

# FROM THE PERIPHERY TO THE BRAIN: WIRING THE OLFACTORY SYSTEM

## Abstract

The olfactory system represents a perfect model to study the interactions between the central and peripheral nervous systems in order to establish a neural circuit during early embryonic development. In addition, another important feature of this system is the capability to integrate new cells generated in two neurogenic zones: the olfactory epithelium in the periphery and the wall of the lateral ventricles in the CNS, both during development and adulthood. In all these processes the combination and sequence of specific molecular signals plays a critical role in the wiring of the olfactory axons, as well as the precise location of the incoming cell populations to the olfactory bulb. The purpose of this review is to summarize recent insights into the cellular and molecular events that dictate cell settling position and axonal trajectories from their origin in the olfactory placode to the formation of synapses in the olfactory bulb to ensure rapid and reliable transmission of olfactory information from the nose to the brain.

## Keywords

• Olfactory bulb • Ensheathing cells • Olfactory placode • Olfactory epithelium • RT-PCR  
 • Embryo • Mice • Sensory neurons • Molecular signaling • Glomerulus

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## Introduction

Chemical signals are likely the most ancient form of communication, and many behavioural responses are based on olfactory information. The olfactory system is composed of two different but complementary structures: the main olfactory system, which detects volatile odorants in the environment [1], and the accessory olfactory system, which mostly detects the non-volatile components of chemical signals, processing reproductive and social behaviours, through the detection of pheromones [2,3]. Nevertheless, this functional classification is not that rigid, since both systems are capable of responding to each other's stimuli [4-7]. Thus, the olfactory system shows an unusual plasticity rarely seen in other structures within the nervous system. The present review focuses on the development of the main olfactory system.

## Main olfactory system overview

The structure of the main olfactory system was first described at the turn of the nineteenth century by Golgi [8] and Cajal [9,10], as formed

by the olfactory epithelium (OE), olfactory bulb (OB) and olfactory cortex (OC). The OE consists of a pseudostratified columnar epithelium, located at the caudal part of the nasal cavity and formed by supporting cells, globose basal cells and olfactory sensory neurons (OSN). Odorant receptors (OR) are localized on the cilia of the OSN dendrites, directly exposed to the lumen of the nasal cavity. Five million OSN populate the OE, and each one expresses just one gene [11,12] of the about 1300 OR genes, from which about 20% are predicted to be pseudogenes [13-15]. Humans have fewer OR coding genes (640), of which 46% are pseudogenes [16]. Those OR are not spread over the entire OE but are restricted to four zones based on the expression pattern of a few OR [17,18]. OSN expressing the same OR target the same glomeruli [19-21]. Deletion or replacement of an OSN receptor results in an OSN axon misguidance into the glomeruli [21,22]. OSN expressing their receptor coalescence at embryonic day 12 (E12) when the first homotopic axonal segregation takes place through the cribose plate [23]. Then, OSN axons in the olfactory bulb join into glomeruli, establishing stereotypic connections in the glomerular layer with periglomerular cells and

dendrites of mitral/tufted (M/T) cells. M/T cells are the primary OB projection neurons and project to different brain areas forming the lateral olfactory tract (LOT) [24-29], relaying odorant-elicited signals to the olfactory cortex. As occurs in the projections from OE to OB, the LOT shows a spatial rearrangement of the axons depending on the projection neurons' position within the OB [30]. This topographical organization is dependent on the neurogenesis of the projection neurons [30,31] and within the LOT, also serves as a spatial rearrangement of the cortical connections.

Unlike other sensory systems, some of the neuronal populations in both OE and OB are replaced throughout the whole life span of the animal [32]. In the OE, newly generated OSN [33] are able to project to the correct glomeruli [34]. In the OB, newly generated interneurons migrate either from the SVZ, in postnatal ages, to repopulate the glomerular and granular layers of the OB [35-39] or, from the lateral ganglionic eminence, at embryonic stages [40]. Interestingly, the percentage of adult-born neurons in the granular layer remains constant at approximately 10%, while the proportion of adult-born neurons in the glomerular layer

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increases over time [41]. In fact, almost all classes of periglomerular interneurons undergo adult neurogenesis, including glutamatergic short axon cells [42]. Thus, the olfactory system shows a solid and well-structured relationship between the central and peripheral nervous systems and, at the same time, a high level of plasticity that allows the integration of new neurons and sensory axons, thus preserving its ability to process the olfactory information.

In this work, we will focus on the generation and development of the afferent sensory fibres into the OB, since it constitutes not only the first relay station in the transduction of the olfactory information into the CNS, but also the source of molecules and cell populations implicated in the growth and guidance of the incoming sensory axons into the OB, in both stages, during development and in adulthood.

### Early differentiation of the olfactory sensory neurons — how does the connectivity work?

The OE is the first structure that develops during embryogenesis; it arises from the

olfactory placode (OP), a thickening of the surface ectoderm, visible at E9 in the rostral prosencephalon of mice. One day later, the placode invaginates to form the olfactory pit from which several cell populations migrate to the telencephalic vesicle, both in rodents [43-47] and chick [48]. These cells form a heterogeneous population expressing a different variety of markers as they migrate towards the CNS (see below). During these early stages, and despite the absence of a fully anatomically differentiated OB, projection neurons have already undergone generation and differentiation processes [49-52]. The close developmental timing between OE and OB raises the question whether the early OE axons play an inductive role in the OB protrusion and projection neurons' generation [53,54]. Nevertheless, in the *Pax-6<sup>Sey-Neu</sup>* mutant mice, where the OE is absent, exists a structure called olfactory bulb-like structure (OBLS) [51,55]. This structure has projection neurons which give rise to a lateral tract where their axons project to cortical caudo-ventral areas. As we will describe in depth, the OSN axons play

an important role in the OB development; however, they do not induce the generation of the projection neurons (first OB neurons generated in the development), since they reach to the OB primordium at E12 in mouse [56] when the first projections neurons are already generated (Figure 1).

