

# Expression, Subcellular Localization, and Regulation of Sigma Receptor in Retinal Müller Cells

Guoliang Jiang,<sup>1</sup> Barbara Mysona,<sup>1</sup> Ying Dun,<sup>1</sup> Jaya P. Gnana-Prakasam,<sup>2</sup> Navjotsin Pabla,<sup>1</sup> Weiguo Li,<sup>1</sup> Zheng Dong,<sup>1</sup> Vadivel Ganapathy,<sup>2</sup> and Sylvia B. Smith<sup>1,3</sup>

**PURPOSE.** Sigma receptors ( $\sigma$ Rs) are nonopioid, nonphencyclidine binding sites with robust neuroprotective properties. Type 1  $\sigma$ R1 ( $\sigma$ R1) is expressed in brain oligodendrocytes, but its expression and binding capacity have not been analyzed in retinal glial cells. This study examined the expression, subcellular localization, binding activity, and regulation of  $\sigma$ R1 in retinal Müller cells.

**METHODS.** Primary mouse Müller cells (MCs) were analyzed by RT-PCR, immunoblotting, and immunocytochemistry for the expression of  $\sigma$ R1, and data were compared with those of the rat Müller cell line (rMC-1) and the rat ganglion cell line (RGC-5). Confocal microscopy was used to determine the subcellular  $\sigma$ R1 location in primary mouse MCs. Membranes prepared from these cells were used for binding assays with [<sup>3</sup>H]-pentazocine (PTZ). The kinetics of binding, the ability of various  $\sigma$ R1 ligands to compete with  $\sigma$ R1 binding, and the effects of donated nitric oxide (NO) and reactive oxygen species (ROS) on binding were examined.

**RESULTS.**  $\sigma$ R1 is expressed in primary mouse MCs and is localized to the nuclear and endoplasmic reticulum membranes. Binding assays showed that in primary mouse MCs, rMC-1, and RGC-5, the binding of PTZ was saturable. [<sup>3</sup>H]-PTZ bound with high affinity in RGC-5 and rMC-1 cells, and the binding was similarly robust in primary mouse MCs. Competition studies showed marked inhibition of [<sup>3</sup>H]-PTZ binding in the presence of  $\sigma$ R1-specific ligands. Incubation of cells with NO and ROS donors markedly increased  $\sigma$ R1 binding activity.

**CONCLUSIONS.** MCs express  $\sigma$ R1 and demonstrate robust  $\sigma$ R1 binding activity, which is inhibited by  $\sigma$ R1 ligands and is stimulated during oxidative stress. The potential of Müller cells to bind  $\sigma$ R1 ligands may prove beneficial in retinal degenerative diseases such as diabetic retinopathy. (*Invest Ophthalmol Vis Sci.* 2006;47:5576–5582) DOI:10.1167/iovs.06-0608

Sigma receptors ( $\sigma$ Rs) are nonopiate, nonphencyclidine binding sites<sup>1</sup> whose ligands have robust neuroprotective properties. The endogenous ligand and the physiological function of  $\sigma$ R1 have not been elucidated.  $\sigma$ Rs consist of several subtypes distinguishable by biochemical and pharmacologic means.<sup>2</sup> Among these, type 1  $\sigma$ R ( $\sigma$ R1) is best characterized.

The cDNA encoding  $\sigma$ R1 was cloned originally from guinea pig liver<sup>3</sup> and subsequently from human, mouse, and rat.<sup>4–7</sup> The  $\sigma$ R1 cDNA predicts a protein of 223 amino acids ( $M_r$ , 25–28 kDa).<sup>3</sup> Initial hydropathy analysis of the deduced  $\sigma$ R1 amino acid sequence suggested a single transmembrane segment.<sup>3,4,6</sup> Recently, Aydar et al.<sup>8</sup> showed that, when expressed in *Xenopus laevis* oocytes,  $\sigma$ R1 has two transmembrane segments with the NH<sub>2</sub> and COOH termini on the cytoplasmic side of the membrane.

