Sensing via intestinal sweet taste pathways

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The detection of nutrients in the gastrointestinal (GI) tract is of fundamental significance to the control of motility, glycemia and energy intake, and yet we barely know the most fundamental aspects of this process. This is in stark contrast to the mechanisms underlying the detection of lingual taste, which have been increasingly well characterized in recent years, and which provide an excellent starting point for characterizing nutrient detection in the intestine. This review focuses on the form and function of sweet taste transduction mechanisms identified in the intestinal tract; it does not focus on sensors for fatty acids or proteins. It examines the intestinal cell types equipped with sweet taste transduction molecules in animals and humans, their location, and potential signals that transduce the presence of nutrients to neural pathways involved in reflex control of GI motility.

Keywords: sweet taste molecules, small intestine, vagal afferents, gastric emptying, glucose sensing, carbohydrate absorption

INTRODUCTION

The ability to sense the chemical composition of ingested material and respond appropriately is critical to individual survival. The gastrointestinal (GI) tract possesses specific sensors throughout its length which are tuned to detect the wide array of luminal stimuli present, and in response, direct neural and humoral signals that control motility, digestion, and absorption. Knowledge of the primary chemosensory mechanisms used by the GI tract is limited, and this will need to be addressed since it is fundamental to being able to manipulate digestive physiology in patients with a spectrum of disorders involving the gut, including obesity, malnutrition, diabetes mellitus, and those with functional GI syndromes.

It is well established that exposure of the proximal small intestine to the products of carbohydrate digestion generates signals that powerfully influence motor and sensory function of the GI tract. This small intestinal feedback is critical to the regulation of gastric emptying, which is tightly controlled in health in the range 1–3 kcal/min (Brener et al., 1983), and to the generation of GI symptoms and the secretion of peptide hormones that induce satiation or influence glycemic control. While a number of these responses are triggered via true humoral pathways upon interaction of nutrients with specialized gut epithelial cells, many result from paracrine activation of the intrinsic and extrinsic nerve pathways of the GI tract (see Cummings and Overduin, 2007; Raybould, 2010). There is strong functional evidence for vagus nerve pathways mediating chemosensory responses from the intestine. Early studies revealed that disruption of vagal pathways via subdiaphragmatic vagotomy or afferent denervation largely or completely blocked inhibition of gastric emptying, gastric acid secretion, and food intake in response to intestinal nutrient exposure in animals and humans (Wilkinson and Johnston, 1973; Roze et al., 1977; Raybould and Holzer, 1992; Schwartz et al., 1993). Over subsequent years, considerable progress has been made in understanding the activation and connectivity of vagus nerve pathways involved in nutrient reflexes, and their subsequent control of glycemia, food intake, and satiety (Raybould, 1991, 2010; Zittel et al., 1994; Lavin et al., 1996; Imeryuz et al., 1997; Broberger and Hokfelt, 2001; Blackshaw et al., 2007). It is now known that in the presence of intestinal nutrient, vagal afferent endings are activated by signals from epithelial cells and convey sensory information to the central nervous system for processing. In response to these signals, gastric vagal motor neurons mediate an “accommodation” response, involving relaxation of the proximal stomach (Azpiroz and Malagelada, 1987; Horowitz and Dent, 1991) together with suppression of antral contractions and the stimulation of tonic and phasic pyloric pressures (Figure 1). The net effect of these actions is to slow gastric emptying to a rate matched to the absorptive capacity of the small intestine (Raybould, 1998). The sensory arm of this reflex is critically dependent on the length and region of small intestine exposed to nutrient and prior nutrient exposure, while the motor events powerfully influence glycemic control (Lin et al., 1989; Horowitz et al., 2002; Little et al., 2006).

Vagal sensory innervation of the intestine arises from afferent neurons with cell bodies in the nodose and jugular ganglia, with endings concentrated in the upper GI tract. Anterograde nerve tracing studies have shown that vagal afferent nerves terminate within the parenchyma of mucosal villi in close contact with the basal lamina, but not with the epithelial surface (Berthoud et al., 1995). This anatomy largely precludes a direct action of luminal nutrient on vagal afferents and suggests that specialized epithelial cells within the intestinal mucosa serve as primary sense cells for nutrients. Specialized intestinal epithelial cells represent less than 1% of the intestinal cell population but collectively from the largest endocrine organ of the body. These regionally distributed mucosal cells possess receptors and transduction machinery to sense luminal stimuli, and in turn, collectively release over 20 different gut
peptides or bioactive molecules throughout the GI tract (see Rindi et al., 2004; Cummings and Overduin, 2007). As a consequence, the intestine is likely to sense nutrients at multiple sites and recruit diverse and cascading sensing strategies. In support of a paracrine action involving the enteroendocrine cells, vagal afferent nerves are known to express receptors for a number of key hormones released by gut epithelial cells, including those released in response to luminal nutrients such as carbohydrates (Kakei et al., 2002; Raybould et al., 2003; Nakagawa et al., 2004; Vahl et al., 2007; Bucinskaite et al., 2009). However, although paracrine mediators and vagal pathways involved in intestinal nutrient signaling are beginning to be revealed, the molecular identity of transduction mechanisms involved in nutrient detection in the GI tract remain poorly defined. In contrast, the identities of molecules involved in detecting sweetness in the tongue have been elucidated much more thoroughly (Nelson et al., 2001; Zhao et al., 2003).

