

Identification of a Pathway from the Retina to Koniocellular Layer K1 in the Lateral Geniculate Nucleus of Marmoset

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Three well characterized pathways in primate vision (midget-parvocellular, parasol-magnocellular, bistratified-koniocellular) have been traced from the first synapse in the retina, through the visual thalamus (lateral geniculate nucleus, LGN), to the visual cortex. Here we identify a pathway from the first synapse in the retina to koniocellular layer K1 in marmoset monkeys (*Callithrix jacchus*). Particle-mediated gene transfer of an expression plasmid for the postsynaptic density 95–green fluorescent protein (PSD95-GFP) was used to label excitatory synapses on retinal ganglion cells and combined with immunofluorescence to identify the presynaptic bipolar cells. We found that axon terminals of one type of diffuse bipolar cell (DB6) provide dominant synaptic input to the dendrites of narrow thorny ganglion cells. Retrograde tracer injections into the LGN and photofilling of retinal ganglion cells showed that narrow thorny cells were preferentially labeled when koniocellular layer K1 was targeted. Layer K1 contains cells with high sensitivity for rapid movement, and layer K1 sends projections to association visual areas as well as to primary visual cortex. We hypothesize that the DB6–narrow thorny–koniocellular pathway contributes to residual visual functions (“blindsight”) that survive injury to primary visual cortex in adult or early life.

Key words: bipolar cells; ganglion cells; koniocellular; parallel pathways; primate retina

Introduction

Parallel visual pathways are established in the retina and remain segregated within the dorsal lateral geniculate nucleus (LGN) of the thalamus (Lee et al., 2010). Midget ganglion cells send their axons to the parvocellular (P) and parasol cells send their axons to the magnocellular (M) layers of the LGN (Leventhal et al., 1981). The P pathway serves high-acuity spatial vision and red–green color vision, feeding into the form recognition (“what”) pathway. By contrast, signals traveling on the M pathway serve motion and spatial analysis (the “where” pathway). The P and M pathways are established at the first synapse in the visual pathway. In the P pathway cones contact midget bipolar cells which in turn contact midget ganglion cells (Kolb and DeKorver, 1991; Calkins et al., 1994); in the M pathway, diffuse bipolar (DB) cells receive input from multiple cones and provide output to parasol cells (Boycott and Wässle, 1991;

Jacoby and Marshak, 2000; Puthussery et al., 2013). A third retino-thalamic pathway involves the koniocellular (K) layers of the LGN (Hendry and Reid, 2000; Casagrande and Xu, 2003). Projections to K layers include ganglion cells called small bistratified cells, which display blue ON/yellow OFF responses (Dacey and Lee, 1994; Martin et al., 1997; Szmajda et al., 2008).

In addition to these three well characterized pathways, at least 12 low-density ganglion cell types that project to the LGN have been identified (Dacey et al., 2003; Crook et al., 2008; Szmajda et al., 2008). These ganglion cell types are difficult to target, and their connectivity within the retina and exact projection within the LGN are not known. Here we trace a pathway from cone photoreceptors through bipolar and ganglion cells to the ventral-most koniocellular layer K1 in the LGN.

Materials and Methods

Animals. For ganglion cell labeling by particle-mediated gene transfer two female adult marmosets (*Callithrix jacchus*) were obtained from CLEA Japan and maintained in the National Institute for Basic Biology Bioresource Center. Procedures adhered to the National Institute for Physiological Sciences (NIPS) code of practice for care and use of animals and were approved by the institutional ethics committee of NIPS (No. 11A172) and the National Institute of Natural Sciences (Nos. 08B005, 10B001).

Tracer injections were performed on five (three male, two female) adult marmosets obtained from the Australian National Health and Medical Research Council (NHMRC) combined breeding facility. Procedures were performed according to NHMRC guidelines for care and use of animals and were approved by the institutional animal ethics committee.

Labeling of ganglion cells by particle-mediated gene transfer. Animals were anesthetized with medetomidine (0.01 mg/kg, i.m.; Domitor

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Orion) and ketamine (10 mg/kg; Ketalar, Daiichi Sankyo Espha) then overdosed with sodium pentobarbital (80–150 mg/kg; Somnopentyl, Kyoritsu). Organotypic culturing and transfection of retinas was performed as described previously (Koizumi et al., 2007; Jakobs et al., 2008; Moritoh et al., 2013). In brief, quadrants of retina were mounted ganglion cell side up onto a 0.4 μm Millicell tissue culture insert (Millipore), and then transfected via particle-mediated transfer using CMV-PSD95-GFP fusion plasmid coated gold microcarriers. The plasmid was a gift from Dr Masaki Fukata (NIPS). The ratio of DNA to the gold microcarriers (1.6 μm) was 1.5 μg plasmid/1 mg gold. The retina was kept in culture for 3 d.

Tracer injections, photofilling, and LGN reconstruction. Tracer injections followed by photofilling were performed according to methods developed by Dacey et al. (2003). The retinas were those used in Szmajda et al. (2008) and Percival et al. (2009, 2011, 2013) where detailed procedures are described. The focus of these previous studies was on ganglion cells projecting to layer K3. To study the injection sites in the LGN, the animal was perfused with 0.9% saline followed by 4% paraformaldehyde (PFA) in 0.1 M PB and 10% glycerol in 4% PFA in 0.1 M phosphate buffer (PB). Serial coronal sections were cut through the thalamus using a freezing microtome. Alternate sections were mounted directly in Mowiol or processed with cresyl violet. Injection sites were reconstructed by outlining the LGN layers and tracing the dense core of the injection site over serial sections.

Immunohistochemistry of gene-gunned retinas and image acquisition. Retinas were fixed in 4% PFA in PB for 60 min at 4°C, and processed for immunofluorescence as described previously (Percival et al., 2011). We used antibodies against CD15 (mouse monoclonal; 1:6; gift from Prof Mai, University of Düsseldorf, Germany) and against the C-terminal binding protein (CtBP2, mouse monoclonal, 1:10,000; Catalog #612044; BD Biosciences). Secondary antibodies were obtained from Jackson ImmunoResearch.

Images of PSD95-GFP cells were acquired on a confocal microscope (Zeiss LSM 700) equipped with 488, 555, and 639 nm lasers using Plan-Apochromat, 20 \times /0.8 objective (2048 \times 2048 pixels). Images of retrogradely labeled cells were obtained on a Zeiss Axioplan 2 microscope equipped with a digital camera (Axiocam, Zeiss) using a 40 \times (Plan-Neofluar, 40 \times /1.3 oil differential interference contrast, DIC), or 63 \times (Plan-Apochromat, 63 \times /1.4 Oil DIC) objective. Micrographs of the LGN were acquired as described previously (Szmajda et al., 2008; Percival et al., 2013). Brightness and contrast of digital images were adjusted using Axiovision (Zeiss), Zen (Zeiss), Adobe Photoshop CS4 (v11.0.2), or Imaris (Bitplane) software.

Analysis. Cells were traced in three dimensions from image stacks (Imaris). The number of PSD95-GFP puncta located in a volume immediately surrounding the bipolar axons, and the proportion of the puncta located within a distance of 1 μm from the surface area of the bipolar axons, were determined (see Fig. 2I–K). As control, one image channel was flipped around the y -axis and the proportion of puncta close to the surface was determined in this pseudorandom fashion. Voxel colocalization (x, y : 0.156 μm ; z : 1.065 μm) was determined using Imaris.

Results

DB6 cells provide input to narrow thorny ganglion cells

The focus of our study is on a population of ganglion cells named narrow thorny cells (Dacey et al., 2003; Yamada et al., 2005). Narrow thorny cells have fine, densely branching dendritic trees decorated with numerous dendritic spines. They resemble the G16 (Kolb et al., 1992) or thorny monostratified ganglion cells (Peterson and Dacey, 1999) in human retina. Based on their stratification, narrow thorny cells are likely to receive input via the DB6 cell (Fig. 1A). In marmoset retina, DB6 cells are immunoreactive to antibodies against CD15 (Chan et al., 2001) and immunoelectron microscopy reveals that DB6 axons make synapses with ganglion and amacrine cells (Fig. 1B; Jusuf et al., 2004). The DB6 axon terminals tile the retina in a “loose” mosaic (Fig. 1C). We analyzed the synaptic output of CD15 immunoreactive DB6 cells in preparations containing gene-gunned cells. The bright fluorescent puncta in gene-gunned cells mark excitatory synaptic input (Fig. 2A)

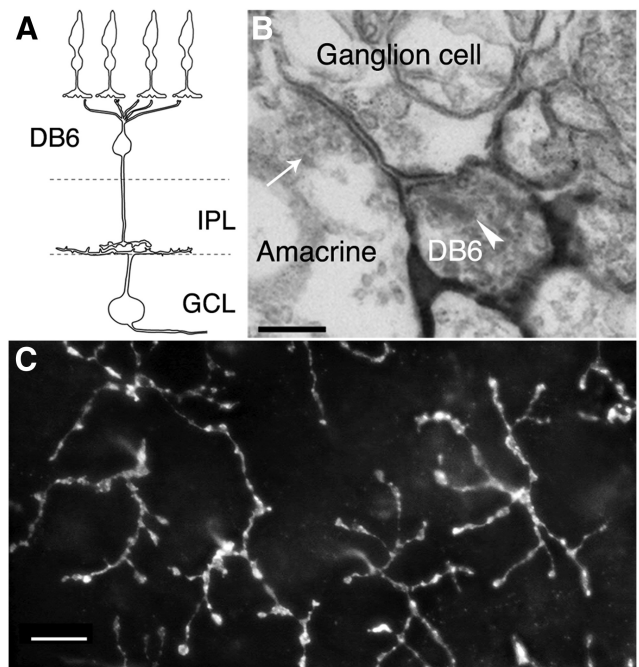


Figure 1. DB6 cells in marmoset retina. **A**, Stratification of DB6 cells in the inner plexiform layer (IPL). GCL, Ganglion cell layer. **B**, Electron micrograph of a CD15 immunoreactive DB6 axon terminal presynaptic to an amacrine (AC) and a ganglion cell (GC) process. Immunoreactivity is located in the membrane of the cell. Arrowhead indicates synaptic ribbon, arrow points to a conventional synapse from an AC profile onto a GC. **C**, Fluorescence micrograph showing axon terminals of CD15 immunoreactive DB6 axons in whole mount. Scale bars: **B**, 0.2 μm ; **C**, 10 μm .

and fainter background labeling arises from PSD95-GFP in transit (Jakobs et al., 2008; Morgan et al., 2008). We confirmed that the PSD95-GFP puncta are the sites of bipolar input by labeling a gene-gunned retinal quadrant with antibodies against CtBP2, which marks bipolar ribbon synapses (tom Dieck et al., 2005). The majority of PSD95-GFP puncta (average $56.9 \pm 39.8\%$, $n = 35$ regions from 13 ganglion cells) was located within 1 μm of a CtBP2 punctum (Fig. 2L–N). Pseudorandom colocalization reduced this percentage to $39.5 \pm 29.7\%$ ($p < 0.02$, Wilcoxon rank sum test). Consistent findings were obtained previously in other species (Jakobs et al., 2008; Morgan et al., 2008; Bleckert et al., 2013).

We analyzed the DB6 inputs to five narrow thorny cells located at ~ 6 mm distance from the fovea. These cells had dendritic field diameters of 245–340 μm and encompassed 24–31 DB6 axon terminals. All DB6 axon terminals showed regions of close association with the dendritic tree of narrow thorny cells (Fig. 2B) and DB6 axons and ganglion cell dendrites were seen to run together (Fig. 2C–H). For three cells we used neighborhood analysis to find local intensity maxima (“spot” detection) and counted the PSD95-GFP puncta located within 1 μm of a DB6 axon terminal (Fig. 2I–K). On average, these PSD95-GFP puncta made up 36.8% (SD 19.8, $n = 25$) of the PSD95-GFP puncta in the immediate vicinity of an individual axon terminal (Table 1). For two further cells we determined the colocalized pixels between the PSD-GFP and the CD15 immunofluorescence channel (Fig. 2E). We then used spot detection on the colocalization channel and counted colocalized spots relative to all PSD-GFP spots in the immediate vicinity of the DB6 axon terminal. We found that the colocalized spots made up an average 43.3% ($\pm 18.8\%$ SD) of the PSD-GFP spots in this region ($n = 14$ axon terminals). On average there were 21 (± 18.1 SD) colocalized areas per axon terminal ($n = 30$).

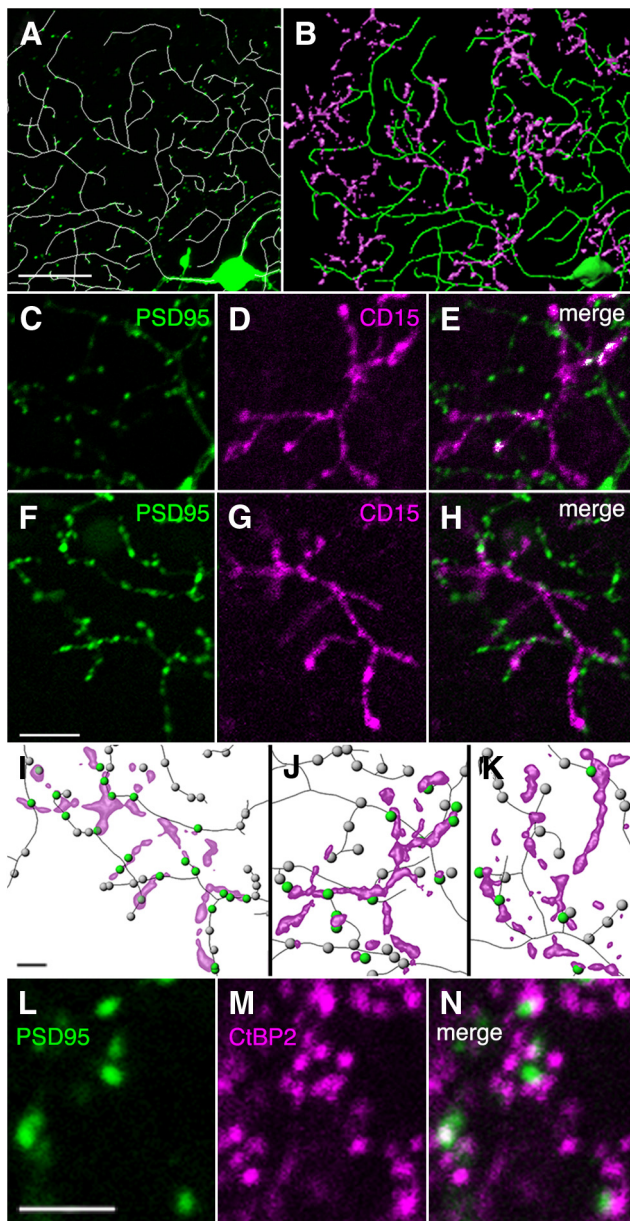


Figure 2. Narrow thorny ganglion cells are connected to DB6 cells. **A**, PSD95-GFP puncta on dendrites (white lines) of a narrow thorny ganglion cell. **B**, Partial reconstruction of this cell together with DB6 axon terminals. **C–H**, High-resolution images of CD15 labeled DB6 axon terminals (magenta) together with dendritic tree of narrow thorny cells. Single slices of a stack of confocal images. Synaptic zones labeled by PSD95-GFP appear as green puncta. **E, H**, Superimpositions of the two channels, revealing the close association between ganglion cell dendrites and DB6 axons. **E**, Colocalized pixels are white. **I–K**, Reconstructed DB6 axons (magenta) associated with dendrites of narrow thorny cells (black lines). PSD95-GFP puncta located in a volume surrounding an axon terminal are represented as green or gray dots (not to scale). Green dots represent PSD95-GFP puncta that are located within $1\ \mu\text{m}$ of the dendrites. **L–N**, PSD95-GFP (postsynaptic, green) puncta of a ganglion cell dendrite are located opposite presynaptic ribbons (magenta) in gene-gunned tissue processed with antibodies against ribbon marker CtBP2. Scale bars: **A**, $30\ \mu\text{m}$; **F**, $10\ \mu\text{m}$; **I**, $5\ \mu\text{m}$; **L**, $3\ \mu\text{m}$.

For one narrow thorny cell we measured 12 DB6 axon terminals using both methods; across axon terminals spot counts showed high correlation (correlation coefficient 0.716, $p < 0.02$) indicating both methods give comparable results. For all cells analyzed, the number of colocalized spots was lower when one channel was flipped ($p < 0.02$, Wilcoxon rank sum test).

We compared the connectivity in the DB6 narrow thorny pathway with the well established connectivity in the midget

Table 1. Analysis of bipolar to ganglion cell connectivity

	No. 1	No. 2	No. 3
Narrow thorny ganglion cells			
DB6 axons analyzed	12	9	4
Percentage of PSD95-GFP puncta located within $1\ \mu\text{m}$ of axon terminal (mean, SD)	38, 12.6	20.9, 9.8	69.3, 10.9
Midget ganglion cells			
Midget bipolar axons analyzed	3	5	1
Percentage of PSD95-GFP puncta located within $1\ \mu\text{m}$ of axon terminal (mean, SD)	51.7, 14.7	43.7, 6.0	56.8

Well labeled bipolar axons that overlapped completely with the dendritic field of the ganglion cell were reconstructed; PSD95-GFP puncta in the immediate vicinity of an axon terminal were detected and the percentage of the puncta located $\leq 1\ \mu\text{m}$ of the ganglion cell dendrite was determined.

pathway by taking advantage of the fact that OFF midget bipolar cells are labeled in addition to DB6 cells with the antibody against CD15 (Chan et al., 2001). For three midget ganglion cells in peripheral retina, the PSD95-GFP puncta located within $1\ \mu\text{m}$ of an OFF midget bipolar axon terminal on average made up 47.8% (SD 9.9, $n = 9$) of the PSD95-GFP puncta in the immediate vicinity of this axon terminal (Table 1).

Identification of narrow thorny cells after retrograde labeling

Tracer injections into the LGN resulted in retrograde labeling of a variety of ganglion cells. Narrow thorny cells (Fig. 3A–C) identified at eccentricities between 2.5 and 5 mm had dendritic field diameters between 100 and $150\ \mu\text{m}$, which is similar to that of parasol cells (Ghosh et al., 1996). Indeed, we initially classified some of the narrow thorny cells as parasol cells. Closer analysis revealed (Fig. 3D, I) that narrow thorny cells, have a higher average dendritic density (dendritic length in $\mu\text{m}/\text{mm}^2$; narrow thorny $424.4 \pm 70.6\ \mu\text{m}$, $n = 9$; parasol $273.5 \pm 70.2\ \mu\text{m}$, $n = 7$, $p < 0.02$, t test), and a higher number of dendritic terminals (including thorns) per $100\ \mu\text{m}^2$ compared with parasol cells (narrow thorny, 3.7 ± 1.4 , $n = 9$; parasol 1.2 ± 0.5 , $n = 7$; $p < 0.02$, t test). The majority (6/9, 67%) of the narrow thorny cells stratified in the ON sublamina of the inner plexiform layer, two cells stratified in the OFF sublamina, and one cell was bistratified with processes in the ON and the OFF sublamina.

Depending on the site of the tracer injection different types of ganglion cells were retrogradely labeled (Szmajda et al., 2008). Parasol cells dominated when the injection site was centered on the magnocellular layers, whereas narrow thorny cells dominated when the injection was centered on layer K1 (Fig. 3E–H). In two additional experiments, where a small injection into layer K1 and a larger injection into layer K3 were made, we encountered eight parasol, eight small bistratified, two narrow thorny, and four other wide-field cells. Parasol cells were always labeled because the injecting electrode must pass through the M layers to reach layer K1. One narrow thorny cell was labeled in an experiment where the injection was centered on K3 but the electrode track included layer K1. We conclude that narrow thorny ganglion cells will be preferentially labeled, if the tracer is injected into layer K1, but projection of narrow thorny cells to other LGN layers cannot be ruled out. Likewise it appears that additional ganglion cell types project to layer K1 (Szmajda et al., 2008).

The DB6 cells contact all cones in their dendritic field with bias against short-wave sensitive cones (Lee and Grünert, 2007). Thus, narrow thorny cells stratifying in the ON sublamina are likely to display achromatic ON responses. Consistent with this prediction, when we collated results from our previous recording studies in LGN (White et al., 2001; Solomon et al., 2002; Tailby et al., 2008) we found that the great majority of receptive fields

recorded in layer K1 (18/22, 82%) show ON-type or ON-OFF-type responses ($p < 0.05$, χ^2 test). By contrast, ON- and OFF-type responses were evenly balanced in layer K3 ($p > 0.5$, χ^2 test), and blue-ON responses were less common in K1 (1/23, 4.3%) than in K3 (77/228, 33.7%, $p < 0.05$, χ^2 test).

Discussion

Narrow thorny ganglion cells have been identified morphologically in the retinas of a variety of primate species but no electrophysiological recordings from these cells exist, thus their functional role can only be inferred from their retinal circuitry and the properties of their geniculate targets in layer K1.

Our results show that a significant proportion of the bipolar input to narrow thorny cells derives from DB6 cells. We suggest that the DB6 input is the dominant bipolar input to narrow thorny cells because the connections of the other bipolar types stratifying in S4 and S5 are well understood: blue cone bipolar cells contact small bistratified cells (Calkins et al., 1998), rod bipolar cells contact amacrine cells (Grünert and Martin, 1991), and invaginating midget bipolar cells contact ON midget ganglion cells (Kolb and DeKorver, 1991; Calkins et al., 1994).

Our results further indicate that DB6 cells in addition to amacrine cells preferentially contact narrow thorny cells. Other outputs of DB6 cells go to large sparse (Percival et al., 2011, their Fig. 3D) and inner stratifying melanopsin-expressing ganglion cells (Grünert et al., 2011, their Fig. 3) but these cells had on average only 2–3 presumed contacts per DB6 terminal and thus did not show the same tight connectivity as do narrow thorny cells (21 contacts per terminal). The only other ganglion cell type stratifying at the level of the DB6 cells is the small bistratified cell and this cell is known to receive dominant input from blue cone bipolar cells (Calkins et al., 1998).

A large proportion of the PSD95-GFP puncta expressed by narrow thorny cells were located close to DB6 axons; a similar relationship was seen for OFF midget ganglion/OFF midget bipolar cells. A likely explanation for the nonassociated puncta is that synaptic output from weakly labeled DB6 axons was underestimated. Some nonassociated puncta may arise from overexpression of PSD95-GFP.

Electrophysiological recordings show diverse response properties in koniocellular pathways (Irvin et al., 1986; White et al., 2001; Xu et al., 2001; Solomon et al., 2002). Specifically, in marmoset and owl monkey limited recordings from neurons in layers K2 and K1 showed receptive fields with high contrast sensitivity, large diameter, high sensitivity to high temporal frequency flicker and negligible responses to short-wave cone stimulation (White et al., 2001; Xu et al., 2001; Solomon et al., 2002). Thus, layers K1

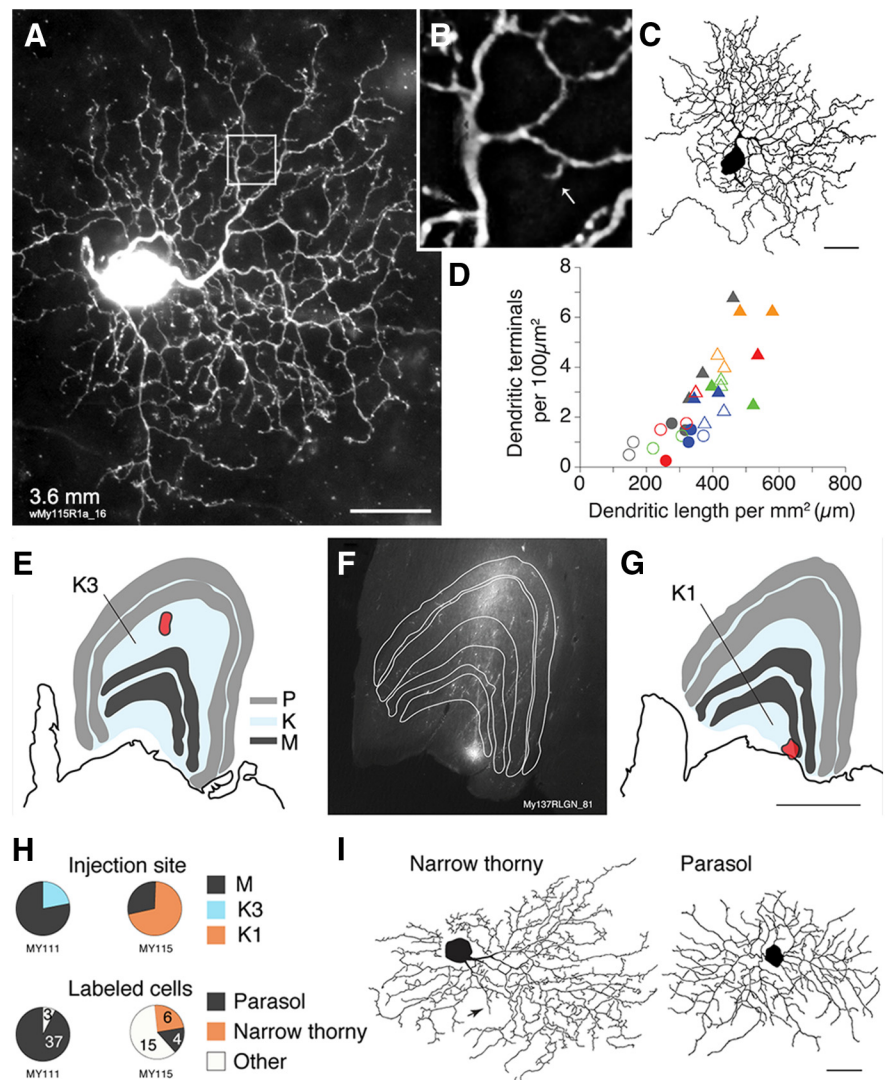


Figure 3. Narrow thorny ganglion cells labeled after injection of koniocellular layer K1. **A**, Fluorescence micrograph of a narrow thorny cell from 3.6 mm eccentricity. Box indicates the region shown in **B**. Arrow points to a dendritic thorn. **C**, Narrow thorny cell from 3.5 mm eccentricity. **D**, Scatter plot showing morphological differences (dendritic length per area, number of dendritic thorns and/or terminals per area) between narrow thorny and parasol cells. Data are based on nine narrow thorny (triangles) and seven parasol (circles) cells. Same symbols indicate different regions of dendritic tree of a specific cell. **E–G**, Marmoset LGN has two P, two M, and four K layers. **F**, Fluorescent micrograph of a coronal section, outlines of LGN are superimposed. Dense core of the injection into K1 can be seen. **E, G**, Reconstruction of the same LGN. Injection sites are indicated in red. **H**, Comparison of injection sites in LGN and number of retrogradely labeled ganglion cells for two experiments. **I**, Drawings of parasol cell from 3.31 mm (magnocellular injection) and narrow thorny cell from 4.85 mm (K1 injection). Scale bars: **A, C, I**, 20 μm ; **G**, 1 mm.

and K2 may be part of a primitive motion-processing pathway, which has been supplanted (or dominated) by the magnocellular pathway in course of primate evolution.

We conclude with the following speculation. It has been established that the K layers are distinct from P and M layers in their main projection sites within the primary visual cortex (Fitzpatrick et al., 1983; Solomon, 2002; Chatterjee and Callaway, 2003; Casagrande et al., 2007). Furthermore, axons from LGN layers K1/K2 preferentially target more superficial cortical layers whereas axons from LGN layers K3–K6 preferentially terminate closer to the granular (M and P recipient) layers (Xu et al., 2001; Casagrande et al., 2007). Layer K1 also projects directly to the motion-selective medial temporal area (MT), bypassing V1 (Sinich et al., 2004). These facts may implicate a role of the narrow thorny-koniocellular K1 pathway in sensitivity of MT receptive

fields to moving stimuli following removal of V1 input in adult (Girard et al., 1992) or following neonatal lesions to V1 (Yu et al., 2013), and more broadly in the residual visual functions (“blindsight”) that survive injury to primary visual cortex in adult or early life (Cowey, 2010).

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