$\sigma$ R1 distribution has been analyzed in brain, and its association with neurons is well established.<sup>9</sup> The receptors modulate ion channel activities at the plasma membrane, neuronal firing, and release of certain neurotransmitters. These receptors are of interest because of their profound capacity to prevent neuronal cell death. They inhibit ischemia-induced glutamate release,<sup>10,11</sup> attenuate postsynaptic glutamate-evoked Ca<sup>2+</sup> influx,<sup>12,13</sup> depress neuronal responsiveness to NMDA receptor stimulation,<sup>14–16</sup> and reduce NO production.<sup>17</sup>

In retina, the role of  $\sigma$ R1 recognition sites in ischemia-reperfusion injury in controlling intraocular pressure and in protection against glutamate-induced neurotoxicity has been reported.<sup>18–20</sup> Binding assays making use of bovine retinal membranes suggested the presence of  $\sigma$ Rs,<sup>21,22</sup> but the studies did not disclose in which retinal cell types  $\sigma$ Rs were present, nor did they establish unequivocally the molecular identity of the receptor. Recently, we used molecular and biochemical methods to study  $\sigma$ R in mouse retina and reported widespread expression of  $\sigma$ R1.<sup>23</sup> RT-PCR analysis amplified  $\sigma$ R1 in neural retina, RPE-choroid complex, and lens. In situ hybridization studies revealed abundant expression of  $\sigma$ R1 in the ganglion cell layer, inner nuclear layer, inner segments of photoreceptor cells, and RPE cells. Immunohistochemical analysis confirmed these observations. Subsequent studies focused on ganglion cells because of their vulnerability in diabetic retinopathy and revealed that  $\sigma$ R1 continues to be expressed under hyperglycemic conditions and during diabetic retinopathy,<sup>24</sup> making it a promising target for neuroprotection against cell death. More recent work using the rat ganglion cell line RGC-5 showed that (+)-pentazocine, a  $\sigma$ R1-specific compound, can block RGC-5 cell death induced by homocysteine and glutamate.<sup>25</sup>

Despite evidence that  $\sigma$ R1 is present in RGCs and other retinal cell types, no studies have been published of  $\sigma$ R1 binding activity in isolated retinal cells. In this study, we examined  $\sigma$ R1 binding in retinal Müller cells (MCs) using the rat cell line (rMC-1), primary MCs, and RGC-5 cells. The rationale for studying  $\sigma$ R1 binding activity in Müller cells is that recent studies have demonstrated a possible role for  $\sigma$ R in glial cell maintenance. Pharmacology studies showed that C6 glioma cells have  $\sigma$ R binding sites.<sup>26</sup> Immunohistochemistry studies demonstrated that, in addition to neurons,  $\sigma$ R1 is present in oligodendrocytes<sup>27</sup> and Schwann cells.<sup>28</sup> Hayashi and Su<sup>29</sup> confirmed  $\sigma$ R1 presence in oligodendrocytes and localized it to the endoplasmic reticulum (ER) forming galactosylceramide-enriched lipid rafts in the myelin sheet of mature oligodendrocytes. They speculated that  $\sigma$ Rs are important for oligodendro-

From the Departments of <sup>1</sup>Cellular Biology and Anatomy, <sup>2</sup>Biochemistry and Molecular Biology, and <sup>3</sup>Ophthalmology, Medical College of Georgia, Augusta, Georgia.

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Corresponding author: Sylvia B. Smith, Department of Cellular Biology and Anatomy, Medical College of Georgia, 1459 Laney-Walker Boulevard, CB 2820, Augusta, GA 30912-2000; sbsmith@mail.mcg.edu.

cyte differentiation and may play a role in the pathogenesis of certain demyelinating diseases.

Müller cells, the key retinal glial cell, span the retinal thickness, contacting and ensheathing neuronal cell bodies and processes. They are crucial role for neuronal survival and provide trophic substances and precursors of neurotransmitters to neurons.<sup>30</sup> Most retinal diseases are associated with reactive MC gliosis, which may contribute to neuronal cell death; hence, we sought to characterize  $\sigma$ R in these cells. Our earlier work with the rat MC line rMC-1 suggested that  $\sigma$ R1 mRNA was present in these cells.<sup>24</sup> In this study, we confirmed that finding and extended the analysis to MCs isolated from mouse retina (primary cell culture). We analyzed the subcellular localization of  $\sigma$ R in primary mouse MCs. Additionally, we used rMC-1, RGC-5, and primary mouse MCs to analyze the binding characteristics of  $\sigma$ R1. Our study represents the first comprehensive analysis of  $\sigma$ R1 in MCs and the first information about the binding characteristics of this receptor in any isolated retinal cell type.

