

# Dog and mouse: toward a balanced view of the mammalian olfactory system

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Ignacio Salazar, Unit of Anatomy and Embryology, Department of Anatomy and Animal Production, Faculty of Veterinary, University of Santiago de Compostela, Av Carballo Calero s/n, 27002 Lugo, Spain e-mail: ignacio.salazar@usc.es Although the most intensively studied mammalian olfactory system is that of the mouse, in which olfactory chemical cues of one kind or another are detected in four different nasal areas [the main olfactory epithelium (MOE), the septal organ (SO), Grüneberg's ganglion, and the sensory epithelium of the vomeronasal organ (VNO)], the extraordinarily sensitive olfactory system of the dog is also an important model that is increasingly used, for example in genomic studies of species evolution. Here we describe the topography and extent of the main olfactory and vomeronasal sensory epithelia of the dog, and we report finding no structures equivalent to the Grüneberg ganglion and SO of the mouse. Since we examined adults, newborns, and fetuses we conclude that these latter structures are absent in dogs, possibly as the result of regression or involution. The absence of a vomeronasal component based on VR2 receptors suggests that the VNO may be undergoing a similar involutionary process.

Keywords: olfactory epithelium, olfactory subsystems, morphology, evolution, dog

#### **INTRODUCTION**

The dog is increasingly appreciated in biomedical research as a species that, unlike purpose-bred laboratory animals, shares the genetic, and clinical variety of human patients (Karlsson and Lindblad-Toh, 2008). Its nasal cavity has been studied from various points of view. Anatomically, Graeger's paper (Graeger, 1958) is considered the classical reference. Physiologically, attention has recently focused on nasal airflow patterns (Craven et al., 2007, 2010). Clinically, the presentation, diagnosis, prognosis, and treatment of tumors is a major current concern (McEntee, 2004; Turek and Lana, 2012; Mason et al., 2013), but the olfactory mucosa of the nasal cavity has also attracted clinical interest because of its unique maintenance of a population of basal cells supporting the continual regeneration of olfactory sensory neurons (OSNs) (Graziadei and Monti-Graziadei, 1979); the intraspinal implantation of cells derived from autologous olfactory mucosa cultures has in fact recently been successful in ameliorating the effects of spinal cord injuries in companion dogs (Granger et al., 2012).

In the mouse, the animal in which the mammalian olfactory system has been most intensively studied, the nasal mucosa features four separate olfactory areas: the main olfactory epithelium (MOE), the septal organ (SO), the ganglion of Grüneberg (GG), and the vomeronasal sensory epithelium [VNsE, not to be confused with the vomeronasal organ (VNO) of which it forms a part] (Breer et al., 2006; Munger et al., 2009; Ma, 2010; Barrios et al., 2014). These four sensory areas can be considered as the points of entry to four olfactory subsystems (OSbS), the integration of which at higher levels is a focus of current research. However, this four-subsystem scheme is by no means exhibited by all mammals, or even by all macrosmatic mammals (Salazar and Sánchez-Quinteiro, 2009). In the work described here we examined its validity for the dog, a notoriously macrosmatic animal. We found that the dog has no GG or SO, and that its VNO shows signs of similar involution.

# MATERIALS AND METHODS

#### ANIMALS

The dogs studied were 31 male or female mesaticephalic adults, 17 newborn males or females from four different litters, and 16 fetuses obtained on days 30, 35, or 40 of gestation. Most were German Shepherds or mongrels derived therefrom, and all were mesaticephalic. All were obtained legally through the dissecting and post-mortem rooms and Department of Clinical Science of our faculty and were treated in accordance with Spanish and EU legislation for the care and handling of animals in research (RD 223/1998, 86/609/EEC) and with the guidelines of the University of Santiago de Compostela Bioethical Committee. The heads of all animals were intact and showed no clinical or post-mortem evidence of neurological disease; all were processed as specified below as soon after death as was possible.

#### **PROCESSING OF SAMPLES AND TISSUE SECTIONS**

Using traditional anatomical techniques, 22 adult heads were carefully prepared by dissection and micro-dissection—from outside to inside—to afford views of the lateral and medial walls of the nasal cavity and lateral and medial views of the turbinates. Views were systematically recorded in photographs and drawings, and the chromatic characteristics of the mucosa were noted. The separate components of the turbinate complex were then dissected and prepared.

Seven puppy heads were similarly prepared except for the final dissection of the turbinate complex.

Transverse sections of uniform thickness were cut on a polystyrene block from two adult heads that had been washed and frozen following appropriate cleaning, fixation by immersion in 10% formaldehyde for 96 h, and removal of the mandibles and associated structures.

