

Heterogeneity in the Accessory Olfactory System

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Abstract

The mammalian accessory olfactory bulb (AOB) is chemoarchitecturally heterogeneous in that it stains differentially with a number of markers; the receptor cells that project to the AOB are similarly heterogeneous. What is the significance of this heterogeneity? We have found that the AOB of the gray, short-tailed opossum, *Monodelphis domestica*, stains differentially with a number of 'markers': antibodies to olfactory marker protein (OMP) and the α subunit of the G protein G_{i2} , the lectin of *Vicia villosa* and NADPH-diaphorase. These markers stain the rostral AOB more strongly than the caudal AOB whereas, the G protein subunit $G_{\alpha\alpha}$ is located predominantly in the posterior subdivision of the AOB. This heterogeneity in the chemoarchitecture of the AOB may reflect a fundamental organizational dichotomy within the vomeronasal system that corresponds to a functional dichotomy. The vomeronasal sensory epithelium also exhibits a chemoarchitectural heterogeneity: receptor cells in the basal third are $G_{\alpha\alpha}$ -immunoreactive whereas the cells in the middle third are $G_{i2\alpha}$ -immunoreactive. Tracing studies using WGA-HRP demonstrate that the neurons in the middle third of the vomeronasal sensory epithelium project their axons to the anterior AOB whereas those in the basal third appear to project to the posterior AOB.

The main olfactory and accessory olfactory systems are represented in the periphery by sensory epithelia containing bipolar neurons that terminate in glomeruli of the main (MOB) and accessory (AOB) olfactory bulbs respectively. One of the prominent current hypotheses concerning chemosensory coding in the main olfactory system proposes that sensory receptor cells with different complements of receptor proteins on their distal processes send specific projections to glomeruli localized in particular sectors of the MOB. In the vomeronasal system, where the glomeruli of the AOB are less distinct, no mechanism of coding similar to that in the main olfactory system has been proposed.

It is generally believed that stimulus coding in the main olfactory system is a reflection of the topographical relationship between the receptor cells of the olfactory epithelium and the glomeruli of the MOB where their axons terminate (Figure 1). However, it has remained a contentious issue whether the glomeruli of the MOB represent a locus for the segregation of information derived from primary sensory neurons responding to distinct odorants or classes of odorants (Kauer, 1980, 1987; Kauer *et al.*, 1994; Schoenfield and Buck, 1994). One approach to this issue is to determine if there are clear differences in the olfactory bulb glomeruli that are reflective of differences in groups of receptor cells.

Characterization of biochemically defined subsets of olfactory neurons may result in identification of neuronal subsets with similar sensitivity spectra to odorants (Schwob, 1992). In rat and mouse olfactory systems bipolar neurons expressing distinct putative receptors are segregated topographically, to a certain degree, in the epithelium (Buck, 1993; Ressler *et al.*, 1993; Vassar *et al.*, 1993). However, within each topographic zone several different types of putative receptors may be expressed and the neurons expressing these different receptors are distributed, apparently at random, within the circumscribed zone. Thus, in the olfactory epithelium a partial pattern of spatial segregation based on receptor type is present (partial topographic correspondence; Figure 1). However, the role of this spatial pattern for the encoding of particular odor stimuli has yet to be determined.

In recent years a chemoarchitectural heterogeneity has been observed in main and accessory olfactory systems in both receptor cells and the nerve/glomerular layers of their respective bulbs. The heterogeneity in the accessory olfactory bulbs of rat, mouse, rabbit, hamster and opossum has been demonstrated with monoclonal antibodies raised to specific carbohydrate moieties (Mori *et al.*, 1987; Schwarting and Crandall, 1991; Schwarting *et al.*, 1992 a,b, 1994), lectin staining (Takami *et al.*, 1992; Taniguchi *et al.*, 1993; Shapiro *et al.*, 1995a), antibodies to G proteins

