

# Histochemical Investigation of the Modal Specificity of Taste

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*The taste mechanism was investigated in a primate (Macaca mulatta). Based on the hypothesis that intracellular enzymes contribute to the transduction of tastes to electric impulses by taste cells, a histochemical survey of the activity of several enzymes was made on taste buds from regions of the mouth associated with sweet, salt, sour, and bitter tastes. Considerable differences were noted among the modalities, which confirmed the hypothesis. An exclusively bitter enzyme was identified.*

The sense of taste is unique among the "chemical" senses, in that it is associated with discrete receptor organs of non-neural derivation, the taste buds. Their existence in humans has been recognized for more than a hundred years,<sup>1</sup> but their exact mode of function remains obscure.

Classically, taste has been subdivided into four modalities, sweet, salt, bitter, and sour (acid). Each of these modalities has been localized to a discrete region on the surface of the tongue: sweet, to the fungiform papillae of the tip of the dorsum; salt, to the fungiform papillae of the lateral aspects of the anterior two thirds of the dorsum; bitter, to the circumvallate papillae of the posterior third of the dorsum; and sour, to the foliate papillae of the lateral aspects of the posterior third of the dorsum (Fig 1).<sup>2</sup>

In 1951, Baradi and Bourne<sup>3</sup> proposed a theoretical taste mechanism wherein the substance being tasted either inhibited or accelerated the rate of an on-going biochemical reaction in a taste region by alter-

ing the efficacy of the enzyme that catalyzed the reaction. A net change in a reaction, then, was the source of the taste impulse, and the fact that enzymes are highly specific accounted for the specificity of the taste impulse. If this hypothesis is valid, the taste buds from a region of predominantly one modality should display an enzyme complement different from that of a region of another modality.

More recent research<sup>4</sup> has shown that taste cells have differing enzyme complements at different stages in their fetal development; perhaps this reflects the developing, and therefore changing, level of function.

It is not unlikely that taste buds of differing modalities also have differing enzymatic compositions, although currently this is a topic of conjecture.

The enzyme complements of the taste buds of the four classical modalities were assayed histochemically, and their specificities, localizations, and intensities were determined.

## Materials and Methods

Adult rhesus monkeys (*Macaca mulatta*) were used. All monkeys were killed by vascular perfusion of physiologic isosmotic saline while they were under anesthesia. Tongues were removed by incision posterior to the circumvallate papillae. Immediately after excision, regions containing taste buds were removed from well within the boundaries of the areas classically assigned to each of the modalities (Fig 2). The tissue then was frozen in liquid nitrogen-isopentane, sectioned at 12 micrometers on a cryostat,\* and mounted on microslides. With each modality, the tissue blocks were oriented so that the microtome cut the taste

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Hatton award: 1st place, predoctoral division.

\* American Optical "Cryo-Cut," American Optical Co., Instrument Division, Buffalo, NY.

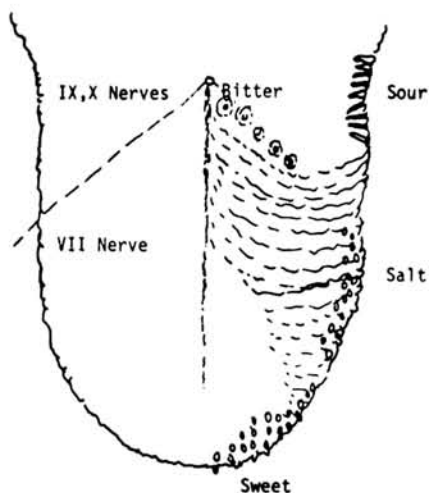


FIG 1.—Classical taste modality localization on the tongue.

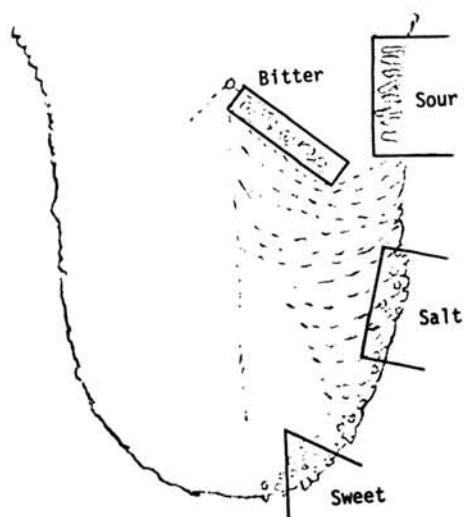


FIG 2.—Excision site for modality assays.

