Types of Parvalbumin-Containing Retinotectal Ganglion Cells in Mouse

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The calcium-binding protein parvalbumin (PV) occurs in the retinal ganglion cells (RGCs) of various vertebrate species. In the present study, we aimed to identify the types of PV-containing RGCs that project to the superior colliculus (SC) in the mouse. We injected retrograde tracer dextran into the mouse SC to label RGCs. PV-containing RGCs were first identified by immunocytochemistry and then neurons double-labeled with dextran and PV were iontophoretically injected with a lipophilic dye, DiI. Subsequently, confocal microscopy was used to characterize the morphologic classification of the PV-immunoreactive (IR) retinotectal ganglion cells on the basis of dendritic field size, branching pattern, and stratification within the inner plexiform layer. Among the 8 different types of PV-containing RGCs in the mouse retina, we found all 8 types of RGCs projecting to the SC. The RGCs were heterogeneous in morphology. The combined approach of using tracer injection and a single cell injection after immunocytochemistry on a particular protein will provide valuable data to further understand the functional features of the RGCs which constitute the retinotectal pathway.

Key words: parvalbumin, retinal ganglion cell, superior colliculus, cell type, cell injection

I. Introduction

Visual information undergoes a first step of processing in the retina, and retinal ganglion cells (RGCs) project to particular visual targets in the brain. Thus, with the advent of modern techniques, the morphological and functional properties of RGCs have been studied in variety of species. The 3 most common types of RGCs have been distinguished in the monkey retina: Large magnocellular (parasol), smaller parvocellular (midget) and small bistatified ganglion cells [11, 33]. Recently, however, at least 8 previously unknown RGCs that project to the lateral geniculate nucleus were identified by using an *in vitro* photostaining method [12] and at least 15 types of wide-field RGCs were identified by microinjection with neurobiotin [48]. In the cat, 2 classes of morphologically distin-

guished RGCs (small beta and large alpha cells) have been shown to correspond to the physiologically defined X and Y cells. Similarly, these retinas have a gamma-type ganglion cell corresponding to a W cell [5, 37, 46]. However, at least 9 types of RGCs were seen anatomically to be non-beta, non-alpha cells, by using intracellular staining [17, 30]. Eleven different types of RGCs were distinguished in the rabbit retina, by using a combination of sophisticated anatomic techniques [32].

The mouse retina is a valuable model for vision research because the mouse is a mainstay of transgenic technology. Therefore, Sun *et al.* [39] have used DiI particle bombardment to describe 14 types of RGCs. Kong *et al.* [21] and Coombs *et al.* [9] have analyzed 11–14 RGCs clusters in the mouse retina using various methods to label cells. Badea and Nathans [2] have analyzed 12 RGC types in the mouse using a genetically directed reporter. Recently, Völgyi *et al.* [44] have morphologically analyzed 22 RGC subtypes based on soma-dendritic parameters in the mouse. Furthermore, our previous studies have shown the morphology of RGCs with the expression of parvalbumin (PV)

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[20] and calretinin [22] by using a newly developed single-cell injection method following immunocytochemistry.

Among the many calcium-binding proteins, at least 3 of the EF-hand calcium-binding proteins, calbindin D28K, calretinin and PV have been applied as good markers to distinguish subpopulations of neurons [3]. Calcium plays a crucial role in cell signal transduction, neurotransmitter secretion and muscle contraction. Calcium-binding proteins are thought to control and modulate these cellular mechanisms of calcium. Misregulation of calcium-binding proteins is closely related to many neurodegenerative diseases [15]. In particular, the expression of parvalbumin (PV) in the nervous system is restricted to the neurons, and mostly to a GABAergic subpopulation of neurons [8]. Recently, neural circuits with PV-immunoreactive (IR) neurons have been studied in the brain and gradually were shown to function in the neural circuitry. Although the systematic study of RGC types with its expression of a particular protein has been previously accomplished for PV and calretinin in the mouse [22, 26], the types of RGCs that project to the superior colliculus (SC), and their expression of specific proteins, have not been studied.

In general, many RGCs project to the lateral geniculate nucleus in the thalamus and SC in the midbrain. The SC is the center of visual motor integration, and controls saccadic movement of eyes. A number of studies have been performed to correlate the distribution and various physiological types of RGCs with the SC [10, 27, 31, 43, 45].

