



Constitutively expressed Protocadherin- α regulates the coalescence and elimination of homotypic olfactory axons through its cytoplasmic region

Sonoko Hasegawa¹, Takahiro Hirabayashi¹, Takahiko Kondo¹, Ken Inoue¹, Shigeyuki Esumi^{1,2}, Atsushi Okayama¹, Shun Hamada^{1,3} and Takeshi Yagi^{1*}

¹ KOKORO-Biology Group and CREST-JST, Laboratories for Integrated Biology, Graduate School of Frontier Biosciences, Osaka University, Osaka, Japan

² Department of Morphological Neural Science, Graduate School of Medical Sciences, Kumamoto University, Kumamoto, Japan

³ Department of Nutrition and Health Sciences, Fukuoka Women's University, Fukuoka, Japan

Edited by:

Alistair N. Garratt, Max Delbrück Center for Molecular Medicine, Germany

Reviewed by:

Alistair N. Garratt, Max Delbrück Center for Molecular Medicine, Germany

Andrew Chess, Mount Sinai School of Medicine, USA

*Correspondence:

Takeshi Yagi, Graduate School of Frontier Biosciences, Osaka University, 1-3 Yamadaoka, Suita, Osaka 565-0871, Japan.
e-mail: yagi@fbs.osaka-u.ac.jp.

Olfactory sensory neuron (OSN) axons coalesce into specific glomeruli in the olfactory bulb (OB) according to their odorant receptor (OR) expression. Several guidance molecules enhance the coalescence of homotypic OSN projections, in an OR-specific- and neural-activity-dependent manner. However, the mechanism by which homotypic OSN axons are organized into glomeruli is unsolved. We previously reported that the clustered protocadherin- α (Pcdh- α) family of diverse cadherin-related molecules plays roles in the coalescence and elimination of homotypic OSN axons throughout development. Here we showed that the elimination of small ectopic homotypic glomeruli required the constitutive expression of a Pcdh- α isoform and Pcdh- α 's cytoplasmic region, but not OR specificity or neural activity. These results suggest that Pcdh- α proteins provide a cytoplasmic signal to regulate repulsive activity for homotypic OSN axons independently of OR expression and neural activity. The counterbalancing effect of Pcdh- α proteins for the axonal coalescence mechanisms mediated by other olfactory guidance molecules indicate a possible mechanism for the organization of homotypic OSN axons into glomeruli during development.

Keywords: olfactory, axon, protocadherin, Pcdh, convergence, elimination, neural circuit, neuron

INTRODUCTION

The olfactory system can recognize and discriminate an enormous number of odor molecules in the external environment. In this system, individual olfactory sensory neurons (OSNs) in the olfactory epithelium (OE) express only one type of functional odorant receptor (OR) from ~ 1000 genes in mice (Buck and Axel, 1991; Chess et al., 1994; Malnic et al., 1999). The OSNs expressing one type of OR project and coalesce their axons into specific glomeruli that are spatially arranged on the surface of the olfactory bulb (OB) (Ressler et al., 1994; Vassar et al., 1994; Mombaerts et al., 1996).

The coalescence of OSN axons into glomeruli is directed by the expressed OR and is dependent on neural activity (Feinstein and Mombaerts, 2004; Feinstein et al., 2004; Mombaerts, 2006; Serizawa et al., 2006). In this process, guidance molecules including cell adhesion molecules and receptors organize the OSN axon projections; these guidance molecules include ephrin/Eph, Semaphorin/neuropilin, Plexin, BIG-2, and Kirrel2/3 (Schwartz et al., 2000, 2004; Walz et al., 2002; Cutforth et al., 2003; Taniguchi et al., 2003; Imai et al., 2006; Serizawa et al., 2006; Col et al., 2007; Kaneko-Goto et al., 2008; Takeuchi et al., 2010). The expression levels of these guidance molecules in the OSNs are regulated by OR-derived signals and the anatomical location of the OSNs in the OE, resulting in the coalescence of homotypic OSNs. Thus, the distinct expression levels and

combinations of guidance molecules in the OSNs constitute one basic mechanism for the approximate projection and coalescence of OSN axons.

The protocadherin- α (Pcdh- α) genes belong to the clustered Pcdh families, which encode cadherin-related proteins with distinct domains derived from multiple variable exons: six extracellular cadherin domains, a transmembrane domain, and a short cytoplasmic domain (Kohmura et al., 1998; Wu and Maniatis, 1999; Yagi, 2012). They also have a common cytoplasmic tail (type A or B) derived from three or four constant exons (Kohmura et al., 1998; Sugino et al., 2000). The diverse Pcdh- α molecules are required for the coalescence of homotypic OSN axons into the OR-specific glomeruli of the OB (Hasegawa et al., 2008).

To address the functional significance of the diversity of Pcdh- α proteins, here we produced mutants in which exons $\alpha 2$ - $\alpha C 2$, i.e., all but exon $\alpha 1$, were deleted in the variable region of the Pcdh- α cluster. Surprisingly, in the mutant mice, the remaining $\alpha 1$ isoform compensated for the others, and was constitutively expressed in the neurons including OSNs. The coalescence of OSN projections in these mutants looked normal. In contrast, loss of the common cytoplasmic region from the Pcdh- α proteins disrupted the axonal coalescence. These findings indicate that constitutively expressed Pcdh- α proteins provide a cytoplasmic signal to regulate axonal coalescence and eliminate ectopic glomeruli. We also showed that the expression and function of

Pcdh- α were not dependent on either OR-specific signaling or OSN-derived neural activity. Based on these results, we propose that Pcdh- α has an activity that causes repulsion (or elimination) in homotypic OSN axons, that is different from the coalescence activity provided by other olfactory guidance molecules.

MATERIALS AND METHODS

ANIMAL EXPERIMENTS

All the experimental procedures were in accordance with the Guide for the Care and Use of Laboratory Animals of the Science Council of Japan and were approved by the Animal Experiment Committee of Osaka University.

GENERATION OF Pcdh- α MUTANT MICE

Similar to produce *Pcdha* ^{Δ CR/ Δ CR} mice (Hasegawa et al., 2008), we generated *Pcdha* ^{Δ CR2/ Δ CR2} mice using the CAG-Cre transgene and two targeted mutations: one G1loxP allele and another ΔA allele (Katori et al., 2009). The *Pcdha* ^{Δ CR2/ Δ CR2} mice expressed no Pcdh- α protein similar to *Pcdha* ^{Δ CR/ Δ CR} mice (data not shown).

By mating G16Neo mutant mice, in which two *loxP* sites were inserted between exons $\alpha 1$ and $\alpha 2$ (Noguchi et al., 2009), G1loxP mice in which a *loxP* site was inserted between exons $\alpha c 2$ and $\alpha CR 1$ (Hasegawa et al., 2008), and synaptosomal complex protein1 (*Sycp*)-Cre transgenic mice (Noguchi et al., 2009), male mice carrying the G16Neo allele, G1loxP allele, and *Sycp*-Cre transgene were generated. These mice were crossed with C57BL/6 females, and the genotypes of pups were determined by Southern blotting using probes amplified by PCR with Probe-G16-F (5'-GGAGGACATGCACAAGTCATG-3') and Probe-G16-R (5'-TTGTGGTGTACAGCGACACC-3') primers, and by PCR using G16 primer (5'-GGCTATCCTGTGCTACAGAAC-3'), G16-GTP-R2 primer (5'-CCAATTAATATTGAGATTCATCCCC-3'), and G1-GTP-R primer (5'-GCCCAGGATGGCTCAAATTC-3'). Some pups carried the *Pcdha* ^{Δ (2-c2)} or the *Pcdha*^{*dup*(2-c2)} allele generated by trans-allelic targeted meiotic recombination (TAMERE) in the testis (Herault et al., 1998) (Figure 5). *Pcdha* ^{ΔA / ΔA} mutant mice with a truncated Pcdh- α protein were described previously (Katori et al., 2009).

RT-PCR

Total RNA from mouse whole brain was extracted with TRIzol Reagent (Life Technologies), and the cDNAs were synthesized using SuperScriptIII reverse transcriptase (Life Technologies), according to the manufacturer's protocol. Forward primers were $\alpha 1$, 5'-GTGACCACGCAGAAGTAAAT-3'; $\alpha 2$, 5'-GAAGAGAGACAACCACCTT-3'; $\alpha 3$, 5'-GACAACTGGTTGGAGACAT-3'; $\alpha 4$, 5'-CAATTGCAGTCTGCAGAGGA-3'; $\alpha 5$, 5'-ACCTCAGGGACCCAGCTCTA-3'; $\alpha 6$, 5'-GCATCAGGATTGAACGACG-3'; $\alpha 7$, 5'-CCTACCTCAGGGTCCCAGCT-3'; $\alpha 8$, 5'-CCATCTGTTTCTTTGGACTC-3'; $\alpha 9$, 5'-GGAAAGTCA TTCTGTTGGAG-3'; $\alpha 10$, 5'-GGTTCTGGAGATAGTGGAGT-3'; $\alpha 11$, 5'-GGAAAGACAGGAGTCAGAGT-3'; $\alpha 12$, 5'-GTCA GAGAGAAAGGCAGGTA-3'; $\alpha c 1$, 5'-GGGGATCATTCAA TGTGGA-3'; $\alpha c 2$, 5'-CCGGGAACCCTGATTATCCTA-3', and reverse primer was $\alpha CR 5'$ -GACTGTTTGGGGTTGCC-3'. Quantitative PCR analysis was performed with the SYBR

Green Master Mix (Life Technologies) using ABI 7900HT (Life Technologies) with primers for $\alpha 1$, 5'-CCCAGTTTGAAC ATAGGC-3' and 5'-CGAGGCAGAGTAGCGCC-3'; for αCR , 5'-AGAGCAGGCATGCACAGC-3' and 5'-GACTGTTTGGGG TTGCC-3'; *GAPDH*, 5'-GACTTCAACAGCAACTCCCAC-3' and 5'-TCCACCACCCTGTTGCTGTAG-3'. For statistical analysis of the qPCR data, One-Way ANOVA, Tukey's test was used.

SINGLE CELL RT-PCR

Single cell RT-PCR analysis of OSNs was performed as described previously (Esumi et al., 2005). Briefly, single OSNs were obtained from the OE of adult OMP-GFP mice (a generous gift from Dr. Mombaerts). We picked up single GFP-positive OSNs with a glass capillary and placed them in individual PCR tubes. After cDNA synthesis from the single cells, the first multiplex PCR was performed using a 5' PCR primer designed to contain a consensus sequence common to all the variable exons of the *Pcdh*- α family. Next, we amplified each *Pcdh*- α isoform using a specific primer pair and semi-nested second PCR. Finally, we directly sequenced the PCR-amplified products to identify which *Pcdh*- α mRNAs were expressed in individual cells.

IMMUNOBLOT ANALYSIS

Mouse whole brains were homogenized in 0.32 M sucrose containing 1 mM EDTA and protease inhibitors, with a Dounce homogenizer. The homogenates were spun at 800 \times g for 10 min. The supernatants were spun at 20,000 \times g at 4°C for 30 min. The pellets were lysed with lysis buffer (20 mM Tris-HCl, pH 7.5, 150 mM NaCl, 1 mM EDTA, 1% Triton X-100, protease inhibitors) and spun at 20,000 \times g at 4°C for 30 min. The supernatants were subjected to SDS-PAGE followed by immunoblot analysis using a rabbit anti-Pcdha antibody (Murata et al., 2004).

In situ HYBRIDIZATION HISTOCHEMISTRY

In situ hybridization histochemistry was performed as described previously (Katori et al., 2009; Noguchi et al., 2009). Briefly, 10 μ m-thick fresh-frozen sections were prepared and hybridized with a constant-region cRNA probe for $\alpha 4$ cDNA (nucleotides 2518–4559) to detect all the *Pcdh*- α mRNA members. To detect certain variable isoforms of the *Pcdh*- α mRNAs, probes for $\alpha 1$ cDNA (nucleotides 396–1134), and $\alpha 11$ cDNA (nucleotides 398–2367) were used.