## MATERIALS AND METHODS

### Reagents

Materials were obtained as follows: DMEM/F12, reagent (TRIZOL) and penicillin-streptomycin (Gibco-Invitrogen Corp., Grand Island, NY); fetal bovine serum (FBS; Atlanta Biologicals, Atlanta, GA); [1,3-<sup>3</sup>H] (+)-pentazocine (specific radioactivity, 37 Ci/mmol), [2,3-<sup>3</sup>H]-(+)-3-(3-hydroxyphenyl)N-(1-propyl)piperidine [<sup>3</sup>H]-(+)-3-PPP (specific radioactivity, 92.4 Ci/mmol) (DuPont-NEN, Boston, MA); IgG-horse radish peroxidase (HRP; Santa Cruz Biotechnology, Santa Cruz, CA); Cy-3 IgG, 488-IgG, and 568-IgG (Alexa Fluor; Invitrogen, Carlsbad, CA); monoclonal anti-lamin A and monoclonal anti-PDI (Abcam Inc., Cambridge, MA); polyclonal antivimentin (Chemicon International, Temecula, CA); enhanced chemiluminescence (ECL) detection kit (Pierce Biotechnology, Inc., Rockford, IL); Bio-Rad Protein Assay Reagent (Bio-Rad, Hercules, CA); commercial blocker (PowerBlock; BioGenex, San Ramon, CA); protease inhibitors (Complete Mini Protease Inhibitor Cocktail tablets; GeneAmp RNA PCR Kit; Applied Biosystems/Roche Molecular System, Branchburg, NJ); DNA polymerase (TaKaRa *Taq*; Takara Bio Inc., Otsu, Shiga, Japan); sense and antisense primers for rat and mouse  $\sigma$ R1 (Integrated DNA Technologies, Inc., Coralville, IA); (+)-3-PPP, carbapentane, 1,3-di-(2-tolyl)guanidine (DTG), haloperidol, and phenytoin (Research Biochemicals, Natick, MA); (+)-pentazocine, anti- $\beta$ -actin, 3-nitroso-N-acetylpenicillamine (SNAP), S-nitrosoglutathione (SNOG), 3-morpholininosydnonimine (SIN-1), hydrogen peroxide 30% (wt/wt) solution, xanthine, xanthine oxidase (X/XO), and all other chemicals (Sigma-Aldrich Chemical Co., St. Louis, MO). rMC-1 cells and RGC-5 cells were kind gifts, respectively, of Vijay P. Sarthy and Neeraj Agarwal. Anti-CRALBP was a generous gift of John Saari.

### Animals and Isolation of Müller Cells

C57Bl/6 mouse breeding pairs (Harlan Sprague-Dawley, Indianapolis, IN) were maintained in our colony in accordance with the ARVO Statement for the Use of Animals in Ophthalmic and Vision Research. Müller cells were isolated from 6- to 7-day-old mice following our published methods,<sup>31</sup> which were adapted from those of Hicks and Courtois.<sup>32</sup> Briefly, eyeballs were removed, placed in Dulbecco modified Eagle medium (DMEM) with gentamicin, and soaked for 3 hours at 25°C in the dark. Then they were rinsed in PBS and were incubated in buffer containing trypsin, EDTA, and collagenase. Retinas were removed from eyeballs (taking care to avoid contamination by pigmented RPE), placed in DMEM supplemented with glucose, FBS, and penicillin/streptomycin, and gently pipetted into small aggregates at a density of 10 to 16 retinas per dish. Isolated cells were detected within 1 to 3 days. By 3 to 5 days, substantial cell growth ensued. Cultures were washed vigorously with medium until only a strongly adherent flat cell population remained. Cells were passaged 1 to 3 days after washing

and were seeded into culture flasks (50,000 cells/cm<sup>2</sup>); culture media were changed three times per week. Purity of the cultures was verified using antibodies that are known markers of Müller cells (CRALBP, vimentin, glutamine synthetase, GLAST). Glial fibrillary acidic protein (GFAP), typically considered a marker of Müller cells under stress, was detected only at a low level.<sup>31</sup> Immunocytochemical studies using markers for neurons (neurofilament-L, a major component of neuronal cytoskeleton) and RPE (RPE-65) showed minimal detection.

### Cell Culture

Primary mouse MCs, rMC-1, and RGC-5<sup>31,33,34</sup> were cultured in DMEM/F12 supplemented with 10% FBS, 100 U/mL penicillin, and 100  $\mu$ g/mL streptomycin and were maintained at 37°C in a humidified chamber of 5% CO<sub>2</sub>. Medium was replaced every other day. On confluence, cultures were passaged by dissociation in 0.05% (wt/vol) trypsin in PBS.