In the present study, we have extended the systematic morphological analysis approach to the study of mouse RGCs with their expression of specific proteins. Our study aimed to identify morphological types PV-containing RGCs with their retinotectal pathways by using advanced approaches. The advanced approach we chose involved triple labeling techniques: retrograde tracer injection and single-cell injection of lipophilic dye, following immunocytochemistry.

II. Materials and Methods

Animals and retrograde tracer injections

Twenty-five adult mice (C57/BL6) were anesthetized intramuscularly with a mixture of ketamine hydrochloride (30–40 mg/kg) and xylazine (3–6 mg/kg), supplemented as needed to maintain anesthesia. The animals were mounted in a stereotaxic apparatus. A small craniotomy was then produced, and dextran (tetramethylrhodamine, 3000 MW, 1–3 μ l of a 10% solution in 0.1 M phosphate buffer (PB) (pH 7.4); Invitrogen Corp., Eugene, OR, USA) was injected using a 5 μ l Hamilton microsyringe attached to a specially designed injection device. All dextran injections into SC were identified by stereotaxic coordinates obtained from mouse brain atlas [16]. Animals were then allowed to recover and survive 48 hr before perfusion. The National Institute of Health guidelines for the use and care of animals were followed for all experimental procedures. All efforts were made to minimize animal suffering as well as the number of animals used.

Perfusion and tissue processing

Mice were deeply anesthetized with a mixture of ketamine hydrochloride (30–40 mg/kg) and xylazine (3–6 mg/kg) before perfusion. In each experiment, the eyes were quickly enucleated after a reference point was taken to label the superior pole, and they were immersed in 0.1 M PB (pH 7.4). Proparacaine HCl (100–200 μ l) was applied to the cornea to suppress blink reflexes. All animals were transcardially perfused with 4% paraformaldehyde and 0.3–0.5% glutaraldehyde in 0.1 M PB (pH 7.4) containing 0.002% added calcium chloride. Following a pre-rinse with approximately 30 ml of phosphate-buffered saline (PBS, pH 7.2) over a period of 1–3 min, each mouse was perfused with 20–30 ml of fixation solution for 5–10 min via a syringe needle inserted through the left ventricle and aorta. The head was then removed and placed in the fixation

Fig. 1. Single cell injection into parvalbumin (PV) and dextran co-localized neurons on the whole-mounted retina. (A) PV-containing neurons were labeled with a monoclonal antibody against PV. (B) Retrogradely injected tracer dextran-labeled neurons in the retina. (C, D) A DiI-filled micropipette targeted double-labeled neurons. (E–H) A DiI-Filled neuron revealed the distinctive morphological pattern. Note the axon of targeted cell in (E). FITC-labeled PV-containing neurons (A–D) were viewed with a water-immersion lens, using a 100 W mercury source and Zeiss filter set 09. The DiI-filled cells (E–H) were viewed using another filter set 20.

solution for 2–3 hr. The brain was removed from the skull, stored for 2–3 hr in the same fixation solution, and left overnight in 0.1 M PB (pH 7.4) containing 8% sucrose and 0.002% calcium chloride. The SC was removed, mounted onto a chuck, and cut into 50 μ m thick sections with a Vibratome. The tissues were coverslipped with Vectashield mounting medium (Vector Laboratories, Burlingame, CA, USA).

Immunocytochemistry

After enucleation, the retinas were carefully isolated from the eye cup in 0.1 M PB (pH 7.4) and mounted onto a black, nonfluorescent filter membrane (0.45 mm, Black, HABP 47 mm; Millipore, Bedford, MA, USA) with the ganglion cell layer upper. The filter membrane and the attached retina were fixed for 30 min in a dish containing 4% paraformaldehyde in 0.1 M PB (pH 7.4). After 3 rinses in 0.1 M PB, we used conventional immunocytochemical techniques that we previously described in detail [18]. Briefly, the tissues were labeled with monoclonal antibodies against PV (1:100–200, Sigma-Aldrich, Temecula, CA, USA), in 0.1 M PB for 2 hr. The secondary antibody used was FITC-conjugated horse anti-mouse IgG (1:50; Vector Laboratories) in 0.1 M PB for 1 to 2 hr.