IMMUNOHISTOCHEMISTRY

Immunohistochemistry was performed as described previously (Hasegawa et al., 2008). We used antibodies against mOR-EG and MOR28, kindly given by Dr. Yoshihara, and a rabbit anti-Pcdha CR antibody, a generous gift from Dr. Watanabe.

β -GALACTOSIDASE HISTOCHEMISTRY

β -galactosidase histochemistry was performed as described previously (Hasegawa et al., 2008). Littermates from *Pcdha*^{+/ Δ CR}, *Pcdha*^{+/ Δ CR2}, *Pcdha*^{+/ Δ (2-c2)}, and *Pcdha*^{+/ ΔA} heterozygous parents were examined to compare WT and mutant mice that were also homozygous for M71-IRES-taulacZ, MOR23-IRES-taulacZ, or P2-IRES-taulacZ (kindly provided from Dr. Mombaerts). For this study, we mainly used *Pcdha* ^{Δ CR/ Δ CR} mice, except for

counting the P2 glomeruli at P7 and P30, for which we used *Pcdha* $\Delta^{CR2/\Delta CR2}$ mice. The phenotypes described in this study were similar for these strains (see **Table 1**).

UNILATERAL NARIS OCCLUSION

Unilateral naris occlusion was performed as described previously (Philpot et al., 1997).

IMAGE ANALYSIS

Fluorescent digital images were captured using a fluorescence microscope (Olympus BX51) equipped with a DP50 CCD camera (Olympus).

STATISTICAL ANALYSIS

Statistical analysis was conducted using StatView J-4.5 (SAS Institute, Cary, NC). Comparison of the number of glomeruli was performed by the Mann–Whitney test. Values in graphs were expressed as the mean \pm SEM.

RESULTS

ABNORMAL AXONAL COALESCENCE OF THE OSNs IN *Pcdh*- α MUTANTS

The diverse *Pcdh*- α family is required for the normal coalescence of OSN axons into glomeruli and for the elimination of ectopic glomeruli in the OB. In *Pcdh*- α -deficient mice, abnormal, small ectopic glomeruli are observed for OSNs expressing the M71, M72, and MOR23 ORs (Hasegawa et al., 2008). To confirm whether *Pcdh*- α is involved in the axonal coalescence of homotypic OSNs in the glomeruli in ventral and dorsal positions of the OB, we examined the axonal coalescence of OSNs expressing the ORs MOR28 (ventral side) and mOR-EG (dorsal side). The glomerular position of MOR28, but not of mOR-EG, is greatly influenced by the disruption of *Neuropilin-2* or *Plexin-A3* (Takeuchi et al., 2010). The disruption of *BIG-2* impairs the

axonal coalescence of MOR28 and mOR-EG OSNs to different degrees (Kaneko-Goto et al., 2008). Immunostaining with OR-specific antibodies showed increased numbers of glomeruli in the OSNs at P30 in the *Pcdh*- α -deficient (*Pcdha* $\Delta^{CR/\Delta CR}$) vs. wild-type (WT) mice, for mOR-EG (averages, lateral 1.6 vs. 1.0; medial 1.4 vs. 1.0, respectively) and MOR28 (lateral 2.4 vs. 1.2; medial 1.3 vs. 1.0) (**Table 1**, **Figure 1A**), similar to the previous results for M71, M72, and MOR23 (Hasegawa et al., 2008).

The organization and projections of axons expressing the OR P2 are well-studied; P2 OSNs possess distinct characteristics from other OR-expressing OSNs, such as a lower sensitivity to the loss of neural activity (Lin et al., 2000; Zheng et al., 2000) and a distinct zonal distribution within the OE. Therefore, we next examined the P2 glomeruli in WT and *Pcdh*- α -deficient (*Pcdha* $\Delta^{CR/\Delta CR}$ and *Pcdha* $\Delta^{CR2/\Delta CR2}$) mice with a P2-IRES-*tau*lacZ locus. As seen for the other ORs, the number of P2 glomeruli increased in the *Pcdh*- α -deficient mice (**Figure 1B**). In whole-mount preparations, the lateral half-bulb showed a mean of 1.9 labeled glomeruli in WT mice ($n = 16$ half-bulbs), and 4.0 glomeruli in the *Pcdh*- α -deficient mice ($n = 20$ half-bulbs). In the medial half-bulb, there were 2.1 glomeruli in the WT ($n = 17$ half-bulbs), and 4.1 glomeruli in the *Pcdh*- α -deficient mice ($n = 20$ half-bulbs) (**Table 1**). Thus, all the homotypic OSNs examined showed a similar phenotype of increased ectopic glomeruli in *Pcdh*- α -deficient mice. These results suggested that *Pcdh*- α 's function contributes to axonal coalescence and the elimination of ectopic glomeruli for all kinds of homotypic OSNs.

GLOMERULAR FORMATION IN *Pcdh*- α MUTANT MICE AT LATE-EMBRYONIC AND NEONATAL STAGES

As described above, the *Pcdh*- α -deficient mice had multiple, small, extraneous glomeruli for all the OSN axons examined. These ectopic glomeruli persist until adulthood (Hasegawa et al.,

Table 1 | Number of glomeruli per half-bulb in sectional analysis.

	Age	(n)	Lateral	(Min–Max)	(n)	Medial	(Min–Max)
10 μM-THICK SECTION							
mOR-EG	P30						
WT		(10)	1.0	(–)	(12)	1.0	(1.0–3.0)
$\Delta CR/\Delta CR$		(10)	1.6	(1.0–4.0) $P = 0.0301$	(09)	1.4	(1.0–2.0) $P = 0.0139$
MOR28	P30						
WT		(10)	1.2	(1.0–2.0)	(10)	1.0	(–)
$\Delta CR/\Delta CR$		(10)	2.4	(1.0–4.0) $P = 0.0053$	(10)	1.3	(1.0–3.0) $P = 0.1468$
50 μM-THICK SECTION							
P2	P30						
WT		(16)	1.9	(1.0–3.0)	(17)	2.1	(1.0–4.0)
$\Delta CR/\Delta CR$		(20)	4.0	(3.0–6.0) $P < 0.0001$	(20)	4.1	(2.0–6.0) $P < 0.0001$
P2	P30						
WT		(13)	2.7	(2.0–4.0)	(14)	2.2	(1.0–4.0)
$\Delta CR2/\Delta CR2$		(09)	5.3	(5.0–6.0) $P < 0.0001$	(09)	3.8	(2.0–6.0) $P = 0.0036$
MOR23	P30						
WT		(12)	1.3	(1.0–2.0)	(12)	1.0	(–)
$\Delta CR2/\Delta CR2$		(12)	2.0	(1.0–3.0) $P = 0.0099$	(12)	1.6	(1.0–3.0) $P = 0.0059$

$n =$ half-bulb, Mann–Whitney U -test.

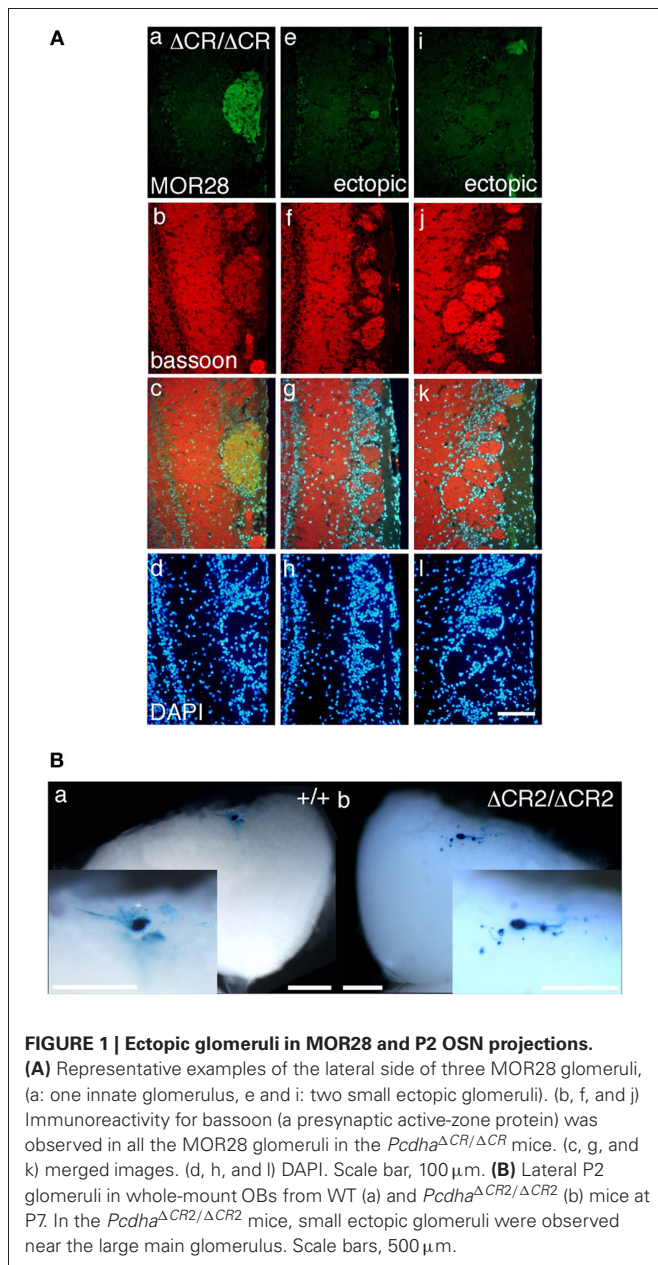


FIGURE 1 | Ectopic glomeruli in MOR28 and P2 OSN projections.

(A) Representative examples of the lateral side of three MOR28 glomeruli, (a: one innate glomerulus, e and i: two small ectopic glomeruli). (b, f, and j) Immunoreactivity for bassoon (a presynaptic active-zone protein) was observed in all the MOR28 glomeruli in the *Pcdha* $\Delta CR/\Delta CR$ mice. (c, g, and k) merged images. (d, h, and l) DAPI. Scale bar, 100 μ m. **(B)** Lateral P2 glomeruli in whole-mount OBs from WT (a) and *Pcdha* $\Delta CR2/\Delta CR2$ (b) mice at P7. In the *Pcdha* $\Delta CR2/\Delta CR2$ mice, small ectopic glomeruli were observed near the large main glomerulus. Scale bars, 500 μ m.

2008). To determine whether the abnormalities in the *Pcdh*- α -deficient olfactory system resulted primarily from an inability of homotypic OSN axons to coalesce, we next analyzed the formation of glomeruli during early development using mice from crosses between M71-IRES-taulacZ or P2-IRES-taulacZ and WT or *Pcdha* $\Delta CR/\Delta CR$ mice (Hasegawa et al., 2008). The glomerular structures are initiated on embryonic day (E) 15–16, when both OSN fibers and mitral cell dendrites contribute to the formation of glomerulus-like structures (Blanchart et al., 2006). In the *Pcdha* $\Delta CR/\Delta CR$ mice, we observed some stray marked fibers that projected to inappropriate regions distant from the target site, and many more M71 and P2 glomerulus-like structures than in WT mice, even on postnatal day 0 (P0) (arrowheads in Figures 2A, 3A).