Taste receptor cells in the oral cavity form the frontline of sensory detection of the GI tract and provide immediate and critical input to direct innate affective behaviors (to high caloric food sources) or aversive behaviors (to noxious substances). Taste buds comprise a heterogeneous group of 50–150 taste receptor cells, which are located throughout the tongue and soft palate. Taste receptor cells are classified on an ultrastructural basis into four cell classes – type I (dark) cells are candidate taste cells for salty taste and likely to have a support role akin to glial cells (Lawton et al., 2000; Vandenbeuch et al., 2008), type II (light) cells frequently possess apical microvilli and are individually receptive to either sweet, bitter, or umami (amino acid) stimuli (Yang et al., 2000; Clapp et al., 2004; DeFazio et al., 2006). Type III (presynaptic) cells are candidate sour taste cells and the only taste cell to form synapses with gustatory nerves (Huang et al., 2008; Kataoka et al., 2008), while type IV (basal) cells are presumed to be progenitor cells (Stone et al., 2002). Differences in the expression of taste receptors and transduction molecules in type II and III cells have also been reported between circumvallate and fungiform papillae, suggesting that these cells display differential sensitivities according to their physical location (Kim et al., 2003; Shigemura et al., 2008).

The attractive taste modalities, sweet and umami, are detected by a small family of class-C, G-protein coupled receptors (GPCR) – T1R1, T1R2, and T1R3 – which are related to metabotropic glutamate, extracellular calcium-sensing, and type B γ-aminobutyric-acid receptors. Subsets of T1R have been shown to heterodimerize to form a umami receptor (T1R1 + T1R3), responsive to the L-amino acids glutamate and aspartate, or to heterodimerize to form a broadly tuned sweet taste receptor (T1R2 + T1R3) responsive to sugars, D-amino acids, sweet proteins and sweeteners. It is also proposed that T1R2 or T1R3 may homodimerize in some circumstances to form low-efficiency sweet taste receptors (Hoon et al., 1999; Nelson et al., 2001, 2002; Li et al., 2002; Damak et al., 2003; Zhao et al., 2003; Delay et al., 2006). T1R receptor dimers have been shown to be expressed in individual taste receptor cells, equipping them to detect a single taste modality (Adler et al., 2000; Nelson et al., 2001; Yoshiida et al., 2009). Indeed, persuasive evidence now exists of a labeled-line model of taste discrimination which occurs at the level of the taste receptor cell (Chandrashkekar et al., 2006; Yarnolinsky et al., 2009).

T1R receptors possess a characteristic large extracellular N-terminus domain linked to a seven-transmembrane C-terminal domain by a short cysteine-rich domain. Bitter taste, in contrast, is mediated by a diverse family of around 30 class-C GPCR – the T2R – which possess short N-terminus domains and which are frequently co-expressed in bitter responsive taste cells (Chandrashkekar et al., 2000; Lindemann, 2001). A common intracellular signal transduction pathway is activated by T1R or T2R stimuli in individual taste receptor cells, involving activation of Gα, Gβγ, and Gβγ-subunits of the taste-specific G-protein, gustducin, followed by a signal transduction cascade involving phospholipase Cβ2 (PLCβ2) and inositol trisphosphate (IP3). This cascade results in the release of intracellular Ca2+ and gating of the taste-specific cation channel TRPM5 (Wong et al., 1996; Perez et al., 2002; Chandrashekar et al., 2006; Clapp et al., 2008). It is also well described that α-gustducin reduces intracellular levels of cAMP through activation of phosphodiesterases (Clapp et al., 2008). Depolarization then occurs, and mediators are released from taste receptor cells to activate gustatory nerves in a paracrine manner; norepinephrine, glutamate, acetylcholine, cholecystokinin (CCK), vasoactive intestinal peptide, neuropeptide Y, adenosine 5′-triphosphate (ATP), serotonin (5-HT), and the incretin hormone glucagon-like peptide-1 (GLP-1) have been identified as taste receptor cell mediators thus far (Nagahama and Kurihara, 1985; Nagai et al., 1998; Finger et al., 2005; Shen et al., 2005; Zhao et al., 2005; Roper, 2007; Martin et al., 2009).

PARALLELS BETWEEN TONGUE AND SMALL INTESTINAL TASTE MECHANISMS

There is unequivocal evidence that the presence of carbohydrate in the proximal small intestine slows gastric emptying in a response that is nutrient-specific, and not secondary to an increase in osmolality detected by intestinal osmoreceptors (Lin et al., 2001). The notion that taste mechanisms exist throughout the GI tract is drawn from similarities between specialized intestinal epithelial cells and...
lingual taste cells. Both cell types are polarized, possess apical microvilli that are exposed to the chemical environment of the lumen and are equipped to release neuromediators (Figure 2). Evidence from studies focused on carbohydrate absorption has also added support, with functional studies underpinning the actions of a specific glucose sensor on the luminal surface (Dyer et al., 2003; Margolskee et al., 2007).

A key advance in understanding sensory systems of the GI tract was the work of Höfer et al. (1996) and Höfer and Drenckhahn (1998), who identified mRNA and protein expression of \(\alpha\)-gustducin in brush cells of the gastric cardia, duodenum, and pancreatic duct of rats. This finding built on earlier work of this group in which GI brush cells were shown to express high levels of nitric oxide synthase (Kugler et al., 1994) and raised the possibility of nitric oxide as the first putative mediator of this transduction pathway. Subsequent studies in a number of rodent intestinal cell lines and tissues revealed that sweet taste receptors T1R2 and T1R3, along with their signal transduction partner \(\alpha\)-gustducin, were expressed in the small intestine at mRNA and protein level (Dyer et al., 2005; Margolskee et al., 2007). Our work extended this key finding to reveal that immunolabeling for \(\alpha\)-gustducin was expressed throughout the GI tract with peak expression in epithelial cells of the mid-jejunum (Sutherland et al., 2007a), where carbohydrate-induced reflexes are likely to be initiated (Lin et al., 1989). This study also explored the phenotype of intestinal epithelial cells in mice that expressed \(\alpha\)-gustducin (Figure 3); the majority of taste cells in the mid-jejunum were identified as brush cells (57%) or GLP-1-containing L-cells (27%) or GLP-1-containing L-cells (15%; Sutherland et al., 2007a).

