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## Role of the immune modulator programmed cell death-1 during development and apoptosis of mouse retinal ganglion cells

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### Abstract

**PURPOSE**—Mammalian programmed cell death-1 (PD-1) is a membrane-associated receptor regulating the balance between T cell activation, tolerance and immunopathology, however its role in neurons has not yet been defined. We investigate the hypothesis that PD-1 signaling actively promotes retinal ganglion cell (RGC) death within the developing mouse retina.

**METHODS**—Mature retinal cell types expressing PD-1 were identified by immunofluorescence staining of vertical retina sections; developmental expression was localized by immunostaining and quantified by Western analysis. PD-1 involvement in developmental RGC survival was assessed *in vitro* using retina explants and *in vivo* using PD-1 knockout mice. PD-1 ligand gene expression was detected by RT-PCR.

**RESULTS**—PD-1 is expressed in most adult RGCs, and undergoes dynamic upregulation during the early postnatal window of retinal cell maturation and physiological programmed cell death (PCD). *In vitro* blockade of PD-1 signaling during this time selectively increases survival of RGCs. Furthermore, PD-1 deficient mice show a selective increase in RGC number in the neonatal retina at the peak of developmental RGC death. Lastly, throughout postnatal retina maturation, we find gene expression of both immune PD-1 ligand genes, PD-L1 and PD-L2.

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**CONCLUSIONS**—These findings collectively support a novel role for a PD-1-mediated signaling pathway in developmental PCD during postnatal RGC maturation.

## Introduction

Modulation of signaling elicited by cell-cell interactions is critical for the formation and remodeling of neuronal synaptic networks in development and learning, as well as for the generation of immunity towards environmental and endogenous antigens.<sup>1,2</sup> In some cases, molecules involved in such selection may be shared by both the immune and central nervous systems. For example, the MHC class I ligand and its receptor component CD3 $\xi$ , a canonical receptor-ligand immune recognition pair, are required to establish functional connections between the retina and brain during development<sup>3</sup>, and the initiating complement protein C1q marks neural retina synapses for elimination in both development and degenerative disease<sup>4</sup>.

In the immune system, cell-cell interaction molecules are critical for regulating lymphocyte function. PD-1 (CD279) is a key immunoregulatory receptor, inducibly expressed on T cells, B cells, NK T cells, activated monocytes, and dendritic cells.<sup>5</sup> PD-1 transduces an inhibitory signal when engaged in combination with the T cell receptor (TCR). These immunoinhibitory signals regulate the extent of T cell activation, attenuate anti-microbial immunity, facilitate chronic viral infections, and provide inhibitory signals that regulate both central and peripheral T cell tolerance.<sup>2</sup> During the establishment of central tolerance, PD-1 is expressed on developing thymocytes as they progress through several maturational stages, where PD-1 signaling modifies signaling thresholds in thymocytes during both positive and negative selection stages of maturation.<sup>6,7</sup> In addition, PD-1 regulates both the induction and maintenance of peripheral T cell tolerance by limiting mature self-reactive T cell function.<sup>5</sup>

We recently observed constitutive neuronal expression of PD-1 in retinal ganglion cells (RGCs)<sup>8</sup>, suggesting that PD-1 may also provide inhibitory signals important for physiologic loss of neurons during retinal maturation. In this study, we tested whether PD-1 has a parallel role in negative selection during neuronal network formation in the developing and adult retina, an organ with well-defined cytological architecture that has been an important model for investigating molecular mechanisms of neurogenesis.

## Materials and Methods

### Animals

Mice were purchased from Charles River Laboratory unless otherwise noted. Embryonic and adult CD1 mice were used for PD-1 blocking experiments. C57BL/6 mice were used for PD-1 immunoblotting and PD-1 ligand gene expression studies. For the PD-1<sup>-/-</sup> characterization, PD-1<sup>-/-</sup> mice were constructed in the C57BL/6 background, as previously described<sup>9</sup>, and wildtype C57BL/6 age-matched mice were purchased from the Charles River Laboratory. All animal experiments were reviewed and approved by the UCLA Chancellor's Animal Research Committee in adherence to the ARVO Statement for the Use of Animals in Ophthalmic and Vision Research.

### Tissue Preparation

Eyes were rapidly enucleated from adult and embryonic mice. Posterior eye cups were fixed in 4% paraformaldehyde at room temperature for one hour, followed by cryoprotection in 30% sucrose/PBS and OCT embedding. Cryostat sectioning was performed at a thickness of 6-8  $\mu$ m. The sections were selected to represent the equivalent regions of the globes and

taken just adjacent to the optic nerve plane. In order to assure validity of analysis, sections were used for quantification only if the retina architecture represented a vertical section through the retina.

### RGC Isolation

Positive selection of RGCs was performed using magnetic beads coated with a Thy-1.1 monoclonal antibody (Millipore/Chemicon), as previously described<sup>10</sup>.

### Retinal Explants

Neural retinas were dissected from newborn P0 mice and cultured as previously described<sup>11</sup>. Briefly, retinas were placed on polycarbonate filter discs (Millipore) and cultured for 24 hours at 37°C/5% CO<sub>2</sub> in DMEM/F12 (Mediatech) containing 10% fetal bovine serum (Omega Scientific), and 1 × N2 supplement (Gibco). For the antibody blocking experiment, the following functional grade antibodies were used at a final concentration of 7 µg/ml: anti-mouse PD-1 (clone J43, hamster IgG, eBioscience) for which specificity<sup>12</sup> and ability to block interaction with PD-L1 and PD-L2<sup>13</sup> have previously been described, or isotype control hamster IgG (eBioscience). After 24 hours in culture, explants were fixed, cryoprotected, embedded in OCT, and cryosectioned, as described above.

### Immunostaining

Sections were incubated at 4°C overnight with the following primary antibodies: anti-mouse PD-1 (clone 29F.1A12, rat IgG2a) for which specificity has previously been described<sup>14</sup>; RGC markers anti-Brn3a (clone 5A3.2, mouse IgG1, Millipore/Chemicon)<sup>15</sup> and anti-NeuN (mouse IgG1, clone A60, Millipore/Chemicon)<sup>16</sup>; amacrine cell marker anti-mouse AP2α (3B5, mouse IgG2b, Developmental Studies Hybridoma Bank)<sup>17</sup>; or apoptotic cell marker rabbit anti-mouse activated caspase-3 (R&D Systems)<sup>18</sup>. For immunohistochemistry, the Rat IgG Vectastain ABC Kit, AEC substrate kit, and hematoxylin QS counterstain (Vector) were used according to product instructions. For immunofluorescence staining, AlexaFluor 488 or AlexaFluor 594 conjugated secondary fluorescent antibodies and 1.4 µM DAPI counterstain (Invitrogen) were used.

### Quantification of Immunofluorescence Staining

Retina sections were imaged at 200X magnification, using a Nikon E800 microscope and a SPOTII digital camera, yielding a retina image width of 0.63mm. Within the GCL, immunostained nuclei or cells and DAPI positive nuclei were counted, for each retina section, by two investigators masked to the identity of samples and experiment. Values are reported as the mean and standard error of the mean (S.E.M.). The Mann-Whitney, non-parametric *t*-test was used to analyze differences between experimental groups, and *p*<0.05 was considered significant. Prism 5 (GraphPad) was used for graphing and statistical analysis. For the PD-1 blocking study, each of three experimental repeats examined 4 retinal explants; for each explant, 5 sections (n=5) were co-stained with caspase-3, Brn3a, and DAPI. For the PD-1<sup>-/-</sup> study, 3 animals were examined at each time point, for both PD-1<sup>-/-</sup> and wildtype; at least 3 retina sections (n≥3) animal were stained with Brn3a or AP2, and DAPI.

### Western Blot Analysis

Retinas were isolated from C57BL/6 and PD-1<sup>-/-</sup> mice. Retinas were sonicated on ice in lysis buffer (30 mM Tris-HCl, 10 mM EGTA, 5 mM EDTA, 1% Triton X-100, 250 mM sucrose)<sup>19</sup> with complete protease inhibitor cocktail (Roche), for 2 min (4 × 30 second pulses). 30 µg of each lysate was analyzed by SDS-PAGE as previously described<sup>20</sup>. These primary antibodies were used: rat anti-mouse PD-1 (clone 29F.9A2, rat IgG2a,k) or mouse

anti- $\beta$ -actin (clone 2A2.1, mouse IgG1, US Biologicals). PD-1 and  $\beta$ -actin were detected with anti-rat IgG-HRP and anti-mouse IgG-HRP (Southern Biotech), respectively.

## RT-PCR

Neural retinas were dissected or primary thymocytes were isolated. Retinas from 3 animals were pooled for each time point; 3 experimental repeats were performed. Tissues or cells were stabilized in RNA later (Qiagen). Total RNA was extracted using an RNeasy Mini Kit (Qiagen) and purified using the RNeasy MinElute Cleanup Kit (Qiagen). 0.5 $\mu$ g total RNA was used as a template for reverse transcription using the SuperScript III First-Strand Synthesis SuperMix for qRT-PCR (Invitrogen) and amplified using Platinum Quantitative PCR SuperMix-UDG (Invitrogen) on an ABI 7500 platform. The following 20X Taqman assays (Applied Biosystems) were used: Mouse GAPDH Endogenous Control, FAM-MGB (GenBank Accession No. NM\_008084.2); PD-L1-FAM (Mm00452055\_m1, GenBank NM\_021893.2); PD-L2-FAM (Mm01208507\_m1, GenBank NM\_021396.1).

## Results

### PD-1 is Expressed in RGCs and Displaced Amacrine Cells of the Adult Mouse Retina

In the adult retina, PD-1 expression was detected mainly in the ganglion cell layer (GCL) and a sub-population of cells in the inner nuclear layer (INL)<sup>8</sup>. Since the mouse retina consists of six functionally distinct neuronal cell types distributed in three cell layers, we were interested to determine the identity of the cell populations that express PD-1. Immunofluorescent double labeling with antibodies directed against specific retinal cell makers revealed colocalization of PD-1 with a specific RGC marker, Brn3a, and an amacrine cell marker AP2 $\alpha$ , hereafter referred to as AP2 (Fig. 1A). To further validate RGC-specific PD-1 expression, we used another well-established RGC marker Thy-1<sup>21</sup>. Thy-1.1 conjugated magnetic beads were used to isolate RGCs, followed by PD-1 immunostaining (Fig. 1B), and our results confirm that Thy-1 positive cells are PD-1 positive. Quantification of co-staining demonstrated that over 95% of the Brn3a positive cells were PD-1 positive, and among the PD-1 positive cells in GCL, 68% were Brn3a positive RGCs and 28% were AP2 positive amacrine cells (Fig. 1C). To assess the possibility that PD-1 expression is coupled to Brn3a expression, we assessed colocalization of PD-1 with an independent RGC marker NeuN<sup>16</sup> (Fig. 2A). Amongst NeuN positive RGCs, we found 99 $\pm$ 0.2% PD-1 positive and 1.2 $\pm$ 0.2% PD-1 negative cells (Fig. 2B). Thus, we conclude that authentic PD-1 is expressed by mature retinal neurons, predominantly in the RGC population.

### Dynamic Developmental Expression of PD-1 in the Retina

We next characterized the regulation of PD-1 expression in developing retina. RGCs are the first retinal neurons, emerging after embryonic day 11 (E11) in mice and completing maturation after a postnatal retinal maturation process<sup>22</sup>. We therefore examined the spatio-temporal expression pattern of PD-1 at defined times during the period of retinal development. Low levels of PD-1 expression were initially detected in the inner and outer neuroblast layers by immunohistochemistry at E14 (Fig. 3A), around the peak period of RGC genesis. Weak expression of PD-1 appeared throughout both the GCL, containing RGCs, and the neuroblast layer, containing migrating RGC precursors and retinal progenitors (Fig. 3A, E17). By postnatal day 0 (P0), relatively high levels of PD-1 were present at the inner retina, where postmigratory RGC precursors undergo differentiation; substantial PD-1 expression was also detected in a subset of proliferating neuroblasts (arrows) occupying the ventricular zone (vz). In contrast to the embryonic and adult time points, PD-1 expression appeared greatest at P0 and P13 (Fig. 3A). In the adult retina, the PD-1 expression appeared decreased, with expression retained in the GCL and in a subset of

INL neurons (Fig. 3A, P24, arrows). These observations by immunohistochemistry were confirmed through quantification of postnatal PD-1 protein expression by Western blot analysis (Fig. 3B). Retinal PD-1 expression was greatest during P0-P4, coinciding with a very active time of neuronal culling<sup>23</sup>; total PD-1 expression decreased dramatically, 85% by P9, and 96% P42, as compared to P0 (Fig. 3C). Together, these findings indicate that PD-1 is present in the developing retina, especially in the differentiating RGCs, and the dramatic increase in PD-1 expression coincides with the period of retinal cell maturation and synaptogenesis.

### PD-1 Signaling Modulates Apoptosis of Neonatal RGCs

During murine postnatal retina maturation, 50% of RGCs undergo programmed cell death (PCD)<sup>23</sup>, representing one of the key mechanisms regulating cell-number homeostasis. To test whether functional PD-1 signaling is involved in apoptosis of neonatal RGCs, we performed a study on postnatal retinal explants, a culture system which preserves neuronal connections and retinal cytological architecture and serves as an *in vitro* model to study the developing retina. Neonatal retinal explants were treated with an antibody that blocks PD-1 function and compared to both isotype control antibody treated and untreated explants. After a 24-hour incubation, explants were evaluated for expression of cleaved caspase-3, Brn3a, and AP2, as markers for apoptosis, RGCs, and amacrine cells, respectively.

When retinal explants were cultured in the presence of a PD-1 blocking antibody, there was a significant decrease in caspase-3-mediated apoptosis in the GCL of retinal explants ( $p < 0.0001$ ; Fig. 4A, B). In contrast, isotype control antibody and untreated controls had indistinguishable levels of apoptosis ( $p = 0.25$ , Fig. 4B). Brn3a, a specific marker for RGCs, is observed to be downregulated or absent during RGC apoptosis<sup>24</sup>. A significant increase in the number of Brn3a positive cells was observed in the explants that were exposed to the PD-1 functional blocking antibody ( $p < 0.0001$ ; Fig. 4A, C), concordant with the decreased apoptosis in the GCL. We also assessed the effect of blocking PD-1 in amacrine cells, a subset of which express PD-1 (Fig. 4D). We considered amacrine cells as a specificity control, since unlike RGCs, amacrine cells are resistant to apoptosis during the first two days after birth<sup>25</sup>. Amacrine cells did not undergo apoptosis in our *in vitro* system (Fig. 4D), and there were no significant changes in numbers of amacrine cells after administration of antibodies ( $p = 0.48$ ; Fig. 4D). Taken together, the data suggest that PD-1 signaling is important in murine postnatal retinal maturation, where active signaling through the PD-1 axis may be critical for caspase-3-mediated RGC apoptosis during this developmental period.

### Absence of PD-1 Selectively Increases RGC Cellularity During the Peak of Developmental RGC Apoptosis

To test the physiological importance of PD-1 in eye development, we compared retinal structure and cellular composition in the developing PD-1<sup>-/-</sup> mouse retina to aged-matched wildtype C57BL/6 controls. To assess GCL composition, immunofluorescent staining was performed using the RGC and amacrine cell specific nuclear markers Brn3a and AP2, respectively (Fig. 5A). Total cell number in the GCL, as measured by the number of DAPI positive cells/mm, was not significantly changed in the absence of PD-1 at P2 (Fig. 5B). However, during this critical developmental window the PD-1<sup>-/-</sup> retina had a significant increase in the fraction of Brn3a positive RGCs ( $p < 0.0001$ ) within the GCL, where “fraction” refers to Brn3a/DAPI and will hereafter be referred to as “RGC fraction”. In contrast, there was no significant change in the fraction of AP2 positive amacrine cells ( $p = 0.15$ ) within this layer (Fig. 5B). In the PD-1<sup>-/-</sup> animals, at P2 and P4, the peak of naturally occurring PCD, there was an increase in number of Brn3a positive cells in the GCL ( $p < 0.0001$  for each, Fig. 5C); RGC fraction was also significantly increased at P2 and

P4 ( $p \leq 0.0003$  for each, data not shown). The early developmental increase in RGC number did not persist: by P9 and subsequent time points, the PD-1<sup>-/-</sup> retina showed no significant difference in either RGC number (P9  $p=0.62$ , adult  $p=0.052$ , Fig. 5C) or RGC fraction (P9  $p=0.40$ , adult  $p=0.29$ , data not shown), as compared to wildtype. These data show that the absence of PD-1 is associated with a transiently increased RGC number during the peak of post-natal neuronal culling in the mouse retina, providing further evidence for the importance of PD-1 function during the critical window of retinal maturation.

### PD-1 ligand Genes Are Expressed Throughout Postnatal Retina Maturation

To investigate the cell surface interactions responsible for PD-1 mediated PCD in the developing mouse retina, we wished to characterize expression of the PD-1 ligands present during retina development. There are two known PD-1 ligands in the immune system, PD-1 ligand 1 and ligand 2 (PD-L1 and PD-L2), which upon ligation with PD-1, function to regulate the balance between lymphocyte activation and suppression<sup>5</sup>. In mice, PD-L2 expression is thought to be restricted to hematopoietic cell types, while PD-L1 protein is widely expressed in many organs, including immunoprivileged regions such as placenta<sup>5</sup>, cornea endothelium and ciliary body of the eye<sup>26</sup>, and brain<sup>27</sup>. PD-L1 is transcribed in the mature mouse retina<sup>26</sup>, making it a strong candidate for the PD-ligand functioning during developmental culling. We examined PD-L1 and PD-L2 expression during postnatal development at P0, 2, 4, 7, and 30 by reverse transcription PCR (RT-PCR) and found both ligands to be expressed at all ages studied (Fig. 6), supporting the idea that PD-L1 and/or PD-L2 are the activating signal for PD-1 mediated PCD.

### Discussion

Programmed cell death (PCD) regulates central nervous system (CNS) cell-number homeostasis and formation of functional neuronal networks<sup>28, 29</sup>, through a balance of survival factors and death signals<sup>22</sup>. In particular, it has been well established that neurotrophic factors and electrical activity contribute to postnatal RGC survival<sup>25, 30-34</sup>. This study introduces a distinct hypothesis that PCD may also result from an active negative selection process through the PD-1 pathway. PD-1 protein is transiently up-regulated in the GCL and INL during the critical postnatal period of RGC target-finding and synaptogenesis. This spatio-temporal pattern is consistent with an important role for PD-1 receptor function in RGC culling. Our functional perturbations, using a PD-1 receptor-neutralizing antibody *in vitro* and PD-1 deficient mice *in vivo*, have demonstrated selective increases in RGC survival, thus strongly supporting a role for PD-1 signaling during the peak period of RGC physiological PCD. PD-1 is also developmentally expressed in a subset of cells located within the ventricular zone of the neonatal retina, where retinal progenitors are localized, so it is possible that PD-1 ligation has an additional role in regulating the survival of progenitors. Whether the observed RGC protective effect functions directly through a blockade on RGCs, or indirectly through an effect on progenitors, is an important consideration. Since no overt altered phenotypes have been detected in other postnatally generated cell types, such as amacrine cells, in PD-1 null retinas, it is unlikely that PD-1 plays a significant developmental role in controlling amacrine cell number. The role of PD-1 expression and definition of the amacrine cell subset expressing this molecule requires additional investigation. Our results thus support a direct function for PD-1 signaling in RGC survival. We present the first evidence that PD-L1 and PD-L2 are transcribed throughout postnatal retina maturation. PD-Ligands may be expressed by either neuronal, endothelial, or resident immune (e.g., macrophage or dendritic) cell types; spatio-temporal localization of these ligands will be key to understanding which cell compartments experience the consequences of PD-1 mediated signaling during development. However,

currently we cannot rule out the possibility that, in addition to the known PD-ligands, novel PD-ligand-like molecules could be involved in CNS-specific functions of PD-1 signaling.

Since PD-1 ligation does not directly engage apoptotic pathways in T cells, but rather acts as a co-signaling event, reducing signal strength from the TCR during thymocyte selection<sup>6</sup>, we propose a similar mechanism for PD-1 signaling during RGC culling: PD-1 may act to modulate the strength of neuronal survival signals. We have determined that the TCR subunits TCR $\beta$  and CD3e are not expressed in the developing or adult retina, by RT-PCR and immunostaining (data not shown). Thus, TCR is not the neuronal co-receptor and identification of such molecules will be an important next step. It should be noted that the PD-1<sup>-/-</sup> retina shows only a transient increase in RGCs, with normal cellularity after retinal maturation is complete. This implies the role for additional mechanisms that determine the overall outcome of neuronal culling in the retina, presumably including those previously delineated<sup>25, 30-34</sup>. Since immune PD-ligands are known to have functional interactions with molecules other than PD-1<sup>35</sup>, we cannot rule out the possibility that one or both PD-ligands could be necessary for RGC culling. While normal RGC cellularity is eventually achieved in PD-1<sup>-/-</sup> mice, it will be important to determine if the temporal disorder of RGC formation has consequences for definitive retina organization, retinogeniculate synapse formation, and visual function.

Programmed cell death of RGCs is prominent during the first 11 days after birth, peaking during the first 5 days<sup>28, 36</sup>, with a key role for caspase-3<sup>18, 37</sup>. Our findings are consistent with a PD-1 mediated mechanism for physiological apoptosis through a caspase-dependent pathway, as a PD-1 blockade decreases caspase-3 induction. The retinal explant system used in this study involves experimental axotomy, but axotomy-induced RGC death is an unlikely explanation for our observations since it is independent of caspase activation in the neonatal retina.<sup>38</sup> While the precise PD-1-mediated apoptotic mechanism has yet to be defined, two possible mediators are pro-apoptotic Bax<sup>39</sup> and anti-apoptotic Bcl-2<sup>40</sup>, both known to be critical intracellular effector molecules regulating developmental RGC apoptosis in mouse models<sup>41</sup>. Furthermore, Bcl-2 is negatively regulated by PD-1 mediated signaling during immunological thymocyte maturation<sup>6</sup>.

According to the trophic factor theory for neuronal survival, RGCs die after unsuccessful competition for target-derived trophic factors, with stable connectivity also influenced by neuronal activity.<sup>33, 34</sup> The present study provides evidence for an additional regulatory mechanism: RGC death can also involve an active process of negative selection. It is notable that developmental RGC apoptosis may be similar to the molecular pathogenesis of RGC degeneration.<sup>42, 43</sup> Intriguingly, the PD-1 receptor is constitutively expressed and PD-1 ligands are transcribed in the adult retina, a state where there is no active cell death, and we speculate that PD-1 ligation may trigger RGC death upon optic nerve damage. In addition, since PD-1 is constitutively expressed across neuronal populations in many cerebral compartments<sup>8</sup>, PD-1 signaling might also act to augment neuronal injury in cerebral disease. Accordingly, the present study gives impetus to investigation of the role of PD-1 in neurodegenerative diseases and in specific therapies for neurodegenerative disorders.

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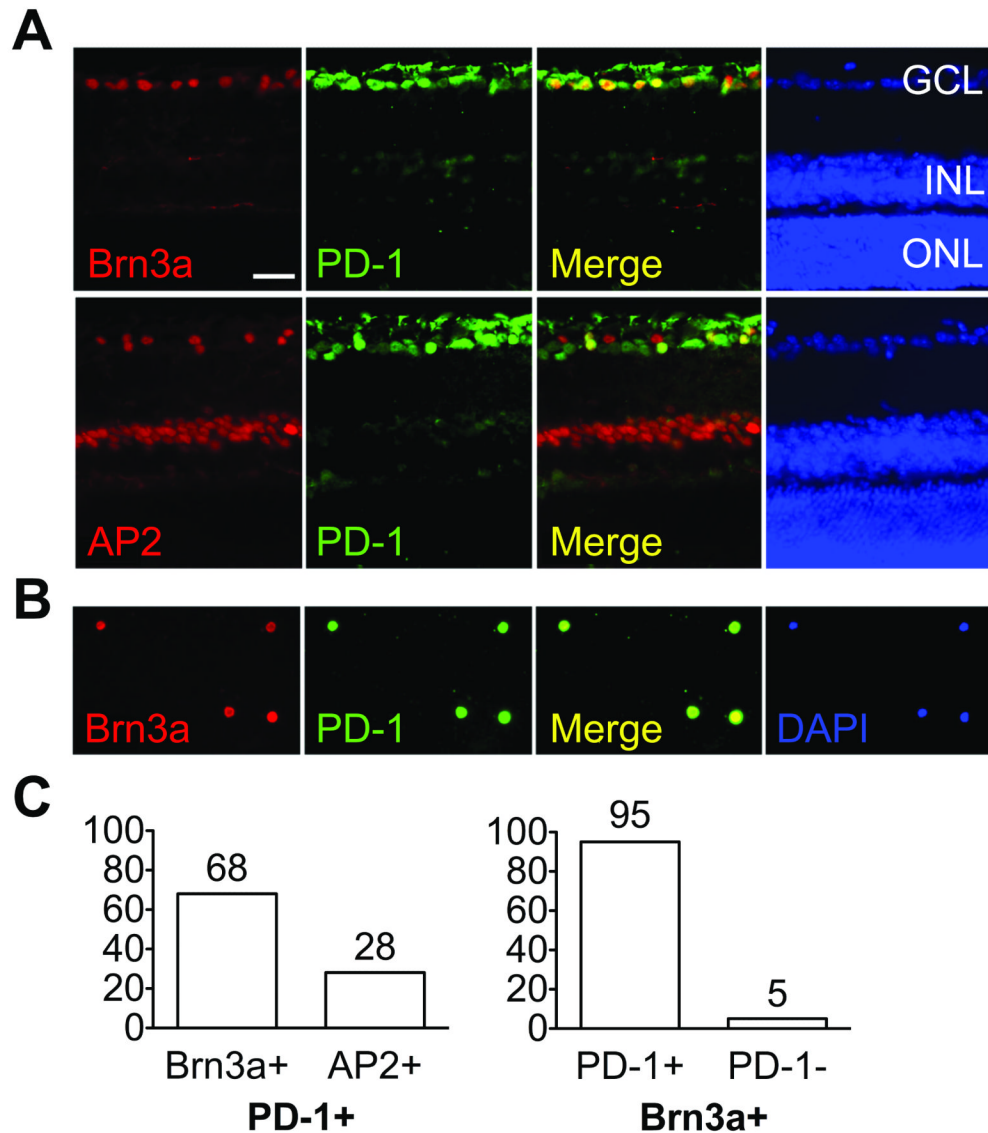
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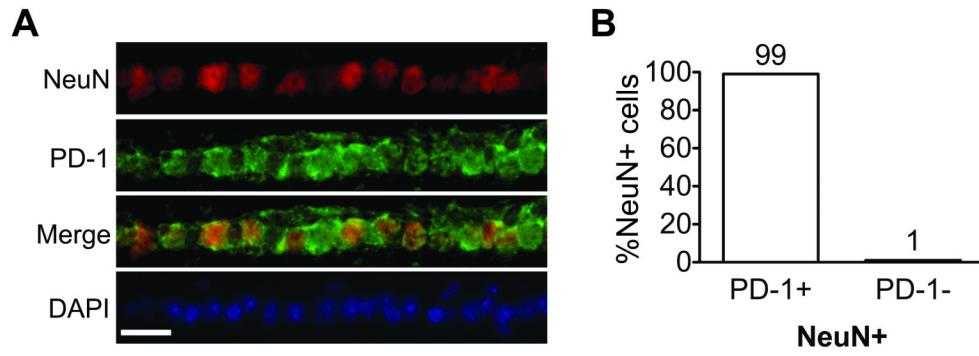


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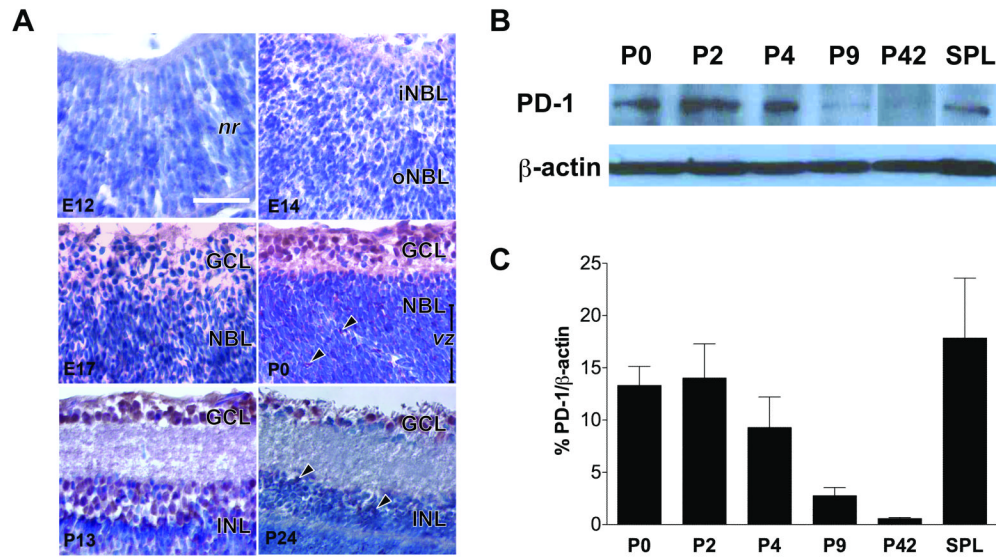


**Figure 1.** PD-1 expression in the adult mouse retina. (A) Double staining of PD-1 with Brn3a or AP2 on vertical retina sections. Left to right panels represent Brn3a or AP2 (red), PD-1 (green), merge (yellow) and nuclear DAPI (blue) staining of retinal sections. (B) Isolated RGCs using Thy-1 coated magnetic beads. Left to right panels represent Brn3a (red), PD-1 (green), and nuclear DAPI (blue). (C) Quantification of PD-1 positive population *in vivo*. Percentage of Brn3a (68%  $\pm$ 11%) or AP2 (28%  $\pm$ 7%) cells among PD-1 positive cells (left). Percentage of PD-1 positive (95%  $\pm$ 3.4%) and PD-1 negative (5%  $\pm$ 1.6%) cells among Brn3a positive RGCs (right). Percentages represent mean  $\pm$ S.E.M. GCL, ganglion cell layer; INL, inner nuclear layer; ONL, outer nuclear layer. Scale bar, 50 $\mu$ m. These findings are representative of four experiments.



**Figure 2.**

Confirmation of PD-1 expression in adult RGCs. **(A)** Double staining of PD-1 with NeuN on vertical retina sections. GCL is shown, with top to bottom panels representing NeuN (red), PD-1 (green), merge (yellow), and nuclear DAPI (blue) staining. Scale bar, 50 $\mu$ m. **(B)** Quantification of NeuN positive population *in vivo*. Percentage of PD-1 positive (99 $\pm$ 0.2%) and PD-1 negative (1.2 $\pm$ 0.2%) cells among NeuN positive RGCs is shown. These findings are representative of three experiments.

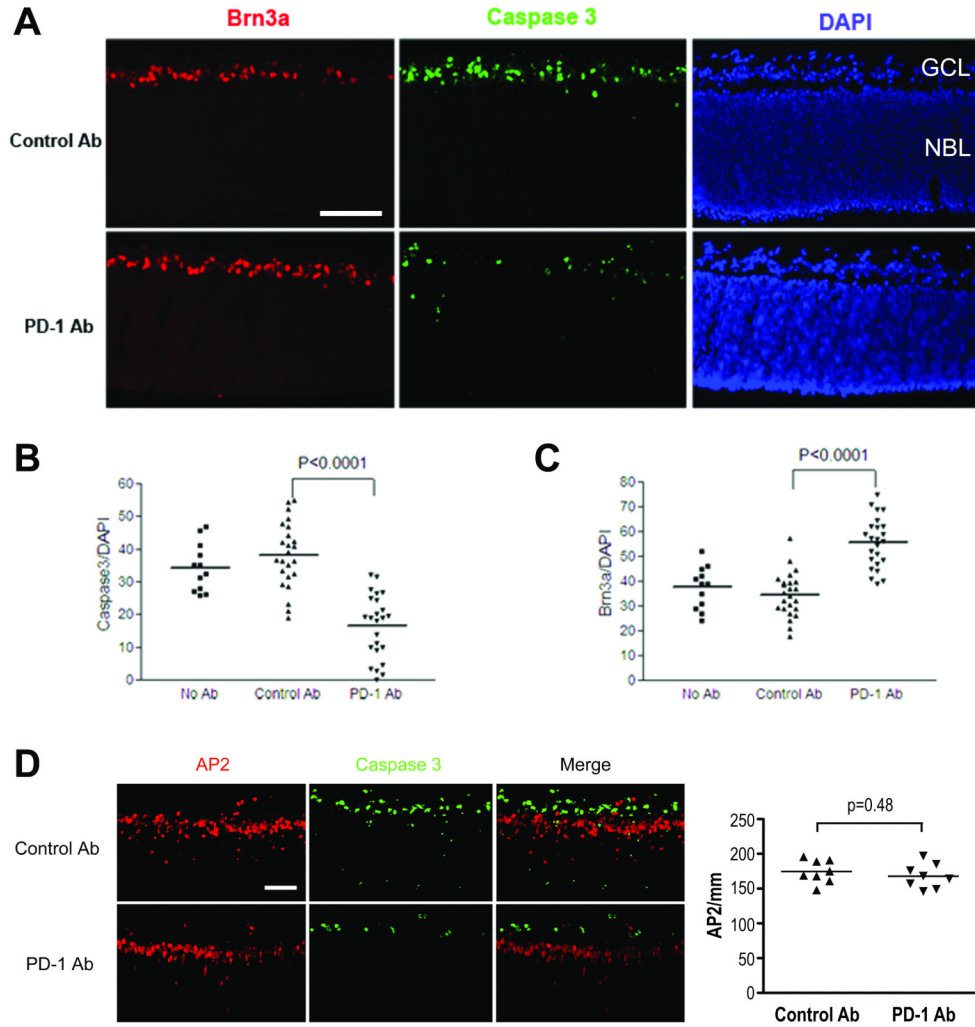


**Figure 3.**

Dynamic regulation of PD-1 expression during retinal development. (A)

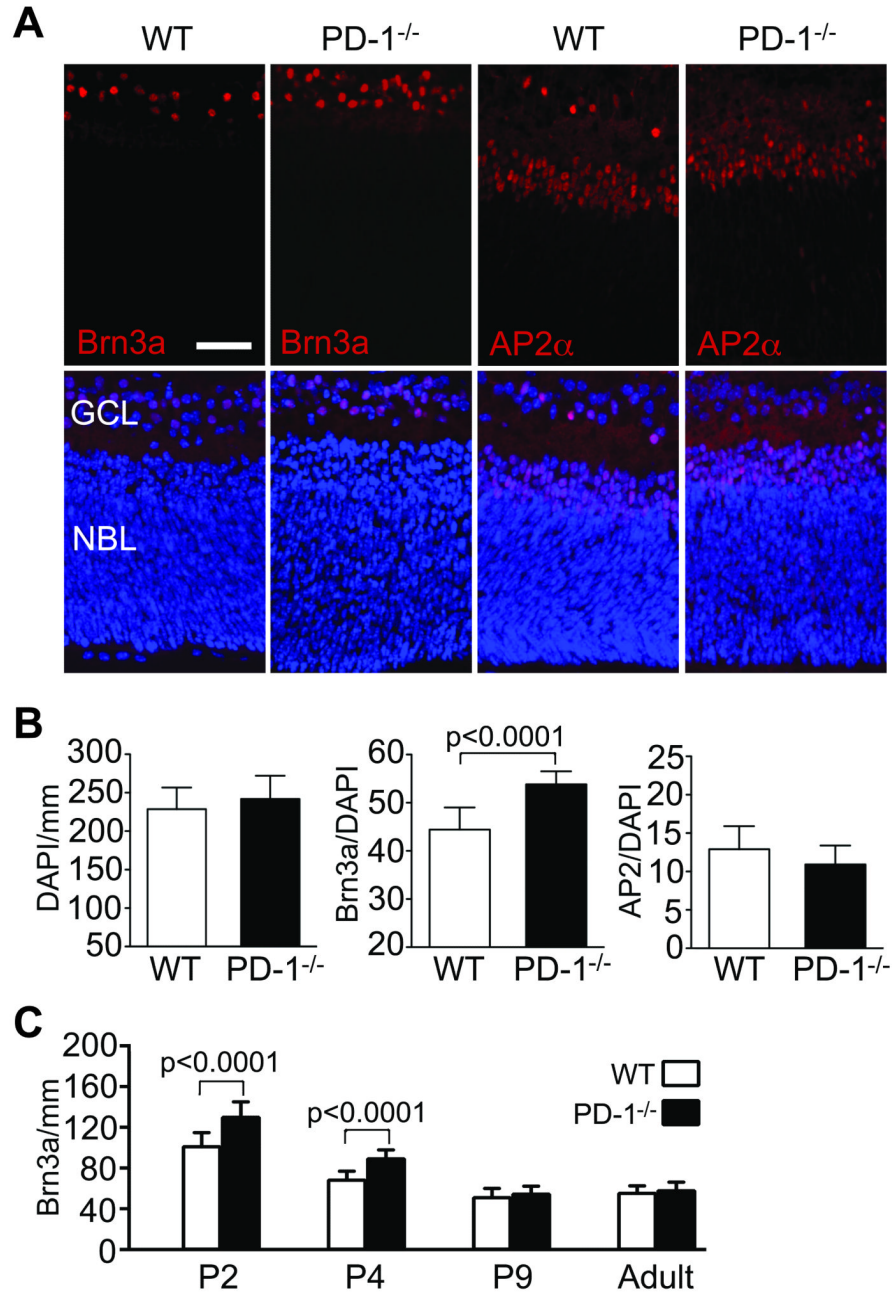
Immunohistochemical staining on retinal sections from embryonic days E12, E14, E17, and postnatal days P0, P13, P24. PD-1 (red) and hematoxylin (blue). *nr*, neural retina; iNBL & oNBL, inner & outer neuroblast layer; GCL, ganglion cell layer; *vz*, ventricular zone; INL, inner nuclear layer. Scale bar, 50 $\mu$ m. These findings are representative of three experiments.

(B) Western blot for PD-1 at P0, P2, P4, P9, P42, and spleen as a positive control.  $\beta$ -actin was used as a loading control. (C) Quantification of PD-1 immunoblots. PD-1 expression decreases by  $80\pm 3.8\%$  by P9, and  $96\pm 0.9\%$  by P42, as compared to P0 expression; values reported as mean $\pm$ S.E.M.  $p=0.03$  by ANOVA analysis across all retina samples. SPL, spleen. Negative controls for reagent (secondary antibody only) and tissue (skeletal muscle) showed no detectable staining (data not shown). These findings are representative of three animals per time point.

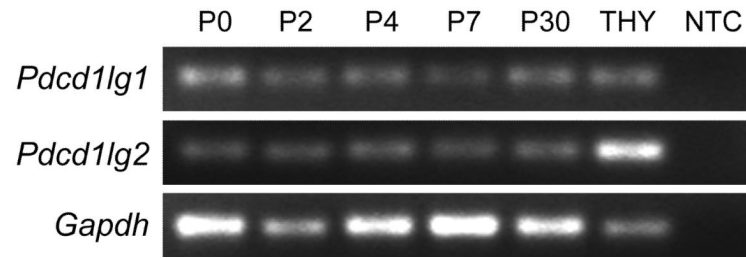


**Figure 4.**

PD-1 blockade inhibits apoptosis of retinal ganglion cells in the developing retina. Neonatal retinal explants were incubated for 24 hours in basal medium containing isotype control antibody (top panel) or PD-1 blocking antibody (bottom panel) (**A** and **D**). (**A**, **D**) Immunofluorescent staining was performed with antibodies against Brn3a (red, **A**) or AP2 (red, **D**), activated form of caspase-3 (green), and DAPI (blue). (**B**, **C**) Quantification of percentage of Brn3a positive cells among DAPI positive cells (**B**) or caspase-3 positive cells among DAPI positive cells (**C**) in the GCL in three experimental groups, as indicated in the figure. (**D**) Amacrine cell survival is not affected by PD-1 blockade. Immunofluorescent staining was performed and the merge image supports lack of colocalization of cleaved caspase-3 and AP2 positive amacrine cells. Quantification of AP2 positive cells also revealed no difference between the anti-PD-1 and control antibody treatments. Scale bars, 100 $\mu$ m. All findings are representative of three experiments.

**Figure 5.**

Increased RGCs during postnatal retinal development in the PD-1<sup>-/-</sup> mouse. (A) Immunofluorescence staining was performed on both wildtype and PD-1<sup>-/-</sup> P2 retina cryosections using antibodies against either Brn3a (red) or AP2 (red). Nuclei were visualized with DAPI (blue). NBL, neuroblast layer. (B) Within the GCL, quantification of total cells (DAPI/mm), Brn3a<sup>+</sup> cells (Brn3a/DAPI), and AP2<sup>+</sup> cells (AP2/mm), at postnatal day P2, in wildtype and PD-1<sup>-/-</sup>. p-values denote significant differences between WT and PD-1<sup>-/-</sup>. (C) Quantification of Brn3a in wildtype and PD-1<sup>-/-</sup> retinal sections from P2, P4, P9, and adult (8 week old) animals was performed as noted. p-values denote significant differences, at P2 and P4, between WT and PD-1<sup>-/-</sup>. Scale bar, 50μm.



**Figure 6.**

Expression of PD-L1 and PD-L2 throughout retina maturation. RT-PCR products showing *Pdc11g1* (PD-L1 gene) and *Pdc11g2* (PD-L2 gene) expression at various ages, with *Gapdh* as an endogenous control and thymus as a positive control. NTC, no template control.