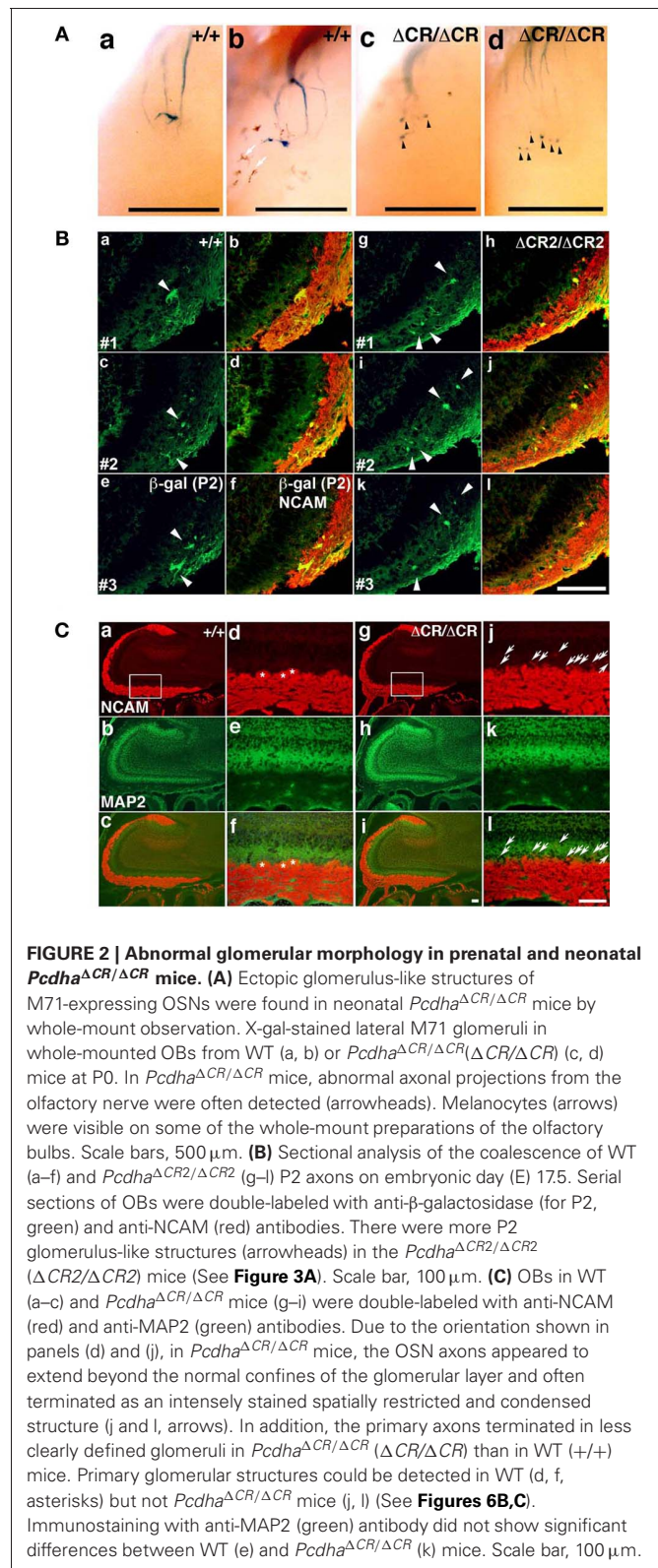
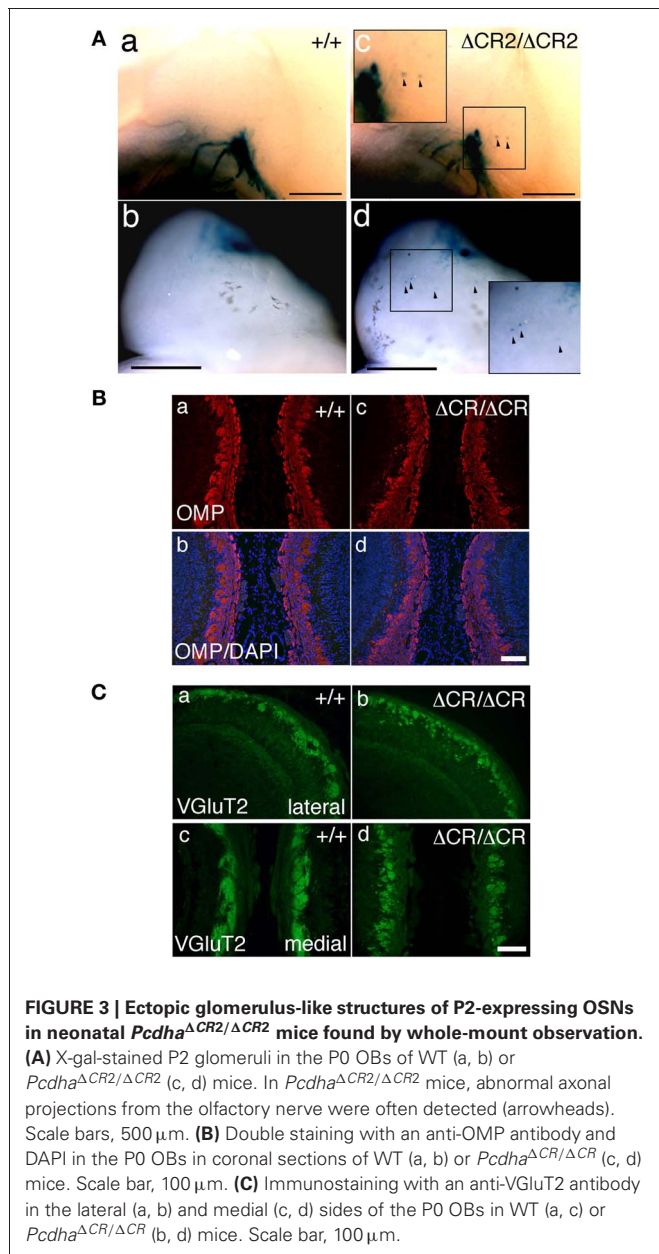


FIGURE 2 | Abnormal glomerular morphology in prenatal and neonatal *Pcdha* $\Delta CR/\Delta CR$ mice.

(A) Ectopic glomerulus-like structures of M71-expressing OSNs were found in neonatal *Pcdha* $\Delta CR/\Delta CR$ mice by whole-mount observation. X-gal-stained lateral M71 glomeruli in whole-mounted OBs from WT (a, b) or *Pcdha* $\Delta CR/\Delta CR$ ($\Delta CR/\Delta CR$) (c, d) mice at P0. In *Pcdha* $\Delta CR/\Delta CR$ mice, abnormal axonal projections from the olfactory nerve were often detected (arrowheads). Melanocytes (arrows) were visible on some of the whole-mount preparations of the olfactory bulbs. Scale bars, 500 μ m. **(B)** Sectional analysis of the coalescence of WT (a–f) and *Pcdha* $\Delta CR2/\Delta CR2$ (g–l) P2 axons on embryonic day (E) 17.5. Serial sections of OBs were double-labeled with anti- β -galactosidase (for P2, green) and anti-NCAM (red) antibodies. There were more P2 glomerulus-like structures (arrowheads) in the *Pcdha* $\Delta CR2/\Delta CR2$ ($\Delta CR2/\Delta CR2$) mice (See Figure 3A). Scale bar, 100 μ m. **(C)** OBs in WT (a–c) and *Pcdha* $\Delta CR/\Delta CR$ mice (d–f) were double-labeled with anti-NCAM (red) and anti-MAP2 (green) antibodies. Due to the orientation shown in panels (d) and (j), in *Pcdha* $\Delta CR/\Delta CR$ mice, the OSN axons appeared to extend beyond the normal confines of the glomerular layer and often terminated as an intensely stained spatially restricted and condensed structure (j and l, arrows). In addition, the primary axons terminated in less clearly defined glomeruli in *Pcdha* $\Delta CR/\Delta CR$ ($\Delta CR/\Delta CR$) than in WT (+/+) mice. Primary glomerular structures could be detected in WT (d, f, asterisks) but not *Pcdha* $\Delta CR/\Delta CR$ mice (j, l) (See Figures 6B,C). Immunostaining with anti-MAP2 (green) antibody did not show significant differences between WT (e) and *Pcdha* $\Delta CR/\Delta CR$ (k) mice. Scale bar, 100 μ m.

Sectional analysis of the OBs on E17.5 clearly showed the abnormal coalescence of P2 axons—multiple small and extraneous P2 glomerular-like structures—during the early formation of glomeruli (Figure 2B). These results indicated that



the ectopic glomeruli found in the *Pcdha*^{ΔCR/ΔCR} mice, which were often maintained until adulthood, were derived from the late-embryonic stage before birth, suggesting that Pcdh- α is involved in the initial axonal coalescence of the homotypic OSNs as well as the elimination of ectopic glomeruli during post-natal development. Ebrahimi and Chess (2000) showed that the successful coalescence of OSN axons at adulthood depends on the population size of OSNs (more OSNs result in better coalescence). On the other hand, small ectopic glomeruli that appear in normal newborn mice completely disappear by adulthood.

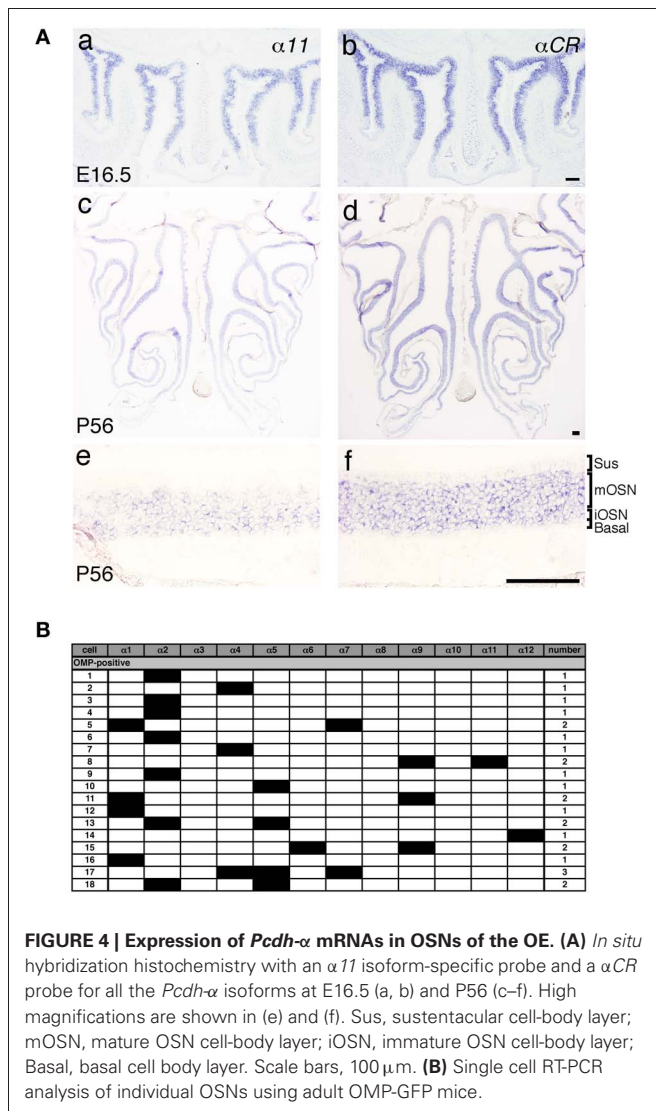
To analyze the dendrites of bulb neurons and OSN axons in more detail, we performed immunostaining for NCAM, an axonal marker and for MAP-2, a dendritic marker. In WT

mice at P0, the outline of individual glomeruli surrounded by periglomerular cells could be recognized in the glomerular layer by NCAM staining (asterisks in **Figure 2C** d, f). In contrast, such glomerular structures were not seen in *Pcdha*^{ΔCR/ΔCR} mice, in which NCAM-positive axons from the OSNs were often found in the external plexiform layer (arrows in **Figure 2C** j, l). These abnormal axonal clusters were confirmed by immunostaining with anti-OMP, a marker of OSNs and with anti-VGLuT2, a presynaptic marker (**Figures 3B,C**). These results suggested that abnormal axonal coalescence was a common feature of all the homotypic OSNs in the *Pcdha*^{ΔCR/ΔCR} mice. These observations support the idea that Pcdh- α helps to determine the precise organization of homotypic OSN axon projections during all developmental stages.