These data support the hypothesis that lingual taste pathways are conserved throughout the alimentary tract and employed in primary intestinal chemosensation. However, they also indicate that the majority of intestinal taste cells in mice employ mediators other than GLP-1 or 5-HT in intestinal signaling. A further key finding was that, like lingual taste receptor cells, multiple components of the sweet taste GPCR pathway were expressed in individual intestinal taste cells. This evidence was provided at protein level by Mace et al. (2007), who showed co-localization of T1R2 and T1R3 in jejunal epithelial cells in rats using immunolabeling. Quantitative molecular approaches also revealed that T1R, T2R, and \(\alpha\)-gustducin were co-expressed within TRPM5-expressing duodenal epithelial cells in mice, demonstrating that a complete sweet taste transduction cascade could operate within these cells (Bezencon et al., 2007).

Expression of sweet taste molecules in the human GI tract was first shown by Rozengurt et al. (2006), who demonstrated immunolabeling for \(\alpha\)-gustducin in colonic epithelial L-cells that also expressed GLP-1 and peptide YY (PYY). Follow-up molecular studies from our group and others have since revealed expression of T1R2, T1R3, \(\alpha\)-gustducin, and TRPM5 throughout the human GI tract, noting prominent expression in the proximal small intestine (Bezencon et al., 2007; Young et al., 2009). One-third of taste cells (expressing \(\alpha\)-gustducin) in the duodenum have been shown to be L-cells, while a smaller proportion were found to be K-cells expressing glucose-dependent insulinoctopic peptide (GIP) or incretin cells that expressed both GLP-1 and GIP (K/L-cells; Theodorakis et al., 2006; Jang et al., 2007). This seminal study also revealed expression of sweet taste receptors and signal transduction partners in individual duodenal L-cells.

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*Figure 2 | Model of intestinal sensing by sweet taste cells and mucosal vagal afferents (adapted, with permission, from Bertrand, 2009).*

(A) Intestinal wall showing villus-crypt and location of sweet taste cells. Different sweet taste cells are shown within the epithelial layer (EPI, alternate colors) indicating the range of intestinal enteroendocrine and brush cells identified with sweet taste machinery. Vagal afferent nerve terminals are shown adjacent the basolateral membrane where they can be activated in response to paracrine signaling (neuromediators), triggering nutrient reflexes that alter behavior (food intake) and slow gastric emptying. (B) Expanded schematic of boxed area in (A), showing key components of the intestinal sweet taste signaling pathway proposed to operate in an enteroendocrine sweet taste cell; (1) Heterodimeric sweet taste receptors comprising the GPRC1R2 and T1R3 detect a wide range of sweet tastants in the intestinal lumen; (2) Upon GPCR binding, the taste-specific G-protein Gustducin is activated, liberating Gxα and G3β3 and Gy13-subunits which are thought to activate PLCβ2. (3) leading to the release of intracellular calcium from IP3-sensitive stores. Ga-gustducin may also reduce intracellular levels of cAMP via activation of phosphodiesterases (not shown). (4) Rising intracellular calcium can then gate the taste-specific cation channel TRPM5, leading to Na+ influx, membrane depolarization, neuromediator release, and (5) nerve terminal activation.
Young Intestinal sweet taste transduction

Youth intestinal sweet taste transduction glucose in anesthetized mice. We showed that transcript levels of T1R2 were markedly reduced in glucose-perfused jejunal loops, in the absence of changes in blood glucose level or levels of other taste molecule transcripts (Young et al., 2009). We also reported that the copy numbers of T1R2, T1R3, α-gustducin, and TRPM5 transcripts in human duodenal biopsies were inversely correlated with blood glucose levels at the time of endoscopy (Young et al., 2009). Taken together, these findings indicate that levels of sweet taste molecule transcripts are modulated in the intestine by both luminal and metabolic factors – a feature that may reflect their signaling of acute postprandial nutrients and chronic dietary intake patterns. Importantly, these data highlight the importance of considering prevailing and historic exposure of intestinal sweet taste receptors to their environment in order to accurately gauge their respective mRNA or protein levels.

There is also much interest in recent news that sweet taste molecules are widely distributed in other glucose sensing structures. T1R and taste G-proteins have now been identified in nutrient-sensitive forebrain structures in mice, where they register circulating glucose levels in a metabolism-independent manner and display similar transcriptional control mechanisms to sweet taste molecules of the intestine (Ren et al., 2009). Mouse pancreatic β-islet cells have also recently been shown to express T1R2, T1R3, α-gustducin, and TRPM5 transcripts in human duodenal biopsies were inversely correlated with blood glucose levels at the time of endoscopy (Young et al., 2009). Taken together, these findings indicate that levels of sweet taste molecule transcripts are modulated in the intestine by both luminal and metabolic factors – a feature that may reflect their signaling of acute postprandial nutrients and chronic dietary intake patterns. Importantly, these data highlight the importance of considering prevailing and historic exposure of intestinal sweet taste receptors to their environment in order to accurately gauge their respective mRNA or protein levels.

Much awaited functional evidence of intestinal sweet taste mechanisms was provided simultaneously in cell lines and in knockout mice deficient in α-gustducin. Firstly, mouse endocrine cells of the GLUTag line showed markedly increased GLP-1 and GIP secretion upon exposure to the sweetener sucralose, which was subsequently blocked by the sweet taste inhibitor gurmarin (Margolskee et al., 2007). Similar results were obtained for sucralose stimulated GLP-1 release in the human L-cell line NCI-H716, which was blocked by either RNA interference targeting of α-gustducin or the human sweet taste antagonist lactisole (Jang et al., 2007). Second, knockout mice lacking α-gustducin displayed a major deficit in glucose-stimulated GLP-1 release, along with a corresponding delay in reaching peak circulating insulin levels (Jang et al., 2007). Together, these results strongly linked the function of sweet taste molecules to intestinal incretin hormone release.