Cell injection

The fundamental procedure for single-cell injection following immunocytochemistry on whole mount retina has been previously described in detail [19, 20, 22] and therefore will be presented in a brief outlined. The FITC- and dextran-labeled neurons were viewed with a microscope equipped with a Zeiss 40X Plan Achromplan (NA 0.80) water-immersion lens and Zeiss filter set 09 (excitation, 450–490 nm; emission, 515 nm). The co-localized cell was filled iontophoretically by passing a positive current of 5–20 nA and 0.2–0.5 mV with 1% lipophilic dye, DiI (1% DiI in ethanol; Molecular Probes, Eugene, OR, USA) [19]. The optimal filling time was 15–30 min (Fig. 1). When the selected cell was filled, the grid reticle was moved laterally and another cell was injected into the near center of the grid. After several cells in the retina were filled, the tissue was rinsed in 0.1 M PB (pH 7.4), and fixed for 2 hr in 4% paraformaldehyde. To counterstain the tissue, the retina was immersed for 1 hr in a nucleic acid dye SYTO13 (10–20 nm; Molecular Probes). The tissue was coverslipped with Vectashield mounting medium (Vector Laboratories).

Data analysis

The cells were imaged using a confocal scanning module (LSM 700, Carl Zeiss Meditec Incorporation, Jena, Germany) mounted on a fluorescence microscope (Axio Observer.Z1, Carl Zeiss) equipped with a C-Apochromat

Fig. 2. Determining the level of dendritic stratification of PV-filled retinal ganglion cells (RGCs). Through-focus series of DiI-injected cell (left column), and SYTO 13 (right column) were matched. INL, inner nuclear layer; GCL, ganglion cell layer. Bar=50 μ m.

40×/1.20w Korr M27 for small cells and an Apochromat 20×/1.20w Korr M27 for large cells. Two sets of images were collected of each injected cell. A through-focus of the DiI-filled cells was imaged using an emission filter (580DF32; Chroma Technology Corp., Rockingham, VT, USA). Subsequently, a through-focus of the nucleic acid-stained (SYTO13; Molecular Probes) nuclei was collected using an emission filter (515DF30), to measure the thickness of the inner plexiform layer (IPL) and to determine the level of stratification of the DiI-filled cells. The ganglion cell layer (GCL)/IPL boundary was defined as 100% and the IPL/inner nuclear layer (INL) boundary was defined as 0%. The z-coordinate of each dendritic branch was measured from the distance between GCL/IPL and the IPL/INL boundaries (Fig. 2). The dendritic field area and diameter were computed with a ZEN 2009 program (Carl Zeiss Microimaging GmbH 2006–2009) by connecting the distal most tips of the dendrites [33]. Images were adjusted in terms of brightness and contrast (Photoshop CS; Adobe Systems, Mountain View, CA, USA).

III. Results

Retrograde tracer injection into SC

In the present study, 25 mice were used to precisely detect RGCs which projected to the SC. In most cases, retrograde tracer (dextran, tetramethylrhodamine, 3000 MW) injections were applied to visual layers of the SC (Fig. 3A). However, some tracers diffused into other layers of the SC where retinal fibers have been known to terminate relatively sparsely. Forty-eight hours after these injections, retrogradely labeled cells were obviously identified as retinotectal RGCs in mouse retinas. Following experimental procedures, retrogradely labeled RGCs and PV-containing RGCs in isolated retinas were confirmed by a microscope equipped with a water-immersion lens (Fig. 3B and 3C). In present study, we also found many PV-IR RGCs were not labeled by tracer in the area where tracer filled many PV-IR RGCs. Therefore, Fig. 3D (arrows) shows double-labeled cells of the retinotectal pathway. These cells were applied to single-cell injection of lipophilic dye.

Topography of RGCs that project to SC

The location and extent of the dextran labeling in the retina differed in most cases. These are caused by the differences in location and extent of the injections in the SC. Dextran-labeled RGCs located at intermediate retinal positions projected to corresponding intermediate positions along the cranial-caudal axis of the SC. Thus, RGCs in the nasal retina projected to targets at the far caudal portion of the SC, and RGCs in the temporal retina project to targets at the far rostral end of the SC [4, 13, 23]. After injecting dextran into the rostral SC on the right side, the retinal portions of retrogradely labeled cells were seen to be located in the temporal area (Fig. 4A). In all cases, the labeled cells were irregularly spaced within the retinal region of labeled cells, and a wide range of ganglion cells projected to the SC (Fig. 4). Our finding in mouse retina was similar to the results of studies in rodent species [27, 31, 43].