### Role of the migratory mass in olfactory bulb development — ensheathing cells, the pillars of axonal growth

Another striking event that follows the development and differentiation of the OE is the generation and migration of several cell populations towards the rostral telencephalic vesicle (Figure 2A, B). Here we focus on the migratory mass (MM) [44,57], formed by a mix of glia and neuronal migrating cells, closely related to the OSN axons and contributing to the OB development. The nature of these cells remains controversial, since they are mainly identified by the expression of neuronal markers such as GAP-43 [58],  $\beta$ -III-tubulin [47] and the vesicular glutamate transporter type 2 [59]. An elegant and acute analyses of the MM

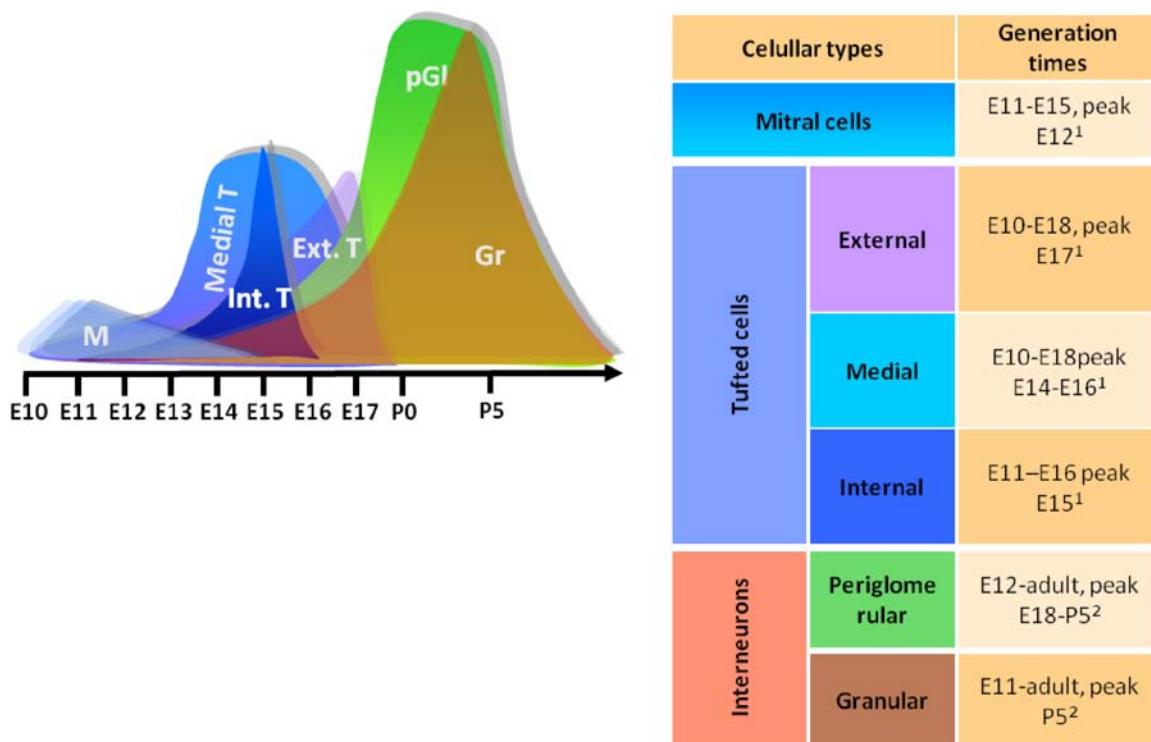
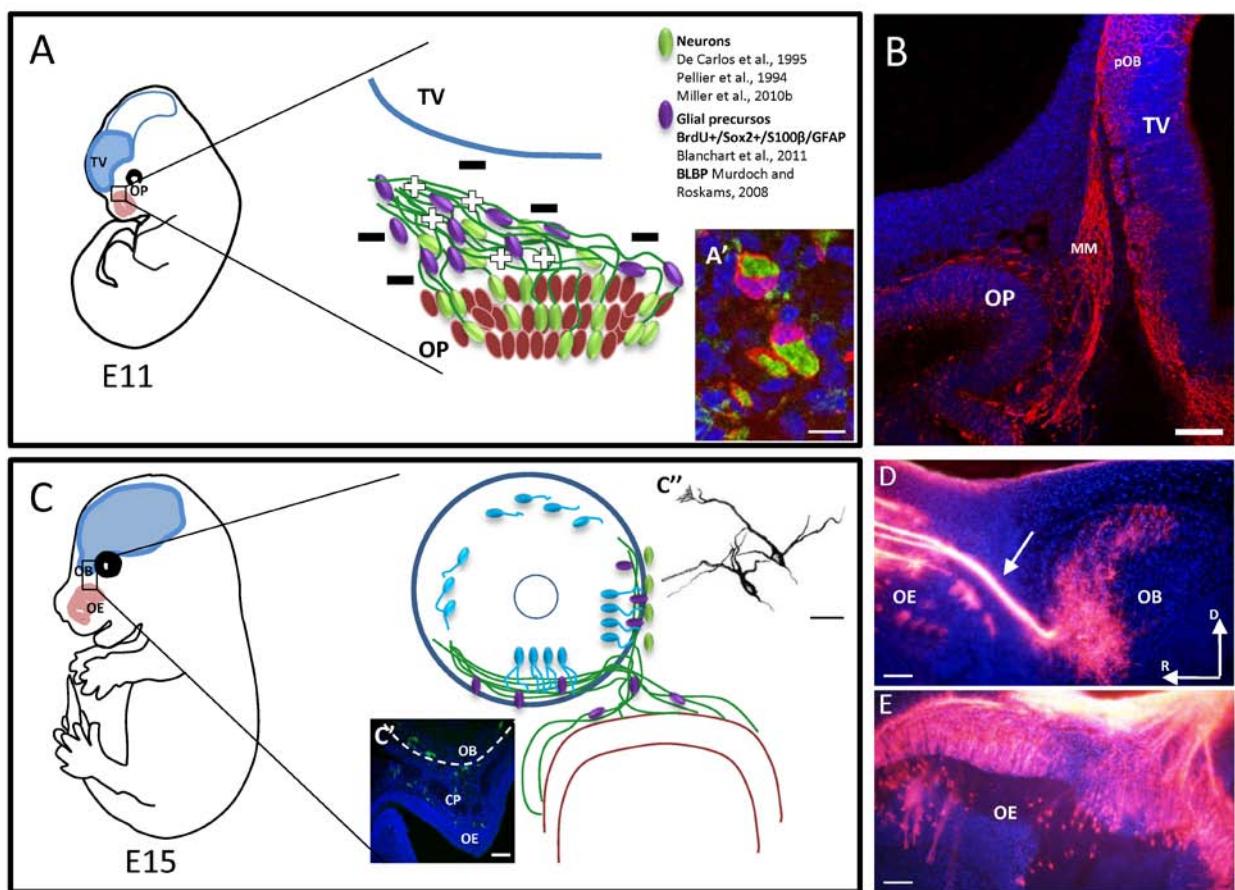