It is now well established that chronic changes in luminal glucose exposure modify GI response to glucose; for example, chronic increases in dietary carbohydrate reduce satiety satisfaction and accelerate gastric emptying in response to subsequent carbohydrate intake in healthy humans (Horowitz et al., 1996), while starvation is associated with delayed gastric emptying of glucose (Corvilain et al., 1995). Correspondingly, intestinal sweet taste molecules undergo dynamic regulation of expression in response to luminal or systemic glucose, which appears to be an essential feature of this system. Mace et al. (2007) first showed that exposure of rat jejunal membrane fractions to high concentrations of glucose or sucralose led to rapid trafficking of T1R2, T1R3, and α-gustducin protein away from the brush border. Our group subsequently identified transcriptional control of T1R2 by intestinal glucose in anesthetized mice. We showed that transcript levels of T1R2 were markedly reduced in glucose-perfused jejunal loops, in the absence of changes in blood glucose level or levels of other taste molecule transcripts (Young et al., 2009). We also reported that the copy numbers of T1R2, T1R3, α-gustducin, and TRPM5 transcripts in human duodenal biopsies were inversely correlated with blood glucose levels at the time of endoscopy (Young et al., 2009). Taken together, these findings indicate that levels of sweet taste molecule transcripts are modulated in the intestine by both luminal and metabolic factors – a feature that may reflect their signaling of acute postprandial nutrients and chronic dietary intake patterns. Importantly, these data highlight the importance of considering prevailing and historic exposure of intestinal sweet taste receptors to their environment in order to accurately gauge their respective mRNA or protein levels.

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**FIGURE 3 | Co-expression of α-gustducin within mucosal cell populations in mouse jejunum (from Sutherland et al., 2007a).**

Immunolabeling for α-gustducin “taste” cells in mouse jejunal mucosa (red fluorescence, A,E,I) was present in cells colabeled for 5-HT (green, B, GLP-1 (green, F), or brush cell-like labeled by the plant lectin Ulex europaeus agglutinin-1 (UEA-1; blue, J). Composite images (yellow, C,G; purple, K) depict localization within the same cells. The majority of α-gustducin cells in mouse jejunum were brush cell-like (57 ± 4% of all α-gustducin cells, L), while smaller proportions were 5-HT producing enterochromaffin cells (27 ± 2%, D) or GLP-1 producing L-cells (15 ± 2%, H). Scale bar = 50 μM.
fine-tuning systems responses for glucose homeostasis, and support the notion of multiple sensing strategies for carbohydrates (discussed further below).

**TASTE MECHANISMS IN SPECIALIZED INTESTINAL EPITHELIAL CELLS**

Understanding the phenotype of intestinal epithelial cells that bear sweet taste receptors is an important focus of a number of research groups, and candidate cells have been identified. Here, evidence of specific endocrine and brush cells involved in intestinal sweet taste transduction is reviewed, with reference to expression in cell culture, rodents and humans (Table 1).

**L-CELLS**

Much research effort has been focused on L-cells due to their ability to detect digestion products of carbohydrates, fats, and proteins and respond by secreting the incretin hormone GLP-1. L-cells are distributed throughout the GI tract, with greatest density in the distal intestine (ileum, colon); yet not insignificant numbers are present in the proximal small intestine in most species (Bryant et al., 1983; Eissele et al., 1992). These cells use alternative post-translational processing of proglucagon to produce the bioactive peptides GLP-1, GLP-2, and oxyntomodulin, although most attention has focused on the actions of GLP-1 (Holst, 2007). GLP-1 is known to be released rapidly in proportion to caloric load and the length of small intestine exposed to carbohydrate, and acts to increase satiation, stimulate insulin release, suppress glucagon secretion and slow gastric emptying (Schirra and Göke, 2005; Horowitz and Nauck, 2006; Little et al., 2006). In addition, complex carbohydrates that reach the distal intestine may be fermented to short chain fatty acids, which can be interacted with L-cells via other nutrient sensors (for review, see Reimann, 2010). Effects of GLP-1 on slowing of gastric emptying are mediated largely via a vagal reflex pathway; GLP-1 receptors have been localized on abdominal vagal afferents, on nodose ganglion neurons and in brainstem neurons (notably within the dorsal vagal complex; (Drucker and Asa, 1988; Goke et al., 1995; Imeryuz et al., 1997; Merchenthaler et al., 1999; Kakei et al., 2002; Nakagawa et al., 2004). While GLP-1 has also been shown to excite vagal afferents directly, the gastrointestinal inhibition achieved may occur in part via a central site of action on vagal motor neurons (Imeryuz et al., 1997; Kakei et al., 2002; Nagell et al., 2006; Nakade et al., 2006; Bucinskaite et al., 2009; Holmes et al., 2009; Ruttimann et al., 2009). However, the relative importance of a central site of action of endogenous GLP-1 is uncertain, due to the rapid degradation of GLP-1 that occurs within minutes at the site of peripheral release and within the circulation and liver by the ubiquitous enzyme dipeptidyl peptidase-IV (Deacon et al., 1996; Holst, 2007). The functional importance of GLP-1 in mediating reflex actions of intestinal carbohydrate on gastric emptying rate is now well established – genetic variation around the GLP-1 allele strongly correlates with changes in gastric emptying rate in mice strains (Kumar et al., 2008) while the GLP-1 antagonist exendin (9–39) accelerates gastric emptying in both rodents and humans (Imeryuz et al., 1997; Schirra et al., 2009; Deane et al., 2010).