Types of PV-containing RGCs that project to SC

Our previous study has revealed that 28.98% (13,003/44,587) of RGCs in the mouse express PV [20]. In the present study, 108 RGCs from 25 retinas were analyzed by a laser scanning confocal microscope. Ganglion cells were identified by the presence of an axon after DiI injection. For simplicity and to avoid controversy regarding ganglion cell types or clusters within the mouse and between species [9, 21, 32, 39], we did not rely on pre-existing names. Instead, PV-containing RGCs with their retinotectal pathway were classified by dendritic field size, branching pattern, and stratification within IPL. We also used a simple name, as in the previous study, PV cell types with their retinotectal pathways 1 through 8 (PV1-8; Table 1). However, each cell type corresponds to the recent classification of Sun *et al.* [39]. A description of cells follows the order of their dendritic field diameter.

PV1 cells possessed a small dendritic field (Fig. 5), with dendrites exhibiting abrupt bends and turns along their course. The field diameter of the cells was $166 \pm 33 \mu\text{m}$. The dendritic arbors branched in the outer part of the IPL ($33.68 \pm 2.33\%$, Fig. 8). As previously reported [20], the

Fig. 3. Retrograde fluorescent tracer dextran injection into the mouse superior colliculus (SC). (A) Injection site of the SC. (B) PV-immunoreactive (IR) cells in the ganglion cell layer of the retina. (C) Dextran containing cells in the same focus as (B). (D) Merged image of (B) and (C). Some cells (arrowheads) are clearly co-localized by anti-parvalbumin antibody and dextran in the mouse retina. SGL, superficial gray layer; OL, optic layer; IGL, intermediate gray layer. Bars=100 μm (A), 50 μm (D).

Fig. 4. (A) Distribution of retrogradely labeled ganglion cells after injections into the SC. High magnification of dextran labeled cells in the whole mount mouse retina: (a) peripheral region, (b)–(c) mid central regions and (d) central region. Bars=1 mm (A), 200 μ m (a–d).

dendritic arbors contained many short branches and were often asymmetric. It was obvious that PV1 cells belong to type B4 (Table 1).

PV2 cells possessed a sparse dendritic field that was curvy and recursive (Fig. 5). PV2 cells came in ON and OFF varieties. The field diameter of the ON variety was 227 ± 35 μ m, whereas the dendritic field of the OFF variety was 191 ± 33 μ m. These cells were stratified in the inner IPL ($64.74 \pm 5.25\%$) or the outer IPL ($30.35 \pm 4.95\%$). As previously reported [20], the dendritic branches were moderately wavy and were mostly longer than those of PV1. PV2 cells appeared to belong to type B3 [39].

PV3 cells possessed medium and medium-dense dendritic fields (Fig. 6). These cells were bistratified in

the IPL with at a depth of $57.21 \pm 8.29\%$ and $29.01 \pm 8.54\%$ with a dendritic field diameter of 193 ± 37 μ m. They comprised 21.3% of our sample. PV3 cells had thin, recursive, loop-forming dendrites. They were very similar to the ON-OFF direction selective cells that have been extensively described in the rabbit [49] and mouse retinas [47]. PV3 cells appeared to belong to type D2 [39].

PV4 cells possessed medium-to-large and medium-dense dendritic fields (Fig. 6). The field diameter of the cells was 226 ± 18 μ m, and they stratifying in the outer IPL ($34.57 \pm 6.47\%$). They had 3–5 primary dendrites that tapered from the soma to the periphery. The dendritic arbors covered the dendritic field evenly and were often asymmetric. PV4 cells corresponded to type C5 [39].

Table 1. Quantitative data of parvalbumin-containing mouse RGC in this study and their homologues from previous data

Cell Type	Dendritic Field Position (% of IPL Depth; Mean $\pm$ SD)	Number in Sample	% of Total Ganglion Cell Filled	Dendritic Field Diameter (μ m; Mean $\pm$ SD)	Dendritic Field Area (μ m ²)	Sun <i>et al.</i> [39]		
						Type	% of Total RGCs	Classification Scheme (Dendritic Field Size/density)
PV 1	33.68 ± 2.33	5	4.6	166 ± 33	19707	B4	5.7	Small/dense
PV 2*	64.74 ± 5.25	9	19.4	227 ± 35	37635	B3	10.6	Small to medium/sparse
	30.35 ± 4.95	12		191 ± 33	25306			
PV 3†	57.21 ± 8.29	23	21.3	193 ± 37	27456	D2	15.5	Medium/medium
	29.01 ± 8.54							
PV 4	34.57 ± 6.47	12	11.1	226 ± 18	38838	C5	11.2	Medium/medium
PV 5	28.93 ± 4.57	15	13.8	219 ± 46	35198	C4	5.9	Medium/dense
PV 6*	65.11 ± 6.10	7	11.1	273 ± 30	51381	C2	10.2	Medium to large/sparse
	35.07 ± 4.33	5		257 ± 22	47762			
PV 7	57.02 ± 1.43	5	4.6	248 ± 35	48186	C1	3.2	Medium to large/medium
PV 8	65.5 ± 4.62	15	13.8	297 ± 52	67007	A1	4.5	Large/sparse
Total		108	100					

* Monostratified cell type with ON and OFF population.