Figure 1. Generation time curves of the different olfactory bulb cell populations during development. First generated cells are the projection neurons and then the interneurons. Interneuron generation continues through the whole life. <sup>1</sup>[49,51] <sup>2</sup>[49,153-155]

described the existence of a heterogeneous neuronal labelling of DCX, OMP, DNER, GnRH, and Lhx2 [60]. However, we recently recognized that heterogeneous neuronal labelling could also correspond to growing axons [61] and a wide range of glial markers, such as S100 $\beta$ , GFAP or BLBP, are expressed in the MM [62,63]. Moreover, GFP-retrovirus-infected MM cells, colabelled with the neurogenic marker BrdU, revealed the existence of cell proliferation within the MM. In addition, MM-cells coexpress the neural stem cell markers, Sox2 and nestin, which enhances the idea of the MM as a niche of glial and progenitor cells that give rise to different glial cells along the OSN pathway,

both in the mesenchyme and in the glomerular layer of the OB. Furthermore, these GFP-infected MM cells express the glial markers GFAP, p75, S100 $\beta$  and BLBP, but none express neuronal marker [61]. In addition, these GFP-infected MM cells were always intimately related to the glomeruli and axons bundles across the mesenchyme, both common features of olfactory ensheathing cells (OEC) [43,64,65]. This fact raises the possibility that MM is a niche of newly generated ensheathing cells at early embryonic stages. This hypothesis supports the *in vitro* idea that OECs facilitate the growing of the OSN axons through the mesenchyme and their implication in the

axonal segregation into the glomeruli [66-75]. Furthermore, activation of Wnt/ $\beta$ -catenin signalling, in Wnt reporter TOPgal transgenic mice, labels a subpopulation of OECs in the most external part of the OB, in a dependent way [76], through its canonical pathway [77], playing an important role in the sorting of OSN axons and glomerulus formation. To sum up, the MM is not just a bulk of cells migrating with the axons, but rather a necessary axonal scaffold to grow towards the OB. Whether the OB sends or produces some type of molecular signal for these migrating cells still remains to be elucidated. So far, possible chemoattractive role of the OB on the migrating olfactory cells

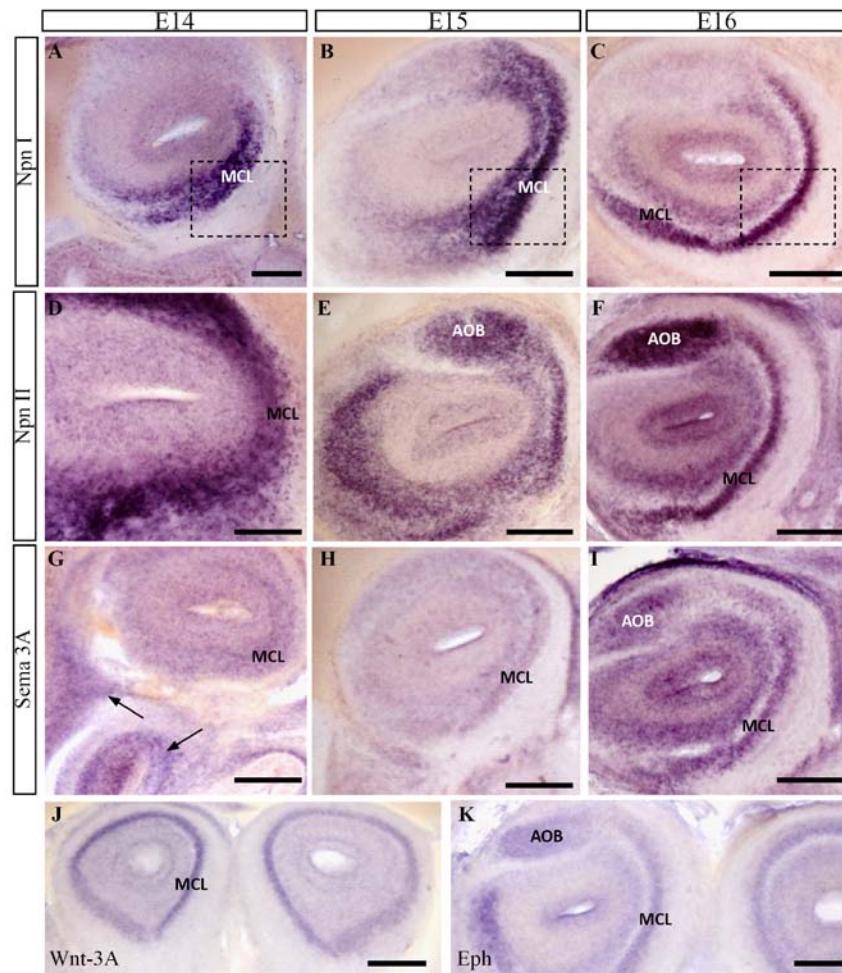


**Figure 2.** Schematic representation of the earlier stages in the olfactory system development. A. At early stages olfactory placode gives rise to the olfactory pit (OP). From hereafter several cell populations from the OP migrate together with olfactory axons towards the telencephalic vesicle (TV), forming the Migratory Mass (MM). Neurons (green) and glial precursors (purple), follow a pathway defined by the expression of molecular cues from mesenchyme, the MM cells and from target regions in the TV. A' Many of these glial precursors processed (S100 $\beta$ + cells, red) envelop the axon bundles (TuJ1, green). B. At E11, TuJ1 expression (red) labels the OP, MM and the prospective OB in the TV. C. At E15, MM cells are still present in both the mesenchyme and the Olfactory Bulb (purple and inset C', GFP cells), while the axons start to innervate the OB from the ventromedial part to the laterodorsal regions. This event will trigger the reorientation of projections neurons achieving its final position (C''). Simultaneously, the number of synaptic contacts will increase following the same innervation pattern. Still, at this age LHRH, GnRH cells are migrating towards diencephalic areas (green). D, E. Dil crystal into the OE (D) shows the OSN axons (D, arrow) reaching the medial and ventral parts OB, while a Dil injection into the OB (E) labels OSN in the OE. E14, sagittal sections. OB, olfactory bulb; CP, cribriform plate; OE, olfactory epithelium; OP, olfactory pit; MM, migratory mass; TV, telencephalic vesicle, pOB, prospective OB. Scale bars, A' = 10  $\mu$ m, B and C' = 100  $\mu$ m, C'' = 50  $\mu$ m, D and E = 100  $\mu$ m.

was described, where OEC migrate along a gradient concentration of soluble factors released by the OB [78]. Nevertheless, this effect seems unlikely to occur at the beginning of the migration since the mesenchyme situated between OB and OE, contains a wide range of clue molecules that affect both the OSN axonal growth and the OE migrating cells. Thus, the possibility of the OB exerting an attractive effect could most likely occur at the final phase of cell migration into the OB.

### The sensitive phase is a key component of connectivity — spatiotemporal development of axonal connections and neurons

The MM fuses with the rostral part of the mouse telencephalon at E12/E13. This event will trigger the initial innervation of OSN axons into the OB, although this process does not occur simultaneously. First, in the sensitive phase, olfactory axons innervate the ventromedial part of the OB and, thereafter, extend to lateral and dorsal regions (Figure 2C-E; [52]). During this phase, between E15/E16, the protoglomeruli are formed; while at E14, the OSN axons stay in the ONL without entering into the inner OB layers. Furthermore, this axonal innervation occurs in parallel with the reorientation of the OB projection neurons [79]. But most importantly, these effects seem to be only exerted at E14-E15, corresponding to the reorientation cell process [51]. This process shares some common features with the development of the olfactory system in *Drosophila*, where the projection neurons are differentiated before the arrival of the olfactory axons [80]. This could be due to molecular signals, derived from either the OE [81] or from the mesenchyme, such as retinoic acid, FGF8, BMPs or Sonic [82]. The lack of projection neuron orientation in the OBLS could be due to the absence of retinoic acid in Pax-6<sup>Sey-Neu</sup> [83]. Nevertheless, the question whether the OE axons either exert a trophic effect or secrete certain molecular clues, as inducers of neuronal reorientation, remains unclear. Both the first synaptic contact and its spatiotemporal pattern occur simultaneously with the axonal innervation pattern and reorientation of the OB projection neurons



**Figure 3.** Unchanged expression pattern of several genes implicated in axonal guidance. A-C. ISH for *Npn1* in the OB. *Npn1* expression is mainly restricted to the MCL during the sensitive phase, decreasing the expression intensity level throughout the ages. Boxes represent the OB region selected for the quantitative RT-PCR analyses. D-E. Expression pattern of *Npn2* mRNA. *Npn2* expression is similar to *Npn1*, with the highest expression in the MCL. G-H. *Sema3A*, the Npn receptor, is expressed in both the MCL and mesenchyme (arrows, G). J. At E16, *Wnt-3A* is also expressed in the MCL without any changes during the sensitive phase. K. *Eph* receptor expression is also restricted to the MCL. ISH, in situ hybridization; MCL, mitral cell layer; OB, olfactory bulb. Scale bars, A-C and E-K = 200  $\mu$ m; D = 100  $\mu$ m.

[63]. These data raise the question whether these newly established synaptic contacts contribute to the refinement of the dendritic tree of the projection neurons. In fact, dendritic stability is higher within the olfactory glomeruli upon exposure to odor-enriched environment [84]; and the maturation process of the dendritic tree is heavily influenced by excitatory neurotransmission. Moreover, an interaction of  $\text{Ca}^{2+}$  with some intracellular enzymes has been proposed, as well as the synthesis of trophic factors, or simply

the synaptic activity itself contributes [84-90]. Thus, the sensitive phase represents a unique event during OB development, which follows a well-established spatiotemporal pattern by the establishment of synaptic connections and the initial steps towards the differentiation of M/T cells. As those processes turn out to be critical in the information "flow" from the PNS to the CNS, this stage requires the participation of cellular, axonal and chemical clues to set up the right axonal and cellular wiring.

**Molecular barriers and spatiotemporal gene expression gradients — a better understanding of the sensitive phase**  
 So far, we focused on the cellular compartments during the sensitive phase, the relationship between mesenchyme, OE/OB and the presence of different molecular clues during the development of the olfactory pathway is conclusive. To investigate the differential profiles of these structures, we performed a quantitative RT-PCR analyses of a list of genes involved in migration, axon guidance and extracellular matrix molecules (Table 1). We used three representative embryonic ages, and the tissue was limited to the ventro-medial part of the OB (Figure 3A-C). That is the place where the first OE axons contact with the telencephalic vesicles and it is also the location for the first synapses [52,63,91].

Comparing gene expression between the three developmental stages of the sensitive phase, 21 out of 99 selected genes showed significant differences (Table 2). Only one of the analyzed genes showed significant differences between E14 and E15; seven genes between E14-E15; and 13 genes between E14-E16. It is noteworthy that the unique gene with a differential expression between E15 and E16 is Neuropilin-2 (Nrp-2). This protein, which constitutes the main receptor of the secretable semaphorin 3B, 3F and 3C proteins [92-94], is one of the main candidates in the guidance of OE axons towards the OB, both in chicken [95] and rodents [94,96,97]. In fact, olfactory axons of the Nrp2-/- mice [98] or of those with expression of dominant negative Nrp1, the Sema3A receptor [99], failed to converge into the correct glomeruli, reaching to the inner OB layers. This suggests that the expression of different semaphorins in the mitral cell layer exerted a barrier effect on the incoming axons, preventing them from growing further into the OB as occurs in the Nrp2 mutant mice. While the expression pattern of the *Nrp-2* mRNA is similar during the three selected ages (Figure 3D-F), RT-PCR analyses revealed different transcript levels. This change could be related to the presence of a ligand in the mitral cell layer, which is important in the targeting of the olfactory axons into specific glomeruli. Furthermore, quantitative RT-PCR

analyses also shows significant differences between E14/E15 in the mRNA levels of Sema3C and Sema3E, and between E14/E16 in these two *nrp* ligands. Sema3C mRNA is localized in the OB [56,98,100,101], but not in the AOB [94,100,102,103]. Moreover, although the expression mRNA pattern did not show changes during the sensitive-phase (Figure 3), the mRNA levels decreased from E14 to E15 (Table 2). This suggests the Neuropilin family and their ligands, the semaphorins, as one of the candidates in the OB innervation, and could partially explain the sharp “behaviour” of OSN axons during this phase. Together with these two proteins, significant differences between the selected ages were also detected in the mRNA expression of the following four molecules: i) Aggrecan (*Acan*), a large aggregating proteoglycan or chondroitin sulphate proteoglycan 1, which is the major proteoglycan in the articular cartilage providing a hydrated gel structure, and it also mediates chondrocyte-chondrocyte and chondrocyte-matrix interactions [104]. As a member of a chondroitin sulfate family, *Acan* could be involved in the migration of different cell subtypes and axons within the mesenchyme during the sensitive phase; ii) Brevican (*Bcan*) belongs to the chondroitin sulphate family proteins [105], and it was recently described as one of the ECM proteins produced and released by astrocytes and involved in synapse generation in the hippocampus [106]. Moreover, *Bcan* is also expressed in the astrocytes ensheathing cerebellar glomeruli [107]. Thus, due to the significant increase in synapses during the OB sensitive phase [63], the up-regulation of the *Bcan* mRNA suggests an active role in the OB synaptogenic process; iii) Collagen A3 (IV) chain, first described in humans [108], both in kidney glomeruli formation [109] and kidney disease Alport syndrome [110]; iv) NRCAM or neuron-glia-CAM-related cell adhesion molecule, implicated in the guidance of thalamocortical axons by the interactions with the *Nrp-2* which is critical for Sema3F-induced guidance [111]. All these proteins are implicated and related to ECM and axonal growth, thus their expression variation or abundance in the OB could trigger changes within the environment, allowing

the axon entrance into the external plexiform layer and the formation of the first glomerulus [52,112-115], giving rise to the glomerular layer at E16.

On the other hand, molecules such as neuroligin 1 (NLgn1), implicated in regulation of synaptogenesis in hippocampal cultures [116] and in presynaptic differentiation [117], could be potential candidates in the OB synaptogenesis, as between E14/E15 the highest increase in the number of synapses. Another potential molecule is Presenilin 1 (Psen1), required for gamma secretase function *in vivo* [118] and neural progenitor cell differentiation in the brain [119]. Finally, between E14-E16, we observed a significant variation of mRNA in the following genes: C-C chemokine receptor type 5 (Ccr5), a member of the beta chemokine receptors family of integral membrane proteins implicated in numerous processes during brain development [120] and inflammation [121]; chemokine (C-X-C motif) ligand 1 (CXCL1), a small cytokine belonging to the CXC chemokine family, described in ovarian cancer [122], inflammation [123], precursor cell proliferation and migration [124-126], and neuronal migration [127]; Deleted in Colorectal Carcinoma (DCC), a transmembrane receptor, first discovered in colorectal cancer [128], and receptor of the netrin molecule and therefore a clue protein for axon guidance [129]. Genetic studies in mice and *C. elegans* showed that DCC is a receptor that mediates netrin1 attraction of commissural axons [130,131]. DCC has both attractive and repulsive effects [132], being also implicated in regulation of cell death [133] and in surveillance of developing retinal cell types [134]. Moreover, its role in cell apoptosis could be related to the presence of apoptotic events when the MM reaches the OB [135]. The neurotrophic tyrosine kinase, receptor type 3 (Ntrk3) is another example of a molecule implicated in the proliferation and differentiation of neurons during development, and in growth and survival in adult [136], playing an important role in survival and cell death [137]. The differential gene expression of both DCC and Ntrk3 during the sensitive phase might partially explain the balance between cell proliferation [61] and apoptosis [135] during the cell migration of the migratory

**Table 1.** List of all genes analyzed during the sensitive phase. Differential gene expression was analyzed by quantitative RT-PCR of RNA extracted from the ventro-medial part of the olfactory bulb, where the first OSN axons start to innervate the OB [52]. All genes were selected based on their function in development, axon migration, cytoskeleton modifications, apoptotic and/or differentiation effects and ECM protein interactions, including axon guidance, repulsion and attraction.

Gene	Name	Known function	Reference
<i>Acan</i>	Aggrecan, large aggregating proteoglycan, or chondroitin sulfate proteoglycan 1	Major component of extracellular matrix of cartilagenous tissues.	[156]
<i>Adam10</i>	Disintegrin and metalloproteinase domain-containing protein 10, also known as ADAM10 or CDw156 or CD156c	Responsible for the proteolytic release of several other cell-surface proteins, including heparin-binding epidermal growth-like factor, ephrin-A2 and for constitutive and regulated alpha-secretase cleavage of amyloid precursor protein (APP).	
<i>Akt</i>	Protein Kinase B (PKB)	Controls the tempo of the newborn neurons integration during adult neurogenesis, including correct neuron positioning, dendritic development and synapse formation	
<i>Arf6</i>	ADP-ribosylation factor 6	Regulates endocytic recycling and cytoskeleton remodeling	
<i>Arg1</i>	Arginase type I	Possible role in nitric oxide and polyamine metabolism.	
<i>Arg2</i>	Arginase Type II		
<i>Astn1</i>	Astrotactin 1	Neuronal adhesion molecule required for glial-guided migration of young postmitotic neuroblasts in cortical regions of developing brain, including cerebrum, hippocampus, cerebellum and olfactory bulb	[157]
<i>Bcan</i>	Brevican	Major chondroitin sulfate in the adult brain. Mediates interactions with ECM.	[105,106]
<i>Bdnf</i>	Brain-derived neurotrophic factor		
<i>Bgn</i>	Biglycan	ECM	
<i>Bmp7</i>	Bone morphogenetic protein also known as osteogenic protein-1 or OP-1	Transformation of mesenchimal cells	
<i>Boc</i>		Cell surface receptors of the immunoglobulin (Ig)/fibronectin type III	
<i>Ccl5</i>	Chemokine (C-C motif) ligand 5	Chemotactic cytokine or chemokine	
<i>Ccr5</i>	C-C chemokine receptor type 5	Integral membrane proteins	[158]
<i>Ccrl1</i>	C-C chemokine receptor type 11	G protein-coupled receptor family, and is a receptor for C-C type chemokines	
<i>Cdc42</i>	Cell division control protein 42 homolog	GTPase of the Rho-subfamily, which regulates signaling pathways that control diverse cellular functions including cell morphology, migration, endocytosis and cell cycle progression	
<i>Cdh11</i>	Cadherin-11	Integral membrane protein that mediate calcium-dependent cell-cell adhesion	
<i>Cdh13</i>	T-cadherin	Cell-cell contacts, dynamic regulation of morphogenetic processes in embryos and tissue integrity in adult organisms	
<i>Cdh2</i>	Cadherin-2	Nonclassic-type cadherins-contact-mediated communication, with cadherin domains acting as homophilic binding regions and the EGF-like domains involved in cell adhesion and receptor-ligand interactions	
<i>Celsr1</i>	Cadherin EGF LAG seven-pass G-type receptor 1		[159]
<i>Chek1</i>	Serine/threonine-protein kinase Chk1	Phosphatase in cell cycle control, particularly for entry into mitosis	