DIFFERENTIAL EXPRESSION OF EACH *Pcdh*- α ISOFORM IN SINGLE NEURONS INCLUDING OSNs

Pcdh- α mRNAs are extensively expressed in almost all of the OSNs from E11.5 to adulthood, and Pcdh- α proteins are enriched in the OSN axons and their terminals in the glomeruli (Hasegawa et al., 2008). However, the mechanisms by which the diverse Pcdh- α family contributes to the axonal coalescence into glomeruli remain unclear. To address this issue, we examined the expression pattern of various *Pcdh*- α isoforms in the OSNs. First, *in situ* hybridization histochemistry with probes for isoform-specific $\alpha 11$ and isoform-common αCR was performed for the OSNs of the OE. Extensive staining was observed at E16.5 for both the $\alpha 11$ and αCR probes (**Figure 4A** a, b). At P56, their expressions were widely observed in the OE, and appeared in both the immature and mature OSN cell body layers but not in the sustentacular or basal cell layers (**Figure 4A** c, d). In contrast to the αCR probe, the isoform-specific $\alpha 11$ probe showed a mosaic staining pattern (**Figure 4A** e). Similar mosaic patterns were observed with other α isoform-specific probes (data not shown, and see $\alpha 1$ in **Figure 6E**). The differential expression of *Pcdh*- α isoforms in single neurons is also observed in periglomerular cells (Kohmura et al., 1998) and mitral/tufted cells (data not shown) in the OB. These results suggested that individual OSNs and neurons in the OB differentially express *Pcdh*- α isoforms, similar to Purkinje cells (Esumi et al., 2005).

A previous single-cell RT-PCR analysis of Purkinje cells revealed strong evidence for the stochastic and combinatorial expression of *Pcdh*- α isoforms in individual neurons (Esumi et al., 2005). We therefore performed single-cell RT-PCR analysis of the OSNs to determine the expression pattern of Pcdh- α isoforms in these neurons at the single-cell level. Using OMP-GFP mice, we picked up GFP-positive cells. Of 30 GFP-positive OSNs, 18 were OMP positive and also had a product representing at least one of the 12 *Pcdh*- α variable isoforms (**Figure 4B**). Each single OSN expressed a different combination of *Pcdh*- α isoforms, except for four $\alpha 2$ -positive and two $\alpha 4$ -positive cells. Although this one-tube single-cell RT-PCR method has more experimental limitations than the split single-cell RT-PCR method, these results indicated that individual OSNs differentially express *Pcdh*- α isoforms, similar to Purkinje cells.



AXONAL COALESCENCE IN MUTANT MICE IN WHICH THE $\alpha 2$ TO $\alpha C2$ EXONS IN THE VARIABLE REGION OF THE *Pcdh-α* CLUSTER ARE DELETED

Our findings indicated that Pcdh- α proteins are essential for the homotypic axonal coalescence of OSNs; however, it was not known whether all or some specific Pcdh- α isoforms were necessary for this coalescence. To address this question, we produced and analyzed a *Pcdh-α* variable exon-deletion mutant line: *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice (Figure 5), in which the $\alpha 2$ - $\alpha C2$ exons were deleted, and only $\alpha 1$ remained, in the variable region of the *Pcdh-α* cluster (Figure 6A). In the *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice, the remaining $\alpha 1$ isoform was highly expressed in the brain (Figure 6B). The expression level of the $\alpha 1$ gene in the *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mouse brain increased by approximately 70-folds compared to WT (Figure 6C), while the total level of *Pcdh-α* expression detected by the common cytoplasmic region was similar between the *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ and WT mice. In immunoblots of brain extract samples, broad bands were immunostained by an anti-Pcdh- α CR antibody (Murata et al.,

2004) that recognized a common cytoplasmic region of Pcdh- α isoforms in WT mice; in contrast, a sharp band was strongly stained in the *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice (Figure 6D). We previously reported that in deletion mutants of the variable region of the *Pcdh-α* cluster, the missing exons are efficiently compensated for by the remaining variable exons (Noguchi et al., 2009). Therefore, to examine the compensation of *Pcdh-α* expression by the remaining $\alpha 1$ gene in the OSNs of *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice, we performed *in situ* hybridization analysis of the OE of WT and *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice with $\alpha 1$ and αCR probes (Figure 6E). Interestingly all the OSNs in the OE of *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice extensively expressed the $\alpha 1$ isoform similar to that of αCR , while the OSNs of WT mice rarely expressed $\alpha 1$. However, the expression pattern and level of the total *Pcdh-α* isoforms detected with the αCR probe were not markedly changed (Figure 6E). Immunostaining with the anti-Pcdh- α CR antibody also showed that the distribution pattern and level of the Pcdh- α proteins were not markedly different between the WT and *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice (Figure 6F). In *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice, the $\alpha 1$ protein was extensively distributed throughout the OSN axons and glomeruli. These results indicated that the $\alpha 1$ isoform completely compensated for the expression of the other *Pcdh-α* isoforms in all the OSNs and neurons of the OBs.

We next examined the axonal coalescence of homotypic OSNs of the *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice. We crossed the *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice with M71-IRES-*taulacZ* or MOR23-IRES-*taulacZ* mice. In whole-mount OB preparations of M71-IRES-*taulacZ* or MOR23-IRES-*taulacZ* mice at P30, there was typically one labeled glomerulus per half-bulb at the lateral and medial side in both WT and *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice (Table 2) (Figure 7A). Sectional analyses of the MOR23 glomeruli showed that the average number of glomeruli in both the lateral and medial sides of the half-bulbs were not significantly different between the WT and *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice (Table 2). Whole-mount analyses of the M71 glomeruli also showed no significant difference between the WT and *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice in the average number of glomeruli in both the lateral and medial sides of the half-bulbs (Table 2). The coalescence of M71 and MOR23 axons appeared normal in the *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice, in which the $\alpha 1$ protein was extensively enriched in all the OSN axons and glomeruli (Figure 7B). In addition, immunostaining of adjacent sections with the anti-Pcdh- α CR antibody revealed that the $\alpha 1$ protein was distributed in both the lateral M71 and medial MOR23 glomeruli of the OB in the *Pcdha* $^{\Delta(2-2)}/\Delta(2-2)$ mice (Figure 7B). These results indicated that a diversity of Pcdh- α isoforms in OSNs is not always required for the axonal coalescence of M71 and MOR23 homotypic OSNs into glomeruli. Instead, constitutive expression of the $\alpha 1$ isoform in neurons including OSNs was sufficient for the normal coalescence and elimination of OSN projections.

REQUIREMENT OF THE COMMON CYTOPLASMIC REGION OF *Pcdh-α* PROTEINS FOR THE NORMAL COALESCENCE AND ELIMINATION OF ECTOPIC OSN PROJECTIONS IN GLOMERULI

Pcdha $^{\Delta CR/\Delta CR}$ mice are presumptive null mutants of the *Pcdh-α* locus; no Pcdh- α proteins are seen in the brain of these mice (Hasegawa et al., 2008). Another *Pcdh-α* mutant mouse,

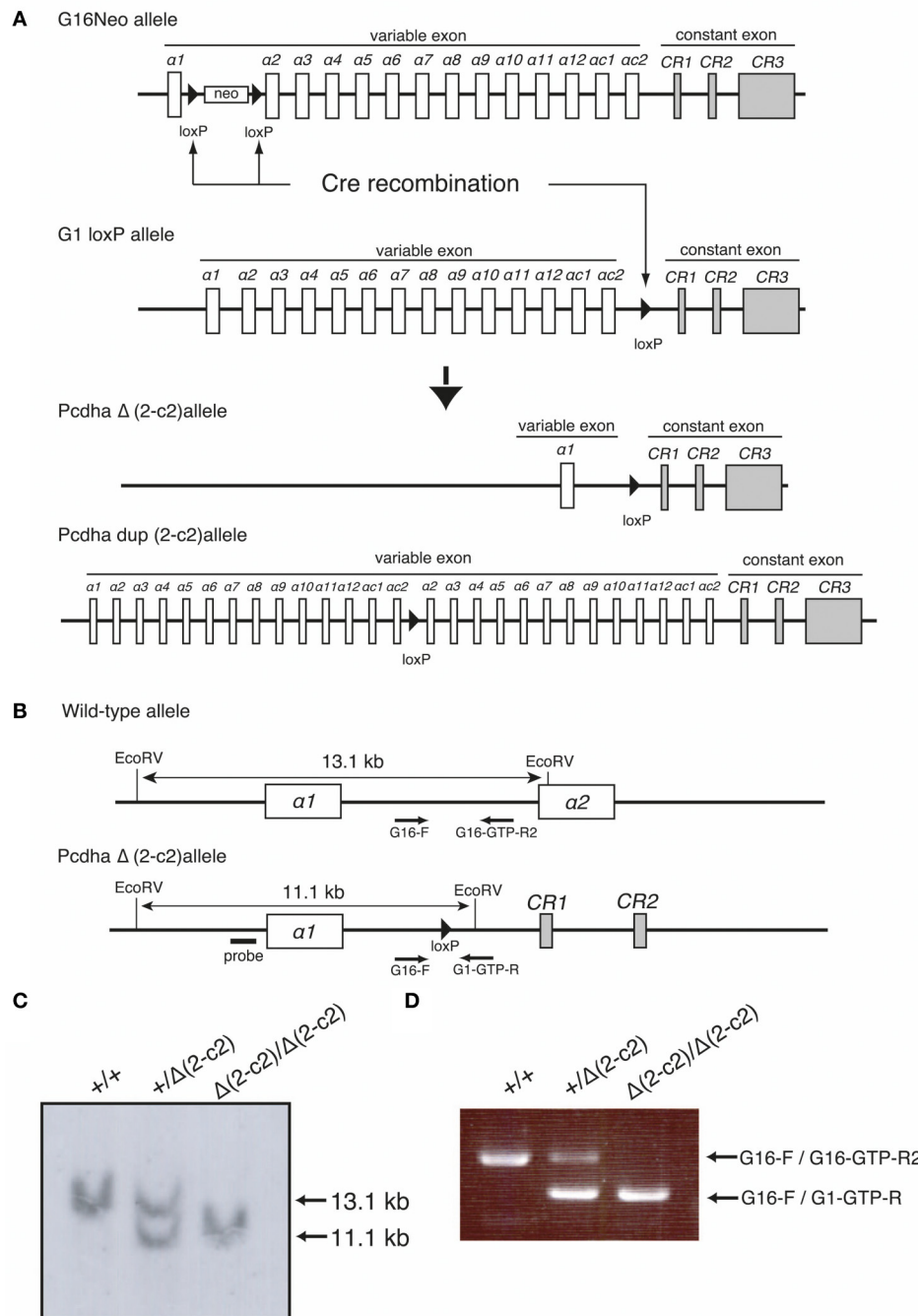


FIGURE 5 | Generation of the $Pcdha^{\Delta(2-c2)}$ and $Pcdha^{dup(2-c2)}$ alleles in mice. (A) $Pcdha^{\Delta(2-c2)}$ and $Pcdha^{dup(2-c2)}$ mice were generated by mating G16Neo mice, G1 loxP mice, and Sycp-Cre transgenic mice created by synaptosomal-Cre (TAMERE) system in the testis. This Cre is expressed

during meiotic crossing-over. (B) Partial genomic structures of the WT and $Pcdha^{\Delta(2-c2)}$ allele. (C,D) Genotyping of WT (+/+), $Pcdha^{+/\Delta(2-c2)}$ [+/ $\Delta(2-c2)$], and $Pcdha^{\Delta(2-c2)/\Delta(2-c2)}$ [$\Delta(2-c2)/\Delta(2-c2)$] by Southern blot and PCR analyses (see "Materials and Methods").

$Pcdha^{\Delta A/\Delta A}$, lacks the common cytoplasmic region (56 amino acids) of Pcdh- α A-type isoforms, and expresses a truncated Pcdh- α protein that lacks the A-type specific cytoplasmic tail (Katori et al., 2009). $Pcdha^{\Delta A/\Delta A}$ mice show disrupted and diffuse serotonergic axon projections, similar to those of the

$Pcdha^{\Delta CR/\Delta CR}$ mice (Katori et al., 2009). To further address how the Pcdh- α proteins control the axonal coalescence of homotypic OSNs expressing specific ORs, we analyzed the axonal coalescence in $Pcdha^{\Delta A/\Delta A}$ mice with the M71-IRES-taulacZ locus. We found a perturbed coalescence of M71 axons in the $Pcdha^{\Delta A/\Delta A}$ mice at

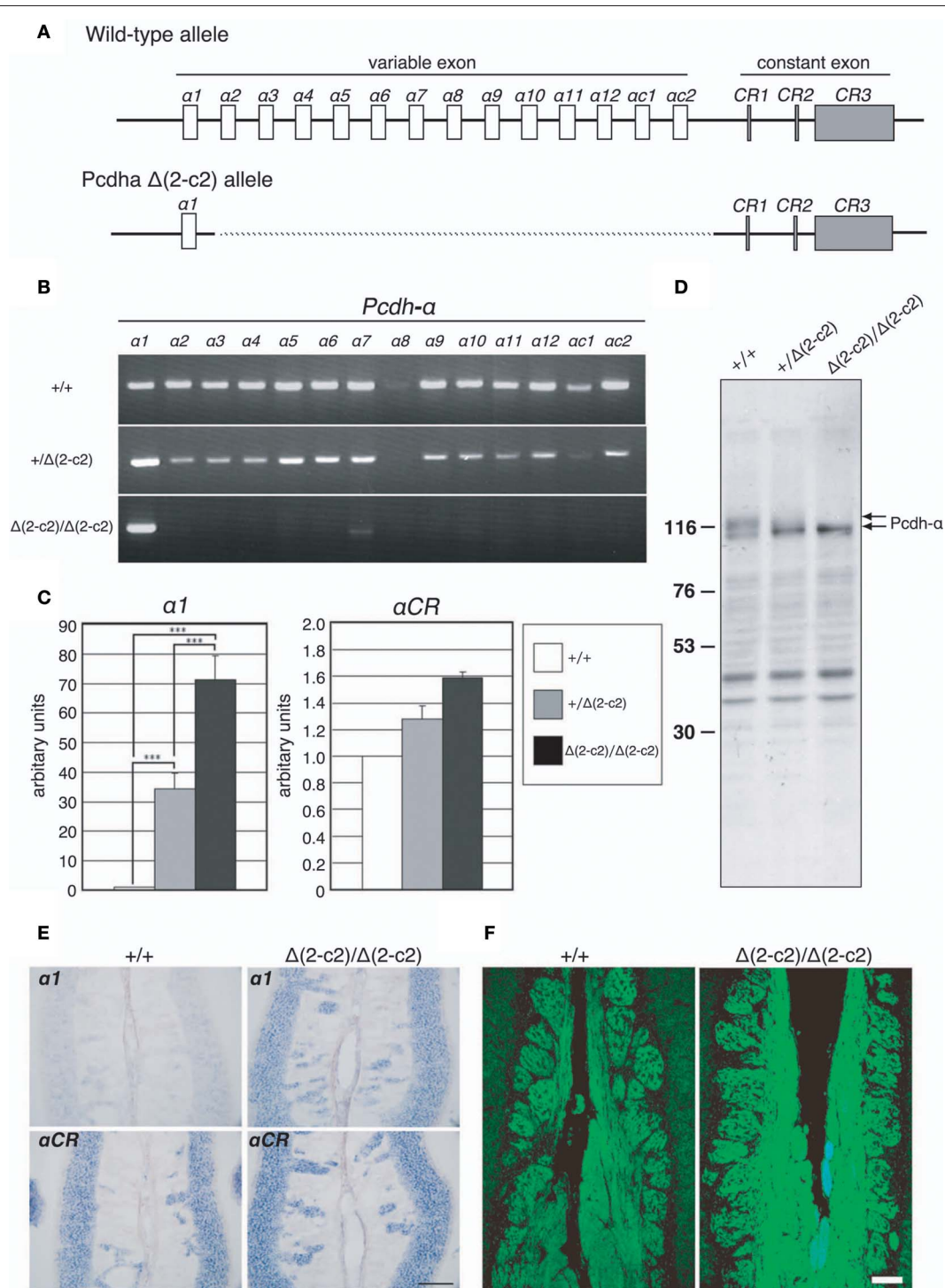


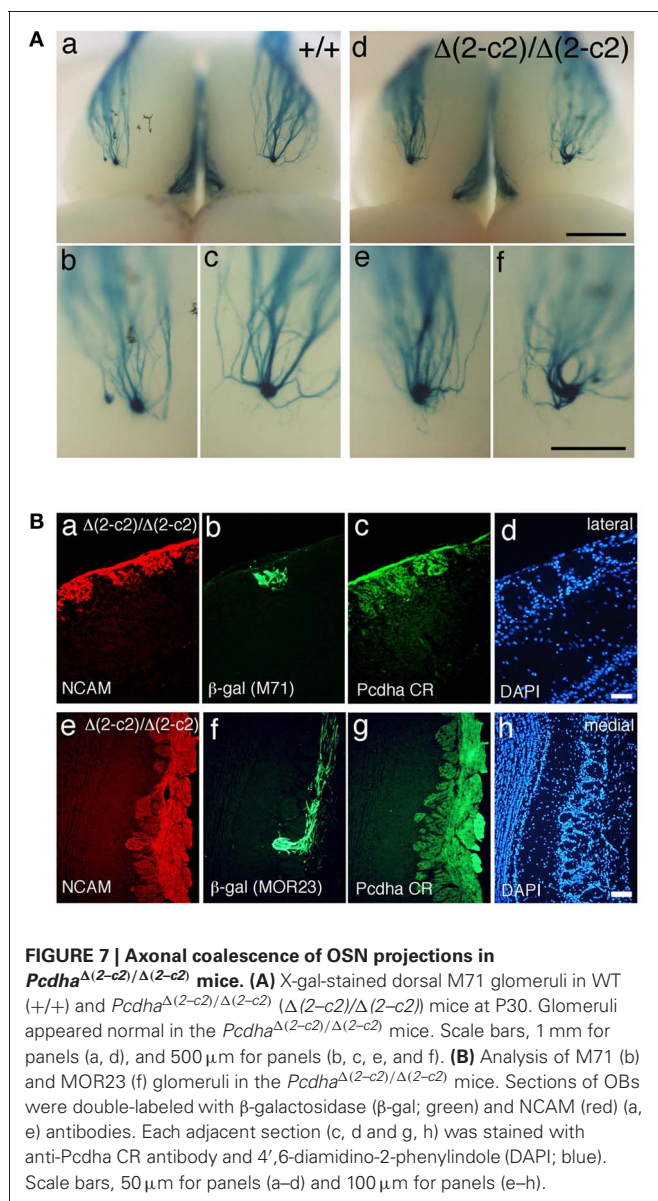
FIGURE 6 | *Pcdha* $\Delta(2-c2)/\Delta(2-c2)$ deletion mutant mice. (A) Wild-type *Pcdha* genes consist of variable-region ($\alpha 1$ to $\alpha 12$, $\alpha c 1$ and $\alpha c 2$) and constant-region (CR1–CR3) exons. The individual variable exons are transcribed from their own promoters. A *Pcdh- α* transcript is produced from one variable exon and three or four constant exons by splicing. In the *Pcdha* $\Delta(2-c2)/\Delta(2-c2)$ mice, exons $\alpha 2$ – $\alpha c 2$ were deleted, leaving only exon $\alpha 1$ in the variable region. (B) RT-PCR analysis of brain extracts of WT (+/+), *Pcdha* $^{+/\Delta(2-c2)}$ [+/ $\Delta(2-c2)$], and *Pcdha* $\Delta(2-c2)/\Delta(2-c2)$ [$\Delta(2-c2)/\Delta(2-c2)$] mice. (C) qRT-PCR analysis of $\alpha 1$ transcripts in the brain of WT (+/+), *Pcdha* $^{+/\Delta(2-c2)}$ [+/ $\Delta(2-c2)$], and

Pcdha $\Delta(2-c2)/\Delta(2-c2)$ [$\Delta(2-c2)/\Delta(2-c2)$] mice. *** $P < 0.0001$, vs. WT and vs. *Pcdha* $^{+/\Delta(2-c2)}$. Data are shown as the mean \pm S.D. (D) Immunoblotting analysis of brain lysates with an anti-Pcdh α CR antibody. (E) Expression of $\alpha 1$ and αCR transcripts in OSNs of the OE were examined by *in situ* hybridization histochemistry. Constitutive expression of $\alpha 1$ transcripts was seen in the OSNs of *Pcdha* $\Delta(2-c2)/\Delta(2-c2)$ [$\Delta(2-c2)/\Delta(2-c2)$] mice. Scale bar, 100 μ m. (F) Pcdh- α immunoreactivity with an anti-Pcdh α CR antibody was strong in the OSN axons and glomeruli of both WT (+/+) and *Pcdha* $\Delta(2-c2)/\Delta(2-c2)$ [$\Delta(2-c2)/\Delta(2-c2)$] mice. Scale bar, 100 μ m.

Table 2 | Number of glomeruli per half-bulb.

	Age	(n)	Lateral	(Min–Max)	(n)	Medial	(Min–Max)
50 μM-THICK SECTION							
MOR23	P30						
WT		(12)	1.0	(-)	(12)	1.0	(-)
$\Delta(2-c2)/\Delta(2-c2)$		(16)	1.0	(-)	(16)	1.1	(1.0–2.0) $P = 0.2120$
WHOLE-MOUNT							
M71	P30						
WT		(14)	1.4	(1.0–3.0)	(14)	1.5	(1.0–3.0)
$\Delta(2-c2)/\Delta(2-c2)$		(16)	1.7	(1.0–3.0) $P = 0.4620$	(16)	1.1	(1.0–2.0) $P = 0.0191$
M71	P30						
WT		(10)	1.2	(1.0–2.0)	(10)	1.1	(1.0–2.0)
$\Delta A/\Delta A$		(10)	2.1	(1.0–3.0) $P = 0.0150$	(10)	1.6	(1.0–2.0) $P = 0.0223$

n = half-bulb, Mann–Whitney U -test



P30 (**Figure 8A**) and P7 (**Figure 8B**). Similar abnormalities were found in the $Pcdha^{\Delta CR/\Delta CR}$ mice (**Figure 8B**, Hasegawa et al., 2008). Whole-mount analysis of the M71-IRES-*taulacZ* mice typically showed one labeled glomerulus per half-bulb at the lateral and medial side and in rare cases a second glomerulus; thus, there was an average of 1.1–1.2 glomeruli per lateral or medial side of the half-bulb at P30 in WT mice. In contrast, $Pcdha^{\Delta A/\Delta A}$ mice showed a perturbed coalescence of M71 axons and significantly higher numbers of M71 glomeruli compared to WT mice (**Table 2**). These results indicated that the common cytoplasmic region among Pcdh- α proteins is required for both the initial coalescence of OSN axons and the elimination of glomeruli during development.