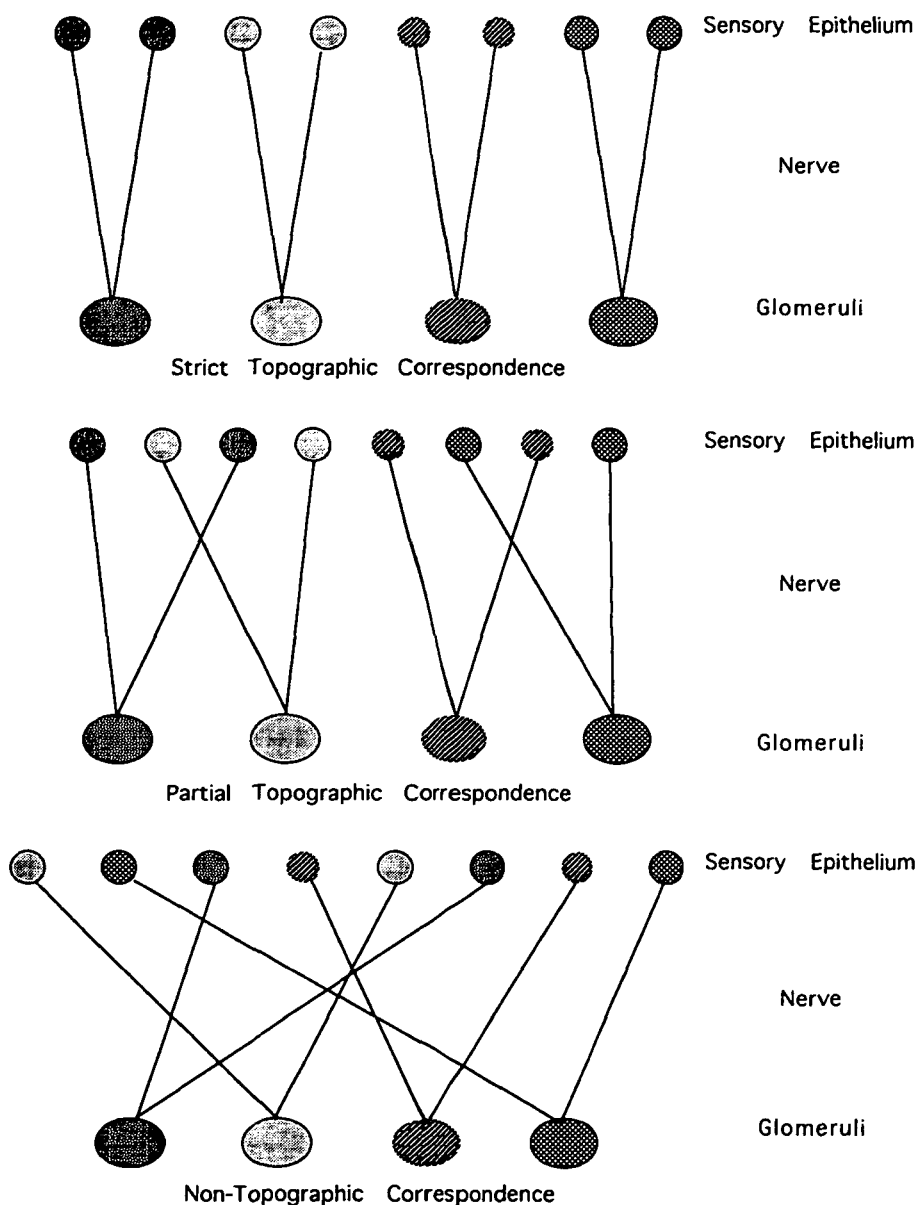


Figure 1 Three models of topographic correspondence between chemically identified cells in the sensory epithelium of chemoreceptive organs and their termination sites in glomeruli of the main or accessory olfactory bulb.

(Shinohara *et al.*, 1992; Halpern *et al.*, 1995), antibodies to olfactory marker protein (Shnayder *et al.*, 1993) and NADPH diaphorase histochemistry (Shnayder *et al.*, 1994; Halpern and Jia, 1995). Most recently, in rat, mouse and opossum two populations of vomeronasal receptor cells have been identified using antibodies to G proteins (Halpern *et al.*, 1995; Jia and Halpern, 1995a,b, 1996; Berghard and Buck, 1996) and each of these two populations appear to project to one of the two chemoarchitecturally defined subdivisions of the accessory olfactory bulb (Shapiro *et al.*, 1995b; Jia and Halpern, 1995a,b, 1996). In addition, an *in situ* hybridization study using probes to $G_{\alpha_{i2}}$ and G_{α_o}

confirmed the immunohistochemical findings of differential localization of receptor neurons expressing these proteins in the vomeronasal sensory epithelium (Berghard and Buck, 1996).