Ten puppy heads were prepared for histological examination as follows. Eight heads (group 1) were fixed by immersion in neutral buffered formaldehyde, where they remained until use, and two (group 2) were immersed in Bouin's fixative for 24 h and then transferred to 70% alcohol. For examination of the whole nasal cavity (mainly with a view to delimiting the MOE), two group 1 heads were decalcified in Shandon TBD-1 rapid decalcifier (Thermo, Pittsburgh, PA) and embedded in one piece, appropriately oriented, in paraffin wax, after which serial transverse sections 8-10 µm thick were cut. 710 alternate sections from one of these heads, and 710 corresponding sections from the other, were transferred to slides and stained with hematoxylin-eosin (HE). The remaining eight heads (six from group 1 and the two group 2 heads) were prepared, sectioned and stained in the same way, except that before sectioning the nasal cavity was divided transversally into three blocks of more or less equal length.

All fetal heads were processed for histological examination in the same way as the puppy heads, with the main goal of scrutinizing the regions in which the GG and SO were expected to be found, if present. The posterior nasal cavity was also studied.

Owing to the difficulty of histological preparation of the whole adult nasal cavity, the following subregions of the remaining seven adult heads were excised, decalcified, and embedded in paraffin wax for sectioning and subsequent staining of the sections with HE or antibodies (see below): (i) four levels of the nasal septum; and (ii) the individual turbinates.

#### **IMMUNOHISTOCHEMISTRY**

Immunohistochemical studies were performed using antibodies against olfactory marker protein (OMP) (Wako Chemicals, 1:500 dilution) and the G-proteins Gai2 (Santa Cruz Biotechnology, 1:100) and  $G_{\alpha o}$  (Santa Cruz Biotechnology and Medical & Biological Lab Co., 1:100). Sections were dewaxed in xylene, rehydrated, and successively incubated (1) for 30 min at room temperature in PBS containing 5% normal horse serum and 2% bovine serum albumin, (2) for 24 h at 4°C in primary antibody solution, (3) for 1 h in biotinylated secondary antibody solution, and (4) for 2 h in a solution of avidin-biotin-horseradish peroxidase complex (ABC Vectastain reagent); after which standard procedures for visualization of the horseradish peroxidase complex with 3,3-diaminobenzidine were followed, and the sections were dehydrated through alcohols, cleared in xylene, and coverslipped. Sections of formalin- or Bouin-fixed canine and murine olfactory bulbs were used as control tissues.

#### IMAGE ACQUISITION AND PROCESSING

Digital images were captured using a Karl Zeiss Axiocam MRc5 digital camera. When necessary, Adobe Photoshop 6.0 (Adobe Systems, San Jose, CA) was used to adjust contrast and brightness to equilibrate light levels, and/or to crop, resize, and rotate the images for presentation; no additional digital image manipulation was performed.



FIGURE 1 | Lateral view of the nasal septum of an adult dog, showing the difference in color between sensory (yellow-brown) and respiratory (red-orange) mucosa. The rectangle frames the vomeronarasal organ. Scale bar: 2 cm (see Figure S1).

# RESULTS

The nasal cavities form a bilaterally symmetric pair flanking the nasal septum. **Figure 1** and Figure S1 show views of the lateral and medial cavity walls and of the turbinate complex within the cavity; **Figure 2** lateral and medial views of the turbinate complex without the walls of the cavity; and Figure S2 lateral and medial views of each separate component of the turbinate complex. This comprises six lateral ectoturbinates, four medial endoturbinates, and the ventral concha; the dorsal concha is a rostral projection of endoturbinate I, and the middle concha part of endoturbinate II. Parts of ectoturbinates 2 and 3 project to the frontal sinus. Transverse sections (**Figure 3** and Figure S3) clearly show the aptness of the term "complex."

In the mucosa lining the nasal cavity, color differentiates between the respiratory and olfactory epithelia, the former exhibiting shades of red or orange, the latter shades of yellow or brown (Figure 1 and Figure S1). Histological examination of selected zones of the nasal septum and of the isolated turbinates 2, 3, II, and III (Figure 4 and Figure S4) shows the transition between the two types of epithelia (Figure 4A), and also differences among different areas of sensory mucosa in regard to the thicknesses of epithelium and lamina propria and the relative thicknesses of the layers of the three epithelial cell types (sustentacular, basal, and neuronal) (Figures 4B-E and Figure S4). Immunohistochemically, the mature neurons and their apical projections of these zones were stained by anti-OMP, and the nerve bundles in the lamina propria by both anti-OMP and anti- $G_{\alpha 0}$ , but anti- $G_{\alpha i2}$  stained nothing specifically (Figure 5 and Figure S5). Staining consistent with these results was observed in the olfactory bulbs of dogs and mice used as controls (results not shown).

The VNO, enveloped in a cartilaginous lamina, lies adjacent to the nasal septum (**Figure 1**). Since the VNSE is located internally, forming the central levels of the medial wall of the VNO duct, its macroscopic display would be difficult, requiring a precise, somewhat curvilinear longitudinal section through the cartilage and its content (the duct and the other soft tissues that surround it). Histologically, the VNsE is composed of basal cells, receptor neurons and supporting cells, while the anterior part of the VNO duct features stratified squamous epithelium and the posterior



part simple columnar epithelium (**Figure 6** and Figure S6). The immunohistochemical features of the VNsE have recently been reported elsewhere (Salazar et al., 2013) and are commented on below in the Discussion.

The above descriptions all refer to adult specimens. In newborns the frontal sinus and ectoturbinate 6 are missing, and ectoturbinate 1 is very small (**Figure 7**). The MOE cannot usually be distinguished chromatically from respiratory epithelium, and must be identified histologically. Figure S7 shows the territory it occupies as determined by examination of series of transverse sections, which also show the simplicity and small area of the neonatal turbinates in comparison with those of adults (Figure S8).

In neither adults nor newborns did we find any structure with the characteristics defining the SO and GG in mice (Barrios et al., 2014). Among our 16 fetal specimens (**Figure 8** shows sections



**FIGURE 3 | Transverse frozen section of the nasal cavity.** D, dorsal nasal concha. Ectoturbinates are identified by Arabic numerals (1–5) and endoturbinates by roman numerals (I–IV). Scale bar: 2 cm (see Figure S3).



FIGURE 4 | (A) Hematoxylin-eosin-stained longitudinal section showing the transition (arrowed) from the respiratory to the sensory epithelium.
(B–E) Hematoxylin-eosin stained transverse sections showing the mucosa of ectoturbinate 2 (B), ectoturbinate 3 (C), endoturbinate III (D), and the nasal septum (E). Yellow bars indicate sensory epithelium; blue bars, lamina propria; and white arrows, bone. 1, supporting cells; 2, neurons; 3, basal cells. Scale bars: (A) 2 mm; (B–E) 100 μm (see Figure S4).



from one) there was one specimen in which the possible existence of GG cells was suggested by the presence of a nerve surrounded by vessels in the region homologous to the mouse GG, but OMP failed to label any cells in this area.

#### DISCUSSION

Among mammalian olfactory systems, the most extensively, and intensively studied has been that of the mouse-a circumstance it shares with many of the other characteristics of this animal (Paigen, 1995). It comprises four major subsystems (OSbS) stimulated via four different areas of the nasal mucosa: the MOE, the VNsE, the SO, and the GG (Barrios et al., 2014). A five-subsystem scheme emerges if the difference between vomeronasal subsystems based on semiochemical receptor types VR1 and VR2 is taken into account (Barrios et al., 2014). Although numerous studies of the canine nasal mucosa have been published (Lauruschkus, 1942; Müller, 1955; Neuhaus, 1955; Adams and Hotchkiss, 1983; Kavoi et al., 2010), we are unaware of any previous systematic search for these four or five OSbS in the dog. The findings reported above, together with our previous observation that the canine vomeronasal system binds anti- $G_{\alpha i2}$  but not anti- $G_{\alpha o}$  antibodies (Salazar et al., 2013), show that the dog has only the main and vomeronasal



subsystems, and that the latter is in all probability entirely VR1-based.

Our observations of the MOE show different areas to differ considerably, especially as regards the thickness of the neuron layer and hence the surface density of neurons, even on a single turbinate. However, we were unable to organize the observed variation in a classification of mucosal types, or to relate our observations to the two-type classification proposed previously (Bock et al., 2009) on the basis of morphological characteristics, neuron turnover, and immunoreactivity.

Whereas the murine vomeronasal system expresses the VR1associated protein  $G_{\alpha i2}$  in the apical layer of the VNsE and the anterior half of the accessory olfactory bulb (AOB), and the VR2associated protein  $G_{\alpha o}$  in the basal layer of the VNsE and the posterior half of the AOB (Barrios et al., 2014), the canine system does not express  $G_{\alpha o}$ , and expresses  $G_{\alpha i2}$  throughout the AOB as well as in the VNsE (Salazar et al., 2013). This behavior is in keeping with reported failure to find functional VR2 genes in the canine genome (Young and Trask, 2007). Given their presence in opossum (Young and Trask, 2007) and rodents (Ishii and Mombaerts, 2011), their absence from the dog (and cow and primates) (Young and Trask, 2007) must be the result of an involutionary process (Salazar and Sánchez-Quinteiro, 2009).