L-cell lines of mouse and human origin were amongst the first to show expression of T1R2, T1R3, α-gustducin and to release GLP-1 via sweet taste receptor mechanisms (Reimann and Gribble, 2002; Dyer et al., 2005; Jang et al., 2007; Margolskee et al., 2007). Immunolabelling studies in mouse and human intestine subsequently revealed populations of L-cells equipped with α-gustducin that ranged from 15% (mouse mid-jejunum) up to 90% of all L-cells (human duodenum; Rozengurt et al., 2006; Jang et al., 2007; Sutherland et al., 2007b). Half of small intestinal L-cells in weaned piglets were also recently shown to be immunolabeled for T1R (Moran et al., 2010b). Much awaited functional evidence of intestinal taste signaling was provided by Jang et al. (2007), who showed that mice deficient in expression of α-gustducin

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**Table 1 | Sweet taste molecules identified in small intestinal cell populations.**

| Cell type | Major transmitters | Cell culture | Rodents | Human |
|-----------|-------------------|-------------|---------|-------|
| Enteroendocrine | | | | |
| L-cell | GLP-1 | T1R2, T1R3, GD | T1R2, T1R3, GD | T1R2, T1R3 GD, TRPM5 |
| | GLP-2, PYY, oxyntomodulin | | | |
| K-cell | GIP | T1R2, T1R3, GD | GD | |
| L/K-cell | GLP-1, GIP | | | |
| EC cell | 5-HT | T1R3 | GD | |
| | chromogranin-A, ATP, melatonin, GABA, uroguanylin, dynorphin | | | |
| I-cell | CCK | T1R2, T1R3, GD | | |
| Brush cell | Nitric oxide* | | T1R1, T1R3 GD, TRPM5 | |
| Enterocyte | None known | – | – | – |

*Gastrointestinal brush cells in rats express neuronal nitric oxide synthase (Höfer et al., 1996), which is not expressed in mouse intestinal bush cells (Sutherland et al., 2007b). Cell culture includes mucosal cells cultured from rodent and human origin. GD, α-gustducin; –, no expression; blank cell, not determined. Bold text indicates major transmitters for each cell type, unbolded text indicates minor transmitters.**
possessed impaired glucose-stimulated GLP-1 secretion, the first such evidence linking sweet taste signaling to incretin hormone responses. However, despite these multiple lines of evidence a number of studies have provided contrary views. Examination of highly purified L-cell populations from transgenic mice revealed only marginal levels of transcript for sweet taste molecules and an inability of sucralose to stimulate GLP-1 release from primary small intestinal cell cultures (Reimann et al., 2008). Whether these findings reflect the small proportion of L-cells known to express α-gustducin in rodents (Sutherland et al., 2007b), the prevailing cell isolation conditions or sensitivity of transcript detection in these studies is unclear. In addition, while a significant α-gustducin/L-cell population is reported to exist in human duodenum (Jang et al., 2007), our own studies indicate that primarily separate populations of cells express GLP-1 or α-gustducin in duodenal biopsies from healthy humans (Young et al., 2010). Whether this difference is due to the use of postmortem tissue in the studies by Jang et al. (2007) compared to our own studies using fresh tissue sources is unclear. It should also be noted that in all studies to date, less than half of all α-gustducin cells in the proximal small intestine of animals and humans co-express GLP-1, suggesting other mediators are likely to be important in signaling sweet and/or other taste modalities. Finally, while sweet taste receptor mechanism have been strongly linked to L-cell responses, the presence of other glucose sensing pathways are known in these cells, including electrolyte glucose transporters and metabolism-dependent mechanisms (Tolhurst et al., 2009; Reimann, 2010); these are discussed in turn below. The exact roles of sweet taste receptors in L-cells are not established, and key experiments are awaited to resolve their contribution to glucose-stimulated release of GLP-1 in animals and humans.

**K-CELLS**

Glucose-dependent insulinoentre-secreting K-cells are located within the proximal small intestine, and are responsive to digestion products of both carbohydrates and fat, as well as certain amino acids (Baggio and Drucker, 2007). GIP is rapidly released postprandially in proportion to caloric load (Pilichiewicz et al., 2007), and acts to stimulate insulin release and to promote lipid storage in adipocytes, the latter an emerging role that may link incretin hormone release in animals and humans co-express GLP-1, suggesting other mediators are likely to be important in signaling sweet and/or other taste modalities. Finally, while sweet taste receptor mechanism have been strongly linked to L-cell responses, the presence of other glucose sensing pathways are known in these cells, including electrolyte glucose transporters and metabolism-dependent mechanisms (Tolhurst et al., 2009; Reimann, 2010); these are discussed in turn below. The exact roles of sweet taste receptors in L-cells are not established, and key experiments are awaited to resolve their contribution to glucose-stimulated release of GLP-1 in animals and humans.

**ENTEROCROMAFFIN CELLS**

Serotonin (5-HT) secreting EC cells are a prominent enteroendocrine cell type in the GI tract and an important target for transducing a wide array of chemical and mechanical cues (for reviews, see Grundy, 2006, 2008; Blackshaw et al., 2007; Raybould, 2010). EC cells are distributed throughout the GI tract and exert paracrine effects via 5-HT receptor (gurmarin, stevioside) however do not appear to influence serotonin release and activation of 5-HT receptors present on intrinsic and extrinsic nerve endings within the submucosa; circulating 5-HT levels are kept low due to specific uptake into platelets. A number of other bioactive compounds have also been identified in EC cell populations, including chromogranin-A, melatonin, ATP, GABA, uroguanylin, and dynorphin although less is known of their actions in the intestine (see Bertrand and Bertrand, 2010). In contrast to other enteroendocrine cells in the GI tract, functional responses of native EC cell to luminal stimuli are less well characterized. It is known that carbohydrates can trigger 5-HT release from EC cell lines and from purified human small intestinal EC cells (Kim et al., 2001; Kidd et al., 2008). Correspondingly, 5-HT receptors are expressed on small intestinal vagal afferents in rodents and 5-HT receptor antagonists block vagal afferent activation, and suppression of food intake, as well as attenuate slowing of gastric emptying in rodents in response to intestinal carbohydrates (Zhu et al., 2001; Glatzle et al., 2002; Raybould et al., 2003; Savastano et al., 2005; Wu et al., 2005). Effects of 5-HT, receptor antagonists on carbohydrate-induced slowing of gastric emptying have not been reported in humans, and although a single study has described an attenuation of solid meal emptying in healthy subjects by a 5-HT receptor antagonist (Akkermans et al., 1988), most indicate no effect (Stacher et al., 1990, 1991; Netzer et al., 2002). Together these data suggest that the involvement of 5-HT3 receptors in gastric emptying reflexes is likely to be species-specific, but with a limited role in humans.