† Bistratified cell type.

Fig. 5. PV-IR RGCs from the mouse SC. The PV1 and PV2 types of PV-containing retinotectal ganglion cells were identified in the present study. Each type corresponded to Sun *et al.*'s classification [39]. Bar=50 μ m.

Fig. 6. PV-IR RGCs from the mouse SC. The PV3 to 5 types of PV-containing retinotectal ganglion cells were identified in the present study. Each type corresponded to Sun *et al.*'s classification [39]. Bar=50 μ m.

Fig. 7. PV-IR RGCs from the mouse SC. The PV6 to 8 types of PV-containing retinotectal ganglion cells were identified in the present study. Each type corresponded to Sun *et al.*'s classification [39]. Bar=50 μ m.

PV5 possessed medium-to large, relatively dense dendritic fields (Fig. 6). The field diameter of the cells was 219 ± 46 μ m, and they stratified in the outer IPL ($28.93 \pm 4.57\%$). As previously reported [20], these cells had 3–6 primary dendrites that were generally smooth, with compact and densely branched dendritic fields. The dendritic arbors overlapped in many fields of the cells. PV5 cells appeared to belong to type C4 [39].

PV6 possessed large and medium-sparse dendritic fields (Fig. 7). PV6 came in ON and OFF varieties. The field diameter of the ON variety was 273 ± 30 μ m, whereas the dendritic field of the OFF variety was 257 ± 22 μ m. These cells were stratified in the inner IPL ($65.11 \pm 6.10\%$) or outer IPL ($35.07 \pm 4.33\%$). As previously reported [20], these cells were characterized by irregular-looking dendritic arbors, which lacked an orderly, radiating pattern. PV6 cells corresponded to type C2 [39].

PV7 cells possessed large and medium-dense dendritic arbors (Fig. 7). The dendritic field diameter of the cells was 248 ± 35 μ m. The dendritic arbor branched in the inner part of the IPL ($57.02 \pm 1.43\%$). These cells had 3–5 primary dendrites, and the dendritic processes rarely crossed one another. They were very similar to the ON-direction selective cells that have been extensively described in mouse retinas [40]. PV7 cells corresponded to type C1 [39].

PV8 cells possessed large, sparse dendritic arbors (Fig. 7). They stratified at $65.5 \pm 4.62\%$ of the IPL, with a

dendritic field diameter of 297 ± 52 μ m. These cells had 3–6 primary dendrites, from which the process extended radially from a soma, emitting several long branches along the way. PV8 cells were similar to the ON variety of alpha cells of most mammalian retinas [26], and appeared to belong to type A1 [39].

IV. Discussion

In the present study, we identified PV-IR retinotectal cells in the mouse retina by using a triple-labeling technique. The newly advanced method is especially useful for matching a neuron's morphology with its protein content and neural pathway.

The recent evidence strongly indicates that the family members of each major RGC type in the retina have all been identified. It is clear that mammalian retinas contain 10–15 different RGC types [28]. Sun *et al.* [39] classified at least 14 RGC types in the mouse. Among these 14 types, our previous study [20] has shown that at least 8 different types expressed PV. These types were A1, B3, B4, C1, C2, C4, C5 and D2 [39]. The present study indicated that all these 8 types of PV-IR RGCs project to the mouse SC. These cells have a small to large dendritic field size. We also have shown that RGCs retrogradely labeled with dextran were found in an orderly retinotopic fashion relative to the SC [4, 13, 23]. Studies in PV-IR RGCs of the

Fig. 8. The branching and levels of stratification of the 8 types of PV-IR mouse RGCs identified in our study. Bar=50 μ m.

various mammalian retinas suggest that PV-IR RGCs, if not all RGCs, most likely belong to a different chemical subpopulation for each type [7, 36].