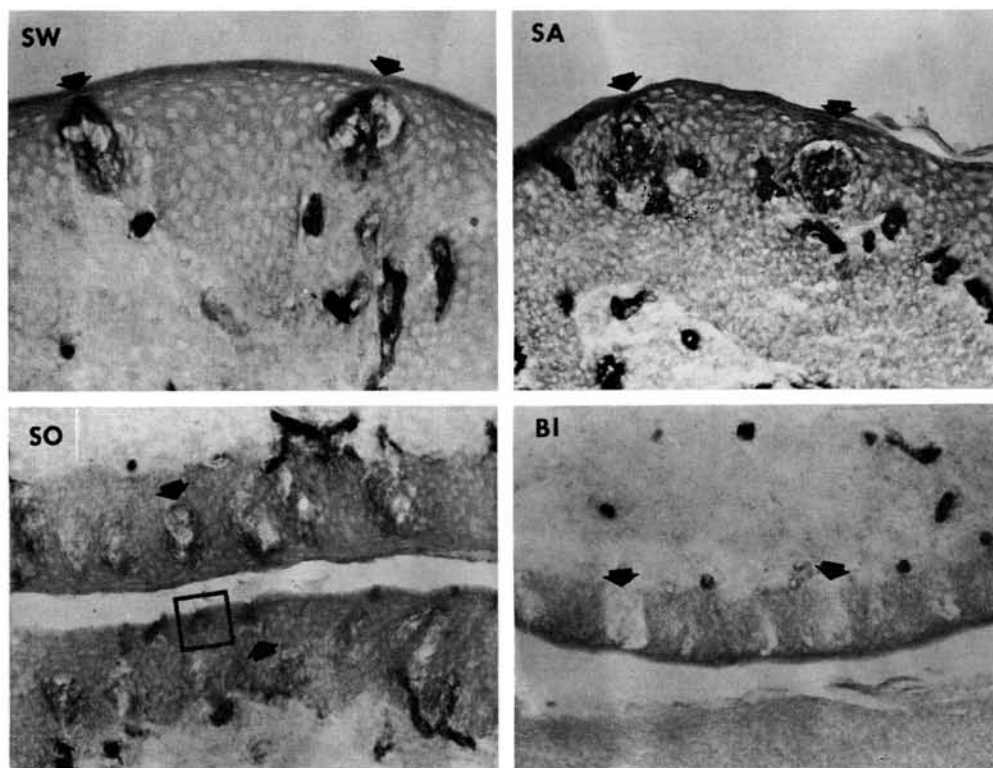


FIG 3.—Alkaline phosphatase. Sweet (*SW*) and salt (*SA*) displayed activity throughout taste buds (*arrows*), but sour (*SO*) and bitter (*BI*) stained only lightly and principally in the region of the taste pore (*square*) where the oral fluids contact the taste cells (orig mag  $\times 200$ , reproduced at 56%).

TABLE  
ENZYMATIC ACTIVITY IN TASTE BUDS

Enzyme	Sweet	Salt	Sour	Bitter
Succinic dehydrogenase	—	—	++	++++
Choline oxidase	+++	+	++++	+++
Cytochrome oxidase	+	+	+	++
Nonspecific esterase	++	—	+	++++
NAD diaphorase	++	+++	++++	++++
NADP diaphorase	+	+	+++	++++
$\alpha$ -Glycerophosphate dehydrogenase	+	+	+++	++++
$\beta$ -Hydroxybutyric dehydrogenase	—	—	—	+++
D-amino acid oxidase	+++	+++	+++	+++
Alkaline phosphatase	++++	++++	+	—
Glucose-6-phosphate dehydrogenase	++	—	++	+
Acid phosphatase	++	++	++++	++++
Adenosine triphosphatase	++	++	+	+++
Nucleotide diphosphatase	++	++	++	+++

buds in the long axis to facilitate comparison.