The presence of sweet taste molecules was first described in mouse intestinal EC cells by our group, where a quarter of midjejunal EC cells demonstrated immunolabeling for α-gustducin (Sutherland et al., 2007b). A small population of intestinal EC
cells in weaned piglets was also recently shown to be immunolabeled for T1R (Moran et al., 2010b). Based on studies in TRPM5-transgenic mice, however, it appears that if taste signaling is functional in EC cells, it is likely to operate independent of TRPM5 (Bezencon et al., 2008; Kokrashvili et al., 2009). Indeed, TRPM4 has been proposed as an alternative taste transduction partner based on lingual studies performed in TRPM5-deficient mice (Ohkuri et al., 2009). Transcripts for T1R3 and T2R have also been identified in purified human small intestinal EC cells, and despite the apparent absence of T1R2 transcript, these cells released 5-HT in response to stimulation with the sweetener sucralose (Kidd et al., 2008). Sucralose stimulation also led to phosphorylation of extracellular signal-related kinase in these EC cells, itself a cardinal feature of sweet taste receptor activation (Ozek et al., 2004; Jang et al., 2007). Human EC cells, however, also expressed the glucose transporter SGLT-1, and 5-HT responses were abolished in the presence of phloridzin (Kidd et al., 2008). Considering the role of T1R in the sweet taste response, it is conceivable that human EC cells may use T1R to conduct a distinct combination of sweet stimuli.

**I-CELLS**

Cholecystokinin-secreting I-cells are located largely within the proximal small intestine and are more responsive to digestion products of fats and proteins, than to carbohydrates (Rehfeld, 1978; Cummings and Overduin, 2007). However, it is well established that CCK is released by intraduodenal glucose infusions in humans, and like GIP release, is independent of the length of small intestine exposure (Liddle et al., 1985; Little et al., 2006). While considerable evidence is gathering for roles of bitter taste receptors and other nutrient receptors in intestinal CCK release, evidence from mouse enteroendocrine STC-1 cells (which secrete CCK) shows expression of transcripts for T1R2, T1R3, and α-gustducin (Wu et al., 2002; Dyer et al., 2005; Rozengurt and Sternini, 2007), indicating that sweet taste signaling may be involved in glucose-induced CCK release. However, STC-1 cells are poorly differentiated and multipotential and it remains to be established if the potential for sweet taste signaling in this cell line translates to CCK release mechanisms in native I-cells.

**BRUSH CELLS (TUFT OR CAVEOLATED CELLS)**

Brush cells are sparse within the epithelium of the respiratory and GI tracts of animals, and are rare in humans (Reid et al., 2005). These cells possess a distinct pear shape and apical microvilli similar to lingual taste cells, but unlike enteroendocrine cells, do not contain secretory granules or transmitter vesicles. The possibility of brush cells participating in nutrient signaling was first recognized by Höfer and Drenckhahn (1996, 1998) and Höfer et al. (1996), who identified solitary brush cells immunopositive for the taste G-protein α-gustducin in the gastric cardia, duodenum and pancreatic duct of the rat, based on cell morphology and immunolabeling for the cytoskeletal markers villin and cytokeratin-18. These brush cells were also immunopositive for neuronal nitric oxide synthase (nNOS) suggesting a possible role for nitric oxide (NO) in taste signaling from brush cells in rats. In our studies, we used binding of the plant lectin Ulex europaeus agglutinin-1 (UEA-1) as a species-specific marker for the glyocalyx of brush cells in mice (Gebert et al., 2000). We revealed that over half of solitary taste cells in the mouse small intestine (immunopositive for α-gustducin) bound UEA-1 (Sutherland et al., 2007b). However, despite the use of multiple antibodies against nNOS, all brush taste cells in mice were immunonegative for nNOS (Sutherland et al., 2007b). This finding reveals that intestinal brush cells are likely to contribute significantly to taste signaling in mice via an as yet unidentified, non-nitrergic mediator. To this end, TRPM5-expressing intestinal cells in transgenic mice were recently shown to resemble brush cells morphologically and immunologically, and to express high levels of the opioids β-endorphin and Met-enkephalin in the apical portion, as well as uroguanylin (Kokrashvili et al., 2009). However, in the same study these cells were also shown to be unresponsive to glucose or a sweetener ex vivo, indicating an apparent lack of sweet taste signaling mechanisms. This is an intriguing finding given that a high proportion of the same cell population identified by another group expressed T1R1, T1R3, and α-gustducin but not T1R2 (Bezencon et al., 2007), and may point toward a signal role of this cell type for umami stimuli. Alternatively, this finding may indicate that TRPM5-independent signaling of sweet taste may operate in subsets of intestinal brush cells (or in endocrine cells) as occurs in different lingual taste receptor cells (Ohkuri et al., 2009). Other signal candidates for brush cells include the gaseous mediators carbon monoxide or hydrogen sulfide (Miller et al., 1998; Bezencon et al., 2008; Linden et al., 2008).

Due to the unmet requirement of a definitive marker of brush cells in humans little data exist for comparative taste molecule expression in human GI brush cells, despite their rarity. In our own recent study GI epithelial cells immunopositive for α-gustducin in humans displayed similar morphology to rodent brush cells (Sutherland et al., 2008), however other data in humans has primarily focused on taste molecule expression in endocrine cells. Ultrastructural studies have however revealed that as in rodents, GI brush cells in humans may be extensively innervated at their basal membrane (Morroni et al., 2007; Bezencon et al., 2008), providing further support of a receptive role for this cell type.

