Human Biology of Taste

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Taste or gustation is one of the 5 traditional senses including hearing, sight, touch, and smell. The sense of taste has classically been limited to the 5 basic taste qualities: sweet, salty, sour, bitter, and umami or savory. Advances from the Human Genome Project and others have allowed the identification and determination of many of the genes and molecular mechanisms involved in taste biology. The ubiquitous G protein–coupled receptors (GPCRs) make up the sweet, umami, and bitter receptors. Although less clear in humans, transient receptor potential ion channels are thought to mediate salty and sour taste; however, other targets have been identified. Furthermore, taste receptors have been located throughout the body and appear to be involved in many regulatory processes. An emerging interplay is revealed between chemical sensing in the periphery, cortical processing, performance, and physiology and likely the pathophysiology of diseases such as diabetes.
taste, and pain. Research on filiform papillae has lagged behind taste systems, but will be critical in understanding the influence of texture.

Taste buds are the primary sensory unit of the taste system and are imbedded under the keratinous layer of the papillae with a taste pore exposed to the external milieu. Figure 1 shows that taste buds are composed of 150 to 300 tightly packed cylindrical cells of epithelial origin. At least, 5 types of cells make up a taste bud: type 1, 2, 3 cells, basal cells, and neuronal processes (Figure 2). The various types of taste cells were originally characterized by the presence or absence of dense granules. Evidence now suggests that each taste modality is mutually exclusive to a subset of individual taste cells or 1 taste modality for 1 taste cell. For example, a type 2 sweet sensitive cell would express sweet receptors, but would not express bitter or umami receptors and vice versa. Sour is thought to be located on type 3 cells and sodium on type 1 cells. It has been shown that type 2 taste cells release adenosine triphosphate (ATP) in response to tastant activation. Figure 3 shows the current understanding of type 2 and 3 cell communication. ATP released from stimulated type 2 receptor cells activates P2Y adenosine receptors on nearby type 3 cells, releasing serotonin and stimulating afferent fibers to the CNS.

Although sensory processing at the level of the taste bud is complex, the transfer of information to the CNS seems to be via a labeled line. Three CNs innervate the tongue: the chorda tympani (CN-VII), glossophrangeal (CN-IX), and trigeminal (CN-V). The chorda tympani innervate the anterior fungiform papillae of the tongue. The glossophrangeal innervates the circumvallate and foliate papillae of the posterior portion of the tongue. The trigeminal nerve receives information from the filiform papillae and from various nerve endings throughout the oral cavity. Taste information is projected to the insula of the gustatory cortex, where a gustotopic map has been created. Each individual taste has a “hot spot” in the insular that responds to a particular taste.

Taste Mechanisms

Sweet

Sweet taste is one of the most hedonically pleasurable senses. The goal of sweet taste is to detect highly caloric saccharides for ingestion. Sweet taste is hardwired into our genes known as tas1R2 and tas1R3. Even newborn children will show positive stereotypical behavior when exposed to sugar solutions.

Figure 4A shows that sweet responsive type 2 taste
cells express the C class receptors, G protein–coupled receptors (GPCRs) T1R2 + T1R3. 15 See Urwyler, 2011, for an excellent review of C Class GPCRs. 16 GPCRs are a ubiquitous class of proteins that function to detect extracellular signals and transmit that information to the cell. 17 It is estimated that GPCRs make up 1% of the human genome and that 50% of drugs target these proteins. 17 C class GPCRs are heterodimers with a large extracellular domain, a cysteine rich hinge, region and 7 transmembrane domain (7-TMD). The 7-TMD is composed of 7 α-helices per subunit, which thread through the cell membrane. When a molecule binds to a GPCR, a conformational change occurs in the protein, resulting in the activation of an intracellular heterotrimeric G protein composed of alpha, beta, and gamma subunits that can stimulate multiple downstream pathways. Well-characterized pathways exist via mobilization of calcium stores from the endoplasmic reticulum by activation of phospholipase beta 3, modulation of phosphodiesterase or adenylyl cyclase (AC)/guanylyl cyclase. 18,19 These pathways modulate the second messengers inositol triphosphate and cAMP, cyclic guanosine monophosphate (cGMP) resulting in depolarization of the taste cell and release of neurotransmitter.

The sweet receptor functions as a dimer with an active site described as a venus fly trap module (VFTM). 20 VFTM has been shown to close upon ligand binding and activate their respective G proteins. Small sugars, such as glucose, are thought to bind into the VFTM of the sweet taste receptor. T1R2 and T1R3 each have a VFTM and are thought to bind to different sweeteners. 21 This may be the basis for synergies and allosteric modulation of sweet taste. Furthermore, the large extracellular domain, the cysteine-rich linking region and the 7-TMD collectively allow for the binding of a rich array of modulators, agonists, and antagonists. 16

Evidence has shown that T1R2/3 is expressed more widely than just the taste buds. Sweet taste receptors have been shown to be expressed in the K and L cells of the intestine, the beta-cells of the pancreas, bladder and hippocampus of the brain. 22–25 The functional significance of the sweet taste receptor expression is hypothesized to be the requirement for sugar sensing in these tissues and the implication for a wider role in regulating metabolism for T1R2/3. Initial evidence came from studies with the immortalized enteroendocrine cell line NCI-H716. 26 These cells express T1R2/3 and the machinery to release glucagon-like peptide-1 (GLP-1) in response to sweet stimuli. GLP-1, peptide tyrosine tyrosine (PYY), and gastrointestinal inhibitory peptide are key hormones mediating the incretin effect, or the gut stimulation of insulin release from beta-cells of the pancreas. 27,28 T1R2/3 regulation of incretin secretion was examined in rats and humans. It was shown that enteroendocrine L-cells co-expressed T1R2/3, GLP-1 and PYY. 29 Furthermore, data show that glucose-stimulated GLP-1 and PYY secretion was blocked by the T1R2/3 blocker lactisole. 30 In additional studies, the incretin effect was isolated by intragastric or intraduodenal perfusion of glucose or mixed meal (17% protein, 30% fat, and 53% carbohydrate), again with or without lactisole. 30 Intragastric infusion of glucose showed a significant stimulatory effect on GLP-1 and PYY secretions in vivo and this effect was blocked by lactisole, suggesting a role for T1R2/3. However, no effect was observed for liquid-mixed meal perfusion, suggesting the sweet taste receptor alone was not responsible for hormone release. Additional work is needed to determine if cephalic phase signals primes incretin effects. 30