The following are the enzyme systems examined and the pathways they represent: succinic dehydrogenase<sup>5</sup> (Krebs cycle ac-

tivity); choline oxidase,<sup>6</sup> cytochrome oxidase,<sup>7</sup>  $\alpha$ -glycerophosphate dehydrogenase,<sup>8</sup>  $\beta$ -hydroxybutyric dehydrogenase,<sup>9</sup> and D-amino acid oxidase<sup>6</sup> (electron transport chain activity); nonspecific esterase<sup>10</sup> (mem-

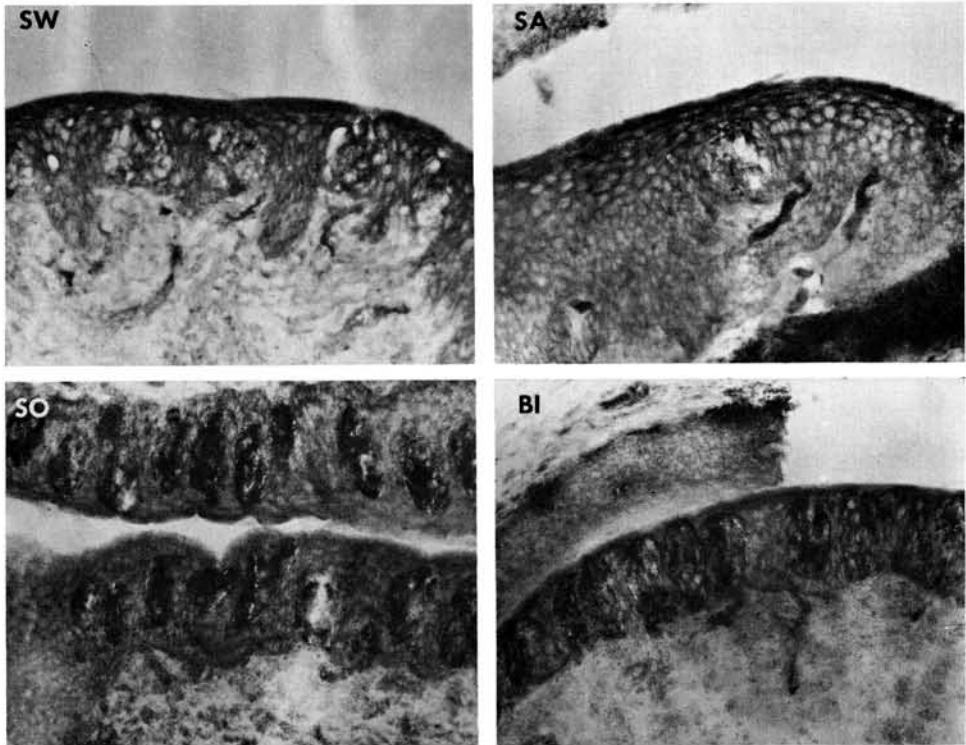


FIG 4.—Acid phosphatase. Sour (SO) and bitter (BI) taste buds stained darkly (great activity) and sweet (SW) and salt (SA) buds showed only minimal activity, also in the region of the pore (orig mag  $\times 200$ , reproduced at 58%).