This chemoarchitectural heterogeneity in the opossum AOB and VNO is illustrated for the G proteins $G_{\alpha_{i2}}$ and G_{α_o} in Figure 2. We conclude from these observations, as well as those of Jia and Halpern (1996) and Berghard and Buck (1996) in the mouse, that the vomeronasal organ contains at least two types of receptor neurons located in different sublayers of the receptor cell layer. The axons of these cells intermingle in the vomeronasal nerve but become

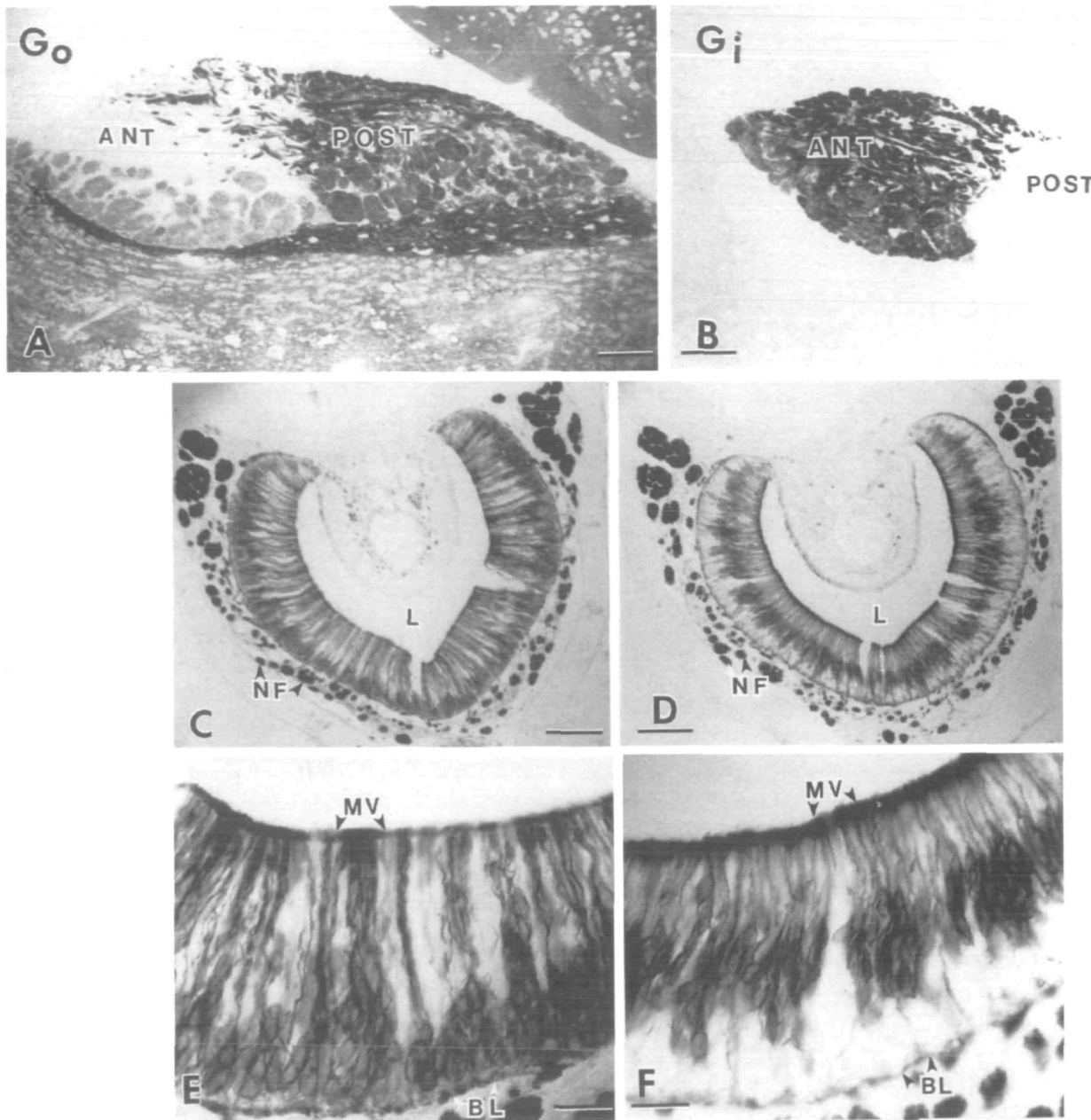


Figure 2 Photomicrographs of sections through the accessory olfactory bulb (**A** and **B**) and vomeronasal organ (**C**, **D**, **E** and **F**) of grey short-tailed opossums stained with antibodies to G_o (**A**, **C** and **E**) and G_i (**B**, **D** and **F**) proteins. Ant = anterior AOB; Post = posterior AOB; L = lumen of vomero-nasal organ; NF = vomeronasal nerve fiber bundles; MV = microvillar border of sensory epithelium; BL = basal lamina of sensory epithelium. Bar = 80 μ m in **A**, **B**, **C** and **D**; 20 μ m in **E** and **F**.

segregated at the level of the accessory olfactory bulb, where they terminate in distinct subdivisions of that structure (Figure 3).

What is the potential significance of this duality in the accessory olfactory system? It is clearly an anatomical substrate for maintaining separate channels of information from the periphery to more central structures. However, if it is the anatomical basis for sensory coding then we are in need of a demonstration that different chemical signals activate the separate channels. The heterogeneity may

represent use of different signal transduction mechanisms within the system. Certainly the universal demonstration, among mammals so far studied, that G proteins are differentially expressed in the two parts of the vomeronasal system supports this notion, as does the demonstration that β NADPH diaphorase reactivity is differentially present in the vomeronasal systems of some mammals. Another possibility is that the differential expression may be related to axon-substrate interactions that are involved in axonal guidance and matching of receptor cells with their