**PURINERGIC CELLS**

A role for purinergic signaling in lingual taste transduction is now well established. Early studies revealed the presence of ionotropic P2X2 and P2X3 receptors in gustatory nerve endings innervating fungiform and vallate taste buds (Bo et al., 1999) and the expression of P2Y receptors coupled to calcium mobilization in taste (Kim et al., 2000; Baryshnikov et al., 2003; Kataoka et al., 2004). P2Y receptors also have been shown to direct paracrine signaling events with 5-HT between taste receptor cell types (Huang et al., 2007). The development of P2X2/P2X3 double knockout mice has provided key information on functional roles of these receptors. These mice display major deficits or complete loss of behavioral and lingual afferent responses to all taste stimuli (Finger et al., 2005). The function of P2X receptors in intestinal nutrient sensing has not been directly investigated, although P2X receptor antagonists have been shown to partially block the motor responses of the small intestine to luminal nutrient in guinea pigs (Gwynne and Bornstein, 2007). Recent developments in lingual
taste research have also led to the identification of the ATP-release hemichannels pannexin-1 and connexin-43 in taste receptor cells responsive to sweet, bitter, and umami stimuli (Romanov et al., 2007; Huang et al., 2009). Moreover, functional evidence of a role for pannexin-1 channels/ATP in lingual taste signaling have been very recently revealed in type II taste receptor cells (Murata et al., 2010); it is likely that markers of pannexin-1 will have future application in probing the phenotype of intestinal taste cells in animals and humans.

**OTHER MECHANISMS OF PERIPHERAL GLUCOSE DETECTION**

**K_{ATP}, CHANNEL-DEPENDENT GLUCOSE SENSING**

ATP-sensitive K_{ATP} channels are archetypal glucose sensors which act as metabolism-dependent ion channels in a number of tissues, most notably in pancreatic β-cells. These channels close in the face of increasing intracellular levels of ATP (and falling MgADP) that accompany glucokinase-dependent glucose metabolism. Due to subsequently reduced background K+ ion flux, small inward currents then facilitate membrane depolarization, which trigger action potentials, Ca2+ entry via voltage-gated Ca2+ channels and insulin secretion (Rorsman, 1997). It is well established that K_{ATP} channels (as well as glucokinase) are expressed in both cell lines and native intestinal L-cells and K-cells in mice and humans (Reimann et al., 2005; Theodorakis et al., 2006; Nielsen et al., 2007; Parker et al., 2009). Despite this, in vivo evidence for a dominant role of K_{ATP} channels in intestinal glucose sensing is currently lacking. Furthermore, it is well established that elevated blood glucose levels per se do not trigger incretin hormone secretion, nor do sulfonlyureas, which effectively inhibit K_{ATP} channels and are used therapeutically to stimulate insulin secretion in human type 2 diabetes (El-Ouaghlidi et al., 2007).

**ELECTROGENIC TRANSPORT (FACILITATIVE GLUCOSE TRANSPORTERS)**

Observations that incretin hormone release was triggered by non-metabolizable substrates of intestinal SGLT-1 in vitro and blocked by its selective inhibitor phloridzin led the concept that SGLT-1 acted as a glucose transceptor, capable of transport and signalization glucose (Sykes et al., 1980; Ritzel et al., 1997; Gribble et al., 2003). SGLT-1 transports glucose across the apical brush border membrane of intestinal epithelial cells with strict stoichiometry to Na+ (1:2) and was shown to generate a net inward current in GLUTag cells sufficient to trigger membrane depolarization and GLP-1 secretion (Gribble et al., 2003). It was noted, however, that inward currents generated in these in vitro studies were very small, and that facilitative background current(s) would be required to trigger electrical activity (Tolhurst et al., 2009), reducing the prospect of SGLT-1 acting as the primary intestinal glucose sensor. However, the first in vivo evidence of glucose-stimulated incretin hormone release that was blocked by phloridzin was recently revealed in mice (Moriya et al., 2009). This finding indicates that SGLT-1 transport may well be a critical determinant of incretin hormone release, at least in this species. SGLT-3, a closely related family member to SGLT-1 that bound with low-affinity but did not transport substrates has also been proposed to serve as an intestinal glucose sensor (Diez-Sampedro et al., 2003; Freeman et al., 2006). However, other studies have indicated that SGLT-3 is not expressed on the intestinal cell surface, and therefore unlikely to be involved in primary glucose responses (Dyer et al., 2007).

A number of animal studies have established that intestinal expression of SGLT-1 is increased in the presence of a broad array of sweet ligands, including artificial sweeteners. This process is now known to occur independent of metabolism and via a sweet taste receptor pathway (Shirazi-Beechey et al., 1991; Dyer et al., 2003; Moran et al., 2010a; Stearns et al., 2010). Critical to this understanding was evidence provided by Margolskee et al. (2007), who showed that intestinal SGLT-1 was not upregulated in response to dietary glucose or sucralose in mice deficient in α-gustducin or T1R3. Along with sweet taste receptors, vagal reflex pathways also appear to play an important role in the control of SGLT-1. Evidence in animals subject to selective vagal afferent ablation has shown that the expected increase in SGLT-1 protein is lost in response to acute and chronic intestinal glucose exposure, and to circadian pattern at anticipated feeding times (Bates et al., 1998; Stearns et al., 2008, 2010). The efferent limb of this vagal reflex is currently unknown, however both cholinergic and opioid receptors may be involved (McFadden et al., 1984; Bastidas et al., 1989). Such a neural reflex may explain the observation that glucose exposure of isolated regions of the GI tract in rats leads to increased SGLT-1 in non-exposed regions (Stearns et al., 2010). It is also that a humoral reflex pathway is triggered by intestinal glucose to modulate distant SGLT-1 expression, for example, GIP release has been proposed as a mechanism linking intestinal sweet taste receptor activation to SGLT-1 regulation in mice (Margolskee et al., 2007; Singh et al., 2008). 5-HT also appears to be involved in this signaling in rats, as the 5-HT3 receptor antagonist ondansetron reduced accumulation of SGLT-1 in response to glucose (Stearns et al., 2010). Together, these observations in animals strongly support the presence of a sweet taste receptor pathway that coordinates intestinal glucose transport capacity in accord with luminal stimuli, and which acts via reflex vagal pathways.