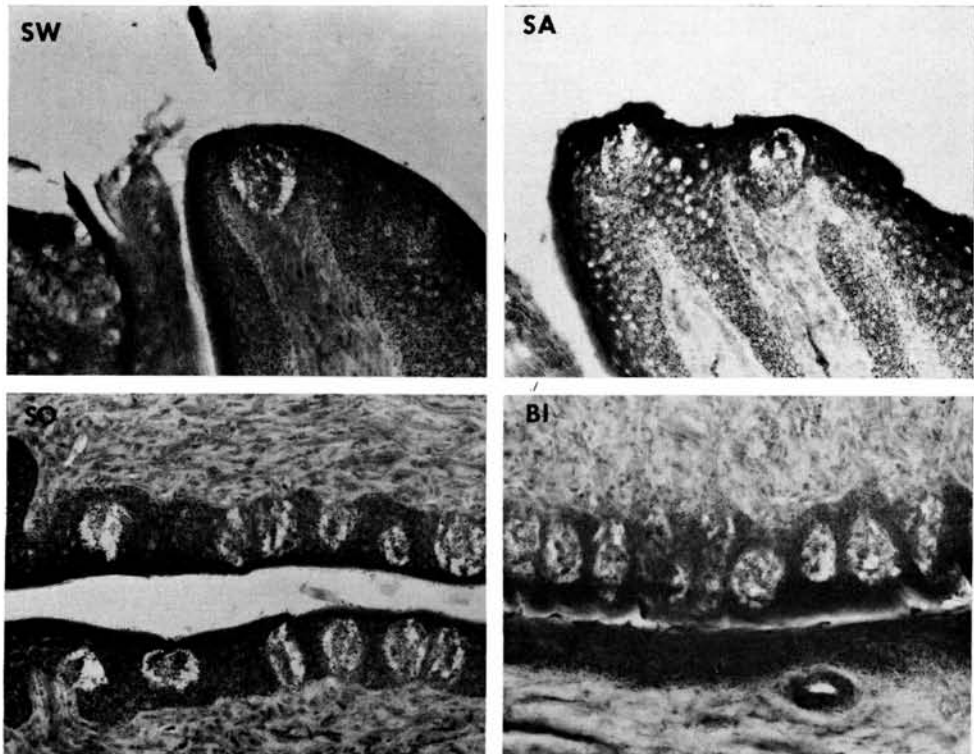


FIG 5.—Succinic dehydrogenase. Sour (SO) and bitter (BI) taste buds displayed more activity than sweet (SW) or salt (SA) buds, but there was no activity in the pore regions of sweet and salt buds (orig mag  $\times 200$ , reproduced at 57%).

brane-bound hydrolytic activity); NAD diaphorase<sup>11</sup> and NADP diaphorase<sup>7</sup> (membrane-bound pentose-phosphate shunt activity); glucose-6-phosphate dehydrogenase<sup>12</sup> (soluble pentose-phosphate shunt activity); adenosine triphosphatase<sup>13</sup> (mitochondrial activity); and nucleotide diphosphatase<sup>14</sup> (Golgi complex activity). Finally, because several investigators have attributed great importance to the role of hydrolytic enzymes in the process of taste,<sup>5,15</sup> alkaline phosphatase<sup>16</sup> (active transport activity) and acid phosphatase<sup>17</sup> (lysosomal activity) were examined. In all instances, the exact methods of the investigators cited were used. As a further control, all four modalities in each assay came from the same monkey, and were incubated simultaneously in the same medium for the same amount of time. In addition, at least two, and as many as six repetitions were performed for each assay until consistent results were obtained. All the data presented here repre-

sent the mean, consistent values obtained for each enzyme. The photomicroscope\* used was set for constant light intensity and exposure values.

### Results

Each of the enzymes studied was present in the taste bud cells, but response in the surrounding epithelium, from which the taste cells are derived, was variable. A synopsis of the observed enzymatic activity in the taste buds is presented in the table. Enzyme activities are indicated by the observed depth of the staining reaction in the range of - (absent) to ++++ (highly active).

The values in the table were obtained by microdensitometric measurement of stain intensity within the taste buds and by comparison of the mean obtained for each modality with the other modality means for each enzyme.

\* Carl Zeiss, Inc., New York, NY.