### Semiquantitative RT-PCR Analysis of $\sigma$ R1 mRNA

Total RNA was prepared from confluent primary mouse MCs, rMC-1, and RGC-5 cells using reagent (TRIZOL; Gibco-Invitrogen Corp.). For rMC-1 and RGC-5, two cell lines derived from rat,<sup>33,34</sup> RT-PCR was carried out with primer pairs specific for rat  $\sigma$ R1<sup>7</sup>: sense, 5'-GTTCT-GACTATTGTGGCGGTGCTG-3'; antisense, 5'-CAAATGCCAGGGTA-GACGGAATAAC-3' (nucleotide positions 80–104 and 567–591; expected PCR product size, 512 bp). For the primary mouse MCs isolated from mouse, RT-PCR was carried out using primer pairs specific for mouse<sup>6</sup>: sense, 5'-CTCGCTGTCTGAGTACGTG-3'; antisense, 5'-AA-GAAAGTGTGGCTGCTAGTGCAA-3' (nucleotide positions 315–333 and 572–593; expected PCR product size, 279 bp). 18S RNA was the internal standard. RT-PCR was performed at 35 cycles, with a denaturing phase of 1 minute at 94°C, an annealing phase of 1 minute at 59°C, and an extension of 2 minutes at 72°C. Twenty microliters of the PCR products were gel electrophoresed and stained with ethidium bromide.

### Immunoblot Analysis of $\sigma$ R1

Immunodetection of  $\sigma$ R1 in RGC-5, rMC-1, and primary mouse MCs followed our published methods.<sup>24,35</sup> Protein samples were subjected to SDS-PAGE and were transferred to nitrocellulose membranes, which were blocked for 1.5 hours with Tris-buffered saline–0.05% Tween-20 containing 5% nonfat milk. Membranes were incubated with anti- $\sigma$ R1 antibody (1:1000)<sup>24</sup> overnight at 4°C, followed by incubation with HRP-conjugated goat anti-rabbit IgG antibody (1:5000). After washing, proteins were visualized with the ECL Western blot detection system. Membranes were washed three times, blocked with 5% nonfat milk for 2 hours, and reprobed with mouse monoclonal anti- $\beta$ -actin antibody (1:5000) as a loading control.

### Immunocytochemical Analysis of $\sigma$ R1

$\sigma$ R1 detection in RGC-5, rMC-1, and primary mouse MCs followed our published protocol.<sup>23–25,35</sup> Cells were incubated with polyclonal anti- $\sigma$ R1 antibody (1:100)<sup>24</sup> and subsequently with Cy-3-conjugated anti-rabbit IgG (1:200). Negative control experiments were performed by incubating the slides without the primary antibody.  $\sigma$ R1 was detected by epifluorescence (Axioplan-2 microscope and AxioVision program; Carl Zeiss, Göttingen, Germany). For  $\sigma$ R1 subcellular localization studies, primary mouse MCs were seeded on coverslips, grown for 24 hours, fixed in ice-cold methanol for 10 minutes, air dried, washed in PBS, blocked (PowerBlock; BioGenex), and incubated overnight at 4°C with the  $\sigma$ R1 antibody (1:100)<sup>24</sup> and either monoclonal anti-lamin A (nuclear membrane marker, 1:25) or monoclonal anti-PDI (ER marker, 1:25). Cells were then incubated for 30 minutes with goat anti-rabbit IgG coupled to dye conjugates and goat anti-mouse IgG (Alexa Fluor 488 and 568; 1:1500; Invitrogen). Negative control sections were treated identically except that PBS replaced the primary antibodies. Coverslips containing the cells were mounted with mounting medium (Vectashield Hardset; Vector Laboratories, Burlingame, CA) on micro-

scope slides and examined under a confocal microscope with imaging software (LSM 510 Meta; Carl Zeiss).

### Preparation of Cell Membranes for Binding Assays

RGC-5, rMC-1, and primary mouse MCs were cultured as described. Cell monolayers were chilled on ice, washed with ice-cold PBS (pH 7.5), and lysed with 5 mM  $K_2HPO_4$ - $KH_2PO_4$  buffer (pH 7.5). The suspension was centrifuged for 30 minutes at 56,800g; the final membrane pellets were rinsed and suspended in 5 mM  $K_2HPO_4$ - $KH_2PO_4$  buffer and homogenized 20 times using a 25-gauge needle. Protein concentration in the final membrane preparation was measured with the protein assay (Bio-Rad Protein Assay Reagent; Bio-Rad).