This mechanism, however, may not extend to humans. A study recently performed by our group to address this possibility in healthy subjects failed to identify any effect of sucralose to enhance acute absorption of intraduodenal infused glucose (Ma et al., 2010). However, the concentration of infused glucose in this study may have saturated available glucose transporters, and therein, masked a possible synergy between sucralose and glucose. It has also yet to be shown whether prolonged exposure to dietary sweeteners can increase small intestinal glucose absorption in humans as it does in mice (Margolskee et al., 2007), or if responses in humans may be sweeteren specific.

The facilitative glucose transporter GLUT2 (which normally transports monosaccharides across the basolateral membrane into circulation) is also regulated by sweet taste receptors in rats. Mace et al. (2007, 2008) have established that intracellular trafficking of GLUT2 and insertion into the apical brush border membrane of intestinal epithelial cells in rats occurs via a sweet taste receptor mechanism. These data imply that, at least in rats, sweet taste receptors link the presence of sweeteners or high levels of glucose in the intestine to increased glucose transport capacity via SGLT-1 and GLUT2. However this mechanism may be species-specific as apical GLUT2 is not detected in intestinal epithelial cells from pigs following carbohydrate supplemented diets (Moran et al., 2010b). Moreover, the rare incidence of glucose–galactose malabsorption in infants (which results from mutations in the SGLT-1 gene; Turk et al., 1991), argues against the presence of a mechanism in humans to translocate GLUT2 to the brush border membrane to otherwise facilitate monosaccharide absorption.
INTESTINAL SWEET TASTE MOLECULES IN DIABETES

Animal models of diabetes and humans with diabetes consistently show altered sensitivity of vagal feedback from the proximal intestine in response to the proximal intestine to carbohydrate. In particular, patients with diabetes exhibit exaggerated upper GI motility reflexes in response to nutrient infusion in the small intestine (Rayner et al., 2002) and up to 50% with longstanding diabetes experience delayed gastric emptying and associated symptoms, such as nausea, abdominal discomfort, and disturbed glycemic control following a meal (Horowitz et al., 2002). Current management of these patients is often suboptimal, due to a lack of understanding of the underlying pathophysiology. It is known that while the prevalence of delayed gastric emptying is high in diabetic patients with autonomic neuropathy, its presence only weakly correlates with disordered cardiovascular autonomic function (Horowitz et al., 1991), suggesting other pathogenic factors are likely to be more important. Correspondingly, an understanding of the regulation of intestinal sweet taste molecules in diabetic patients is likely to be important in assessing their altered sensitivity of nutrient feedback.

Our study was the first to compare intestinal levels of sweet taste molecules in patients with well controlled type 2 diabetes to those without diabetes. This revealed no significant difference in absolute copy number of these transcripts in duodenal biopsies of fasted molecules in patients with well controlled type 2 diabetes to those www.frontiersin.org

Adler, E., Hoon, M. A., Mueller, K. L., references

brain imaging (Nie et al., 2005; Frank et al., 2008). Moreover, mice from sucrose, and shows differences in central activation in functional 2009; Little et al., 2009; Ma et al., 2009). In this respect sucralose is the slowing of gastric emptying (Gregersen et al., 2004; Fujita et al., 2008; Baggio, L. L., and Drucker, D. J. (2007). Biology of incretins: GLP-1 and GIP. Gastroenterology 132, 2131–2157. Baryshnikov, S. G., Rogachevskaja, O. A., and Kolesnikov, S. S. (2003). Calcium signaling mediated by P2Y receptors in mouse taste cells. J. Neurophysiol. 90, 3283–3294. Bastidas, J. A., Yeo, C. J., Schmieg, R. E. Jr., and Zinner, M. J. (1989). Endogenous opiates in the mediation of early meal-induced jejunal absorption of water and electrolytes. Am. J. Surg. 157, 27–32. Bates, S. L., Sharkey, K. A., and Meddings, J. B. (1998). Vagal involvement in die- that lack T1R3 or TRPM5 show a complete loss of taste preference to sucralose, but retain residual responses to sucrose suggesting the presence of T1R3 or TRPM5 independent sweet signaling (Damak et al., 2003, 2006). It is, however, currently unknown whether chronic exposure of the intestine to sweeteners exerts intestinal effects; such an effect, for example, may explain the enhanced GLP-1 release in response to a high fat diet seen following chronic exposure of mice to the non-digestible sugar oligofructose (Cani et al., 2006).

It is also of interest that mouse colonic epithelial cell cultures exposed to sucralose trigger GLP-1 release in a manner that is additive with glucose, suggestive of sweet taste synergism (Reimann et al., 2008). A recent human study has added further support in revealing that ingestion of diet soda (sweetened with sucralose and asaculfame-K) prior to oral glucose load modestly increased GLP-1 release in healthy young volunteers (Brown et al., 2009). These authors concluded that while sweeteners did not release GLP-1 per se, they synergized with glucose to increase GLP-1 release (Brown et al., 2009). It is also well established that multiple allostERIC binding sites exist for T1R2 and T1R3 and that lingual sweet taste synergism readily occurs between combinations of sweeteners known to occupy both orthosteric and allosteric binding sites (Schiffman et al., 1995; Xu et al., 2004; Morini et al., 2005; Zhang et al., 2010). Whether taste synergism translates to in vivo function of intestinal sweet taste receptors is not established, and the potential for interaction between sugars (which bind the orthosteric site) and sweeteners that bind allosteric sites on the intestinal T1R2 receptor is yet to be explored.

CONCLUSION

Based on their relevant distribution and emerging intestinal functions there is much evidence to support sweet taste receptors as the luminal-facing GPCR tuned to carbohydrates and sweeteners in the GI tract. However, key experiments are still needed to determine the signaling mode of sweet taste cells in the human intestine and the contribution of their signals to the triggering of gut hormone release and vagal-mediated nutrient reflexes. In addition, the emerging role of sweet taste receptors in the regulation of intestinal sugar transport capacity remains to be clarified. Despite these caveats, the recent and rapid identification of GPCR involved in lingual and intestinal nutrient detection offers real potential to manipulate digestive physiology for therapeutic gain in patients with disorders of gut motility or sensation, glycemic control, or food intake.

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