

Molecular Mechanisms Underlying Differentiation and Maintenance of High Salt Taste Cells

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Summary

Taste perception is initiated when chemical substances in food are detected by taste cells located within taste buds in the oral epithelium. Humans can recognize five basic taste qualities: sweet, umami, bitter, salty, and sour. Each of the sweet, umami, bitter, and sour tastes is detected by distinct populations of taste cells. Salty taste is mediated by two separate pathways: low-concentration sodium taste and high-concentration salt taste. The former is detected by a unique population of taste cells distinct from those sensing sweet, umami, bitter, and sour, whereas the latter is detected by bitter and sour taste cells. This study focuses on *Eya1*, a transcription factor expressed in bitter taste cells (i.e., taste cells that detect bitter taste and high-concentration salt taste) and undifferentiated taste bud cells, to investigate its role in the differentiation and maintenance of taste cells responsible for bitter and high-salt taste detection.

To analyze the functions of *Eya1* in taste buds, we used *Eya1*-floxed mice in combination with two Cre-expressing lines: one expressing the tamoxifen-inducible CreERT2 recombinase in taste stem cells, and the other expressing Cre in mature sweet, umami, and bitter taste cells. In mice lacking *Eya1* in taste stem cells, bitter/high-salt taste cells were replaced by sweet and umami taste cells, indicating that *Eya1* is essential for the differentiation of bitter/high-salt taste cells. Similarly, in mice lacking *Eya1* in mature taste cells, expression of bitter taste receptors was markedly reduced, and bitter/high-salt taste cells were replaced by sweet and umami taste cells. Behavioral assays revealed that mice lacking *Eya1* in mature taste cells failed to avoid bitter substances such as denatonium and cycloheximide, as well as high concentrations of sodium chloride and calcium chloride. These findings suggest that *Eya1* is required not only for the differentiation but also for the maintenance of bitter/high-salt taste cells. Overall, this study demonstrates that *Eya1* is a key regulator of bitter and high-salt taste cell identity and function and that the *Eya1*-deficient mice generated here serve as a valuable model for investigating the molecular mechanisms underlying high-salt taste perception.