Regressive processes, whether evolutionary or ontogenic, seem also to have led to the absence of both SO and GG from the



canine nasal cavity. Contrary to such frequently made assertions as that "the mammalian nose contains several distinct chemosensory organs, including the... MOE, the... VNO and the SO" (Tian and Ma, 2004), the SO, though present in marsupials, and in mice, rats, and rabbits, is absent from the ferret and the cat as well as the dog, (though reportedly present in fetal cat) (Breipohl et al., 1983; Kociánová et al., 2006; Ma, 2010). Similarly, statements to the effect that Grüneberg found the GG "in all mammals examined, including humans" (Tian and Ma, 2004) ignore the fact that he actually reported definitely observing it in only three of the 13 species he studied: mouse, rat, and hamster (Grüneberg, 1973). In four species (squirrel, guinea pig, shrew, and tarsier-note the two rodents) it seemed to be absent; and in the remaining six, including Homo sapiens, its presence was uncertain. Furthermore, he only studied two species as adults (mouse and shrew) and three more as newborns (rat, mole, and guinea-pig); all his other specimens were embryos or fetuses, and he specifically suggested that in many species the ganglion, though possibly present in these early stages, may regress during prenatal development-a process he definitely asserted of the raccoon on the basis of examination of four fetal stages.

That the dog lacks two of the olfactory epithelia of the mouse (SO and GG), and has an apparently less complex form of a third (the VNsE), may seem surprising in view of its notoriously macrosmatic nature. It may be the case that the larger size of the canine nasal cavity allows physical separation to play a larger part in the dog's discrimination among odors than is possible in the mouse (Schoenfeld and Cleland, 2005); in this respect, it must be borne in mind that our dogs were mainly German

Shepherds or of German Shepherd descent, so the possibility that the SO and GG may be present in other breeds cannot be absolutely ruled out. Another possibility is that in spite of the sensitivity of the dog's sense of smell, and its ability to learn to detect unfamiliar substances by olfaction, its lifestyle prior to its domestication some 15,000 years ago (Leonard et al., 2002; Savolainen et al., 2002) required recognition of a smaller range of olfactory cues than did that of the mouse. In this regard, the dog is reported to have 811 functional olfactory receptor genes as against the 1035 of the mouse (Shi and Zhang, 2009; Niimura, 2012).

Since its domestication, the dog has of course been the object of intensive artificial selection processes that have variously pursued the enhancement of morphological, behavioral and physiological traits (regarding the mechanistic differences among these different types of modification, see Carroll, 2005, 2008; Liao et al., 2010). Creating the approximately 400 breeds of dog now estimated to exist (American Kennel Club, 2006) has altered the genome and its realization in many ways (Wayne and von Holdt, 2012), it is commonly thought that dogs collectively exhibit greater morphological variation than any other land mammal (Ostrander, 2012), and these alterations have been vigorously investigated, especially since the publication of the genome (Lindblad-Toh et al., 2005). Most of these studies have concerned the morphological traits that define breeds-body size, coat, leg length, and width, skull shape, etc. (Boyko et al., 2010; Parker et al., 2010; Schoenebeck and Ostrander, 2013), but others have addressed olfactory receptor genes, finding both marked uniformity at the gene family level, and significant variation at lower levels (Issel-Tarver and Rine, 1996; Olender et al., 2004;



Tacher et al., 2005; Robin et al., 2009; Chen et al., 2012; Derrien et al., 2012; Quignon et al., 2012). This pattern suggests that our main results—the absence of SO, GG, and a VR2-based vomeronasal subsystem—seem likely to be generalizable to breeds other than German Shepherds, but confirmation is required. Light would also be thrown on this issue by a study of wolves to determine whether the absence of these subsystems is a result of domestication.

In the study of olfaction, as in other fields (Karlsson and Lindblad-Toh, 2008), the dog is proving to be a mine of information, and is leading to a much more balanced view of the mammalian olfactory system than was promoted by exclusive attention to the mouse model.

#### **AUTHOR CONTRIBUTIONS**

Ignacio Salazar designed the research and wrote the paper. Arthur W. Barrios, and Pablo Sánchez-Quinteiro performed the work. Arthur W. Barrios, Pablo Sánchez-Quinteiro, and Ignacio Salazar analyzed and discussed the data.

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### **SUPPLEMENTARY MATERIAL**

The Supplementary Material for this article can be found online at: http://www.frontiersin.org/journal/10.3389/fnana. 2014.00106/abstract

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