A Chick Model of Retinal Detachment: Cone Rich and Novel

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Abstract

Background: Development of retinal detachment models in small animals can be difficult and expensive. Here we create and characterize a novel, cone-rich retinal detachment (RD) model in the chick.

Methodology/Principal Findings: Retinal detachments were created in chicks between postnatal days 7 and 21 by subretinal injections of either saline (SA) or hyaluronic acid (HA). Injections were performed through a dilated pupil with observation via surgical microscope, using the fellow eye as a control. Immunohistochemical analyses were performed at days 1, 3, 7, 10 and 14 after retinal detachment to evaluate the cellular responses of photoreceptors, Müller glia, microglia and nonastrocytic inner retinal glia (NIRG). Cell proliferation was detected with bromodeoxyuridine (BrdU)-incorporation and by the expression of proliferating cell nuclear antigen (PCNA). Cell death was detected with terminal deoxynucleotidyl transferase dUTP nick end labeling (TUNEL). In mammalian models of RD, there is shortening of photoreceptor outer segments and mis-trafficking of photoreceptor opsins in areas of RD. Photoreceptor cell death was maximal 1 day after RD, but continued until 14 days after RD. Müller glia up-regulated glial fibrillary acidic protein (GFAP), proliferated, showed interkinetic nuclear migration, and migrated to the subretinal space in areas of detachment. Microglia became reactive; they up-regulated CD45, acquired amoeboid morphology, and migrated toward outer retina in areas of RD. Reactive NIRG cells accumulated in detached areas.

Conclusions/Significance: Subretinal injections of SA or HA in the chick eye successfully produced retinal detachments and cellular responses similar to those seen in standard mammalian models. Given the relatively large eye size, and considering the low cost, the chick model of RD offers advantages for high-throughput studies.

Introduction

Retinal detachment (RD) is a clinically important cause of visual loss; it is common and it is destructive to vision and to the eye itself. Poor visual acuity resulting from RD has been studied in humans and animal models for decades [1]. Such models have included intravitreal injections of dispase, for enzyme disruption of basement membranes [2,3,4], subretinal injection of saline to create a transient RD [5] or hyaluronic acid for a chronic RD [6,7], or intravitreal injection of cells (e.g., fibroblasts, macrophages, retinal pigment epithelial cells) [1,8,9,10]. Currently, the subretinal injection of hyaluronic acid is a prevalent RD model and has helped to explain the cascade of events following RD that can lead to permanent vision loss [6].

Changes to the photoreceptors, glia, and macrophages/microglia appear to be critical in the pathobiology of RD. Specifically, the photoreceptor outer segments (OS) degenerate and many of the photoreceptors apoptose, resulting in thinning of the outer nuclear layer (ONL) [11,12]. This apoptosis is maximal 3 days following a retinal detachment in several mammalian models [13]. Subsequent to photoreceptor damage, Müller glia proliferate, hypertrophy, with up-regulation of intermediate filaments [13,14,15,16], and migrate to the outer nuclear layer (ONL) [17,18,19,20], contributing to the destructive scar formation which is the hallmark of proliferative vitreoretinopathy [6,21]. Müller processes extend beyond the outer limiting membrane (OLM) and limit re-growth of photoreceptor outer segments after the retina is re-attached [18]. In addition, macrophages and microglia become reactive and accumulate in significant numbers in the retina and subretinal space, and contribute to retinal pathophysiology following RD [19,22,23,24,25,26,27].

A wide variety of mammalian species have been used to model retinal detachments and proliferative vitreoretinopathy, including rabbits, cats, mice, and primates [13]. But, other than primates, these species do not have a cone-rich retina needed to model humans. One animal that does possess similar cone density is the ground squirrel (Spermophilus beecheyi) and it has been used as
a model for RD [28,29]. Unfortunately, the ground squirrel model has significant disadvantages in poor availability and difficult handling. For these practical reasons the identification of better cone-rich animal models of RD is important for study of this retinal disorder. In addition, a model which permits our better understanding of the molecular biology of macroglial and microglial cells, and of their responses to retinal damage and to progenitor cells will add special value [30,31,32,33,34,35,36].

