
Peeking into Sigma-1 Receptor Functions Through the Retina

19

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Abstract

This review discusses recent advances towards understanding the sigma-1 receptor (S1R) as an endogenous neuro-protective mechanism in the retina, a favorable experimental model system. The exquisite architecture of the mammalian retina features layered and intricately wired neurons supported by non-neuronal cells. Ganglion neurons, photoreceptors, as well as the retinal pigment epithelium, are susceptible to degeneration that leads to major retinal diseases such as glaucoma, diabetic retinopathy, and age-related macular degeneration (AMD), and ultimately, blindness. The S1R protein is found essentially in every retinal cell type, with high abundance in the ganglion cell layer. Ultrastructural studies of photoreceptors, bipolar cells, and ganglion cells show a predominant localization of S1R in the nuclear envelope. A protective role of S1R for ganglion and photoreceptor cells is supported by in vitro and in vivo experiments. Most recently, studies suggest that S1R may also protect retinal neurons via its activities in Müller glia and microglia. The S1R functions in the retina may be attributed to a reduction of excitotoxicity, oxidative stress, ER stress response, or inflammation. S1R knockout mice are being used to delineate the S1R-specific effects. In summary, while significant progress has been made towards the objective of establishing a S1R-targeted paradigm for retinal neuro-protection, critical questions remain. In particular, context-dependent effects and potential side effects of interventions targeting S1R need to be studied in more diverse and more clinically relevant animal models.

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Keywords

Sigma-1 receptor • Retinal diseases • Retinal ganglion neurons • Müller glia • Photoreceptors • Neuro-protection

19.1 Introduction

Earlier pharmacological profiling revealed two subtypes of sigma receptors (S1R and S2R) [1]. The S1R sequence has been cloned [2], while the identity of S2R remains unknown [3–5]. Despite numerous studies since its discovery [6, 7], S1R remains mysterious. Outstanding questions include the following: (1) *Identity*. No homolog of S1R is found in mammalian genomes. Curiously, the only protein that shares >30 % sequence identity with S1R is the yeast C-8,7 sterol isomerase [8]. But S1R is not found in yeast and it does not possess sterol isomerase activity. While an NMR structure of partial S1R was recently reported [9], an atomic structure of the whole protein has yet to be unveiled. (2) *Function*. In contrast to its unique identity, S1R is ubiquitously distributed, with high abundance in the central nervous system and liver [8]. Paradoxically, while S1R knockout mice do not exhibit overt phenotypes [10], S1R is linked to an array of pathological conditions such as cancer and neurological disorders (see review) [11]. These studies were conducted mostly using S1R ligands with only a handful employing S1R knockout mice. Hence, the S1R specificity of observed ligand functions awaits further investigations in knockout animals. (3) *Endogenous ligands*. Many synthetic ligands bind to S1R, including a few that have been intensively used for investigating S1R functions (see review) [12]. However, the identity of the true endogenous S1R ligand remains unclear. Several naturally occurring compounds show affinity for S1R, including steroids [13], trace amine [14], and lipids [15–17], but their S1R-specific roles are largely unknown.

Recently, there has been a surge of interest in S1R. In particular, important progress has been made to unravel its important role in the nervous system. A potential neuro-protective function of

S1R is found in animal models of major neurodegenerative diseases including Alzheimer's disease [18], Parkinson's disease [19], amyotrophic lateral sclerosis [20, 21], as well as retinal degenerative diseases [22–25]. The retina presents an excellent model system for studying S1R functions in the central nervous system. The main advantages include the following: (1) The retina is integral to, yet isolated from the brain, thus conveniently accessible for experimentation. (2) Animal models are available for major retinal degenerative diseases. (3) Despite being a thin sheet of tissue, the retina contains diverse cell types including neurons, epithelial cells, macroglia and microglia (see review) [26]. (4) Retinal cells are exquisitely organized into distinct layers [26], and hence advantageous for morphological and pathophysiological investigations (for example, see Fig. 19.1). (5) Since the eye is an immunologically privileged organ [27], immunogenic concerns caused by introducing experimental or therapeutic agents are relatively minor. In spite of a limited number of publications on S1R in the retina, progress has been achieved in identifying neuro-protective functions of S1R. While excellent reviews are available for studies of S1R in the nervous system in general [8, 11, 28], an overview is lacking for studies on S1R specifically in the retina. Here we discuss recent findings on the distribution, function, and molecular mechanisms of S1R in the mammalian retina.

19.2 General Molecular Functions of S1R

Mammalian S1R is a protein of 223 amino acids, with two transmembrane helices and a hydrophobic C-terminal region that putatively form a ligand binding pocket(s) (see review) [12]. An N-terminal double-arginine sequence serves as an endoplasmic reticulum (ER) retention motif.