A series of conceptual and experimental advances have made it possible to identify the neural circuits of retinotectal neurons and the connection of RGCs in the retina to their terminations in the SC of midbrain [1, 4, 10, 13, 23, 27, 31]. In general, retinotectal RGCs as non-standard cells have direction selectivity, local edge detector and suppressed-by-contrast receptive field (W type and Y type). They do not include standard cells underlying high resolution vision (X type). Non-retinotectal RGCs have standard concentric receptive fields [6, 29, 41, 42]. Waleszczyk *et al.* [45] has presented a simplified schema of the cat visual system including the retinotectal pathway. All types, except for beta (X)-type ganglion cells, project to the SC. Thus, W type RGCs project to the most superficial cellular layer of the SC, the stratum zonale and the stratum griseum superficiale. Y type RGCs project to the lower portion of the stratum griseum superficiale, and the upper portion of the stratum griseum intermediale [4, 17, 24, 38]. Moreover, distribution of retinotectal RGCs was also revealed in the rabbit [43], rat [1] and ground squirrel [27, 31] by using retrograde staining. Especially, the ground squirrel RGCs were identified in 4 types of retinotectal neurons, including the beta-type ganglion cell (X type). They have a variety of morphological types of RGCs projected to the SC. The present results of retrograde staining showed a variety of morphological types of RGCs in the mouse retina. Although the physiological functions of mouse RGCs types are still not fully understood, our present study provides evidence that PV-IR retinotectal RGCs in the mouse may possibly have W type and Y type RGCs. Sun *et al.* [39] assumed that A1 (PV8) type RGCs are equivalent to the alpha cell type (Y type). Our small PV-IR retinotectal RGCs (PV1 and PV2) could possibly belong to the W type RGCs, but confirmation requires further electrophysiological and ethological studies. However, PV-IR retinotectal RGCs do not have B2 type RGCs, which were assumed morphologically to be the beta cell type (X type). Thus, our results for PV-IR RGCs correspond

with the previous results in most animals.

PV is thought to play a major role in buffering the intercellular calcium levels and a neuroprotective role from calcium overload. Moreover, PV-IR neurons in the brain can possibly modulate the pathway of principal neurons by controlling excitation and inhibition [14, 35]. It may be possible that the PV-IR RGCs especially contribute to neuroprotection and modulation of neural pathway activity. Our results indicated that retinotectal PV-IR RGCs were monostратified to bistratified, possessed a small-to-large dendritic field size, and had sparse-to-dense dendritic arbors. It is conceivable that PV is expressed in different types of RGCs to differentiate their special functions in specific cell types.

Specifically, it is conceivable that PV may be involved in low spatial resolution pattern vision and controlling the reflex direction of the gaze (W) as well as nonlinear spatial summation within the receptive field (Y) [34, 41]. These results may assume that PV in small PV-IR retinotectal RGCs (PV1 and PV2) may contribute to interactions involved in local edge detection, and the PV in large PV-IR retinotectal RGCs (A1) may contribute to interactions of suppression by contrast. Similarly, Y type RGCs have a large dendritic field and they also transmit signals of nonlinear spatial component within a receptive field to the SC in the cat [4] and primate [10]. Crook *et al.* [10] has shown that Y type RGCs (M) have collicular projections and have linear and nonlinear components of the receptive field. These cells conceivably corresponded to type PV8 in our results. However, the precise function of PV in each cell type needs more detailed studies and remains far from obvious.

Retinotectal RGCs also have a directionally selective response. Especially, On-direction selective ganglion cells and On-Off direction selective ganglion cells project to the lateral geniculate nucleus as well as the SC [25, 50]. On direction selective RGCs and On-Off direction selective RGCs corresponded to type PV7 and PV3, respectively in our result. Although the function of PV in direction selective cells remains far from obvious, the presence of PV in these cells may suggest that PV play a certain role in

modulating signals of direction selectivity. These assumptions also require further study.

In conclusion, the combined newly approach of cell morphology and the selective expression of PV within the neural pathway will provide useful data for further correlation of specific neural circuits of RGCs, and also provide a powerful strategy for matching a neuron's morphology with its expression of a specific protein. In the present study, our results have indicated that PV-IR retinotectal RGCs have all 8 types PV-IR RGCs that were heterogeneous in morphology. PV-IR retinotectal RGCs may have functions similar to those of non standard RGCs, which interact with direction selectivity, local edge detection and suppressed-by-contrast. It is possible that PV-IR neurons have particular functional properties for modulating signals of the neural pathway for a specific niche adaptation. The combined method of triple labeling technique will lead to a better insight of the specific neural circuitry of RGCs.

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