The chick has been used to study the development of the visual system [37,38] and, more recently, for studying retinal damage and potential for regeneration [30,31,32,33,34,35,36]. It is a diurnal species with a sophisticated visual system emphasizing color vision. Chick retina contains four single cones responsible for color vision and one double cone, which may mediate achromatic motion perception [39]. The cone types include those that express visual pigments sensitive to long- (L), medium- (M), or short- (S) wavelengths. By convention, the chick L cone photopigment absorption peaks at 517 nm (also known as chicken red), M2 cone photopigment at 508 nm (chicken green), M1 cone photopigment at 435 nm (chicken blue), and S cone photopigment at 415 nm (chicken violet, the most similar to human S cones) [40]. Rods contain rhodopsin visual pigment (absorption around 500 nm) [40,41]. Cones outnumber rods 6:1 in the chick retina [42], while humans have approximately 20 times more rods than cones [43]. Cones are distributed throughout the entire chick retina with the highest density in the rod-free area centralis, which is analogous to the fovea centralis in humans [38,41].

With this study we want to add further value to the use of the chick in biological research. We propose to develop a new chick model of RD using subretinal injection of hyaluronic acid. We hope to use its many modeling advantages which include (1) a cone-rich retina, (2) novel retinal glia [present in non human primates but not in rodents], (3) the wealth of data on micro and macroglial molecular responses to retinal damage and retinal regeneration, (4) the availability of mutant ocular strains, (5) a sequenced genome, (6) low cost, (7) widespread availability, (8) ease of handling, and (9) large eyes amenable to experimental manipulation.

**Results**

Retinal detachments and controls were examined at days 1, 3–4, 7, 9–10, and 14–16. The RPE initially appeared flattened at day 1, with loss of apical processes (Figs. 1a–c). By 9 days after treatment, the detached RPE had a mound appearance (Figs. 1d–f) with signs of hyperplasia in some samples (Fig. 1c). The photoreceptor outer segments were shorter and degenerated by day 1, compared to controls (Figure 1). In addition, we observed thinning of the outer nuclear layer (ONL), suggesting loss of photoreceptors (Fig. 1f). Saline detachments had resolved by the day 3 timepoint (not shown), while the hyaluronic acid detachments persisted.

Immunohistochemical analyses of photoreceptors showed mis-trafficking and disorganization of L/M opsin, S opsin, and rhodopsin in areas of retinal detachment (Figure 2). These changes were similar in both saline (SA) and hyaluronic acid (HA) RDs even though SA RDs had resolved by day 3. Re-distribution of L/M opsin to the photoreceptor cell body was seen using confocal microscopy. Opsin-immunofluorescence was present across the inner segments, cell bodies and axon terminals of stressed photoreceptors in regions of detachment (Figs. 2–f). At approximately RD day 9, there was reduction in rhodopsin and S opsin levels, with relative preservation of L/M opsin and calbindin (Fig. 2). Labeling for S opsin and rhodopsin were reduced in detached regions of retina, whereas labeling for L/M opsin and calbindin remained, but appeared disorganized (Figure 2q–t).

To evaluate the timecourse of photoreceptor death in this model, TUNEL analysis was performed. Cell death was significantly increased in detached retina (ANOVA, p < 0.0001, Figure 3). Cell death was maximal at day 1 after detachment and then decreased over time (day 1 control: day 1 RD (p < 0.0001), day 1 RD: day 3 RD (p < 0.0001), day 1 RD: day 7 RD (p < 0.0001) and day 1 RD: day 14 RD (p = 0.0004)). TUNEL labeling was not significantly different than in controls at day 3, 7, and 14. The labeling appears primarily in the outer nuclear layer (ONL) but some cells were seen in the INL (data not shown). ONL thickness, a measure of photoreceptor cell survival, was unchanged at day one, but decreased significantly at later timepoints following retinal detachment, measuring 99% of paired control at detachment day 1, 84% at day 3, 55% at day 7, and 47% at 14 days following retinal detachment (Figure 3, ANOVA; p = 0.0002). The mean ONL thickness in regions of detachment was significantly less than in paired controls at day 14 (p < 0.0001) and bordering at day 7 (p = 0.0088); ONL thickness in day 1 RDs was significantly greater compared to days 7 and 14 (p = 0.0005 and 0.0033, respectively).

The cellular responses of the Müller glia to RD were evaluated. Similar to findings in mammalian models, the expression of the intermediate filament GFAP increased in regions of detachment (Figure 4) as early as 3 days after detachment. Expression of transitin, the avian homologue of mammalian nestin, was similarly increased (Fig. 4d). Glial processes positive for GFAP, transitin, and Muller glial marker TOPAP (topographic marker expressed along the anterior-posterior axis, detected by antibody 2M6) were seen extending beyond the outer limiting membrane (OLM) in areas of RD as early as day 3 (Figs. 2, 4). Areas consistent with subretinal scar were visualized at day 7 and 14 with large clusters of subretinal glial processes, as well as cell bodies positive for Sox2 and TOPAP (Fig. 4).

Cell proliferation, particularly of Müller glia, has been reported following retinal detachment in mammalian models [15,43,46]. Figure 5 shows proliferating cells present in detached retina; most were Sox2/PCNA-positive Müller glia (Fig. 5). PCNA-positive cells were more numerous than BrdU-positive cells and predominantly co-localized with Sox2, but not the NRG marker homeobox protein Nkx2.2 (Nkx2.2), thus indicating Müller glia proliferation (Fig. 5). Müller glia also exhibited interkinetic nuclear migration in regions of detachment, with migration of nuclei into the outer retina (Figs. 4–5).

A highly localized infiltration and activation of retinal and subretinal microglia and macrophages is characteristic in mammalian models of RD [22,24,25]. In the chick RD eyes, there was a significant accumulation of CD45-positive microglia/macrophages in the detached retina compared to controls (Fig. 6). Quantification of CD45-immunofluorescence indicated significantly increased mean area of signal per field of view from $2.8 \times 10^4$ to $64.2 \times 10^4$ in RD retina at day 1 ($p = 0.00092$), and the CD45 density sum increased from $7.2 \times 10^4$ to $64.2 \times 10^4$ ($p = 0.0165$). Density sum was significantly elevated in day 3 RD retina at $59.0 \times 10^4$ compared to day 3 controls at $4.0 \times 10^4$ ($p = 0.0118$), the other days did not reach significance after Bonferroni correction.

Recently, a novel population of glial cells (nonastrocytic inner retinal glia (NIRG) has been detected in the retinas of multiple species, including chick and primates [34,47], and seen particularly after retinal damage in regions of focal detachment [34]. NIRG cells are known to express Sox2, Sox9, Nkx2.2, transitin, and vimentin, but are negative for GFAP and Pax2 [34].
Immunostaining for NIRG cells revealed an accumulation of NIRG cells above background controls in areas of RD (Figure 7). Surprisingly, NIRG cells were detected in subretinal scars as well as inner retina.

Discussion

To the best of our knowledge, this is the first study utilizing the chick as a model of retinal detachment. There are many biologic, logistic, and practical advantages of the cone rich chick model system. Briefly, they offer advantages in cones, glia, characterized retinal responses, sequenced genome, availability, low cost, ease of handling, and large eyes. While chicks are not generally selected based on genetic strain, there are some ophthalmic mutants available [as reviewed in [48]]. Since the eye is large, intravitreal injections can be performed easily for potential high-throughput analyses of therapeutic agents.

As an alternative to the ground squirrel, the chick compares favorably in terms of availability, handling, and cost. The chick and ground squirrel cones have similarities, but some differences (Table 1). Chick retina serves tetrachromatic vision and contains four single cones (L, M1, M2, and S sensitive) and one double cone [39]. Ground squirrels have dichromatic vision with cones most sensitive to M (518 nm) and S (436 nm) wavelengths, as determined by ERG testing [49]. Similar to chicks, ground squirrels too are diurnal, with a very high cone density: about 7.5 million cones per retina. This is 1.6 times the human cone density (4.6 million cones) [50]. They have a retina in which 85% of the photoreceptors are cones, about 5.7 cones-per-rod [29]. In comparison, the chicks have 6 cones-per-rod [42]. The highest cone density in ground squirrel (49,550/mm²) is in the visual

Figure 1. Altered outer retinal morphology in chick retinal detachment. Toluidine blue staining timecourse of retinal detachment frozen sections days 1, 3, 10, and 14, showing the typical degenerative changes after HA RD as well as variability of the detachments generated. (a) Normal retina with details of outer retina, including oil droplets (predominantly orange-colored spheres) between the photoreceptor inner and outer segments and RPE apical processes (inset). (b–f) There is shortening of the outer segments and flattening of the RPE in areas of RD. Later there is a decrease in cell bodies in the outer nuclear layer (f). RPE hyperplasia is noted in some sections (C). ONL (outer nuclear layer), OPL (outer plexiform layer), PR (photoreceptor), IS (inner segments), OS (outer segments), RBC (red blood cells). Scale bar (50 microns).

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streak, analogous to the fovea centralis, in a horizontal strip of retina 2mm ventral to the optic nerve head [51]. In comparison, the chick area centralis is reported to have between 10,000–16,000 cones/mm², although average cone density is reported higher by other studies [52]. Unlike mammalian cones, chick cones contain an oil droplet that filters out light below wavelengths 575, 520, 497, or 454 nm [39,53]. Chick rods do not contain an oil droplet.

Figure 2. Mistrafficking of photoreceptor opsins in RD. Calbindin (red) and L/M opsin (green) immunostaining in control retina of the untreated fellow eye (a–c) and hyaluronic acid RD at day 3 (d–f) and 9 (g–i). Note decreased intensity of L/M opsin staining at day 9 HA RDs. Confocal microscopy demonstrates early mistrafficking of L/M opsin (green) with staining of the photoreceptor cell body and cone pedicles (arrows, k, l) in SA RDs at day 3 compared to untreated controls. The L/M opsin staining is present in areas consistent with the cone IS/OS as well as the cone ellipsoids in normal chick retina (j). Some transitin-positive glial processes are seen extending beneath the outer limiting membrane (OLM, arrowhead k, l). Similar to day 3 HA RDs, day 3 SA RDs show mistrafficking of rhodopsin (red) and L/M opsin (green, m, n) and S opsin (green, o, p), despite resolution of subretinal fluid. In advanced HA RD (Day 16 RD, q–t) there is redistribution of L/M opsin and calbindin to the cell body with loss of S opsin and rhodopsin (bracket). ONL (outer nuclear layer), OPL (outer plexiform layer), PRL (photoreceptor layer). Scale bar 50 microns. doi:10.1371/journal.pone.0044257.g002
Despite some differences in their cone structure as compared with mammals, chick cones showed similar responses to retinal detachment. The photoreceptor outer segments demonstrated abnormalities in the detached areas as soon as one day after detachment. There was mis-trafficking of photoreceptor opsins and rhodopsin, similar to that in other models of detachment [28,54,55,56,57]. Cone opsins were redistributed to the plasma membrane as the outer segments degenerated. Similar to Sethi et al. [16], we found L/M opsin redistributing to the cell body in retinas with more severe degeneration in HA RDs. We also observed L/M opsin staining on the photoreceptor cell body and cone pedicles in the re-attached day 3 SA RD eyes.