NO EFFECT OF THE NEURAL ACTIVITY OF OSNs ON THE EXPRESSION LEVEL AND DISTRIBUTION PATTERN OF Pcdh- α

Axon guidance molecules showing OR-specific expression are often regulated by the neural activity of OSNs (Serizawa et al., 2006; Kaneko-Goto et al., 2008); therefore, we examined whether a reduction in OSN neural activity would affect the Pcdh- α expression. WT mice were subjected to unilateral naris occlusion at 3 weeks and analyzed by *in situ* hybridization histochemistry after 1 week. Efficiency of the naris occlusion was validated by the loss of tyrosine hydroxylase (TH) signals from the glomerular layer on the closed side of the OB (**Figure 9A a**) (Stone et al., 1990). In contrast, mRNA signals of the isoform common αCR probe in the OE were quite similar between the open and closed sides (**Figure 9A b**). In the OE, the expression levels with the $\alpha 11$ and αCR probes were quite similar between the open and closed sides (**Figure 9A c, d**). Furthermore, immunohistochemistry of the P30 WT mouse after naris occlusion showed that the distribution and intensity of Pcdh- α immunoreactivity in the OB were almost the same between the closed and open sides (**Figure 9B**). These results indicated that the Pcdh- α expression in the OSNs and OB was not altered by the OSN neural activity.

Naris occlusion leads to sensory deprivation, which might inhibit the neural activity-regulated process by which multiple glomeruli are eliminated. Indeed, a previous study showed that this treatment significantly increases the number of glomeruli (Zou et al., 2004). Therefore, we next analyzed the number

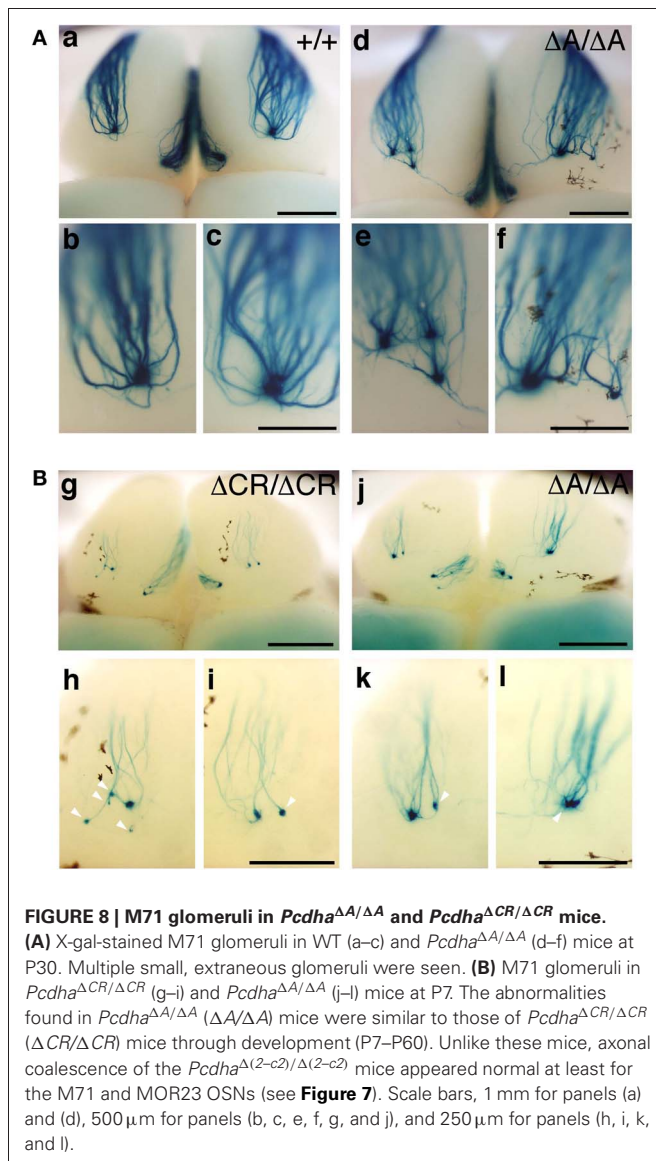


FIGURE 8 | M71 glomeruli in *Pcdh* $^{\Delta A/\Delta A}$ and *Pcdh* $^{\Delta CR/\Delta CR}$ mice. (A) X-gal-stained M71 glomeruli in WT (a–c) and *Pcdh* $^{\Delta A/\Delta A}$ (d–f) mice at P30. Multiple small, extraneous glomeruli were seen. (B) M71 glomeruli in *Pcdh* $^{\Delta CR/\Delta CR}$ (g–i) and *Pcdh* $^{\Delta A/\Delta A}$ (j–l) mice at P7. The abnormalities found in *Pcdh* $^{\Delta A/\Delta A}$ ($\Delta A/\Delta A$) mice were similar to those of *Pcdh* $^{\Delta CR/\Delta CR}$ ($\Delta CR/\Delta CR$) mice through development (P7–P60). Unlike these mice, axonal coalescence of the *Pcdh* $^{\Delta(2-c2)/\Delta(2-c2)}$ mice appeared normal at least for the M71 and MOR23 OSNs (see **Figure 7**). Scale bars, 1 mm for panels (a) and (d), 500 μ m for panels (b, c, e, f, g, and j), and 250 μ m for panels (h, i, k, and l).

of M71 and MOR23 glomeruli between the closed and open sides of the OB to examine the effects of naris occlusion in the *Pcdh* $^{\Delta CR/\Delta CR}$ or *Pcdh* $^{\Delta CR2/\Delta CR2}$ mice. Although the number of ectopic glomeruli was already significantly increased in these *Pcdh*- α -deficient mice, unilateral naris occlusion further increased the number of M71 and MOR23 glomeruli in the WT and *Pcdh*- α -deficient (*Pcdh* $^{\Delta CR/\Delta CR}$ and *Pcdh* $^{\Delta CR2/\Delta CR2}$) mice until they all reached similar levels (**Figure 9C**). Thus, even in *Pcdh*- α -deficient mice, the ectopic glomeruli were further increased by reduced neural activity. Together, these findings further suggest that *Pcdh*- α proteins function continuously to organize the projections of OSN axons and eliminate ectopic glomeruli in a neural activity-independent manner (**Figure 10**).

DISCUSSION

We previously reported that loss of the *Pcdh*- α family (*Pcdh* $^{\Delta CR/\Delta CR}$) disrupts the coalescence of OSN axons into

glomeruli (Hasegawa et al., 2008). However, the relationship between the diverse *Pcdh*- α isoforms and axonal coalescence was not well understood. In the present study, we found that *Pcdh*- α isoform expression and function in the olfactory system are different from those of other known olfactory guidance molecules. First, the phenotypes of axonal coalescence of OSNs in *Pcdh*- α -deficient mice were not restricted to specific OR-expressing OSNs but rather were found in all kinds of homotypic OSNs. Second, the expression of *Pcdh*- α molecules was not altered by the neural activity of OSNs. Third, following *Pcdh*- α deficiency, the numbers of ectopic glomeruli were even further increased by a reduction in neural activity, suggesting that *Pcdh*- α 's functions in coalescing and eliminating abnormal OSN axons are not dependent on neural activity. Fourth, the constitutive expression of *Pcdh*- α protein in neurons including OSNs was essential for normal axonal coalescence, although the diversity of *Pcdh*- α proteins was not always needed. In addition, we found that a common cytoplasmic region among *Pcdh*- α proteins was essential for the axonal coalescence and elimination of ectopic glomeruli. Taken together, we conclude that the *Pcdh*- α family is constitutively involved in mechanisms regulating the coalescence of OSN axons and elimination of small ectopic glomeruli that are independent of OR-specificity and neural activity.

Ebrahimi and Chess (2000) proposed a mechanism for maintaining the axonal coalescence of OSNs, in which the presence of other OSNs expressing the same OR is required, called “interdependence.” The projections of OSNs expressing an ectopic OR always coalesce in newborn mice, although this coalescence is not always maintained in adult mice. The probability of maintaining coalescence in adult mice is positively correlated with the number of OSNs expressing the OR (Ebrahimi and Chess, 2000). Thus, there must be some mechanism for removing the excess glomeruli. In the *Pcdh*- α -mutant mice, even small glomeruli do not disappear in adulthood. The present data showed that the constitutive expression and cytoplasmic region of *Pcdh*- α proteins are required for the elimination of miswired axons.

Pcdh- α MOLECULES MAY FUNCTION IN A NEURAL ACTIVITY-INDEPENDENT MANNER

It has been suggested that neural activity is required for the establishment and maintenance of specific glomeruli (Zou et al., 2004). Blocking the neural activity in OSNs by overexpression of the inward rectifying potassium channel (Kir2.1) induce multiple abnormal glomeruli in P2, MOR28 and MOR23 axons (Yu et al., 2004). On the other hand, in olfactory cyclic nucleotide-gated channel subunit 1 (OCNC1)-deficient mice, M72 axons form multiple abnormal glomeruli, but P2 axons do not (Lin et al., 2000; Zheng et al., 2000). Thus, P2 axons are probably less affected by a reduction in neural activity than other axon types in determining their targeted projections to the OB.

Interestingly, the *Pcdh*- α -deficient mice in our present study clearly showed abnormal multiple glomeruli in all OR-expressing OSNs examined, even in the P2 axons. Although it was already known that the elimination of ectopic glomeruli proceeds through the neural activity of OSNs, our present data suggest that *Pcdh*- α proteins may function in glomerular remodeling in a neural activity-independent fashion.

