

Shh Signaling and Regulatory Gene Expression in Mouse Taste Buds

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Introduction

In mammals, taste buds arise from local epithelium and are maintained by continuous cell renewal (Beidler and Smallman, 1965; Farbman, 1980; Delay *et al.*, 1986; Stone *et al.*, 1995). The life span of taste cells is estimated to be ~10 days. Denervation causes the degeneration of taste buds, indicating that the taste nerves trophically maintain the taste buds. It is assumed that cell proliferation, differentiation and death for taste buds are regulated in a coordinated fashion. However, the mechanism of each process in taste buds remains to be clarified.

Shh signaling in taste buds

Sonic hedgehog (*Shh*) is an evolutionarily conserved intercellular signaling molecule and *Patched1* (*Ptc*) is its receptor (Ingham and McMahon, 2001). *Shh* signaling is known to mediate various inductive events, including cell proliferation and differentiation, in animal development. We have examined the expression of *Shh* signaling-related genes in taste buds of adult mice. The expression of *Shh* was found in the basal cells in taste buds, and *Ptc* expression was observed in the surrounding region of the *Shh* expression (Figure 1; Miura *et al.*, 2001). This *Shh* and *Ptc* expression pattern was observed in every taste papilla (Figure 2). While *Ptc* expression itself can indicate the tissues responding to *Shh* during development (Goodrich *et al.*, 1996; Marigo *et al.*, 1996), the expression of *Shh* target gene *Gli1* was also observed in almost same region where *Ptc* was expressed, indicating that the cells responded to *Shh* signal (Figure 1). Analysis of BrdU incorporation showed mitotic cells were observed mainly in *Ptc*-expressing region, while mitotic cells in and around taste buds have been thought to contain the taste bud progenitors (Beidler and Smallman, 1965; Delay *et al.*, 1986). In addition, the expression of a downstream gene of *Shh*, *Nkx2.2*, was observed in a subset of taste cells (Miura *et al.*, 2003). *Nkx2.2* is a homeobox, which is induced by *Shh* in the ventral region of neural tube and is required for the specification of ventral neuron (Briscoe *et al.*, 2000). These data raise the possibility that *Shh* signal may be

involved in taste cell differentiation and that *Nkx2.2*-expressing cells in taste buds might be derived from the *Ptc*-expressing cells, which received *Shh* signal. The transection of cranial nerve IXth caused the

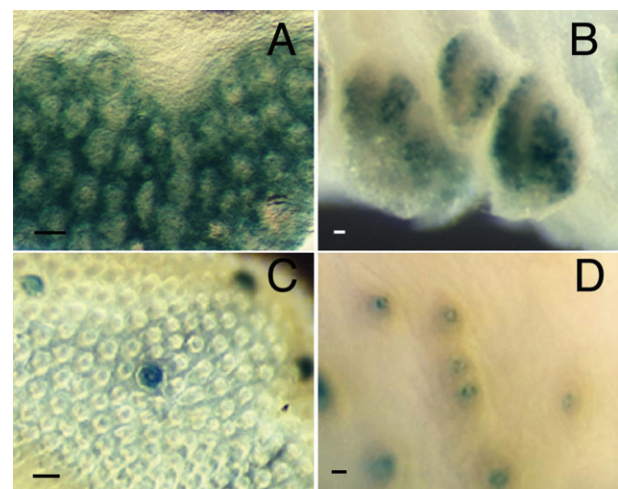


Figure 2 The expression of *Ptc* in taste papillae and soft palate. (A) Circumvallate papilla; (B) foliate papilla; (C) fungiform papilla; (D) soft palate. In order to detect the *Ptc* expression with high sensitivity in whole mount, *Ptc*^{+/-} mice, in which part of *Ptc* exon I and all of exon II were replaced with *lacZ* gene, were used, and the *Ptc* expression was detected by X-gal staining (Goodrich *et al.*, 1996). (A–C) The lingual epithelia were peeled off by collagenase and elastase treatment and used for staining. *Ptc* expression was observed in every taste bud. Scale bars indicate 50 μ m.