### Ligand-Binding Assay

[ $^3H$ ](+)-pentazocine binding to membrane preparations was assayed as described with minor modifications.<sup>36</sup> Samples were incubated with [ $^3H$ ](+)-pentazocine (10 nM) in 250  $\mu$ L of 5 mM  $K_2HPO_4$ - $K_2PO_4$  buffer, pH 7.5, at 25°C for 90 minutes. Binding was terminated by the addition of ice-cold binding buffer, and the mixture was filtered on a Whatman GF/F glass fiber filter, presoaked in 0.3% polyethylenimine. The filter was washed three times with ice-cold binding buffer, and radioactivity associated with the filter was determined by liquid scintillation spectrometry. To study the inhibition of [ $^3H$ ]-pentazocine binding to the  $\sigma$ R1, several competitive inhibitors—(+)-3-PPP, carbapentane, DTG, haloperidol, and (+)-pentazocine—were used. Concentrations for the [ $^3H$ ]-pentazocine and the inhibitors were 50 nM and 1  $\mu$ M, respectively. To investigate the allosteric effects of phenytoin on  $\sigma$ R1, 150  $\mu$ L rMC-1 membrane preparation containing 300  $\mu$ g protein was incubated with 50  $\mu$ L of 125 nM of [ $^3H$ ](+)-3-PPP (final concentration, 25 nM) and 50  $\mu$ L of 250  $\mu$ M phenytoin (final concentration, 50  $\mu$ M) or its solvent (5 mM  $K_2HPO_4$  buffer containing 2.5% DMSO, pH 7.5) for 90 minutes at 25°C. The final volume of the reaction system was 250  $\mu$ L of 50  $\mu$ M unlabeled (+)-pentazocine and was used to define nonspecific binding.

Effects of donor NO and ROS on  $\sigma$ R1 binding activity were examined by treating rMC-1 cells on the second day after seeding for 6 hours with NO donor SNAP (250  $\mu$ M), SNOG (250  $\mu$ M), and SIN-1 (100  $\mu$ M) or ROS donor  $H_2O_2$  (0.00025%) and xanthine (25  $\mu$ M)/xanthine oxidase (10 mU/mL), after which the binding activity was assayed as described.

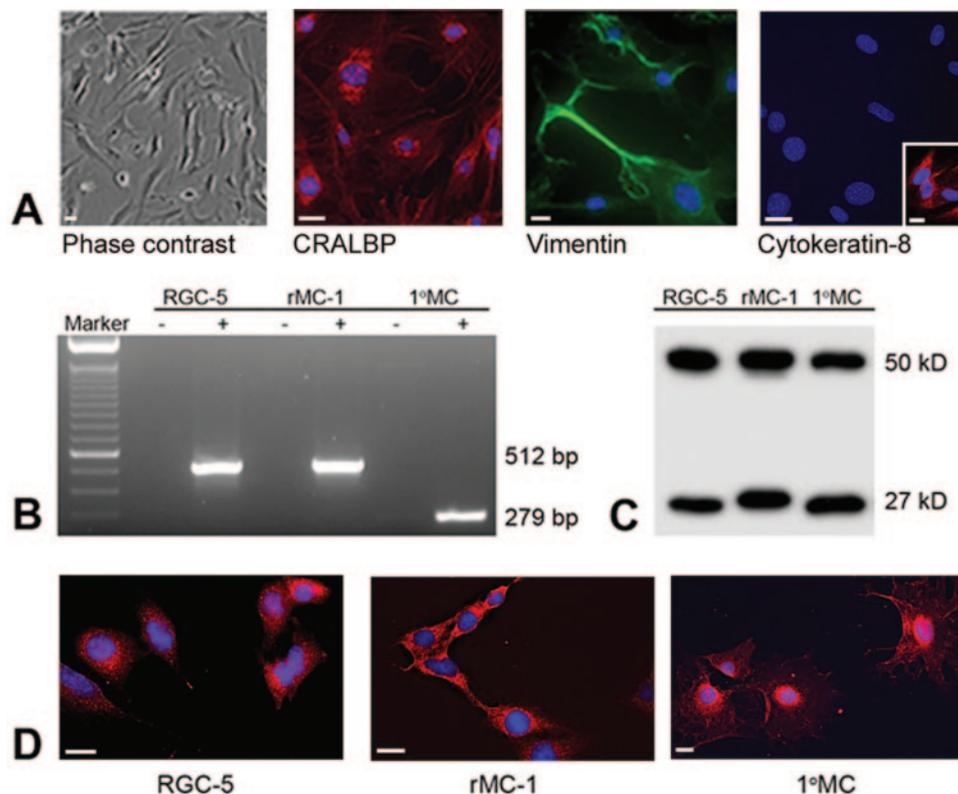
### Data Analysis

Experiments were performed in duplicate or triplicate; each experiment was repeated at least twice. Results are expressed as mean  $\pm$  SE. Equilibrium saturation-binding parameters, dissociation constant ( $K_d$ ), and maximum number of binding sites ( $B_{max}$ ) were calculated by nonlinear regression analysis of the equation for a rectangular hyperbola (SigmaPlot 2001 for Windows, version 7.0; Systat Software Inc., Richmond, CA), and statistical significance ( $P < 0.05$ ) was determined (SigmaStat, version 2; Systat).