The ground squirrel retina has been evaluated with histologic, molecular, and electrophysiologic techniques to determine rod and cone responses to RD. As in other mammalian species, outer segment degeneration began at 1 day. Rex et al. [56,57]...
Figure 4. Müller glia activated in chick RD: Expression of intermediate filaments GFAP (untreated (a), day 5 HA RD (c)) and transitin, the chick homologue of nestin, (day 5 HA RD (d), untreated not shown) are essentially negative in normal control retina and increase after RD by day 3 (not shown). GFAP- and transitin-positive Müller glial processes extend beneath the outer limiting membrane (OLM) in HA RDs (arrows c, d) to the subretinal space. The Müller marker TOPAP (antibody 2M6) is present primarily in a vertical distribution throughout the untreated retina in Müller glia (b) and increases particularly in the outer retina below the OLM in HA RDs, with accompanying undulation of the outer retina (f). Cells in subretinal scars are positive for Müller progenitor marker Sox2 (arrow, e) and 2M6 (arrow, f).

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demonstrated a dramatic loss of mRNA and protein expression of cone opsins early after RD (days 1–3), with a relative preservation of rod opsin expression. Our studies indicate a longer retention of cone markers in the chick RD model with some L/M opsin and calbindin staining detectable in day 14 RDs. However, we note that S opsin-positive cones showed extreme down-regulation of these S photopigments in areas of severe degeneration, suggesting that S cones could be more sensitive to damage from RD. Nork et al. [58] demonstrated significant damage to S cones in human RDs. Experimental RD studies have yielded mixed results regarding whether S cones are more sensitive to RD damage. Future studies of S cone mapping should be done with the chick RD model to answer this question of relative stress sensitivity.

Although there is permanent loss of some photoreceptors following RD, re-attachment studies have demonstrated recovery of rods and cones, including re-expression of cone opsin [56,57].

Sakai demonstrated similar transient depletion of S and M cone opsins in ground squirrel retinas that had been detached for 1 day and then reattached [59]. It is clinically and experimentally important that the time-course of cell death of photoreceptors following RD was similar in the chick model to that seen in human retinas. Dying cells were most abundant at day 1. This is earlier than the day 3 peak in cell death observed in several mammalian RD models [6]. The abundance of dying cells was highly variable between individuals, likely because the size of the detachment was variable. However, outer nuclear layer thickness, reflecting photoreceptor cell death, was initially preserved and did not decline significantly until days 7–14 after detachment. These data fit well with human data suggesting that the interval 7–30 days post-detachment is likely to be most significant for visual loss due to photoreceptor damage [60,61,62,63,64,65]. The mechanism of visual loss in patients with macula-affecting RD is highly

Figure 5. Müller glia proliferate after HA RD. Immunostaining to detect proliferating cells demonstrated more PCNA (blue) than BrdU (green) positive cells at the timepoints tested after RD (day 3, 7, 14). Sox2 (red) labels nuclei of Müller glia and NIRG cells. Merged images of PCNA, Sox2, and BrdU demonstrate a good correlation of PCNA and Sox2, including in subretinal scar. Cells labeling with all three markers are marked with arrows, while Sox2 negative cells labeling with PCNA and BrdU are marked with open arrowheads.

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correlated with both photoreceptor cell death (indicated by thinning of the outer nuclear layer) and by abnormalities in the photoreceptor outer segments (indicated by disruption of the “intermediate line”) on spectral domain optical coherence tomography (SD-OCT) [66]. This disruption can improve in patients over time and fits with improving visual acuity [66]. Future studies in the chick, including use of SD-OCT, will be useful to better evaluate photoreceptor recovery after retinal detachment repair. Here, most experiments used subretinal hyaluronic acid which created a chronic detachment. In saline RDs, the detachments had resolved by day 3. However, in these re-attached saline RDs, responses were similar to those of hyaluronic acid RDs, including mistrafficking of photoreceptor proteins (Fig. 2k–l, m–p) and microglial activation and migration towards the subretinal space (Fig. 6b).

There is a local response of Müller glia in most mammalian models of RD and also in human retinectomy specimens [6,16]. In contrast, the Müller glia in the ground squirrel are non-reactive in areas of RD [28]. This is an important research liability. The chick RD model is a better match for feline and human RD than is the ground squirrel in its prominent Müller glial activation, involving GFAP up-regulation, distal migration and proliferation. Similar to these other models, Müller processes extend beyond the OLM and appear to establish subretinal scars in some eyes. Recently, the Müller glia have been shown to proliferate, extend processes, and migrate beneath the retina to form subretinal scars [21]. Similarly, in the chick model we observed proliferation of Müller glia and interkinetic nuclear migration, with movement toward the outer retina, and extension of processes beyond the OLM. Histological examination of the subretinal scar in the chick indicates the accumulation of Müller glial processes and cell bodies. These results suggest that some Müller progenitor cells detach and migrate wholly into the subretinal space, while some only grow processes into the subretinal space. Such migration patterns and mechanisms are of special importance for translational research. Controlling Müller glial activation and gliosis can be critical to stopping subretinal scars and proliferative vitreoretinopathy.

Chick retina contains a recently-identified special glial cell type termed non-astrocytic inner retinal glia (NIRG). NIRG-like cells have been identified in the retinas of multiple species including dogs and monkeys, but not in the retinas of rodents [34,47]. NIRGs activate and proliferate in response to chemical retinal injury with NMDA and IGF-1 in the chick, which leads to multiple small detachments of the retina [34]. In these detached

**Figure 6. Microglia are up-regulated in chick SA and HA RD.** SA RDs (b) and HA RDs (c) show up-regulation of microglial marker CD45 (green) in areas of RD compared to control retina (a). The SA RD is resolved by day 3. The microglia acquire amoeboid morphology and migrate to the outer retina/subretinal space. Quantitation of CD45 staining in HA RD over time (d, density sum and e, mean area).

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Mueller glia are absent, but an accumulation of NIRG cells fringe the Muller glia around damaged areas in the inner retina, vitread to where the Muller glia typically reside [34]. Little is known about the function of NIRG cells in response to retinal damage. To determine whether NIRG cells were prominent in the model of RD created here, we evaluated NIRG populations in the detached areas. These studies demonstrated a small increase in numbers of NIRG cells in the inner retina. However, Nkx2.2-negative Muller glia, expressing Sox2 and Sox9, were the predominant glial cell type in detached retina in this model. Surprisingly, some NIRG cells were detected in subretinal scars, along with Muller-glia derived cells (negative for Nkx and positive for Sox-9 as well as Sox-2, and TOP\textsuperscript{Ap}). These studies support the notion that macroglial cells contribute to subretinal scar formation after retinal detachment [21].

Areas: Figure 7. NIRG cells are mildly up-regulated after chick HA RD. NIRG cells (arrows) are positive for Nkx2.2 (green, a–f) and Sox9 (orange, c, d) and are increased after HA RD compared to controls. The up-regulation of NIRGs is less than that of Muller glia (Sox9+/Nkx- cells in the inner nuclear layer and outer retina). NIRGs were identified in subretinal scars (f, arrow; green = Nkx, blue = DAPI).

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Similar to other studies of mammalian models, the macrophage/microglial response to RD in the chick is dramatic. Microglia are resident tissue macrophages of the CNS. In the chick eye they are identified by immunostaining with CD45 [67]. In untreated eyes quiescent microglia are typically located in the plexiform layers of the retina. After RD, microglia migrate to the outer retina, up-regulate levels of CD45, acquire amoeboid morphology and enter the subretinal space. Nakazawa et al. [22] identified the importance of the macrophage product MCP-1 in decreasing photoreceptor survival in a murine model of RD. While the chick does not have a known MCP-1 homologue, it will be valuable to determine whether other specific macrophage products can reduce photoreceptor survival similar to those in mammalian models [22], and whether there is a role for microglia contributing to Muller glial proliferation and subretinal scar formation.
In conclusion, the chick model of RD shares many similarities with human, and advantages over other mammalian models. It is a cone-rich and research friendly model, and it could offer added insights into the pathobiology of retinal detachment. It could be helpful for high-throughput studies of neuroprotective or anti-proliferative agents that would ameliorate RD.

### Materials and Methods

**Animals**

Leghorn chicks (*Gallus gallus domesticus*) were obtained from the Department of Animal Sciences at The Ohio State University. They were used at postnatal days 7 to 21 (P7–P21) to create the retinal detachment model under an IACUC approved protocol (#2009A0139). Only one eye was treated. The chicks were housed with free access to Purina chick starter and water in a 12 h light-to-dark cycle in a stainless steel brooder.

**Retinal Detachment Surgery**

This research adheres to the principles of the ARVO Statement for the Use of Animals in Ophthalmic and Vision Research. Chicks (P7–P21) were anesthetized with inhaled isofluorane via nonrebreather mask. Pupil dilation was achieved with multiple rounds of drops of tubocurarine hydrochloride (Sigma: 6.7 mg/ml in saline with 0.1% benzalkonium). Tubocurarine, a neuromuscular blocking agent that is a competitive antagonist of nicotinic neuromuscular acetylcholine receptors [68], was used for dilation since chick iris contains prevalent nicotinic cholinergic receptors [69].

Left eyes were prepped with betadine swabs and a pediatric lid speculum was inserted. A lateral canthotomy was created with a sterile razor blade for better exposure of the globe. The conjunctiva was stabilized with 0.12 forceps and then was dissected temporally with a 25 g needle tip to expose the sclera. A 25 g needle on a 5 cc syringe was inserted approximately 3 mm posterior to the limbus and vitreous was aspirated from the central posterior to the limbus and vitreous was aspirated from the central

**Enucleation and Histology**

Eyes were enucleated for analysis at day 1 (n = 7), 3–4 (n = 10), 5–7 (n = 6), 9–10 (n = 3), and 14–16 (n = 6).

**BrdU Labeling**

Intravitreal injections of 2 micrograms 5-bromo-20-deoxuridine (BrdU; 100 microgram/mL Sigma-Aldrich) were delivered approximately 4 hours prior to euthanization (n = 4 chicks at each timepoint: day 1, 3, 7, 14) as previously described [36]. Frozen sections were immunostained using the anti-BrdU antibody (G3G4, Developmental Studies Hybridoma Bank). BrdU positive cells were determined at different levels in the area of the retinal detachment to determine the number of proliferating cells.

**TUNEL Assay**

Apoptosis was evaluated with the TUNEL assay with the In Situ Cell Death Kit (TMR red; 1215679910, Roche Applied Science) per the manufacturer’s instructions.

**Immunohistochemistry**

Immunohistochemistry was performed to detect markers as previously described [34]. Antibodies to rhodopsin (Rho 4D2, mouse, generous gift Dr. R. Molday, University British Columbia), L/M opsin (AB5405, rabbit, Chemicon), S opsin (rabbit polyclonal anti-blue opsin, Millipore) were used to evaluate photoreceptor opsins. Mouse anti-calbindin (300; Swant Immunochemicals, Bellinzona, Switzerland) was used to detect photoreceptor opsins.

### Table 1. Short-wavelength (S), Medium-wavelength (M), Long-wavelength (L), cones per square millimeter (c/mm²).

| Chick        | Cone Vision | Oil Droplet | High Acuity Vision | Fovea | Average Retinal Cone density |
|--------------|-------------|-------------|--------------------|-------|-----------------------------|
| Tetrachromatic | S-M-L       | Yes         | Area Centralis     | No    | 10,000–35,961c/mm² (a)      |
| Dichromatic  | S-M-L       | No          | Visual Streak      | No    | 24,310–49,550c/mm² (b)      |
| Trichromatic | S-M-L       | No          | Fovea Centralis    | Yes   | 2500c/mm² peripheral 6000c/mm² central (c) |

Additional notes:

*Reviewed in [52]. Cone density varied widely between experiments.
*Peak cone density was highest in the visual streak and lowest density in the dorso-nasal periphery [51].
* Determined by histology from donor eyes [72].

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calbindin. Antibodies to BrdU (G3G4, mouse, Developmental Studies Hybridoma Bank) and PCNA (M0079, mouse, DAKO) were used as markers of proliferating cells. CD45 (HIS-C7, mouse, Cedi Diagnostic) was a marker for microglia/macrophages [67], GFAP (Z0334, rabbit, Dako) and vimentin (H5, mouse, DSHB) were markers for Muller glia. Sox-2 (Y-17, goat, Santa Cruz), Sox-9 (AB5535, rabbit, Chemicon), and transitin (EAP3, mouse, Developmental Studies Hybridoma Bank) were used to identify NIRG cells and Muller glia. DRAQ-5 (Biostats Limited) was used as a counter stain for nuclei to determine of outer nuclear layer thickness.

**Image Analysis**

Photomicrographs were obtained using a Leica DM5000B fluorescent microscope and Leica DC500 digital camera. Confocal images were obtained using a Zeiss LSM 510 at the Hunt-Curtis Imaging Facility. Image analysis was previously described [34]. Images were optimized for color, brightness, and contrast, and double-labeled images overlaid by using Adobe PhotoshopTM 6.0. Cell counts were made from at least three different animals at the area of maximal detachment, and means and standard deviations calculated on data sets. To minimize the variability of region-specific differences within the retina, cell counts were made from the same region of retina for control and retinal detachment areas. Immunofluorescence was quantified by using Image-Pro 6.2. Identiﬁcal illumination, microscope, and camera settings were used to obtain images for quantiﬁcation. Fixed areas were sampled from 5.4 megapixel digital images. These areas were randomly sampled across all retinal layers in areas of detachment or similar control retina areas. For CD45 staining, the retina was selected as the region of interest from 20X field of view measuring 376,441 m². The total area of retinal CD45 immunofluorescence was calculated for regions with pixel intensities above a determined threshold and averaged per group. The average density was calculated as the mean pixel value above threshold within threshold-designated regions. The density sum was calculated as the total pixel values for all pixels within threshold-designated regions. For TUNEL measurements, the number of immunofluorescent cells above threshold was determined for regions with pixel intensities above a determined threshold using Image-Pro 6.2. For outer nuclear layer thickness measurements, Image-Pro 6.2 was used to measure the height of the outer nuclear layer averaging evenly-spaced measurements taken per 20X field [71]. These calculations were determined for regions sampled from at least three different individuals for each experimental condition.

**Statistical Analysis**

To determine whether differences in outer nuclear layer thickness, TUNEL-positive cells, and microglia in the detached retina at different timepoints were significant, analysis of variance (ANOVA) was first performed to confirm a difference in measurements with p<0.05 considered significant. Subsequent least squares means testing with Bonferroni correction was performed (p<0.007 was considered significant, with 7 conditions compared for ONL thickness and TUNEL, and p<0.0125 for microglia analysis with 4 conditions compared (RD to controls)). SAS v9.2 from the SAS Institute Inc. (Cary, NC) was used for analyses.

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**Author Contributions**

Conceived and designed the experiments: CMC AJF. Performed the experiments: CMC CPZ MAS AB SR AJF. Analyzed the data: CMC CPZ MAS AB SR AJF. Wrote the paper: CMC AJF.

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