A major question arising from this research is as follows: Do artificial sweeteners alter the incretin effect and if so what are the physiological consequences? Past studies looking at the effect of artificial sweeteners on a variety of hormone levels in the blood have shown little effect from artificial sweeteners in vivo. 31 However, pronounced effects have been reported in vitro. 31 Studies suggest that local activation of sweet taste receptor from L-cells are thought to cause paracrine release of GLP-1 resulting in translocation of glucose transporters of the brush border in preparation for sugar absorption and distribution. Further evidence for the role of T1R2/3 in metabolism comes from the research showing that the loss of the incretin effect is a specific and early marker for type 2 diabetes. 32 Moreover, studies with T1R2 or T1R3 knockout mice showed similar phenotypes to

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Figure 3. Communication between type 2 and type 3 taste receptor cells. Type 2 cells contain the GPCRs for sweet, bitter, and umami. Activation by ligands stimulates a G protein cascade, releasing intracellular stores of calcium that causes release of ATP. ATP then binds to P2Y receptors on the type 3 presynaptic cells, resulting in serotonin release to stimulate afferent neurons.
gastric bypass mice. Since gastric bypass surgery has the potential to reverse type 2 diabetes, the inference is that aberrant T1R2/3 function is a major factor in the progress on type 2 diabetes and that T1R2/3 is a potential drug target. In fact, the Gymnema sylvestre plant from India is a potent sweet taste blocker and has several thousand years of use as a homeopathic treatment for diabetes. The potential role for the sweet taste receptor in obesity and diabetes is provocative and will require much study, but the benefits of preventing or curing these costly diseases is clearly justified.

**Umami**
Umami is the Japanese word for the savory taste of amino acids, such as monosodium glutamate (MSG). Umami taste was first described in Japan by Kikunae Ikeda in 1908. Controversy surrounded the idea of umami as a primary taste until tas1R1 + tas1R3 was shown to code for the umami receptor. Figure 4B illustrates that the umami taste receptor is also a C class GPCR and contains a common subunit T1R3 with the sweet taste receptor and a unique subunit T1R1. Umami receptors show classical allosteric modulation of MSG response by GMP and IMP. The affinity of MSG can be lowered by an order of magnitude by the addition nucleotides, and this fact has been used in the food industries for over a hundred years to enhance the flavor of savory meals. Although the human umami receptor is very promiscuous, responding to all 20 amino acids, it has the highest affinity for glutamate.

**Bitter**
Currently 25 T2R bitter receptors have been identified in humans and belong to the A class GPCRs family. Figure 4C illustrates the A class receptors function as monomers and have small extracellular domains. It is amazing to note that ~25 bitter receptors can identify a seemingly endless array of compounds. Many pharmacologically active compounds are bitter, which can be a major hindrance to oral compliance. Potassium bitter taste may not be mediated by a T2R-dependent mechanism, but by a potassium-sensitive ion channel on type 2 cells. The identity of this channel is currently under investigation.

**Sour**
Previously it was show that the polycystic kidney disease (PKD) channel PKD1L2 + PKD3L1 mediates sour taste. These receptors have been found on type III taste cells and appear to function by allowing protons to traverse the membrane. The depolarization of the taste cell then stimulates the release of neurotransmitter. Cholecystokinin and neuropeptide Y are candidate neurotransmitters for the sour signal transduction. A study has suggested that PKD channels, mediates sour taste from circumvallate and foliate papillae. In a study of PKD knockout mice, sour taste response was inhibited only 25% to 45%, suggesting a secondary mechanism. Acid sensing ion channel (ASIC) were proposed as the mechanism for the fungiform papillae of the tongue. The researchers found...
that ACIS were present on taste cells and were sour active.

Salt
Salt mechanism has been quite controversial over the last decade. Epithelial sodium channel (ENaC) was an ideal candidate for the salt taste channel, and in rodents ENaC plays a central role in sodium taste; however, in humans ENaC seems to play very little role (Figure 4E). The sodium channel blocker, amiloride, effectively blocks sodium-dependent evoke potentials from rodents and blocks sodium preference, but in humans, amiloride has little to no effect in humans. Some have shown evidence for a second salt detection system. This alternative salt detection system mediates sodium and potassium tastes and is thought to be the ASIC and is much less sensitive to salt concentration; however, today no definitive evidence has identified the human protein responsible for salt taste.

In conclusion, an understanding of human taste biology has exploded over the last decade, but many questions still remain. It is remarkable that food quality and intensity can be coded by just 5 basic tastes; however, when combined with retronasal input from the approximately 300 human olfactory receptors, we perceive a large variety of flavors. The rich flavors of our human diet are the sum of taste, olfaction, and trigeminal input, but the synthesis of perception is the sum of peripheral input modulated by emotion, physiological and metabolic state, and learning. It is now becoming clear that taste receptors have a deeper role then just quality control. They also play an important role in maintaining nutrient homeostasis. How these sensory modalities interplay to regulate weight and satiety will be the next challenge in human nutrition.

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