## RESULTS

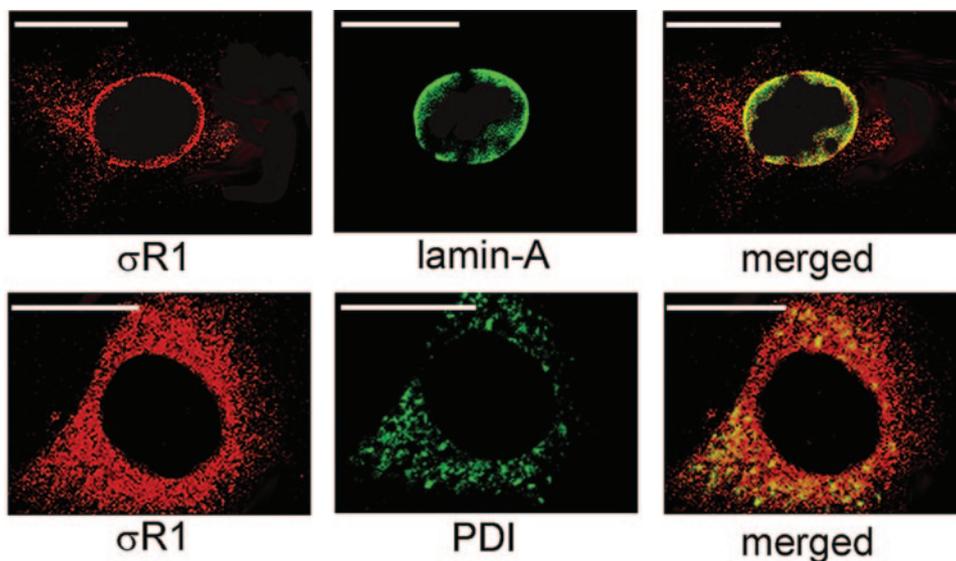
### Analysis of $\sigma$ R1 Expression in Müller and Ganglion Cells

Purity of the mouse Müller cells<sup>31</sup> was confirmed using known antibody markers for Müller cells, including CRALBP and vimentin (Fig. 1A) and glutamine synthetase (data not shown). The cells were not positive for neuronal markers such as NF-L<sup>31</sup> or for cytokeratin-8, which labels RPE cells (Fig. 1A). For expression and ligand-binding studies, primary mouse MCs were used at passage number 5. RT-PCR of the mouse primary mouse MCs amplified the expected PCR product (279 bp; Fig. 1B), indicating that  $\sigma$ R1 is expressed in these cells. RT-PCR analysis of the rat cell lines amplified the expected product (512 bp), consistent with our earlier data using these cell lines.<sup>23</sup> Immunoblotting with an antibody specific for  $\sigma$ R1<sup>24</sup> detected a protein band of the expected size ( $M_r \sim 27$  kDa) in primary mouse MCs, rMC-1, and RGC-5 cells. Immunocyto-



**FIGURE 1.** Analysis of  $\sigma$ R1 expression in retinal cell lines and in primary mouse MCs. (A) Verification of the purity of the primary mouse MCs used in these studies. *Left:* phase-contrast image of cells grown for 3 days in culture. *Middle:* positive immunostaining for CRALBP (red fluorescence) and vimentin (green fluorescence). *Right:* immunolabeling with cytokeratin-8, which is negative in the Müller cells but is positive in ARPE-19 cells (red fluorescence, inset). (B) RT-PCR showing expression of  $\sigma$ R1 mRNA in rat ganglion (RGC-5) and Müller (rMC-1) cell lines (512 bp) and primary mouse MCs (279 bp). For each cell type, PCR was run in the absence (-) or presence (+) of reverse transcriptase. (C) Immunoblotting of RGC-5, rMC-1, and primary mouse MCs showing the detection of  $\sigma$ R1 ( $M_r \sim 27$  kDa) and  $\beta$ -actin ( $M_r \sim 50$  kDa). (D) Cell lines or primary mouse MCs were cultured and subjected to immunocytochemistry to detect  $\sigma$ R1, followed by a Cy-3-conjugated secondary antibody (red). DAPI was used to label nuclei (blue).  $\sigma$ R1 was detected in the cell lines (RGC-5, rMC-1) and in primary mouse MCs. Bar, 15  $\mu$ m.