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Gene	Name	Known function	Reference
<i>Col4a3</i>	Collagen, type IV, alpha 3	Major structural component of basement membranes	
<i>Csnk2b</i>	Casein kinase II subunit beta	Ubiquitous protein kinase which regulates metabolic pathways, signal transduction, transcription, translation, and replication	
<i>Cxcl1</i>	Chemokine (C-X-C motif) ligand 1	Mitogenic properties and it is also implicated in melanoma pathogenesis	[160]
<i>Cxcl12</i>	Chemokine (C-X-C motif) ligand 12	Participates in nervous system development	[161,162]
<i>Cxcr3</i>	Chemokine (C-X-C motif) receptor 3	Induces various cellular responses, most notably integrin activation, cytoskeletal changes and chemotactic migration	
<i>Cxcr4</i>	Chemokine (C-X-C motif) receptor 4	Normal development of the central nervous system, influencing the guidance of both growing axons and migrating neurons	[163,164]
<i>Dab1</i>	Disable homolog-1	Adaptor protein in charge of Reelin signal transduction	[165]
<i>Dcc</i>	Deleted in Colorectal Carcinoma	Bounds to netrin-1, signals are conveyed that can lead to proliferation and cell migration. In the absence of netrin-1, DCC signaling has been shown to induce apoptosis.	[166]
<i>Dcn</i>	Decorin	Small cellular or pericellular matrix proteoglycan closely related in structure to biglycan protein. It is a component of connective tissue, binds to type I collagen fibrils, and plays a role in matrix assembly.	
<i>Fert2</i>	fer (fms/fps related) protein kinase	Required for p38 activation to promote chemotaxis	[167,168]
<i>Fgf2</i>	Basic fibroblast growth factor	Located in basement membranes and in the subendothelial extracellular matrix of blood vessels. Extensively used in stem cell cultures	
<i>Fgfr1</i>	Fibroblast growth factor receptor 1, also known as basic fibroblast growth factor receptor 1, fms-related tyrosine kinase-2 / Pfeiffer syndrome, and CD331	Receptor tyrosine kinase whose ligands are specific members of the fibroblast growth factor family.	
<i>Fgfr3</i>	Fibroblast growth factor receptor 3 or CD333	Participates in the assembly of the extracellular matrix: interacts with type I and type II collagen fibrils and inhibits in vitro fibrillogenesis	
<i>Fmod</i>	Fibromodulin		
<i>Gpc2</i>	Glypican 2, also known cerebroglycan	Integral membrane heparan sulfate proteoglycan found in the developing nervous system. Cerebroglycan participates in cell adhesion and is thought to regulate the growth and guidance of axons. Cerebroglycan has especially high affinity for laminin-1.	[142,169]
<i>Hhip</i>	Hedgehog-interacting protein	Modulates the responses to any Hedgehog signal.	[170]
<i>Igfs8</i>	Immunoglobulin superfamily member 8	Role on integrin-dependent morphology and motility functions. Participate in the regulation of neurite outgrowth and maintenance of the neural network in the adult brain	
<i>Il8rb</i>	Interleukin 8 receptor, beta is a chemokine receptor, is also known as CXCR2	Reported on various cell types, including neurons, astrocytes, microglia, neural progenitor cells, and microvascular endothelial cells. It is related to development of the CNS, neurotransmission, blood-barrier permeability and in several brain pathologies.	[171]
<i>Irf1</i>	Interferon regulatory factor 1	Acts as a tumor suppressor and regulating apoptosis	
<i>Irf8</i>	Interferon regulatory factor 8	Plays a negative regulatory role in cells of the immune system	
<i>Itga3</i>	Integrin, alpha 3 (antigen CD49C, alpha 3 subunit of VLA-3 receptor)	Receptor for fibronectin, laminin, collagen, epiligrin, thrombospondin and CSPG4. Alpha-3/β-1 may mediate with LGALS3 the stimulation by CSPG4 of endothelial cells migration	

continued **Table 1.** List of all genes analyzed during the sensitive phase. Differential gene expression was analyzed by quantitative RT-PCR of RNA extracted from the ventro-medial part of the olfactory bulb, where the first OSN axons start to innervate the OB [52]. All genes were selected based on their function in development, axon migration, cytoskeleton modifications, apoptotic and/or differentiation effects and ECM protein interactions, including axon guidance, repulsion and attraction.

Gene	Name	Known function	Reference
<i>Itpr1</i>	Inositol 1,4,5-triphosphate receptor, type 1	Intracellular channel that mediates calcium release from the endoplasmic reticulum following stimulation by inositol 1,4,5-trisphosphate	
<i>Jag2</i>	Protein jagged-2	Putative Notch ligand involved in the mediation of Notch signaling	
<i>Lmx1b</i>	LIM homeobox transcription factor 1-beta	Dorso-ventral patterning of vertebrate limbs, and ventral brain development	[172]
<i>Lrp2</i>	Low density lipoprotein-related protein 2 or megalin	Multiligand binding receptor found in the plasma membrane.	
<i>Lum</i>	Lumican	Lumican is the major keratan sulfate proteoglycan of the cornea but is also distributed in interstitial collagenous matrices throughout the body	
<i>Myh10</i>	Myosin, heavy chain 10, non-muscle	Plays a role in cytokinesis, cell shape, and specialized functions such as secretion and capping	
<i>Ncan</i>	Neurocan	Modulates neuronal adhesion and neurite growth during development by binding to neural cell adhesion molecules (NG-CAM and N-CAM)	[173]
<i>Ngfr</i>	Nerve growth factor receptor	Mediates cell survival as well as cell death of neural cells	
<i>Nisch</i>	Nischarin	Binds numerous imidazoline ligands that induces initiation of cell-signaling cascades triggering to cell survival, growth and migration	
<i>Nlgn1</i>	Neuroligin-1	Neuronal cell surface protein thought to be involved in cell-cell-interactions by forming intercellular junctions through binding to beta-neurexins.	
<i>Nrcam</i>	Neuronal cell adhesion molecule	Cell adhesion, ankyrin-binding protein involved in neuron-neuron adhesion, axon projection and Schwann cell and axon interactions	[111,174]
<i>Npn1</i>	Neuropilin 1	Implicates in cell migration and axonal targeting.	[175,176]
<i>Npn2</i>	Neuropilin 2		
<i>Ntf3</i>	Neurotrophin 3	Controls survival and differentiation of mammalian neurons	
<i>Ntn1</i>	Netrin 1	Controls guidance of CNS commissural axons and peripheral motor axons. Its association with either DCC or some UNC5 receptors will lead to axon attraction or repulsion	[177,178]
<i>Ntn2l</i>	Netrin-2-like protein precursor	(See Netrin)	
<i>Ntrk2</i>	Neurotrophic tyrosine kinase, receptor, type 2	BDNF modulates synaptic transmission and plasticity primarily through the TrkB receptor, but the molecules involved in BDNF-mediated synaptic modulation are largely unknown.	[179,180]
<i>Ntrk3</i>	neurotrophic tyrosine kinase, receptor, type 3	Receptor for neurotrophin-3. Mediates the multiple effects of this neurotrophic factor, which includes neuronal differentiation and survival.	[137]
<i>Pafah1b1</i>	Platelet-activating factor acetylhydrolase IB subunit alpha	Required for proper activation of Rho GTPases and actin polymerization at the leading edge of locomoting cerebellar neurons and postmigratory hippocampal neurons in response to calcium influx triggered via NMDA receptors	
<i>Pcdhb16</i>	Protocadherin beta-16	Potential calcium-dependent cell-adhesion protein	
<i>Pcdhb21</i>	Protocadherin beta 21	Cell-adhesion protein	
<i>Pex5</i>	Peroxisomal biogenesis factor 5	Essential role in peroxisomal protein import	

continued Table 1. List of all genes analyzed during the sensitive phase. Differential gene expression was analyzed by quantitative RT-PCR of RNA extracted from the ventro-medial part of the olfactory bulb, where the first OSN axons start to innervate the OB [52]. All genes were selected based on their function in development, axon migration, cytoskeleton modifications, apoptotic and/or differentiation effects and ECM protein interactions, including axon guidance, repulsion and attraction.

Gene	Name	Known function	Reference
<i>Prx1</i>	Paired related homeobox protein-like 1	Required for the formation of spatio-temporally appropriate projections from nociceptive sensory neurons to their central targets in the dorsal horn of the spinal cord.	
<i>Psen1</i>	Presenilin-1	Involved in the cleavage of Notch receptors	
<i>Pten</i>	Phosphatase and tensin homolog	The lipid phosphatase activity is critical for its tumor suppressor function. Antagonizes the PI3K-AKT/PKB signaling pathway by dephosphorylating phosphoinositides Dephosphorylates tyrosine-phosphorylated focal adhesion kinase and inhibits cell migration and integrin-mediated cell spreading and focal adhesion formation.	
<i>Ptk2</i>	Protein tyrosine kinase 2	Implicated in signaling pathways involved in cell motility, proliferation and apoptosis.	
<i>Ptpz1</i>	Protein tyrosine phosphatase, receptor-type, Z polypeptide 1	Expressed in developmental CNS and implicated in glioblastoma cell migration	[181,182]
<i>Pxmp3</i>	Peroxisome biogenesis factor 2	The protein is thought to be involved in peroxisomal matrix protein import	
<i>Pxn</i>	Paxillin	Cytoskeletal protein involved in actin-membrane attachment at sites of cell adhesion to the extracellular matrix (focal adhesion)	
<i>Rhoa</i>	Ras homolog gene family, member A	Regulates a signal transduction pathway linking plasma membrane receptors to the assembly of focal adhesions and actin stress fibers	
<i>Robo1</i>	roundabout, axon guidance receptor, homolog 1 (Drosophila)		
<i>Robo2</i>	roundabout, axon guidance receptor, homolog 2 (Drosophila)	Receptors for SLIT1 and SLIT2 which are thought to act as molecular guidance cue in cellular migration, including axonal navigation at the ventral midline of the neural tube and projection of axons to different regions during neuronal development	[145,148,183]
<i>Robo3</i>	roundabout, axon guidance receptor, homolog 3 (Drosophila)		
<i>Sema3A</i>	Semaphorin 3A		
<i>Sema3B</i>	Semaphorin 3B		
<i>Sema3C</i>	Semaphorin 3C		
<i>Sema3D</i>	Semaphorin 3D		
<i>Sema3E</i>	Semaphorin 3E	Secreted and membrane proteins that act as axonal growth cone guidance molecules. They primarily act as short-range inhibitory signals and signal through multimeric receptor complexes. They are usually cues to deflect axons from inappropriate regions, especially important in neural system development	[98,184-186]
<i>Sema3F</i>	Semaphorin 3F		
<i>Sema4F</i>	Semaphorin 4F		
<i>Sema6b</i>	Semaphorin 6b		
<i>Sema6c</i>	Semaphorin 6c		
<i>Shh</i>	Sonic hedgehog homolog	Intercellular signal essential for a variety of patterning events during development: signal produced by the notochord that induces ventral cell fate in the neural tube and somites, and the polarizing signal for patterning of the anterior-posterior axis of the developing limb bud	

continued **Table 1.** List of all genes analyzed during the sensitive phase. Differential gene expression was analyzed by quantitative RT-PCR of RNA extracted from the ventro-medial part of the olfactory bulb, where the first OSN axons start to innervate the OB [52]. All genes were selected based on their function in development, axon migration, cytoskeleton modifications, apoptotic and/or differentiation effects and ECM protein interactions, including axon guidance, repulsion and attraction.

Gene	Name	Known function	Reference
<i>Slit1</i>			
<i>Slit2</i>		Acts as molecular guidance cue in cellular migration, and function appears to be mediated by interaction with roundabout homolog receptors. During neural development involved in axonal navigation at the ventral midline of the neural tube and projection of axons to different regions.	[145,187]
<i>Slit3</i>			
<i>Trl1;Trl6</i>	Transfer RNA leucine 1 (anticodon AAG)		
<i>Tlx3</i>	T-cell leukemia homeobox 3	Transcription factor implicated in differentiation processes	[150]
<i>Tnn</i>	Tenascin N	Sited around migrating cells, at sites of epithelium-mesenchyme interactions, at sites of smooth muscle and connective tissue morphogenesis, and throughout the central and peripheral nervous system, and is often induced under pathological conditions Tenascin-X and tenascin-Y are expressed around developing muscle and blood vessels and tenascin-R expression is limited to the nervous system	[188]
<i>Twist1</i>	Twist homolog 1	Acts as a transcriptional regulator	
<i>Unc5c</i>	Unc-5 homolog C	Receptor for netrin required for axon guidance. Mediates axon repulsion of neuronal growth cones in the developing nervous system upon ligand binding. Axon repulsion in growth cones may be caused by its association with DCC that may trigger signaling for repulsion.	[189]
<i>Vax1</i>	Ventral anterior homeobox 1	Genes of this family are involved in the regulation of body development and morphogenesis	
<i>Vldlr</i>	Very low density lipoprotein receptor	Receptor for the extracellular protein reelin, implicated in migration processes.	[190]

mass towards the OB. Outstanding variations in gene expression during the sensitive phase included the Glycan 2 (GPC2), also known as cerebroglycan, a glycoprophosphatidylinositol-linked integral membrane heparansulfate proteoglycan located on the surface of axons and growth cones [138-141]. GPC2 is also present in the developing nervous system contributing to cell adhesion, growth and guidance of axons [142]. The reported differences may explain the slow, progressive penetration of axons into the developing OB comparing E16 and E14, likely modulated by repulsive effects. Another candidate, the Nischarin (Nsch), is a citoplasmatic protein that binds the citoplasmatic domain of the integrin  $\alpha 5\beta 1$ , inducing  $\alpha 5\beta 1$ -dependent migration in several cell types [143,144]. In our analyses, we found a 34 fold down-regulation of its mRNA during the sensitive phase, which could partially explain the cell migration of the MM throughout all the OB regions, fusing with

the telencephalic vesicle at E13 [61]. Another secreted protein implicated in axon guidance process is Slit-1, a well-known secreted protein in the olfactory system [145-147], as well as spinal motor axons [148] and midbrain [149], which is responsible for growth cones collapse. Levels of *Slit-1* mRNA decreases from E14 to E16, partially supporting its downregulation, and allowing for the incoming OE axons to penetrate the OB and give rise to the glomeruli by E16. Unlike the previous proteins, the T-cell leukemia homeobox 3 (Tlx3) is a transcription factor implicated in multiple differentiation processes [150] and it also mediates by Wnt activation [151]. As mentioned above, activation of the canonical pathway in a subset of OECs [152] seems to be critical for the guidance of OSN axons during the MM migration towards the rostral telencephalic vesicle [77], although its expression pattern does not change during the sensitive phase (Figure 3K). Altogether, these data indicate that

Tlx3 could be implicated in the specification of at least a subpopulation of these migrating cells from the OE, during the early onset of the olfactory connections. In summary, a dynamic balance of multiple attractive and repulsive signal cues during the olfactory development, and specifically during the sensitive phase, are important to allow the formation of the first glomerulus in a time-space dependent manner [52,63]. Moreover, changes in the mRNA levels of a complex array of factors, seem to govern the ability of olfactory axons to undergo guidance, targeting and synaptogenesis.

## Concluding remarks

Regardless of the striking features showed by the olfactory system and the large amount of cell populations and molecular signals involved in these processes, it is still unknown how all these processes cooperate together. Axonal growth and targeting between OE/

**Table 2.** Significant differences in the mRNA levels of different genes implicated in migration, ECM interaction, proliferation, apoptosis and differentiation in the ventro-medial part of the OB during the sensitive phase. Table shows the expression values of each gene, the comparison between the three stages of the sensitive phase (E14, E15 and E16), as well as the fold differences between each stage.

Gene	2-ΔCt		Fold Difference	p-value	Fold Up(+) or Down(-) Regulation
	E14	E15			
<i>Acan</i> ( <i>Aggrecan</i> )	5.33 E-08	4.82 E-07	0.11	0.008	- 9.04
<i>Bcan</i> ( <i>Brevican</i> )	2.68 E-07	1.28 E-06	0.21	0.013	- 4.79
<i>Col4a3</i> ( <i>Collagen, type IV, alpha 3</i> )	1.41 E-08	3.24 E-08	0.44	0.001	- 2.29
<i>Nlgn1</i> ( <i>Neuroligin-1</i> )	1.19 E-08	1.64 E-07	0.07	0.012	- 13.74
<i>Nrcam</i> ( <i>Neuronal cell adhesion molecule</i> )	6.63 E-06	3.07 E-05	0.22	0.028	- 4.63
<i>Psen1</i> ( <i>Presenilin-1</i> )	7.15 E-06	2.44 E-05	0.29	0.001	- 3.41
<i>Sema3c</i> ( <i>Semaphorin 3C</i> )	1.57 E-06	4.25 E-06	0.37	0.032	- 2.72
<b>E15 vs E16</b>					
	E15	E16	<b>E15/E16</b>		
<i>Nrp2</i> ( <i>Neuropilin 2</i> )	5.07 E-06	5.47E-10	9270.94	N/A	+ 9 270.94
<b>E14 vs E16</b>					
	E14	E16	<b>E14/E16</b>		
<i>Acan</i> ( <i>Aggrecan</i> )	5.33 E-08	3.64 E-07	0.15	0.003	- 6.83
<i>Bcan</i> ( <i>Brevican</i> )	2.68 E-07	1.04 E-06	0.26	0.003	- 3.88
<i>Ccr5</i> ( <i>C-C chemokine receptor type 5</i> )	2.14 E-08	5.93 E-08	0.36	0.014	- 2.77
<i>Col4a3</i> ( <i>Collagen, type IV, alpha 3</i> )	1.41 E-08	4.59 E-08	0.31	0.009	- 3.25
<i>Cxcl1</i> ( <i>Chemokine (C-X-C motif) ligand 1</i> )	1.60 E-08	2.20 E-07	0.07	0.039	- 13.76
<i>Dcc</i> ( <i>Deleted in Colorectal Carcinoma</i> )	6.87 E-05	2.82 E-05	2.44	0.001	+ 2.44
<i>Gpc2</i> ( <i>Glypican 2, also known cerebroglycan</i> )	6.26 E-05	2.69 E-05	2.33	0.001	+ 2.33
<i>Nisch</i> ( <i>Nischarin</i> )	6.04 E-05	2.07 E-03	0.03	0.007	- 34.19
<i>Nrcam</i> ( <i>Neuronal cell adhesion molecule</i> )	6.63 E-06	2.59 E-05	0.26	0.016	- 3.92
<i>Ntrk3</i> ( <i>neurotrophic tyrosine kinase, receptor, type 3</i> )	3.92 E-06	2.00 E-06	1.96	0.005	+ 1.96
<i>Sema3e</i> ( <i>Semaphorin 3A</i> )	4.71 E-06	5.12 E-05	0.09	0.018	- 10.89
<i>Slit1</i>	3.77 E-06	2.20 E-06	1.72	0.009	+ 1.72
<i>Tlx3</i> ( <i>T-cell leukemia homeobox 3</i> )	2.14 E-08	3.38 E-07	0.06	0.023	- 15.83

OB during the sensitive phase is a dynamic and comparative process in which cells and axons respond to qualitative and quantitative molecular differences, expressed by neighbouring targets, and make their decisions based on the relative balance of attractive and repulsive forces. Besides, because the well structured spatial organization and specificity in the OB innervations, this system could serve

as a good model to address the important issue involving how the wiring between the PNS and CNS is performed.

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