Current Organizational Scheme for Opossum Vomeronasal System

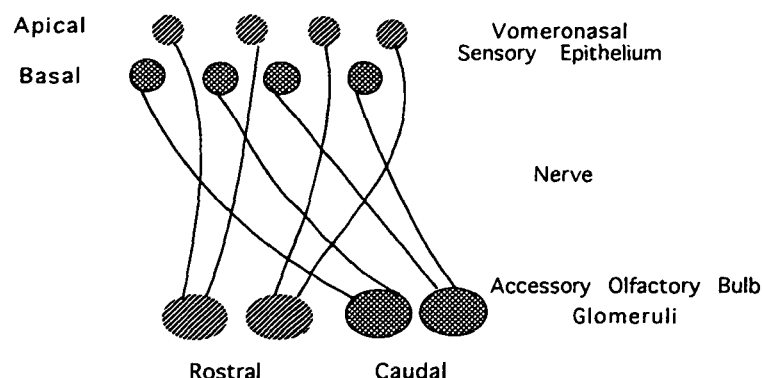


Figure 3 Current model of topographic correspondence between the chemically identified cells in the vomeronasal epithelium and the glomeruli of the accessory olfactory bulb.

appropriate targets in the AOB. For example, the two G proteins may be involved in pathway finding and target selection. G_i and G_o , both sensitive to PTX, have been implicated in neurite-promoting functions of cell adhesion molecules (Doherty *et al.*, 1991; Strittmatter *et al.*, 1991).

Comparisons among mammals of the chemoarchitectonic heterogeneity within the accessory olfactory system reveal some striking similarities and differences (Table 1). For example, whereas G_o is localized in the posterior AOB and G_i is localized in the anterior AOB in the rat, mouse and opossum (Shinohara *et al.*, 1992; Halpern *et al.*, 1995; Jia and Halpern, 1995a,b, 1996), the pattern of lectin staining (Takami *et al.*, 1992; Taniguchi *et al.*, 1993; Shapiro *et al.*, 1995a), olfactory marker protein localization and the pattern of NADPH diaphorase reactivity differs in these three groups of mammals (Davis, 1991; Porteros *et al.*, 1994; Shnyder *et al.*, 1994; Halpern and Jia, 1995). Do these differences between species represent fundamental differences in the organization of the accessory olfactory system or do they represent different methods for solving similar problems of sensory information transduction, coding and segregation? These issues are at present unresolved.

Acknowledgements

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