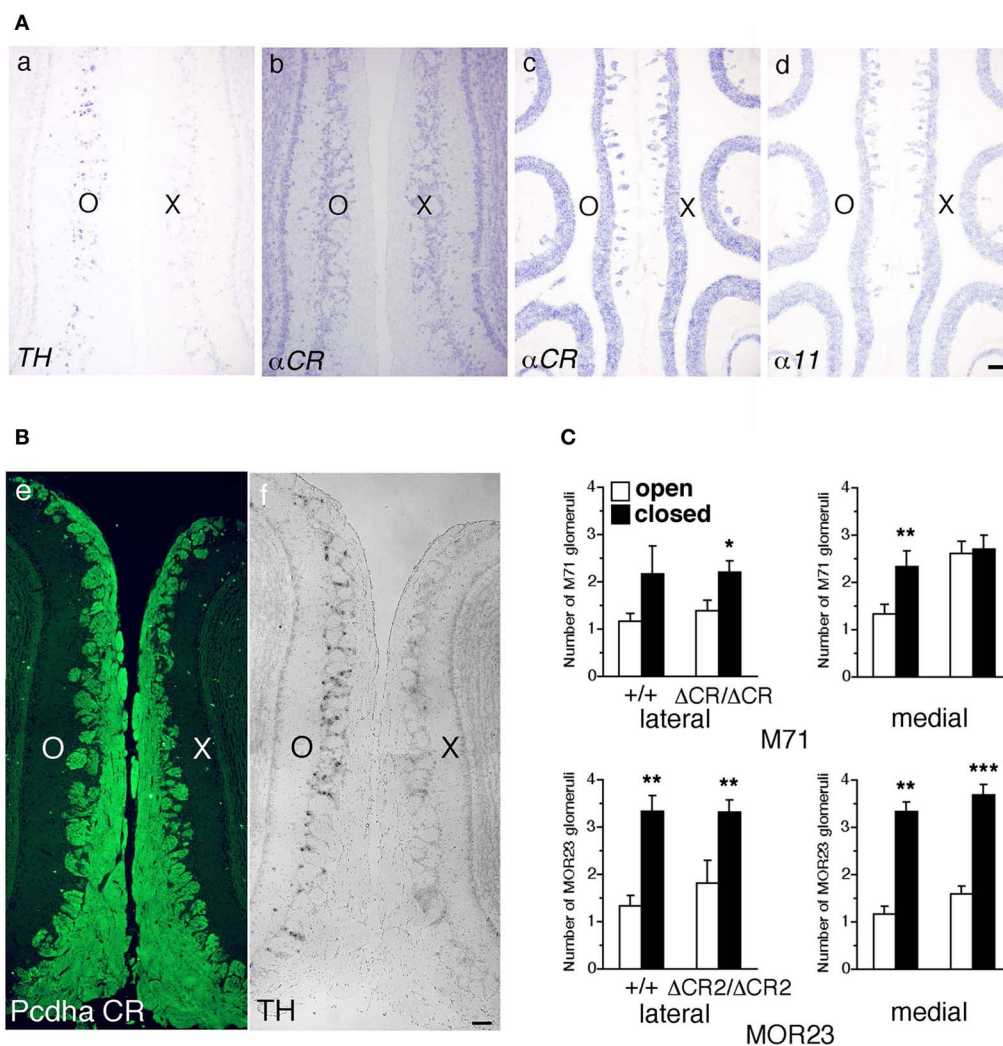


FIGURE 9 | Expression of *Pcdh- α* and the number of M71 and MOR23 glomeruli after naris occlusion of WT and *Pcdh- α* -deficient (*Pcdh α ^{ΔCR/ΔCR}* and *Pcdh α ^{ΔCR2/ΔCR2}*) mice. (A) A WT mouse was subjected to unilateral naris closure at P21 and analyzed after 1 week. The expression of tyrosine hydroxylase (*TH*) (a) and α CR transcripts (b) of the closed (X) and open (O) sides in the OB were examined by *in situ* hybridization histochemistry. The expression of α 11 and α CR transcripts in the OE were also examined (c, d). Scale bar, 100 μ m. (B) A WT mouse was subjected to naris occlusion at P5 and analyzed at P30. *Pcdh- α* immunoreactivity with the anti-*Pcdh α* CR antibody was strong in OSN axons and glomeruli in both the closed (X) and open (O) sides at similar levels (e), in

contrast to the change in *TH* signals (f). Scale bar, 100 μ m. (C) The number of M71 and MOR23 glomeruli per lateral and medial half-bulb of the open (white bars) and closed (black bars) sides after naris occlusion in WT (+/+) and *Pcdh- α* -deficient (*Pcdh α ^{ΔCR/ΔCR}* and *Pcdh α ^{ΔCR2/ΔCR2}*) mice at P30. Although ectopic glomeruli were further increased in *Pcdh α ^{ΔCR/ΔCR}* (Δ CR/ Δ CR) and *Pcdh α ^{ΔCR2/ΔCR2}* (Δ CR2/ Δ CR2) mice by the treatment, the total number of glomeruli after treatment was almost the same in the WT and *Pcdh- α* -deficient (*Pcdh α ^{ΔCR/ΔCR}* and *Pcdh α ^{ΔCR2/ΔCR2}*) mice. The number of MOR23 glomeruli in *Pcdh α ^{ΔCR2/ΔCR2}* mice are shown in **Table 1**. Significant differences at * $P < 0.05$, ** $P < 0.01$, and *** $P < 0.001$ calculated by the Mann-Whitney *U*-test.

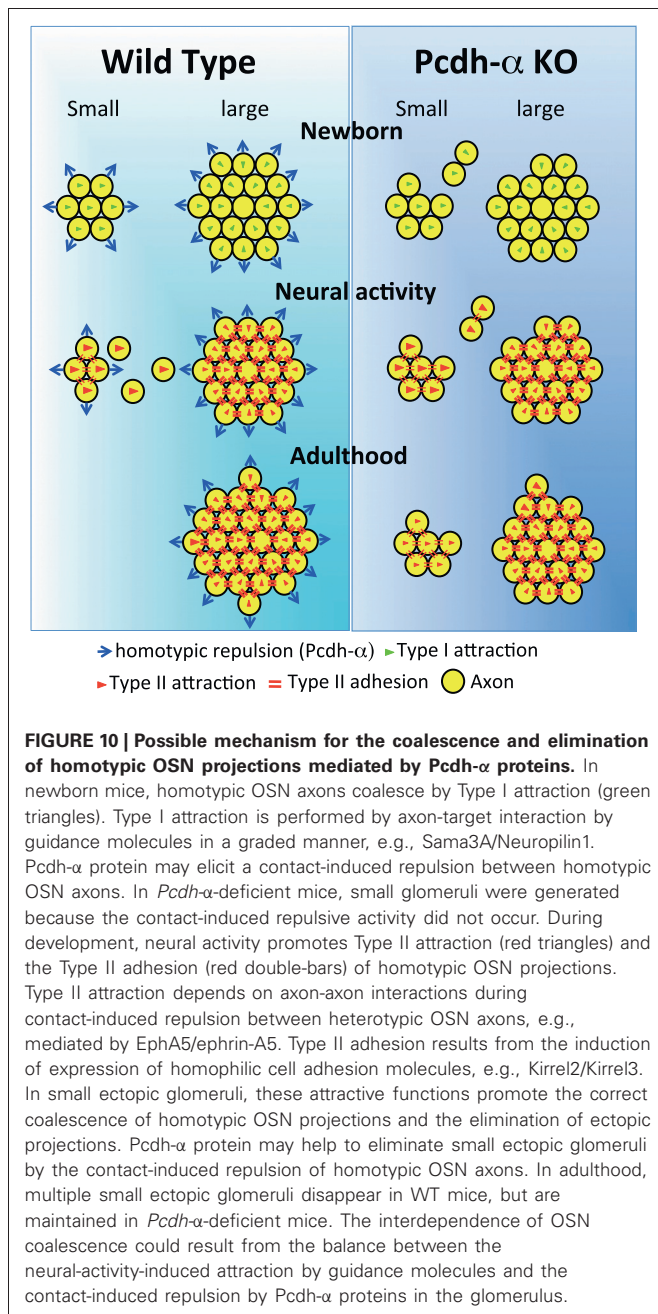
ROLE OF THE CYTOPLASMIC REGION OF *Pcdh- α* PROTEINS

The abnormalities in the axonal coalescence of OSNs in *Pcdh α ^{ΔA/ΔA}* mice were similar to those of *Pcdh α ^{ΔCR/ΔCR}* mice, indicating that the cytoplasmic region of the *Pcdh- α* protein is essential for the axonal coalescence of homotypic OSNs. The cytoplasmic domain of *Pcdh- α* is known to bind cytoplasmic signaling proteins of the focal adhesion kinase (FAK) family, FAK and PYK2 (Chen et al., 2009). The FAK family contributes to signaling cascades that regulate growth cones (Chacon and Fazzari, 2011) and membrane stabilization, with PKC and MARCKS (Garrett

et al., 2012). Interestingly, MARCKS-like and GAP43 proteins down-stream of PKC are extensively expressed in immature OSNs (McIntyre et al., 2010). Although a role for FAK-PKC signaling has not been reported in OSN axons, it may be involved in regulating coalescence and the elimination of glomeruli.

LOSS OF *Pcdh- α* MOLECULAR DIVERSITY: DELETION OF EXONS α 2 TO α 2 IN THE VARIABLE REGION OF THE *Pcdh- α* CLUSTER

In the *Pcdh- α* cluster, the variable region encodes multiple first exons (variable exons) for 14 different *Pcdh- α* isoforms. Each



variable exon is transcribed from its own promoter and *cis*-spliced to the constant region exons, which are common to all the Pcdh- α isoforms. While the $\alpha 1$ – $\alpha 12$ isoforms are expressed randomly, isoforms $\alpha c 1$ – $\alpha c 2$ are expressed constitutively in individual Purkinje cells (Esumi et al., 2005; Kaneko et al., 2006). Here we showed that Pcdh- α isoforms were also constitutively expressed in the OSNs of the OE, in which the $\alpha 1$ – $\alpha 12$ isoforms were randomly expressed, indicating that the mechanism of Pcdh- α gene regulation is similar in Purkinje cells and OSNs. Interestingly, in Pcdh $\alpha^{\Delta(2-c2)/\Delta(2-c2)}$ mice, in which exons $\alpha 2$ – $\alpha c 2$ in the variable region of the Pcdh- α cluster were deleted,

the remaining $\alpha 1$ isoform compensated for the others, and was expressed constitutively in neurons including OSNs (Figure 6). Such compensation of missing Pcdh- α cluster genes is also found in other deletion mutants, Pcdh $\alpha^{\Delta(11-c2)/\Delta(11-c2)}$ and Pcdh $\alpha^{\Delta(2-11)/\Delta(2-11)}$ (Noguchi et al., 2009). In these mice, the total expression level of Pcdh- α isoforms is maintained, and the remaining isoforms compensate for those missing. This compensation can be explained by a mechanism in which a *cis*-element for regulating the Pcdh- α cluster selects one or two gene(s) by binding within the $\alpha 1$ – $\alpha 12$ promoters. In fact, the *cis*-element for the Pcdh- α cluster was identified as HS5-1 (Ribich et al., 2006), and shown to be regulated in Purkinje cells *in vivo* (Yokota et al., 2011; Monahan et al., 2012). In addition, a chromatin factor known as CCCTC-binding factor (CTCF) is essential for gene regulation of the Pcdh- α cluster (Hirayama et al., 2012). Here, in the Pcdh $\alpha^{\Delta(2-c2)/\Delta(2-c2)}$ mice, the same Pcdh- α gene-regulation mechanisms might function in individual OSNs of the OE and in neurons of the OBs. In any case, the Pcdh $\alpha^{\Delta(2-c2)/\Delta(2-c2)}$ mice had normal-looking glomeruli in the OB, indicating that the constitutive expression, but not the diversity, of Pcdh- α protein is essential for the axonal coalescence of OSNs and the elimination of small ectopic glomeruli in the olfactory system.

POSSIBLE MECHANISM FOR THE INVOLVEMENT OF Pcdh- α PROTEIN IN THE COALESCENCE AND ELIMINATION OF HOMOTYPIC OSN AXONS

The development of OSN axon projections and their coalescence into glomeruli from the OE to the OB involves several molecular mechanisms regulated by step-wise processes. The stepwise regulation of OSN projection is categorized into type I for immature OSNs and type II for mature OSNs (Sakano, 2010). Immature and mature OSNs that express the same OR possess similar levels of the same guidance cues, and these homotypic axons project and coalesce to the same glomerulus position in the OB (Sakano, 2010).

Pcdh- α proteins may constitutively function to determine the organization of the OSN axon projections. We propose that the Pcdh- α family uses a novel mechanism for axonal coalescence and the elimination of ectopic glomeruli for all OR-expressing homotypic OSNs. The coalescence activity enhances the specific topography and OR-signaling-dependent neural activity mediated by guidance molecules (Sakano, 2010). Previously identified olfactory guidance molecules are known to contribute exclusively to the coalescence of homotypic OSN axons into rudimentary glomeruli, in a manner regulated by distinct levels of homophilic and repulsive activities of the OR-specific OSNs. However, no guidance molecules providing the homotypic repulsion for axonal elimination had been identified in the olfactory system. Here we propose that Pcdh- α proteins provide a repulsive activity for all homotypic OSNs and have a counterbalancing effect on the axonal coalescence mechanisms mediated by previously known olfactory guidance molecules (Figure 10). This Pcdh- α -mediated signaling provides a novel mechanism for eliminating OR-expressing OSN axons, that is, a repulsive signaling arising from all types of glomeruli that are independent of topography or neural activity.

Consequently, this hypothesis raises the further possibility that Pcdh- α could be a determinant of the specification of a major (large) glomerulus from other minor (small) glomeruli of homotypic OSNs. Pcdh- α protein is extensively expressed in both the presynaptic side of OSNs and the postsynaptic side of mitral/tufted and periglomerular cells, suggesting it may function in the appropriate maintenance and elimination of synaptic connections in homotypic glomeruli. Therefore, we propose a possible mechanism in which the interdependence of homotypic OSNs in axonal coalescence is ensured by a balance between the neural-activity-induced attraction by many guidance molecules and the contact-induced repulsion by Pcdh- α proteins in the glomeruli of the OB (**Figure 10**).