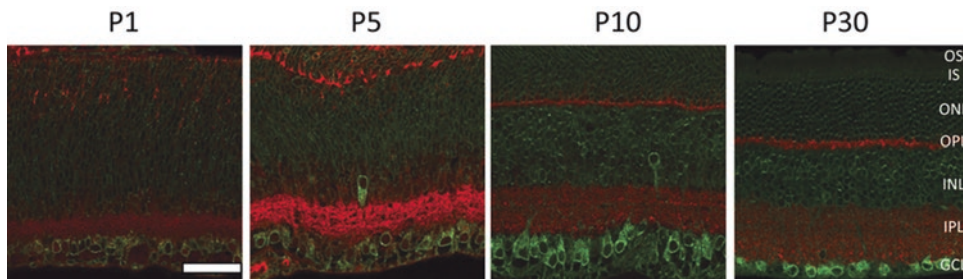


Fig. 19.1 Immunostaining of S1R on mouse retinal sections at different postnatal stages. Green, S1R; red, synaptophysin. OS outer segment, IS inner segment, ONL outer nuclear layer, OPL outer plexiform layer, INL inner

nuclear layer, IPL inner plexiform layer, GCL ganglion cell layer. Scale = 50 μ m (Adapted from: Scientific Reports. 2015.2;5:10689)

Until the discovery of its ligand-operated chaperone function [29], molecular functions of S1R were not known. This study revealed that activated S1R regulates mitochondrial calcium homeostasis by stabilizing the IP3 type 3 receptor at ER/mitochondria contacts. Thus S1R is thought to support cell survival [29]. Follow-up studies suggest that S1R is a multitasking protein involved in a broad range of cellular activities. S1R has been reported to modulate the activity of various, e.g., Na⁺, K⁺, Ca²⁺, Cl⁻, ion channels, likely via direct interactions [30] (see review) [28]. S1R was also reported to interact with G-protein coupled receptors (see review) [12]. Recently, evidence showed that S1R is involved in autophagy [31, 32]. In accordance, S1R also participates in ER stress responses; e.g., S1R interacts with and stabilizes ER stress sensor IRE1 [29, 33, 34]. While interactions of S1R with several lipids were observed earlier (see review) [12], most recently the Su group reported that S1R transports myristic acid to support proper tau phosphorylation and axon extension [17]. They also found that S1R modulates transcriptional activities via interaction with a nuclear envelope protein [35], consistent with the presence of S1R in the nuclear envelope [36, 37].

Despite continuous discoveries about the molecular biology of S1R, it remains an open question as to whether there is a common thread connecting S1R functions. In other words, can all S1R actions be attributed to its chaperone activity? The S1R C-terminal half is believed to be

responsible for its chaperone activity [29, 38]. If the C-terminus of S1R is confined in the ER lumen, how would S1R functionally interact with cytosolic proteins? What are the functions of the other S1R domains, for instance, the central loop region proposed to be cytosolic? It was reported that ligand binding to S1R alters its monomeric/oligomeric states [39–41]. How is this functionally related? In sum, many intriguing questions remain, which would inspire new investigations to help understand S1R functions in the retina as well as other systems.

19.3 Cellular and Sub-cellular Distribution of S1R in the Retina

The neural retina is a sheet of light-sensitive tissue in the back of the eye. Its intricate structure contains three layers of neatly organized neurons: the outer nuclear layer (ONL), the inner nuclear layer (INL), and the ganglion cell layer (GCL). Sandwiched in between are two synaptic layers connecting neurons: the outer plexiform layer (OPL) and the inner plexiform layer (IPL) [26] (Fig. 19.1). The neural retina rests on a nourishing single layer of pigmented cells called retinal pigment epithelium (RPE). ONL is formed by photoreceptors. INL contains bipolar cells, horizontal cells, and amacrine cells. GCL is mainly composed of ganglion cells and displaced amacrine cells. Müller glia cells traverse the entire neural retina and are interconnected with retinal

microglia cells [42]. Vision begins at photoreceptors, which convert light signals into chemical signals and then electrical impulses. Filtered through secondary neurons in INL, the electrical signals are transmitted to ganglion cells, and sent further into the brain through their axons (optic nerve) to be processed into images or other forms of vision.