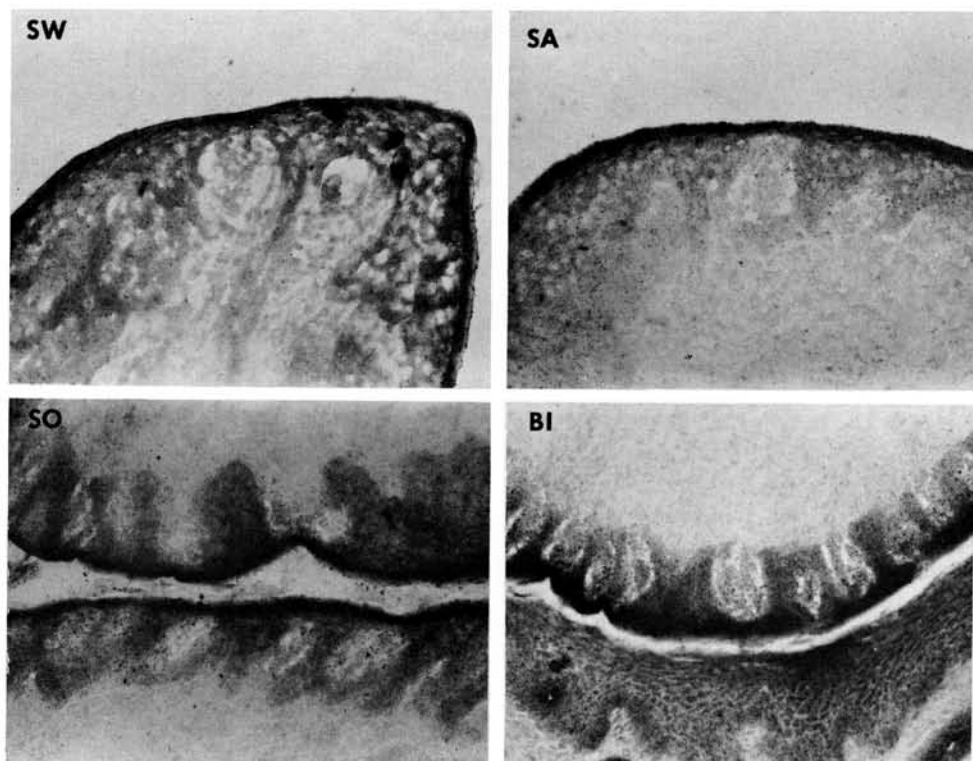


FIG 6.—Beta-hydroxybutyric dehydrogenase. Sweet (*SW*), salt (*SA*), and sour (*SO*) taste buds displayed little activity, but the bitter taste buds (*BI*) showed considerable activity, especially in the surrounding epithelium (orig mag  $\times 200$ , reproduced at 56%).

Alkaline phosphatase (Fig 3) reacted strongly in the sweet and salt modalities, and reacted hardly at all in the bitter and sour modalities. All activity seemed to be concentrated in the region of the taste pore in the sour and bitter buds, whereas the sweet and salt buds stained rather generally and very darkly. This situation is reversed in the acid phosphatase reaction (Fig 4); sour and bitter stained heavily throughout, and sweet and salt showed only slight activity, which is concentrated in the region of the pore. The characteristic staining of the capillary endothelium by the alkaline phosphatase reaction is evident and uniform through the different modalities.

The activity of succinic dehydrogenase (Fig 5) seemed to parallel that of acid phosphatase, with the exception of the pore regions of the sweet and salt buds, which displayed no special activity.

Beta-hydroxybutyric dehydrogenase (Fig

6) appeared to be specific for the bitter modality; all the other modalities displayed no activity. The reaction in the surrounding epithelium was especially notable.

NADP diaphorase (Fig 7) displayed a spectrum of activity that can be graded as least in sweet, progressively more in salt and sour, and most in bitter.

Alpha-glycerophosphate dehydrogenase (Fig 8) showed its strongest activity in the sour and bitter modalities, with heavy epithelial activity around the taste buds. This is especially apparent in the bitter (circumvallate papilla) region, where the heavily stained gustatory epithelium opposed a non-neural epithelium that showed essentially no activity.

Other enzymes studied, but not photographed, showed measurable but somewhat less striking results (Table): Cytochrome oxidase, D-amino acid oxidase, and glucose-6-phosphate dehydrogenase displayed fairly

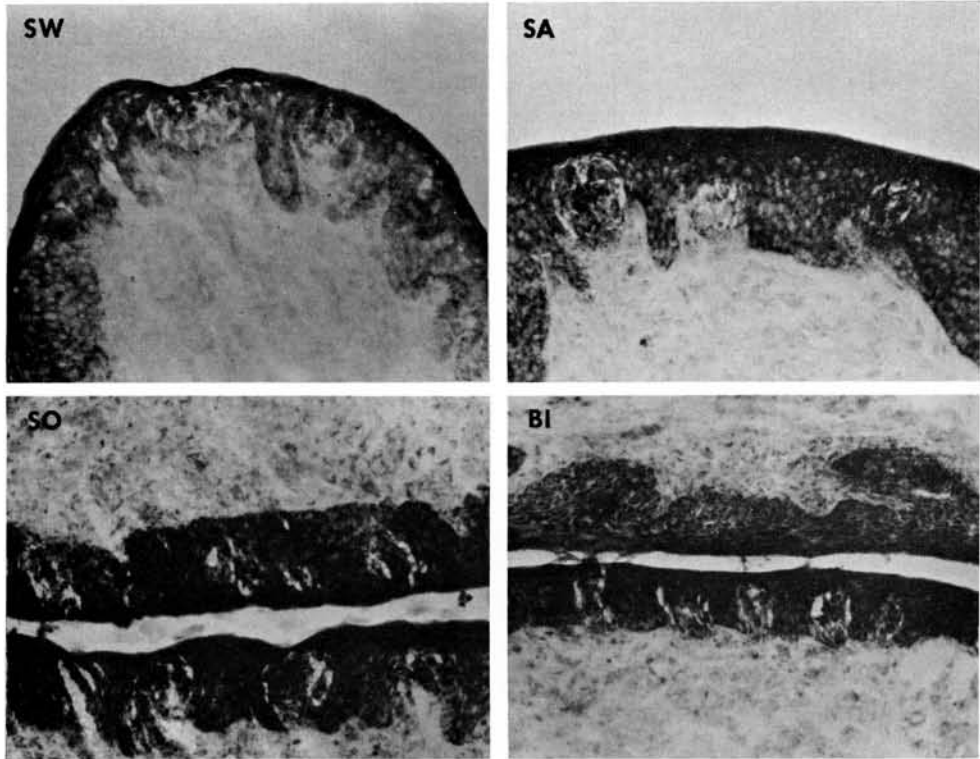


FIG 7.—NADP diaphorase. Taste bud enzyme activity was graded from least to most in the following order: sweet (*SW*), salt (*SA*), sour (*SO*), and bitter (*BI*) (orig mag  $\times 200$ , reproduced at 56%).

even staining from modality to modality. The same is true for adenosine triphosphatase and nucleotide diphosphatase, but sharp intracellular differences were apparent. Adenosine triphosphatase stained the cell membranes of all four modalities sharply, but the pore regions in the sour modality, and especially the bitter modality, displayed intense activity. Nucleotide diphosphatase, however, darkly stained the nuclei of only some of the cells of the buds of all four modalities and the pore regions in all modalities except salt. NAD diaphorase had a graded spectrum of activity that was similar, but not identical to that of NADP diaphorase. Choline oxidase and nonspecific esterase showed activity in all the modalities except salt; greater differences were noted with choline oxidase.

#### Discussion

Succinic dehydrogenase has long been used as a marker for the activity of the

Krebs cycle,<sup>5</sup> especially in the taste buds.<sup>18</sup> Its localization in significant amounts in sour and bitter taste buds, but not in sweet or salt buds, infers that in the former two modalities the missing electrochemical impulse conversion system of Baradi and Bourne,<sup>3,19</sup> is in some way linked to the Krebs tricarboxylic acid cycle.

High acid phosphatase activity in sour and bitter taste buds and low activity in sweet and salt buds is to be expected, because most sour and bitter substances are acidic to some degree,<sup>20</sup> and lysosomes, which contain the enzyme, have been observed in ultrastructural studies of foliate and circumvallate papillae in lower species.<sup>21,22</sup> In addition, it is a lytic phosphate-cleaving enzyme that lends itself to energy mediation.<sup>23</sup> This fact also is true of alkaline phosphatase, which is sweet- and salt-specific; but, in addition, alkaline phosphatase is an energy-mediating enzyme often associated with active transport at the cell-