REFERENCES

- Blanchart, A., De Carlos, J. A., and Lopez-Mascaraque, L. (2006). Time frame of mitral cell development in the mice olfactory bulb. *J. Comp. Neurol.* 496, 529–543.
- Buck, L., and Axel, R. (1991). A novel multigene family may encode odorant receptors: a molecular basis for odor recognition. *Cell* 65, 175–187.
- Chacon, M. R., and Fazzari, P. (2011). FAK: dynamic integration of guidance signals at the growth cone. *Cell Adh. Migr.* 5, 52–55.
- Chen, J., Lu, Y., Meng, S., Han, M. H., Lin, C., and Wang, X. (2009). α - and γ -protocadherins negatively regulate PYK2. *J. Biol. Chem.* 284, 2880–2890.
- Chess, A., Simon, I., Cedar, H., and Axel, R. (1994). Allelic inactivation regulates olfactory receptor gene expression. *Cell* 78, 823–834.
- Col, J. A., Matsuo, T., Storm, D. R., and Rodriguez, I. (2007). Adenylyl cyclase-dependent axonal targeting in the olfactory system. *Development* 134, 2481–2489.
- Cutforth, T., Moring, L., Mendelsohn, M., Nemes, A., Shah, N. M., Kim, M. M., et al. (2003). Axonal ephrins and odorant receptors: coordinate determination of the olfactory sensory map. *Cell* 114, 311–322.
- Ebrahimi, F. A., and Chess, A. (2000). Olfactory neurons are interdependent in maintaining axonal projections. *Curr. Biol.* 10, 219–222.
- Esumi, S., Kakazu, N., Taguchi, Y., Hirayama, T., Sasaki, A., Hirabayashi, T., et al. (2005). Monoallelic yet combinatorial expression of variable exons of the protocadherin- α gene cluster in single neurons. *Nat. Genet.* 37, 171–176.
- Feinstein, P., Bozza, T., Rodriguez, I., Vassalli, A., and Mombaerts, P. (2004). Axon guidance of mouse olfactory sensory neurons by odorant receptors and the β 2 adrenergic receptor. *Cell* 117, 833–846.
- Feinstein, P., and Mombaerts, P. (2004). A contextual model for axonal sorting into glomeruli in the mouse olfactory system. *Cell* 117, 817–831.
- Garrett, A. M., Schreiner, D., Lobas, M. A., and Weiner, J. A. (2012). γ -protocadherins control cortical dendrite arborization by regulating the activity of a FAK/PKC/MARCKS signaling pathway. *Neuron* 74, 269–276.
- Hasegawa, S., Hamada, S., Kumode, Y., Esumi, S., Katori, S., Fukuda, E., et al. (2008). The protocadherin- α family is involved in axonal coalescence of olfactory sensory neurons into glomeruli of the olfactory bulb in mouse. *Mol. Cell. Neurosci.* 38, 66–79.
- Herault, Y., Rassoulzadegan, M., Cuzin, F., and Duboule, D. (1998). Engineering chromosomes in mice through targeted meiotic recombination (TAMERE). *Nat. Genet.* 20, 381–384.
- Hirayama, T., Tarusawa, E., Yoshimura, Y., Galjart, N., and Yagi, T. (2012). CTCF is required for neural development and stochastic expression of clustered Pcdh genes in neurons. *Cell Rep.* 2, 1–13.
- Imai, T., Suzuki, M., and Sakano, H. (2006). Odorant receptor-derived cAMP signals direct axonal targeting. *Science* 314, 657–661.
- Kaneko, R., Kato, H., Kawamura, Y., Esumi, S., Hirayama, T., Hirabayashi, T., et al. (2006). Allelic gene regulation of Pcdh- α and Pcdh- γ clusters involving both monoallelic and biallelic expression in single Purkinje cells. *J. Biol. Chem.* 281, 30551–30560.
- Kaneko-Goto, T., Yoshihara, S., Miyazaki, H., and Yoshihara, Y. (2008). BIG-2 mediates olfactory axon convergence to target glomeruli. *Neuron* 57, 834–846.
- Katori, S., Hamada, S., Noguchi, Y., Fukuda, E., Yamamoto, T., Yamamoto, H., et al. (2009). Protocadherin- α family is required for serotonergic projections to appropriately innervate target brain areas. *J. Neurosci.* 29, 9137–9147.
- Kohmura, N., Senzaki, K., Hamada, S., Kai, N., Yasuda, R., Watanabe, M., et al. (1998). Diversity revealed by a novel family of cadherins expressed in neurons at a synaptic complex. *Neuron* 20, 1137–1151.
- Lin, D. M., Wang, F., Lowe, G., Gold, G. H., Axel, R., Ngai, J., et al. (2000). Formation of precise connections in the olfactory bulb occurs in the absence of odorant-evoked neuronal activity. *Neuron* 26, 69–80.
- Malnic, B., Hirono, J., Sato, T., and Buck, L. B. (1999). Combinatorial receptor codes for odors. *Cell* 96, 713–723.
- McIntyre, J. C., Titlow, W. B., and McClintock, T. S. (2010). Axon growth and guidance genes identify nascent, immature, and mature olfactory sensory neurons. *J. Neurosci. Res.* 88, 3243–3256.
- Mombaerts, P. (2006). Axonal wiring in the mouse olfactory system. *Annu. Rev. Cell. Dev. Biol.* 22, 713–737.
- Mombaerts, P., Wang, F., Dulac, C., Chao, S. K., Nemes, A., Mendelsohn, M., et al. (1996). Visualizing an olfactory sensory map. *Cell* 87, 675–686.
- Monahan, K., Rudnick, N. D., Kehayova, P. D., Pauli, F., Newberry, K. M., Myers, R. M., et al. (2012). Role of CCCTC binding factor (CTCF) and cohesin in the generation of single-cell diversity of protocadherin- α gene expression. *Proc. Natl. Acad. Sci. U.S.A.* 109, 9125–9130.
- Murata, Y., Hamada, S., Morishita, H., Mutoh, T., and Yagi, T. (2004). Interaction with protocadherin- γ regulates the cell surface expression of protocadherin- α . *J. Biol. Chem.* 279, 49508–49516.
- Noguchi, Y., Hirabayashi, T., Katori, S., Kawamura, Y., Sanbo, M., Hirabayashi, M., et al. (2009). Total expression and dual gene-regulatory mechanisms maintained in deletions and duplications of the Pcdha cluster. *J. Biol. Chem.* 284, 32002–32014.
- Philpot, B. D., Lim, J. H., Halpain, S., and Brunjes, P. C. (1997). Experience-dependent modifications in MAP2 phosphorylation in rat olfactory bulb. *J. Neurosci.* 17, 9596–9604.
- Ressler, K. J., Sullivan, S. L., and Buck, L. B. (1994). Information coding in the olfactory system: evidence for a stereotyped and highly organized epitope map in the olfactory bulb. *Cell* 79, 1245–1255.
- Ribich, S., Tasic, B., and Maniatis, T. (2006). Identification of long-range regulatory elements in the protocadherin- α gene cluster. *Proc. Natl. Acad. Sci. U.S.A.* 103, 19719–19724.
- Sakano, H. (2010). Neural map formation in the mouse olfactory system. *Neuron* 67, 530–542.
- Schwartz, G. A., Kostek, C., Ahmad, N., Dibble, C., Pays, L., and Puschel, A. W. (2000). Semaphorin 3A is required for guidance of olfactory axons in mice. *J. Neurosci.* 20, 7691–7697.
- Schwartz, G. A., Raitcheva, D., Crandall, J. E., Burkhardt, C., and Puschel, A. W. (2004). Semaphorin 3A-mediated axon guidance regulates convergence and targeting of P2 odorant receptor axons. *Eur. J. Neurosci.* 19, 1800–1810.
- Serizawa, S., Miyamichi, K., Takeuchi, H., Yamagishi, Y., Suzuki, M., and Sakano, H. (2006). A neuronal identity code for the odorant receptor-specific and activity-dependent axon sorting. *Cell* 127, 1057–1069.

- Stone, D. M., Wessel, T., Joh, T. H., and Baker, H. (1990). Decrease in tyrosine hydroxylase, but not aromatic L-amino acid decarboxylase, messenger RNA in rat olfactory bulb following neonatal, unilateral odor deprivation. *Brain Res. Mol. Brain Res.* 8, 291–300.
- Sugino, H., Hamada, S., Yasuda, R., Tuji, A., Matsuda, Y., Fujita, M., et al. (2000). Genomic organization of the family of CNR cadherin genes in mice and humans. *Genomics* 63, 75–87.
- Takeuchi, H., Inokuchi, K., Aoki, M., Suto, F., Tsuboi, A., Matsuda, I., et al. (2010). Sequential arrival and graded secretion of Sema3F by olfactory neuron axons specify map topography at the bulb. *Cell* 141, 1056–1067.
- Taniguchi, M., Nagao, H., Takahashi, Y. K., Yamaguchi, M., Mitsui, S., Yagi, T., et al. (2003). Distorted odor maps in the olfactory bulb of semaphorin 3A-deficient mice. *J. Neurosci.* 23, 1390–1397.
- Vassar, R., Chao, S. K., Sitcheran, R., Nunez, J. M., Vosshall, L. B., and Axel, R. (1994). Topographic organization of sensory projections to the olfactory bulb. *Cell* 79, 981–991.
- Walz, A., Rodriguez, I., and Mombaerts, P. (2002). Aberrant sensory innervation of the olfactory bulb in neuropilin-2 mutant mice. *J. Neurosci.* 22, 4025–4035.
- Wu, Q., and Maniatis, T. (1999). A striking organization of a large family of human neural cadherin-like cell adhesion genes. *Cell* 97, 779–790.
- Yagi, T. (2012). Molecular codes for neuronal individuality and cell assembly in the brain. *Front. Mol. Neurosci.* 5:45. doi: 10.3389/fnmol.2012.00045
- Yokota, S., Hirayama, T., Hirano, K., Kaneko, R., Toyoda, S., Kawamura, Y., et al. (2011). Identification of the cluster control region for the protocadherin-beta genes located beyond the protocadherin-gamma cluster. *J. Biol. Chem.* 286, 31885–31895.
- Yu, C. R., Power, J., Barnea, G., O'Donnell, S., Brown, H. E., Osborne, J., et al. (2004). Spontaneous neural activity is required for the establishment and maintenance of the olfactory sensory map. *Neuron* 42, 553–566.
- Zheng, C., Feinstein, P., Bozza, T., Rodriguez, I., and Mombaerts, P. (2000). Peripheral olfactory projections are differentially affected in mice deficient in a cyclic nucleotide-gated channel subunit. *Neuron* 26, 81–91.
- Zou, D. J., Feinstein, P., Rivers, A. L., Mathews, G. A., Kim, A., Greer, C. A., et al. (2004). Postnatal refinement of peripheral olfactory projections. *Science* 304, 1976–1979.
- commercial or financial relationships that could be construed as a potential conflict of interest.

Received: 23 August 2012; accepted: 27 September 2012; published online: 16 October 2012.

Citation: Hasegawa S, Hirabayashi T, Kondo T, Inoue K, Esumi S, Okayama A, Hamada S and Yagi T (2012) Constitutively expressed Protocadherin- α regulates the coalescence and elimination of homotypic olfactory axons through its cytoplasmic region. *Front. Mol. Neurosci.* 5:97. doi: 10.3389/fnmol.2012.00097

Copyright © 2012 Hasegawa, Hirabayashi, Kondo, Inoue, Esumi, Okayama, Hamada and Yagi. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits use, distribution and reproduction in other forums, provided the original authors and source are credited and subject to any copyright notices concerning any third-party graphics etc.

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any