While differences may exist among species, S1R is found in all cellular layers in the mouse retina, including GCL, INL, ONL, and RPE, as detected by both in-situ hybridization and immunohistochemistry [23, 37, 43, 44]. The specificity of S1R immunostaining is confirmed by the lack of S1R-positive staining in the retina of S1R knockout mouse [23]. S1R is abundant in GCL in mouse [23, 43], rat [45], monkey, pig, and human retinas [37]. In contrast, staining of S1R is less intense in ONL, unclear in the photoreceptor inner segment, and not detectable in the outer segment [23]. Consistently, using immunoelectron microscopy (EM) we did not observe S1R-positive staining on outer segment membrane discs, or in mitochondria or the ER which are concentrated in the photoreceptor inner segment [37] (Fig. 19.2). Rather, S1R is exclusively localized in photoreceptor nuclear membranes. Similarly in bipolar cells, EM data show that S1R is predominantly present in the nuclear envelope, in both outer and inner nuclear membranes [37]. In ganglion cells, S1R is found not only in nuclear membranes, but also in the ER and lipid droplets (Fig. 19.3). Interestingly, in accordance with S1R localization in nuclear membranes in retinal neurons [37] and Müller cells [36], Tsai et al. reported that S1R influences gene transcription by interacting with the nuclear envelope protein emerin to recruit chromatin-remodeling proteins [35].

In spite of new findings, perplexing questions remain about the distribution of S1R in the retinal neurons. For example, S1R was identified as a molecular chaperone functioning at the ER/mitochondria junction [29]. What is the function of S1R in photoreceptor cells, where it is found neither in the ER or mitochondria? S1R has been reported to interact with multiple ion channels including NMDA receptors in the ganglion cell plasma membrane [46, 47]. However, EM data do not show S1R in the plasma membrane of reti-

nal neurons [37]. Is it possible some channels in the plasma membrane interact with S1R localized in the subsurface ER cisternae [20, 37]? Moreover, S1R expression in the embryonic (E16) mouse retina is barely detectable but continuously increases during development until a mature retina is formed [37]. Is there a possible link between the temporal S1R distribution and retinal development?

19.4 Functions of S1R in the Retina

High-affinity S1R-selective agonists, e.g., (+)-pentazocine, PRE084, SK10047, and antagonists, e.g., NE100, BD1047, BD1063, present convenient pharmacological tools for studying S1R functions in the retina [12]. Using S1R ligands to treat animals (or cells) and whole retina samples for analysis, early studies suggested a neuro-protective role of S1R in the retina [43, 48–52]. Taking advantage of the layered retinal structure that partitions different neurons, in recent studies investigators analyzed cell type-specific S1R functions in the retina (see details in the subsections below). Moreover, S1R knockout mice [10] and retinal disease models provide powerful genetic tools for delineating S1R-specific functions in a given disease or cell type. However, mechanistic studies using isolated retinal neurons, in particular, photoreceptor cells, are challenging, as these highly specialized neurons cannot maintain their physiology and viability in cell culture. While immortalized cell lines are often used to represent corresponding retinal neurons, they are very dissimilar to mature native neurons in morphology and pathophysiology. Moreover, they may be associated with identity complications, e.g., the RGC-5 cell line [53]. Likely because of available methods to culture primary ganglion cells and their high S1R abundance [37, 43, 44], a majority of S1R functional studies in the retina have focused on ganglion neurons and associated disease conditions. Studies have also been extended to other cell types, e.g., Müller glia and microglia. In the following subsections, published studies on each retinal cell type will be discussed.

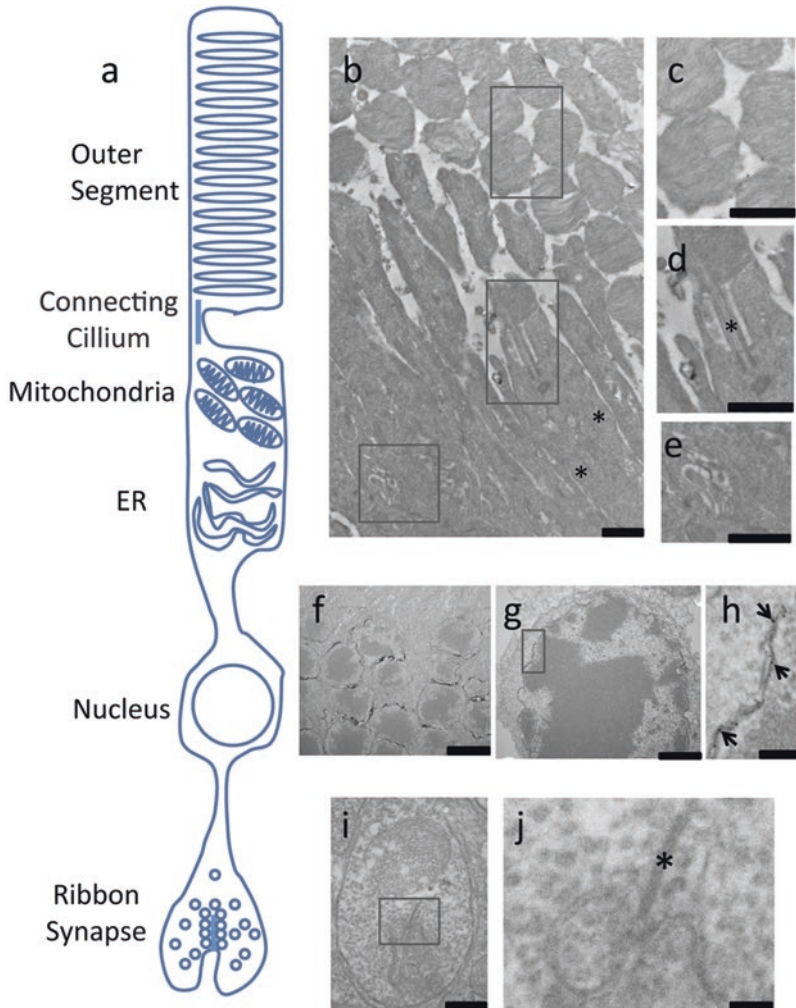


Fig. 19.2 Electron microscopy images showing S1R distribution in the mouse photoreceptor subcellular compartments. (a) Schematic of the compartments in the photoreceptor. (b) Ultrastructure of outer and inner segment. Asterisks label mitochondria. (c–e) Magnified images of the boxed areas in (b), showing the outer segment containing membrane discs, the connecting cilium (asterisk), and the inner segment (including ER), respectively. (f–h), Localization of S1R in the nuclear envelope.

(f) nuclear region of several photoreceptor cells; (g) nuclear envelope of a single cell, (h) magnified box area in (g) showing S1R localization in the outer and inner membranes of the nuclear envelope (pointed to by arrows). (i) and (j), Photoreceptor synaptic terminal. The image in (j) is a magnified box area in (i) revealing the characteristic ribbon (asterisks) and vesicles. Scales: (b–e) and (g), 1 μm ; (f), 3 μm ; (h), 0.2 μm ; (i), 0.5 μm ; (j), 0.1 μm (Adapted from: Scientific Reports. 2015.2;5:10689)

19.4.1 Retinal Ganglion Cells (RGCs)

RGCs make the functional link between the retina and the brain. Approximately 20 subtypes of RGCs process complex visual information collected from bipolar cells and amacrine cells, and then send it to the brain as action potentials along RGC axons (see review) [26]. As such, RGCs play a critical role in vision, and their deteriora-

tion leads to vision loss or impairment. A good example is glaucoma, a prevalent retinal disease characterized by final-stage RGC loss and consequent visual field deficits (see review) [54]. Although no data is available with regard to S1R expression in each specific RGC subtype, it is conceivable that S1R is ubiquitously expressed, based on S1R-positive staining in essentially all GCL cells [23, 37, 43].

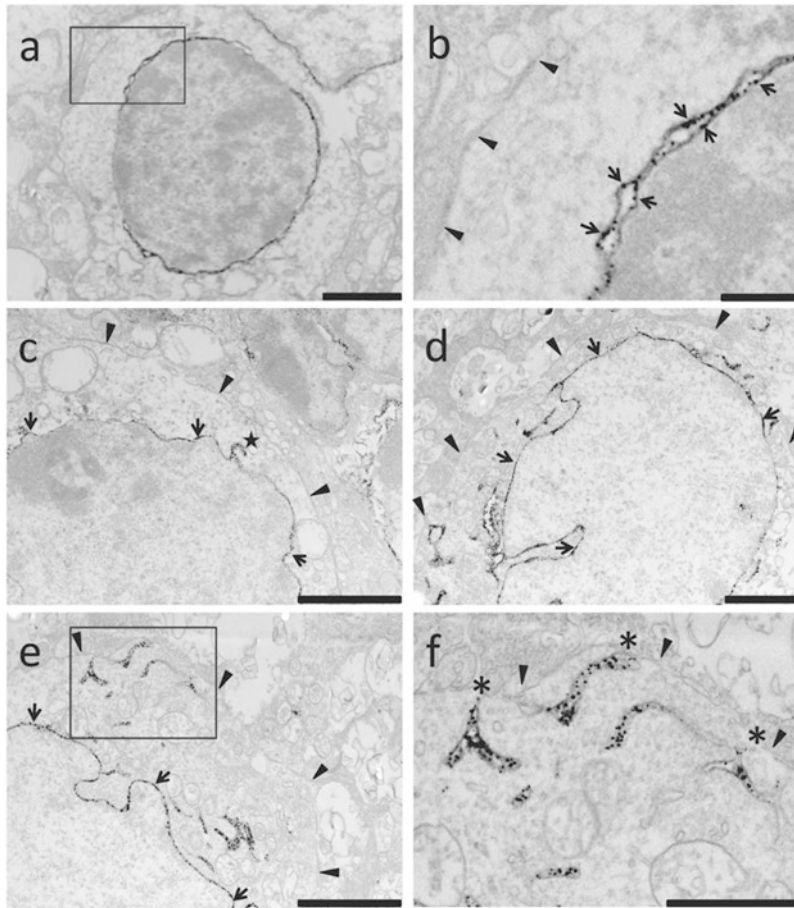


Fig. 19.3 Subcellular localization of S1R in bipolar and ganglion cells of the mouse retina. (a–c), Bipolar cells. (b) shows magnification of the boxed area in (a). Arrows point to S1R immunolabeling in the inner and outer membranes of the nuclear envelope. Arrowheads mark the plasma membrane. (c) shows S1R localization in the ER membrane (*star*) connected to the nuclear envelope (*arrows*). (d–f), Ganglion cells. (d) shows predomi-

nant S1R localization in the nuclear envelope (*arrows*) but not in the plasma membrane (*arrow heads*). (e) highlights the presence of S1R in the ER (*boxed area*). (f) shows the magnification of the boxed area in (e), revealing S1R localization in the ER cisternae (*asterisks*) that are adjacent to the plasma membrane (*arrow heads*). Scales: (a), (c–e), 2 μm ; (b), (f), 0.25 μm (Adapted from: Scientific Reports. 2015.2;5:10689)

In vitro and in vivo studies from several research groups support a pro-survival role of S1R in RGCs. Using both primary mouse RGCs and a RGC-5 cell line, the Smith group showed that the S1R-specific agonist (+)-pentazocine protected against apoptosis induced by homocysteine or glutamate. The mechanism was attributed to the attenuation of excitotoxicity, which was mediated by the NMDA receptor [55, 56]. In a recent study, they observed that (+)-pentazocine also protected RGC-5 cells against oxidative stress; this effect was associated with down-regulation of ER stress proteins [57]. Using

whole-cell patch clamp on RGC-5 cells, the Yorio group found that the S1R agonist SKF10047 promoted cell survival by inhibiting apoptosis-provoking Ca^{2+} influx mediated by the L-type Ca^{2+} channel [58, 59]. In purified rat primary RGCs, they were able to recapitulate the inhibitory effect of S1R activation on Ca^{2+} influx and a possible S1R/L-type channel interaction [60]. Their latest work showed that S1R protected RGCs in vitro against ischemic damage via ERK activation [61]. In an ex vivo study using patch clamp on rat retinal slices, the Yang group observed suppression of NMDA receptor-specific current

responses in both ON and OFF types of RGCs following S1R activation [47]. Their data further suggested that this effect was mediated through a Ca^{2+} -dependent PLC-PKC pathway. In sum, all the foregoing in vitro studies suggest a protective role of activated S1R in RGCs, via attenuation of oxidative stress, excitotoxicity, or Ca^{2+} toxicity involving ion channels. At present it is not clear whether these S1R actions are orchestrated in RGCs under cellular stresses.

In an in vivo study using a spontaneous diabetic retinopathy mouse model, the Smith group

identified a prominent anti-oxidative effect of S1R activation [22]. Treating animals with (+)-pentazocine injection preserved the thickness of IPL and INL, cell number in GCL, as well as organization of Müller glia. Demonstrating a specific role of S1R in ganglion cell neuro-protection, the Guo group reported that cell loss in GCL was significantly faster in S1R knockout mice compared to wild type control after optic nerve crush, an acute glaucoma model [23] (Fig. 19.4). This observation was echoed by another study using S1R knockout mice from the Smith group [62].

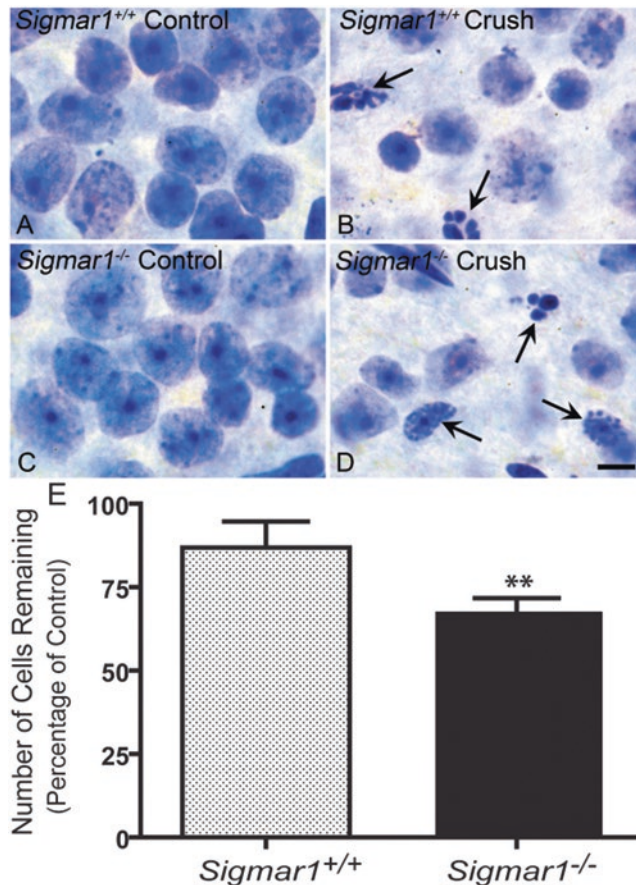


Fig. 19.4 Comparison of the post-crush cell loss in the retinal ganglion cell layer between WT and *Sigmar1*^{-/-} (S1R knockout) mice. (a–d) Nissl-stained retinal whole-mounts from WT (a and b) and *Sigmar1*^{-/-} (c and d) mice. Images were from representative fields (1000×) of the mid-peripheral inferior retinas of 12-month-old mice. For each mouse, while the *right eye* served as untreated control (a and c), the *left eye* was treated by optic nerve crush for 3 s (b and d). Retinal whole-mounts were prepared 7 days after surgery, and the side of the ganglion cell layer was stained. Healthy ganglion cells exhibited larger somas

and nuclei with prominent nucleoli. *Arrows* point to apoptotic cells. (e) Quantification of cells remaining in the retinal ganglion cell layer 1 week after surgery. The number of remaining cells in the experimental eye is represented as a percentage of the untreated control. The data were pooled from three WT and *Sigmar1*^{-/-} pairs of 6-month-old mice and two pairs of 12-month-old mice. There were 86.82 ± 7.90 % (mean ± standard deviation [SD], n = 5) cells remaining in WT mice and 68.31 ± 3.36 % remaining in *Sigmar1*^{-/-} mice. ** *t*-test, p = 0.0013 (Adapted from Mol Vis. 2011;17:1034–1043)

While retinal morphology and electroretinogram (ERG) appeared normal in younger S1R knockout mice, decrease of ERG b-wave amplitudes and GCL nuclei number, as well as disrupted axon structure in the optic nerve head, occurred in S1R knockout mice compared to wild type at 12 months of age. Moreover, using S1R knockout mice, they confirmed a S1R-specific neuro-protective function of (+)-pentazocine in an induced diabetic mouse model [63]. Taken together, these studies support an important role of S1R in alleviating RGC stress and degeneration in RGC disease models.

Given the complexity of the pathophysiology of retinal neuro-degeneration, it is a daunting challenge to delineate the molecular mechanisms of S1R-specific neuro-protection for RGCs. Since primary, mature RGCs do not divide and hence they cannot be expanded in cell culture, it is difficult to perform *in vitro* mechanistic studies using these cells. As RGC-5 which was long used as an RGC line recently proved false [53], it is imperative to establish an appropriate RGC line, for *in vitro* mechanistic research. Moreover, further investigation is needed to better correlate *in vitro* mechanisms to *in vivo* pathophysiology. To better understand the therapeutic potential of targeting S1R for interventions, in particular for treating chronic diseases such as glaucoma, more clinically relevant animal models, e.g., DBA/2J [54], may be utilized. To this end, local drug delivery methods integrating advanced bioengineering technologies would provide new insights and opportunities.

19.4.2 Müller Glia and Microglia

RGCs and Müller glia are closely situated, facilitating their functional interactions in RGC pathophysiology (see review) [42]. In a retinal transcriptome survey, Ha et al. did not find significant changes of ER stress genes in neural retinas isolated from S1R knockout mice compared to wild type control. Interestingly, however, marked expression changes of those genes were observed in Müller cells isolated from knockout versus wild type mice [24]. This finding implies an important role of Müller cells in

previously observed S1R-mediated protection of RGCs.

Müller cells are a major glial cell type in the retina where they serve as anatomical conduits between neurons and their environment [42]. Müller cells are radially oriented, spanning the entire thickness of the retina from the inner limiting membrane to the outer limiting membrane. Studies suggest that Müller cells play essential roles in the retina (see review) [64]. In addition to supporting the structural integrity of the retina, they maintain retinal homeostasis by participating in essential processes such as glucose metabolism, antioxidant production, ion/substrate exchange, and vascular regulation. Müller cells, together with astrocytes and microglia, become reactive in retinal diseases [27].

In a recent report, the Smith group observed an increase of LPS-stimulated secretion of inflammatory proteins from Müller cells isolated from S1R knockout mice versus those from wild type control [65]. Furthermore, (+)-pentazocine treatment of Müller cells inhibited the secretion of inflammatory proteins and NF κ B translocation to the nucleus. In a follow-up study, they found that Müller cells from S1R knockout mice compared to wild type cells manifested more severe oxidative stress, which could be explained by suppressed NRF2 signaling and impaired function of an L-cysteine/L-glutamate antiporter (system xc⁻) [66]. These studies uncovered an essential role of S1R in the suppression of oxidative stress and inflammation in retinal Müller glia. Reporting a different S1R action, Vogler et al. showed that PRE084 mitigated osmotic swelling of Müller cell somas induced by superfusion of rat retinal slices with a hypo-osmotic solution [67]. This S1R effect was likely mediated through activation of a glutamatergic-purinergic signaling cascade known to prevent osmotic Müller cell swelling. Astrocytes are another type of retinal glia that are most abundant in the optic nerve head [64]. To our knowledge, there is no report investigating S1R function in this specific cell type in the retina.

Recently, S1R protein was also found in retinal microglia. Pretreatment of isolated microglia with (+)-pentazocine reduced LPS-stimulated morphological change, intracellular ROS pro-

duction, and secretion of inflammatory cytokines (TNF- α , IL-10, MCP-1). The (+)-pentazocine effects were blocked by S1R antagonist BD1063, suggesting a S1R-specific function [68]. These S1R-mediated responses likely involved suppression of the ERK/JNK MAPK pathway due to S1R activation.

Together, the foregoing reports have brought about a new perspective that S1R may protect RGCs through their functions in Müller glia and/or microglia. They also raise an interesting scenario in which the mechanisms of S1R-mediated retinal neuro-protection are multifactorial, likely involving both neuronal and non-neuronal cells and their interactions. An ensuing question is whether retinal glia or microglia cells can serve as effective therapeutic targets. These cells could play opposite roles. Whereas they are essential for maintaining retinal neuron homeostasis [64], when activated by stress conditions, they may transform into inflammatory cells causing harm to retinal neurons. On the other hand, these cells can be readily isolated from the retina, an advantage for *in vitro* experimental models. Nonetheless, more studies are warranted to understand their role in retinal neuro-degeneration and protection, in the context of specific S1R-associated regulations.

19.4.3 Bipolar Cells, Horizontal Cells, and Amacrine Cells

The nuclei of bipolar, horizontal, and amacrine cells are all in INL, which is situated in between the photoreceptor layer (ONL) and GCL. In mammalian retinas there are approximately a dozen types of bipolar cells, three types of horizontal cells, and 30 types of amacrine cells (see reviews) [26, 54]. Bipolar cells transfer visual signals either directly from photoreceptors to ganglion cells or indirectly through horizontal cells and amacrine cells. Whereas horizontal cells transmit (and modulate) the visual information from photoreceptors to bipolar cells, amacrine cells modulate the signals transmitted from bipolar cells to RGCs. Although S1R distribution in each subtype of the secondary neurons has not been completely delineated [45], immunostain-

ing shows S1R presence in majority of INL cells [23, 43, 45].

Because of a paucity of experimental evidence, the function of S1R in bipolar cells is not known. Vogler et al. reported that S1R activation protects against osmotic swelling of Müller cells, but not of bipolar cells [67]. On the other hand, one-year old S1R knockout mice showed reduced amplitudes of ERG b-wave, which measures the activity of the inner retinal neurons including bipolar cells [62]. Since S1R is found in bovine photoreceptor presynaptic terminals [37], it is tempting to speculate that S1R may modulate neurotransmission to postsynaptic bipolar cells under some circumstances. There is no data available about the function of S1R in horizontal and amacrine cells. Thus, more research is needed to explore S1R functions in these secondary neurons in visual signal transmission. Such information would provide important insight into possible side effects, e.g., disturbance of synaptic transmission, of S1R-targeted interventions.

19.4.4 Photoreceptor Cells

Photoreceptors are highly specialized neurons. Through a biochemical process of phototransduction, they are capable of converting light signals into nerve impulses that eventually lead to vision (see review) [26]. There are two basic types of photoreceptors, rods and cones, each containing four morphologically and functionally distinct compartments. Rods are extremely light sensitive and responsible for night vision; cones respond to bright light and are responsible for day vision and color vision. Photoreceptors are highly susceptible to genetic defects, as well as insults from their environment. There are up to 100 photoreceptor gene loci that cause retinal diseases such as retinitis pigmentosa, a condition characterized by photoreceptor cell death (RetNet). While S1R is found in the nuclear envelope of photoreceptor cells [37], its function is not clear. Most recently, the Hara group demonstrated the importance of S1R for photoreceptors [25]. Using a 661W neuronal cell line, they found that high-affinity S1R ligand cutamesine (named SA4503) attenuated light-induced dis-

ruption of mitochondrial membrane potential and caspase-3/7 activation. Moreover, using a light-induced photoreceptor degeneration model of mice carrying a mutation in RPE65 (an RPE specific protein), cutamesine delivered by intravitreal injection partially rescued light-induced retinal dysfunction (reduced ERG) and ONL thinning. The cutamesine effect could be blocked by S1R antagonist BD1063, suggesting it was S1R specific. As photoreceptors and RPE cells are structurally and functionally dependent on each other [69], it remains unclear which cell type is the primary site of the observed S1R protective function. Of note, mechanisms of photoreceptor degeneration vary in different pathological contexts [70], and so may S1R function. Whether S1R activation is ubiquitously beneficial in the retina awaits more careful testing. Moreover, the predominance of S1R in the photoreceptor nuclear envelope raises an interesting question with regard to possible mechanisms of S1R-specific protection in photoreceptor cells. Therefore, different retinal degeneration models may be used in future experiments to comprehensively understand the role of S1R in photoreceptor pathophysiology.

19.4.5 Retinal Pigment Epithelium (RPE) Cells

RPE is a single layer of cells situated between the light-sensitive outer segments of photoreceptors and the choroid blood supply [26]. RPE possesses many functions essential to the visual process, the chief of which is to maintain photoreceptor homeostasis. Analyses of hereditary types of retinal degeneration reveal a strong dependence of RPE on photoreceptors and *vice versa*. Defects in RPE contribute to initiation and/or progression of AMD in humans (see review) [69]. Characterized by the loss of central vision, AMD is the leading cause of blindness in elderly populations, and no pharmacological treatment is available. Oxidative damage is considered as a major factor for disease onset and progression.

In situ hybridization indicated the presence of S1R mRNA in RPE [43]. However, its protein

abundance and subcellular distribution in RPE cells remain unclear, partly because of the intense auto-fluorescence that masks specific S1R immunostaining. Nevertheless, in an earlier study using a human RPE cell line (ARPE-19) and adult human primary RPE cells, Bucolo et al. were able to reduce H₂O₂-induced DNA damage and cell loss by pre-treatment with PRE084, an effect abolished by S1R antagonists [71]. Most recently, using targeted siRNA screening in a human RPE1 cell line, MacVicar et al. identified S1R as a potential regulator of autophagosome homeostasis involving mitochondrial dynamics [31]. Autophagy is an important stress response pathway responsible for the removal and recycling of damaged or redundant cytosolic constituents. While autophagy is found to be an active process in the RPE in vivo [72], evidence from AMD donors indicates a decline of autophagic flux in the RPE [73]. Although not yet specifically investigated in RPE cells in vivo, S1R has been reported to influence autophagy in vitro [31, 32, 74]. A possible protective role of the S1R via autophagic regulations in RPE cells needs to be further determined experimentally. In light of an anti-oxidative function of S1R and its involvement in lipid metabolism, it appears reasonable that S1R may play a role in maintaining homeostasis of RPE cells, which are situated in a highly oxidative environment to process large amounts of lipids from phagocytosed photoreceptor membrane discs [69].

19.5 Concluding Remarks

The retina, which is composed of diverse cell types, represents a favorable model for studying the functions of S1R. In the past decade, considerable progress has been made in understanding the role of S1R in retinal degenerative diseases. While most efforts have been devoted to retinal ganglion neurons, reports on S1R in other retinal cell types are emerging. These studies generally support a protective role of S1R against stress-induced cell loss. To exploit the therapeutic potential of a S1R-targeted strategy for treating retinal diseases, more studies are required, particularly in the following areas: (1) Investigation using more diverse

pre-clinical retinal disease models for a comprehensive understanding of S1R functions. (2) Delineation of S1R-specific and non-specific effects of S1R-binding drugs, using S1R knockout animals or cells. (3) Determination of the cellular and molecular mechanisms of S1R-mediated retinal neuro-protection. (4) Evaluation of combination therapies using S1R-targeting ligands and drugs targeting other pathways. A deeper understanding of S1R-specific functions and mechanisms in the retina would lead to new therapeutic opportunities, not only for retinal diseases but also other related disorders.

Acknowledgement This work was supported by the National Eye Institute grant R01EY022678 and a Morgridge Institute for Research & the James Christenson Estate Macular Degeneration Research Award (to L-W Guo), and P30EY016665 and S10OD018221 (to the University of Wisconsin Vision Core). The project was also supported by the Clinical and Translational Science Award (CTSA) program, through the NIH National Center for Advancing Translational Sciences (NCATS), grant UL1TR000427. The content is solely the responsibility of the authors and does not necessarily represent the official views of the NIH.

We thank Drs. Miles L. Epstein and Arnold E. Ruoho for critical comments. We also thank Dr. Laura Hogan at the University of Wisconsin Institute for Clinical & Translational Research (ICTR) for editing and proof-reading.

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