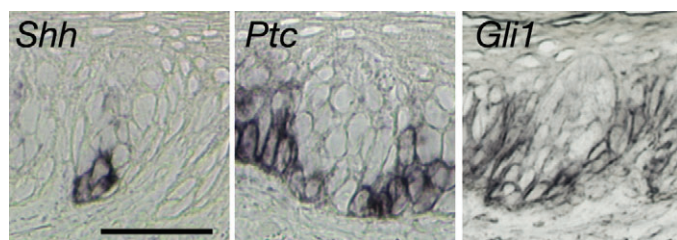


Figure 1 The expression of *Shh*, *Ptc* and *Gli1*. The expression of *Shh* was restricted to the basal cells in taste buds. *Ptc* was observed in the basal side of the epithelial cell around *Shh* expression. The expression of *Gli1* was observed in almost same region where *Ptc* was expressed. Scale bar indicate 50 μ m.

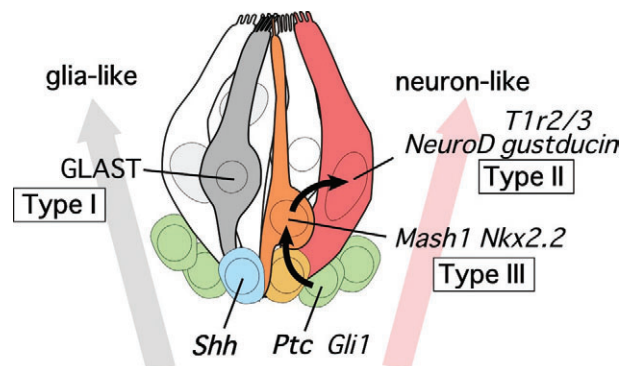


Figure 3 Hypothetical cell lineage in taste buds. *Shh* may be involved in expression of *Ptc* and *Nkx2.2*. Type II cell may be derived from type III cell, while some of type III cell may be in type III cell state throughout cell life. (See text for further details.)

rapid loss of *Shh* expression in the basal cells in taste buds of circumvallate papillae, suggesting the possible involvement of *Shh* signaling in the taste bud maintenance (Miura *et al.*, 2001).

Regulatory genes in taste buds and possible cell lineage

On the other hand, most *Nkx2.2*-expressing taste cells co-expressed *Mash1*, a bHLH transcription factor required for neuronal differentiation in CNS and olfactory epithelium (OE) (Miura *et al.*, 2003). We have previously reported that, in taste buds, the expression of *Mash1* segregated with those of taste reception-related genes, *gustducin* and *Tlr2*, and that *Mash1* expression preceded taste reception-related gene expressions during development (Kusakabe *et al.*, 2002). Based on these data, we speculated that *gustducin*- and *Tlr2*-expressing cells might be derived from *Mash1*-expressing cells. The expression of *NeuroD* in *gustducin*-expressing cells (Suzuki *et al.*, 2002) appears to be consistent with this idea by consideration of the sequential expression of bHLH transcription factors in neuronal differentiation in olfactory epithelium: *Mash1* is expressed in the early phase, and induces the expression of the other bHLH genes including *NeuroD* (Cau *et al.*, 2002). Based on ultrastructural characteristics, the taste bud cells have been categorized into four cell types—basal cell, type I (dark) cell, type II cell and type III (light) cell—and some immunohistochemical markers have been shown to correlate with each cell types: *gustducin* and NCAM are specific for type II and type III, respectively (Yee *et al.*, 2001). However, it is not clear whether these cell types represent separate cell lineages or transient cell states of one cell lineage (Pumplin *et al.*, 1997; Delay *et al.*, 1986). We found almost of all *Mash1*-expressing taste cells were NCAM-positive (unpublished data). Therefore, the hypothesis that *gustducin*-expressing cells derived from *Mash1*-expressing cells proposes the possible cell lineage of type II cell from type III cell. *Mash1* and *NeuroD* are bHLH factors critical for neuronal differentiation, while Type I cell has been considered as a glia-like cell (Lindemann, 1996; Lawton *et al.*, 2000). Recently, the expression of the genes related to Delta-Notch signaling in taste buds has been reported (Seta *et al.*, 2003). This signaling is known to be involved in neuron–glia cell fate determination, and therefore it may be possible for the Delta-Notch signaling to be involved in a cell-fate choice between neuron-like cell and glia-like cell in taste buds. Our hypothesis of taste cell lineage is summarized schematically in Figure 3.

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