**FIGURE 2.** Subcellular localization of  $\sigma$ R1 in primary mouse MCs. MCs were isolated from mouse retina and cultured as described. They were subjected to double-labeling immunocytochemical analysis using a polyclonal antibody specific for  $\sigma$ R1 (red) and monoclonal antibodies (green) that label the nuclear membrane (lamin-A) or the endoplasmic reticulum (PDI). Optical sectioning by confocal microscopy detected colocalization of  $\sigma$ R1 with lamin-A (merged image) and with PDI (merged image). In the merged images, the orange signal was detected when the red and green fluorescence overlapped, indicating colocalization. Bar, 15  $\mu$ m.



chemical analysis of the cells detected  $\sigma$ R1 in RGC-5, rMC-1, and primary mouse MCs (Fig. 1D).

To analyze the subcellular location of  $\sigma$ R1 in primary mouse MCs, markers for the nuclear and ER membranes were used in double-labeling experiments with the  $\sigma$ R1 antibody. As shown in Figure 2A,  $\sigma$ R1 was present in the area, consistent with the nuclear membrane, and the merged image of the  $\sigma$ R1 plus the nuclear membrane marker, lamin-A, showed marked colocalization, suggesting that  $\sigma$ R1 is present on the nuclear membrane. Similarly, optical sectioning at a slightly different cell plane showed intense  $\sigma$ R1 levels in the perinuclear area, consistent with ER localization. The merged image of the  $\sigma$ R1 and the ER membrane protein, PDI (Fig. 2), provided strong evidence that  $\sigma$ R1 was also present in the ER.

### Characterization of [<sup>3</sup>H](+)-Pentazocine Binding in Retinal Cell Membranes

$\sigma$ R1 binding in RGC-5, rMC-1, and primary mouse MCs was characterized using (+)-pentazocine, a high-affinity  $\sigma$ R1 ligand.<sup>37</sup> Binding was saturable over a (+)-pentazocine concentration range of 1.25 to 75 nM (Fig. 3). Apparent  $K_d$  was  $25.0 \pm 5.9$  nM,  $21.5 \pm 2.6$ , and  $18.9 \pm 5.6$  nM for RGC-5, rMC-1, and primary mouse MCs, respectively. Thus, in the cell lines derived from rat and in primary mouse MCs, the affinity constant for the protein was comparable to the cloned  $\sigma$ R1 from rat and mouse. Scatchard analysis of the binding data revealed the presence of a single binding site in each cell type. The binding constants ( $B_{max}$ ) calculated for the RGC-5, rMC-1, and primary mouse MCs were  $1.53 \pm 0.14$ ,  $1.84 \pm 0.08$ , and  $1.32 \pm 0.13$  pmol/mg protein, respectively. These data suggest that the receptor density was comparable between the retinal cell lines and the primary mouse MCs.

The binding of [<sup>3</sup>H](+)-pentazocine (final concentration, 25 nM) to RGC-5, rMC-1 and primary mouse MC membranes was inhibited by several  $\sigma$ R1 ligands (Fig. 4). The order of potency differed among the cell types studied, such that in RGC-5 cells it was (+)-pentazocine > haloperidol > carbetapentane > 3-PPP > DTG and in rMC-1 cells it was haloperidol > (+)-pentazocine > carbetapentane = DTG > 3-PPP. In the primary mouse MCs, the order of potency was carbetapentane = haloperidol > (+)-pentazocine > 3-PPP > DTG.

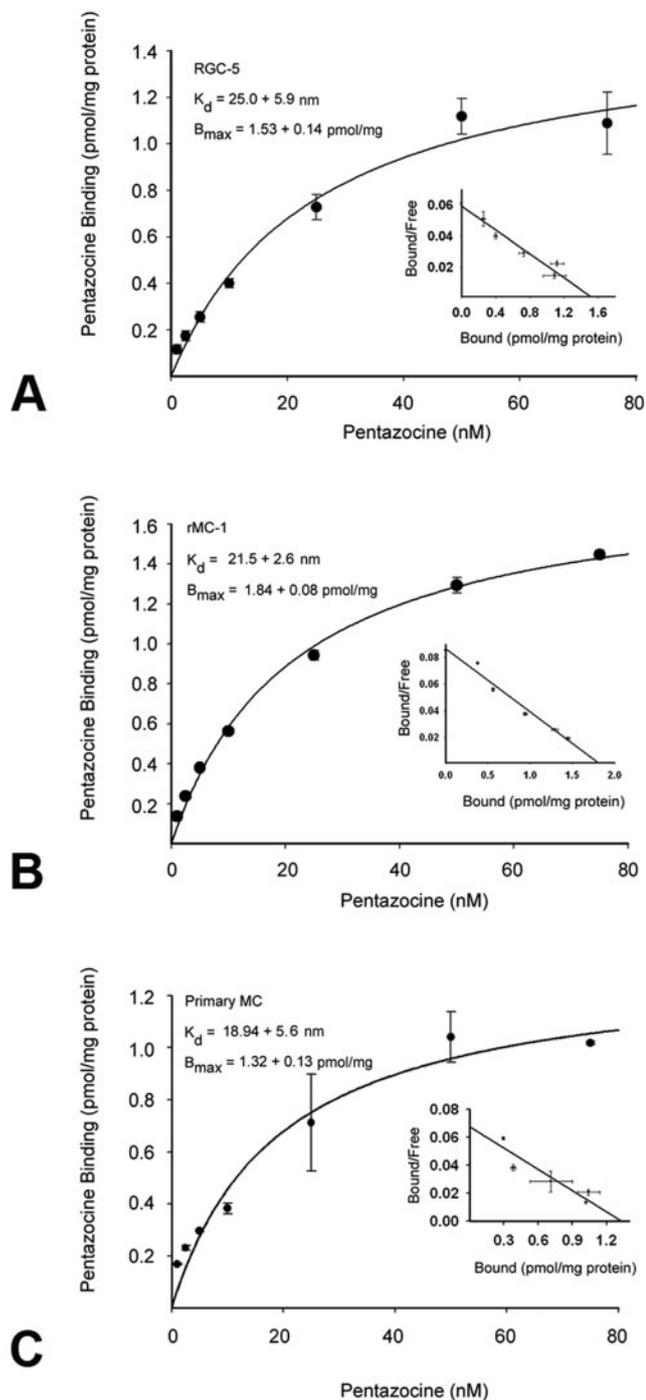
A key pharmacologic characteristic of  $\sigma$ R1, which distinguishes these receptors from type 2 receptors, is the allosteric enhancement of ligand binding to the  $\sigma$ R1 in the presence of phenytoin.<sup>9</sup> Stimulation of binding of the  $\sigma$ R1 agonist (+)-3-

PPP in the presence of phenytoin was reported recently in guinea pig brain.<sup>38</sup> We used primary mouse MCs to confirm that the  $\sigma$ R binding we were studying was specific to type 1 ( $\sigma$ R1). In these studies, the binding of [<sup>3</sup>H](+)-3-PPP was examined in the absence and presence of 50  $\mu$ M phenytoin. There was a 35% stimulation of binding activity in the presence of phenytoin (Fig. 5). The data suggest that the binding activity measured in the isolated Müller cells was specific to  $\sigma$ R1.

The ability to analyze  $\sigma$ R1 binding activity in individual retinal cell types affords the opportunity to investigate the regulation of this activity in the presence of factors implicated in retinal disease. Although it was not the focus of this study to investigate exhaustively the regulation of  $\sigma$ R1, it was important to ascertain whether binding activity was likely to be altered when retinal cells were subjected to stress. To this end, we performed preliminary studies in rMC-1 cells and analyzed  $\sigma$ R1 binding activity when cells were exposed to NO or ROS donors. These stressors were selected because NO<sup>39-42</sup> and oxidative stress<sup>43</sup> have been implicated in the pathogenesis of diabetic retinopathy. The cells were exposed to the NO donors SNAP, SNOG, and SIN-1 or to the ROS donors H<sub>2</sub>O<sub>2</sub> and X/XO for 6 hours, after which the binding activity of  $\sigma$ R1 was assayed (Fig. 6). Treatment of rMC-1 cells with all three NO donors resulted in a marked increase in binding activity, with the greatest effect observed when cells were incubated with SIN-1. There was an approximate threefold increase in binding activity after 6-hour exposure to this NO donor. In experiments using ROS donors, the effects were similarly profound, with a threefold increase in binding activity after 6-hour exposure to H<sub>2</sub>O<sub>2</sub> and a nearly fourfold increase in the presence of X/XO. The findings were observed in three independent experiments.

### DISCUSSION

In this study, we examined  $\sigma$ R1 expression in isolated retinal cell types and comprehensively characterized  $\sigma$ R1 binding activity in these cells. We used molecular techniques and demonstrated that  $\sigma$ R1 is present in primary mouse MCs isolated from the mouse retina. The location of the  $\sigma$ R1 appeared to be perinuclear. We explored this through laser scanning confocal microscopy (LSCM) with antibodies that recognize the nuclear and ER membranes. