may be crucial for survival in the wild, this complete abolition is unexpected, as it would leave no room for any other signaling pathway to be involved in this fundamental behavior. One possible explanation is that it may be an artifact of the highly simplified situation with purified nutritive and non-nutritive solutions directly infused into the intestines, which is far from the normal task to discriminate among a multitude of food items. Thus, studies in more naturalistic environments with more complex food choices should be revealing. Also, given the extensive literature on humoral mediation of gut-to-brain nutrient signaling (e.g., refs. 78, 79), these recent findings suggest that different signaling pathways mediate different behavioral endpoints, such as satiation, appetite, food seeking, and reward generation.

It is also important in this regard to note that previous attempts using highly nonselective total subdiaphragmatic vagotomy or the slightly more selective subdiaphragmatic vagal deafferentation

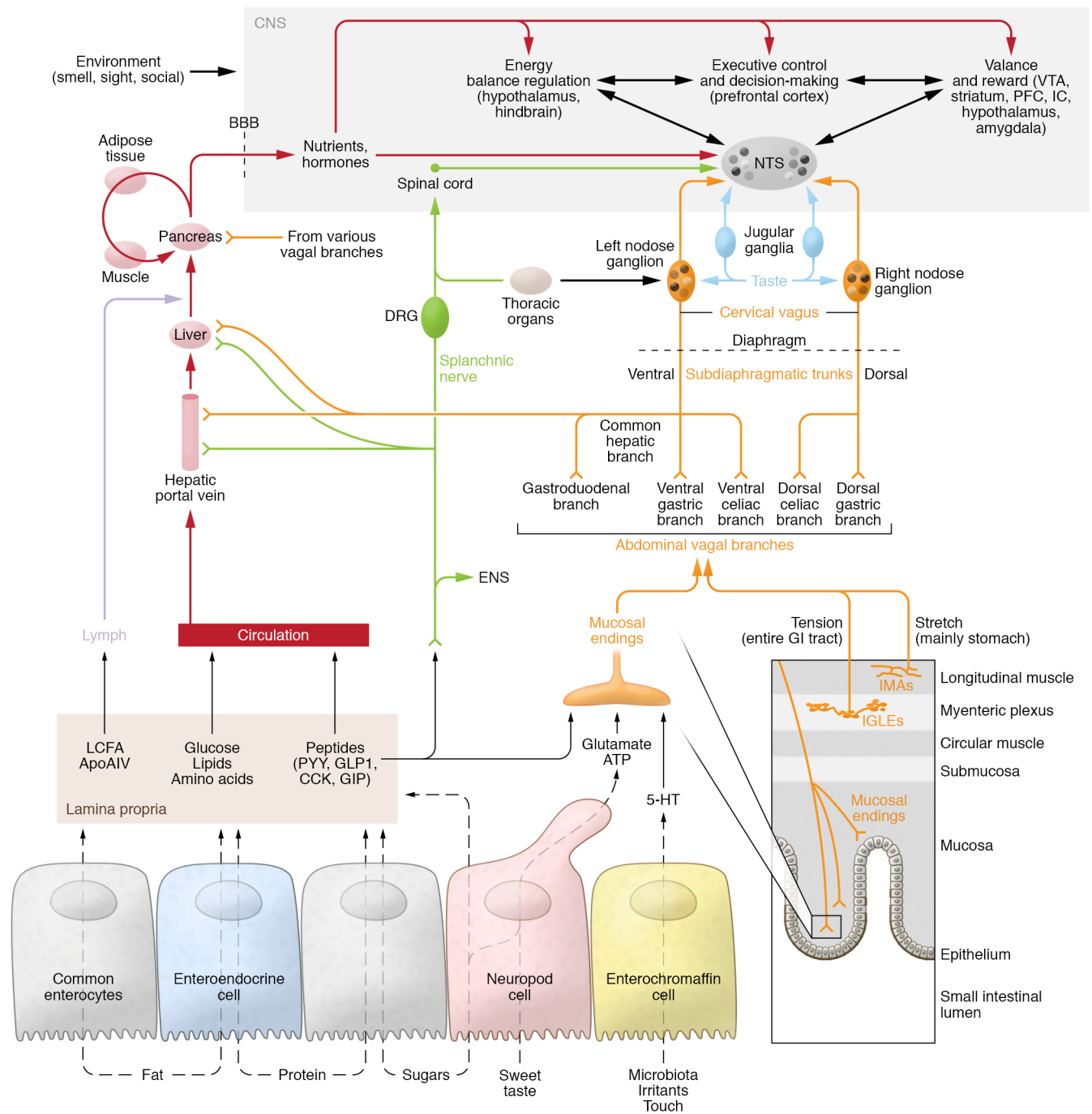


Figure 2. Gut-brain communication relevant to nutrient assimilation. Bottom: Transport mechanisms for dietary nutrients in small-intestinal epithelial cells and transduction mechanisms generating neural and humoral signals. Multistep pathways are shown using dashed lines. The volume and osmotic effects of ingested nutrients also interact with the muscular wall of the alimentary canal and can activate vagal stretch (intramuscular arrays [IMAs] mainly in the stomach) and tension receptors (intraganglionic lamina endings [IGLEs] throughout the gastrointestinal tract). Middle: Signals generated in vagal afferent terminals are projecting through several subdiaphragmatic branches (orange) to the nuclei tractus solitarii (NTS) via the nodose ganglia, which contain the neural cell bodies. Dorsal root afferents can also mediate signals to the brain (green), including the NTS via the spinal cord and spinothalamic tract. The lamina propria is innervated by enteric neurons, including intrinsic primary afferent neurons mediating mucosal signals to the enteric nervous system. Also note that nutrients and hormones in the portal hepatic vein and liver can potentially activate vagal and DRG afferents. Top: The NTS is a hub for sensory neural information originating from the gastrointestinal tract and the oral cavity as well as from humoral inputs via the area postrema and has extensive projections to many brain areas. These include areas involved in the control of food intake (satiety and hunger); homeostatic regulation of energy balance and body weight; the generation of food reward; and executive control and decision-making. Also note that nutrients and hormones taken up into the bloodstream (either directly or after transport through the lymphatic system) can eventually interact with sensors in specific areas of the brain and all other organs. VTA, ventral tegmental area; PFC, prefrontal cortex; IC, insular cortex; BBB, blood-brain barrier; ENS, enteric nervous system; LCFA, long-chain fatty acids; ApoAIV, apolipoprotein AIV; 5-HT, 5-hydroxytryptamine (serotonin).

failed to implicate vagal afferents in nutrient preference learning (80–82), suggesting that at least some critical information is mediated from gut to brain by humoral signals. However, it must also be acknowledged that these nonspecific vagotomies (and similarly manipulations of dorsal root and sympathetic innervation by means of celiac/superior mesenteric ganglionectomy; refs. 81, 83) produce multiple side effects, particularly in the long term, that may occlude the true role of critical, small, and function-specific populations of sensory and/or motor neurons. Regeneration of surgically transected abdominal vagal afferents is another potential confounding problem (84).

The role of other neural pathways

Compared with the vagus nerve, sensory and sympathetic motor fibers innervating the abdominal organs via the spinal cord have received much less attention. This is at least in part due to the challenging anatomical configuration of sensory innervation by dorsal root ganglia (DRG) and the sympathetic nervous system (SNS). There is no “easy target” such as the common hepatic branch for the vagus, and researchers have targeted either the major splanchnic nerve (85, 86) or the celiac/superior mesenteric ganglion complex (81, 83). However, these manipulations are problematic, for the same reasons as are nonselective vagotomies, because there are likely many function-specific populations of SNS efferents. In particular, there are specific sympathetic pre-postganglionic neuron chains of motility-inhibiting, secretion-inhibiting, and vasoconstrictor neurons, which were, at their postganglionic level, defined by their peptide content (87). Also, instead of just two easily accessible nodose ganglia, there are many DRGs harboring a variety of transcriptionally defined subtypes (65) that are even more difficult to manipulate. To date, reporter mice with fluorescently labeled neurons and RNA-Seq have not been applied to the investigation of the innervation patterns in abdominal and other relevant organs, or to investigation of the potential roles of postganglionic SNS and DRG neurons in nutrient sensing and metabolic control. One recent study reported that the duodenal-glucose-induced drop in hypothalamic agouti-related peptide neuron activity was significantly attenuated in mice with prior celiac/superior mesenteric ganglionectomy, while the duodenal-fat-induced drop was attenuated in mice with subdiaphragmatic vagotomy (83), which may be explained by some of the functional differences between vagal and splanchnic glucosensors reported earlier (88). Whether these findings can be confirmed with more selective manipulations remains to be seen.

The potential of viscerofugal enteric neurons to affect food intake and glucose regulation was recently explored with chemogenetic activation of myenteric plexus neurons expressing cocaine- and amphetamine-regulated transcript (CART) located in the microbiota-rich regions, the distal ileum and colon, and projecting to the celiac/superior mesenteric ganglia (89). It was concluded that CART-expressing neurons can modulate blood glucose in a microbiota-dependent fashion via a polysynaptic pathway from the lower intestine to the celiac/superior mesenteric sympathetic ganglia and, in turn, to the pancreas (89). However, the suppressive effects on food intake, glucose tolerance, and insulin secretion were small and rather spurious, and the potential

role of such a signaling pathway in normal physiology awaits further interventional approaches.

Clearly, comprehensive atlases of molecularly distinct, projection-specific postganglionic sympathetic neurons and DRG neurons would greatly facilitate further research into their potential roles in metabolic diseases.

The role of gastrointestinal hormones and other factors

As illustrated in Figure 2, a number of gastrointestinal hormones are secreted from EECs upon stimulation by various nutrients and other factors. Based on studies with systemic administration of such hormones and factors in vagotomized rodents, the generally held view was that most of these hormones are able to change brain function and food intake via both the bloodstream and vagal sensory neurons.

For example, glucagon-like peptide-1 (GLP-1) is produced by EECs located mainly, but not exclusively, in the distal bowel upon carbohydrate and protein absorption (78, 90). GLP-1 and its mechanisms of action have attracted the bulk of attention because stable GLP-1 analogs such as liraglutide and semaglutide are currently among the most effective pharmacological treatments to achieve remission of type 2 diabetes and reductions in food intake and body weight (91–93). In addition, increased GLP-1 signaling is among the leading candidates when it comes to explaining the beneficial effects of bariatric surgeries (94). Our understanding of intestinal GLP-1's exact signaling pathways has been complicated by its short half-life time in peripheral blood, as well as the presence of GLP-1-producing neurons in the NTS and their extensive projections to brain areas involved in energy homeostasis and glycemic control (78, 95). Even after extensive experimentation in a number of species, there remains considerable controversy as to the role of vagal afferents in the physiological and pharmacological effects of GLP-1 and its analogs on food intake and body weight regulation (ref. 95; for an in-depth discussion, see section 13 in ref. 78).

The recent genetics-guided approaches tend to support the conclusion that information from intestinal nutrient sensors does not depend on GLP-1 receptor-bearing (GLP-1R-bearing) vagal afferents. Surprisingly, GLP-1R is mainly expressed in vagal afferents forming IGLEs between the external muscle layers of the stomach and is only expressed in a small subset of vagal afferents innervating the intestinal villi (61, 62). The observation that selective optogenetic activation of GLP-1R-expressing vagal afferents reduces short-term food intake (62) is consistent with the idea that gastric distention contributes to the satiation process and that circulating GLP-1 may modulate their sensitivity (96). However, as pointed out in a recent review (96), the small subset of GLP-1R-expressing vagal afferents innervating the intestinal villi could still be involved in the direct paracrine or synaptic (neuropods) activation of vagal afferents by GLP-1 released from EECs and its upstream effects on satiation via the NTS (95).

A fraction of bile acids escapes the enterohepatic circulation and “leaks” into the peripheral circulation. As bile acids are bona fide ligands for several receptors, this allows for central/peripheral actions of bile acids outside the gastrointestinal tract. Notably, lean animals with bile diversion surgery (to augment peripher-

ally circulating bile acids) have TGR5-dependent reductions in cocaine-induced elevation of accumbal dopamine relative to controls, suggestive of a gut-brain hormonal circuit and potential for treatment for addiction (97).

One problem that arises with focusing on single hormones and factors, as is done in most reductionist experimental settings, is the likelihood that processes like satiation and appetite are regulated by the interaction and synergism between many factors. Although integration of afferent signals controlling food intake at all levels of the neuraxis has been recognized for a long time (e.g., see ref. 98), this issue has recently gained more attention. Thus, the combinatorial actions of GLP-1 and other gut hormones like CCK and PYY, whether mediated by vagal afferents or mediated through the bloodstream, are likely to exceed the individual effects (e.g., see refs. 99, 100), and this concept has found its way into the design of combinatorial polypharmacy for the treatment of metabolic diseases (101).

Gut-brain communication in energy balance regulation and obesity

Long-term energy balance is determined by energy intake, absorptive efficiency, and energy expenditure. Therefore, experimentally induced changes in eating do not necessarily translate into changes in body weight, or more specifically, overeating does not necessarily lead to obesity. This is particularly true for manipulations and challenges that change the satiation process, as changes in meal size are often compensated by changes in meal frequency, with total energy intake unchanged. Here we focus on the far fewer studies of gut-brain communication that provide clear evidence that it causes, prevents, or reverses obesity.

Neural communication pathways

Since vagal afferents have traditionally been mostly implicated in the satiation process, it is not surprising that little evidence for their role in body weight regulation has been provided. Earlier studies using indiscriminate surgical or chemical interruption of vagal afferents through subdiaphragmatic vagal deafferentation in rats or capsaicin treatment in rats and mice found small effects on meal patterns but no overt body weight phenotype (102). Similarly, silencing of vagal afferents using saporin-conjugated CCK injections into the nodose ganglia of rats eliminated CCK- and GLP-1-induced suppression of food intake but had apparently no effect on body weight (103). These observations are reminiscent of observations in rats with meal-contingent, exogenous CCK administration (104) demonstrating that increased meal size is compensated by decreased meal frequency to maintain normal body weight. Also, more selective vagal deafferentation procedures, such as the selective elimination of leptin receptors in vagal afferents in the *Nav1.8/LepR^{fl/fl}* mouse (105) or knock-down of CART transcript in vagal afferent neurons in rats (106), result in a small increase of body weight and adiposity, with slightly increased nighttime food intake and no change in energy expenditure. However, diminished vagal afferent signaling after saporin-CCK injection into the nodose ganglia of rats on a high-fat/high-sugar diet was clearly shown to contribute to excessive body weight gain and adiposity (107). Given that the sensitivity of gut vagal afferents is decreased in obese rodents (108), it will

be important to further distinguish what comes first: obesity, or blunting of vagal afferent sensitivity (108).

The most recent genetics-guided selective manipulations of vagal afferents in mice that yielded specific deficits in nutrient preference learning were not tested for their long-term effects (36, 62, 77). Specifically, it will be interesting to see whether chronic chemogenetic activation of vagal afferent neurons in the two mouse lines that showed significant acute suppression of food intake (*Oxtr^{Cre}* and *Glp1r^{Cre}* mice; ref. 62) also reduces long-term food intake and body weight. Similarly, it will be interesting to see whether chronic activation of neuropod cells or their signaling capacity to vagal afferents (36) or proenkephalin-expressing NTS neurons (77) could be leveraged to suppress long-term food choice, caloric intake, and body weight.

In humans, electrical stimulation of the left cervical vagus nerve, a procedure used to treat refractory epilepsy and depression, has been reported to cause weight loss (109, 110) and increased energy expenditure through BAT thermogenesis (111). A report of reduced food craving upon vagus nerve stimulation (112) was, however, criticized as providing evidence not for reduced craving but rather for confused appetite in these depressed patients (113). A pilot clinical trial revealed beneficial effects on glucose metabolism also with transcutaneous stimulation of the left auricular branch of the vagus nerve (114). Although the vagal branches stimulated in these studies were not cut, it is assumed that the observed effects resulted mainly from activation of vagal afferents and the downstream brain areas (115), not vagal motor outflow. As vagal afferents are relayed by the NTS to various midbrain and forebrain sites, these beneficial effects may be generated at emotional and cognitive levels. However, a loop through the hypothalamus may link stimulation of vagal afferents to efferent autonomic and neuroendocrine channels. Thus, BAT thermogenesis via sympathetic nerves (116, 117), and possibly activation of efferent neurons of the dorsal vagal motor nucleus to the gut and pancreas, are conceivable. Interestingly, the opposite strategy, vagal blockade using high-frequency electrical stimulation of the subdiaphragmatic vagal trunks (vBloc Therapy, EnteroMedics Inc), has also been reported to result in meaningful weight loss (118), reminiscent of total subdiaphragmatic vagotomy recommended in the early days of surgical intervention for the treatment of obesity (119). For all these approaches, it will ultimately be important to identify and verify exactly what function-specific vagal afferents (or efferents) are activated or silenced, in order to generate coherent mechanistic explanations and design more selective treatments.

Finally, the antiinflammatory effects of electrical stimulation of vagal motor outflow in rodents have been widely publicized and its mechanisms discussed (for recent reviews, see refs. 120–122). To what extent the antiinflammatory reflex is linked to the pathogenesis of obesity, and whether it can be successfully translated to treat patients with obesity, remain to be elucidated (123).

Humoral signaling by gut hormones and other factors

As already discussed above, many gut hormones play important roles in the control of food intake and glycemic control. However, only GLP-1 agonists are currently marketed as effective antiobesity and/or antidiabetic drugs, with several others in early clinical trials (92, 93, 124). Given our emphasis on vagal communication, we only

briefly discuss the potential role of gut hormones and other factors directly acting on the brain, and the interested reader should consult the comprehensive recent review by Gimeno et al. (124).

Besides the classical gut hormones, bile acids are increasingly recognized for their actions not only in the gut and liver, but also in many other organs, including the brain, via the widely distributed bile acid receptors FXR and TGR5 (125). Importantly, the gut-restricted FXR agonist fexaramine potently attenuates diet-induced obesity and its associated inflammation and impairments in glucose metabolism in mice over a period of 5 weeks (117). The effect was attributed to increased brown fat thermogenesis and browning of white fat, but not decreased food intake. Furthermore, increased circulating levels of the intestinal hormone FGF15/19 and a shift in the bile acid profile, triggering changes in the microbiome and increased signaling through TGR5, were identified as potential mechanisms (117, 126). Although this would not imply gut-brain communication at first sight, it seems that the specific changes in bile acid profile and gut microbiota can affect glucose homeostasis via an obligatory relay in the dorsal vagal complex (127). In addition, bile acids can modulate GLP-1 secretion from EECs through a complex interaction between inhibitory FXR and stimulatory TGR5 effects (128–130). Aside from the bile acid–receptor interaction itself, bile acids stimulate secretion of FGF15/19 from enterocytes (131), which in turn may act in the brain to suppress food intake and body weight, and improve glucose homeostasis (132, 133).

Gut microbiota

The majority of studies on gut microbiota and obesity are cross-sectional and simply describe associations between specific microbes and obesity. Although such studies are helpful, they cannot provide evidence for causal relationships. Only interventional studies that directly or indirectly manipulate gut microbiota and their signaling pathways can demonstrate the necessity of gut microbiota for the regulation of body weight/adiposity.

The effects of prebiotics (foods that stimulate beneficial microbiota, e.g., inulins, fructo-oligosaccharides), probiotics (live microorganisms such as *Lactobacillus* and bifidobacteria), and synbiotics (synergistic combinations of pre- and probiotics) in patients with obesity and diabetes have been studied in many randomized placebo-controlled trials, with a number of meta-analyses available. Although most of these meta-analyses find small beneficial effects, they also emphasize the poor quality of evidence due to methodological variability and low sample numbers. The largest of these meta-analyses was based on 105 studies comprising a total of 6826 subjects with overweight or obesity (134), and reported mean improvements of -0.94 kg in body weight and -0.55 kg/m² in BMI with probiotic treatment for 3–12 weeks in overweight but not obese subjects. It also found small improvements in fasting blood glucose and glycated hemoglobin (HbA1c) in type 2 diabetics, as well as small improvements in alanine and aspartate aminotransferase in subjects with fatty liver disease (134).

The gold standard for demonstrating a causal role for the gut microbiome in obesity is microbiota transplantation or fecal matter transplantation (FMT). A recent meta-analysis of six placebo-controlled studies with a total of 154 subjects with obesity and dia-

betes found that although mean HbA1c (-1.69 mmol/L) was lower and HDL-cholesterol (0.09 mmol/L) was higher 2–6 weeks after healthy-donor compared with control FMT, there were no significant effects on body weight and BMI after 6–12 weeks (135). Thus, the great promise for gut microbiota manipulations and their resulting changes in gut-brain communication as a cure for obesity has not yet been realized.

Gut-brain communication and bariatric surgery

Bariatric surgery is currently the most effective treatment to achieve sustained weight loss in patients with obesity, and altered gut-brain communication resulting in reduced appetite is the most widely presumed mechanism. Despite intensive preclinical and clinical research, no single mechanism has yet been identified to account for the weight loss, but altered signaling by gut hormones to the brain, either directly or via vagal afferents, has been among the leading candidates. In a large cohort of patients with or without prior vagotomy, Roux-en-Y gastric bypass (RYGB) produced the same weight loss up to 5 years (136). In contrast, selective transection of the celiac vagal branches, which innervate mainly the small and large intestines including the Roux limb, significantly attenuated RYGB-induced weight loss and hypophagia in rats (137). Furthermore, in mice, eating a meal during the first 2 weeks after RYGB leads to exaggerated activation of the brainstem anorexia pathway including the NTS, lateral parabrachial nucleus, and central amygdala, each thought to receive crucial input from vagal afferents (138–140). These observations are consistent with the idea that increased activity of vagal afferents mediates at least some of the early effects of RYGB on food intake and body weight, but further research with more selective sensory vagotomies will be necessary.

Because RYGB — as well as vertical sleeve gastrectomy (VSG) — greatly increases circulating plasma levels of GLP-1 and PYY, gut hormones known to decrease food intake, they were initially thought to play major roles in the beneficial effects on body weight (94, 141). However, subsequent studies in mouse models of bariatric surgery showed that RYGB-induced weight loss is similar in mice with deficient GLP-1R, PYY/Y2R, or combined GLP-1R/Y2R signaling compared with wild-type mice (142–145), suggesting that the critical signaling molecules have not yet been identified. Similarly, bile acid signaling through either TGR5 (146) or FXR (147) is not required for RYGB-induced weight loss, although both TGR5 signaling and FXR signaling play a role in improvements of glycemic control after RYGB (147) and VSG (148–150), and FXR signaling plays a role in VSG-induced weight loss (148). Overall, numerous studies testing single molecules or hormones have failed to identify the driving mechanisms of bariatric surgery. Alternatively, it is much more likely that bariatric surgery alters numerous processes that collectively result in satiety, weight loss, and other beneficial effects that contribute to sustained weight loss.

Conclusions

The vagus nerve is undoubtedly one of the most important links between the gut and the brain when it comes to ingestive behavior and metabolic regulation. The new generation of genetics-based neural manipulations, while fully confirming the distribution and structure, has prompted reassessment of the functional roles of vagal afferent fibers innervating the gut and associated organs.

As molecularly distinct clusters of vagal afferent neurons likely reflect function-specific populations, their selective manipulation should continue to revolutionize their functional anatomy, and the traditional, nonselective vagal manipulations should be abandoned. Similar strategies should also be applied to spinal afferents and autonomic nervous system outflow via the vagus nerve and SNS. It is hoped that a more comprehensive understanding of neural and non-neural pathways of communication will extend the therapeutic repertoire to treat obesity and metabolic disease.

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