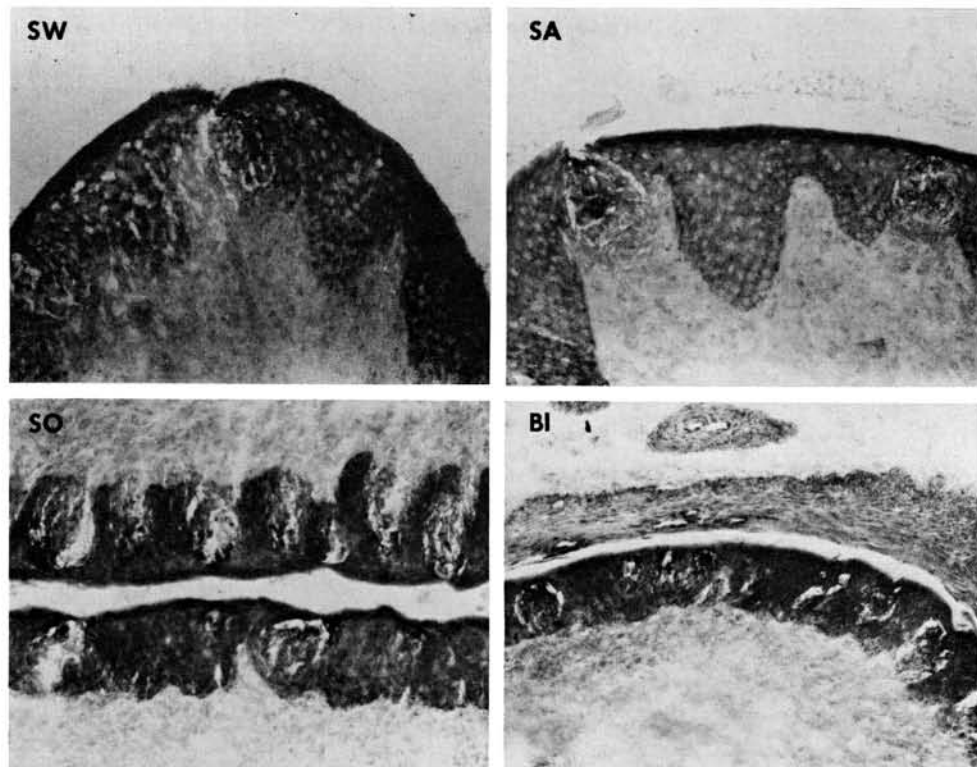


FIG 8.—Alpha-glycerophosphate dehydrogenase. More activity was evident in sour (SO) and bitter (BI) taste buds than in sweet (SW) or salt (SA) buds. Notable activity in epithelium surrounding sour and bitter taste buds was present (orig mag  $\times 200$ , reproduced at 56%).

membrane level,<sup>24</sup> as evidenced by its previously noted activity in capillary endotheliums. Alkaline phosphatase may be the electrochemical converter for sweet and salty tastes.

Choline oxidase,  $\alpha$ -glycerophosphate dehydrogenase,  $\beta$ -hydroxybutyric dehydrogenase, and D-amino acid oxidase are indicators of the so called electron-transport chain,<sup>25</sup> another cellular energy path. The varying results observed in the four modalities for these four enzymes points convincingly toward enzymatic specificity as the determining factor in taste specificity. In addition,  $\beta$ -hydroxybutyric dehydrogenase may well be a true "bitter enzyme."

The other enzymes assayed all showed varying differences both among themselves and among the four modalities. All were linked to either a metabolic pathway (NAD and NADP diaphorase, membrane-bound pentose phosphate shunt<sup>26</sup> and glucose-6-

phosphate dehydrogenase, free cytoplasmic pentose phosphate shunt<sup>27</sup>), or to a biologic function associated with a discrete ultrastructural entity (adenosine triphosphatase, mitochondria<sup>11</sup>; nucleotide diphosphatase, Golgi apparatus,<sup>28</sup> and nonspecific esterase, ribosomes and rough-surfaced endoplasmic reticulum<sup>29</sup>).

This wide variation of enzymatic activity, and wide variation of kinds and types of enzymes involved, easily explains why we are capable of more than 16 tastes (the maximum number of combinations and permutations of the four basic taste modalities). This study only considered a fraction of the enzymes that occur in cells, but variations were found from modality to modality in almost every enzyme studied. One notable exception is cytochrome oxidase, which is an indicator of anaerobic metabolic activity.<sup>30</sup> As expected, the activity is relatively constant from cell to cell; this may imply

that the ultimate transmission (or transduction or both) of the taste signal is by an aerobic pathway, rather than by an anaerobic, pathway.

An overall survey of the data in the table indicates the following generalized subgrouping, on the basis of enzyme activities; sweet with salt and sour with bitter. This may be of developmental significance, since sweet and salt taste buds receive their innervation from the VII nerve, and sour and bitter taste buds from the IX and X nerves (Fig 1), which supply structures derived from different branchial arches.

This fact is of medical interest, because diminutions in taste acuity of these modal subgroups that occur in certain human diseases have been noted and are well-documented.<sup>31</sup> Sweet functions with salt, and sour with bitter, and the two subunits operate quite independently of each other during the course of therapy.

### Conclusions

The mechanism of taste was studied histochemically in a primate. Notable differences in enzymatic activity were demonstrated among taste buds from tongue regions associated with sweet, salt, sour, and bitter taste. The hypothesis of Baradi and Bourne<sup>32</sup> that the specificity and transduction of the taste impulse is due to specific enzymatic involvement was confirmed. Functional association was noted between sweet and salt enzymatic activity, and between sour and bitter enzymatic activity; this agrees with clinical findings. A bitter-specific enzyme,  $\beta$ -hydroxybutyric dehydrogenase, was identified.

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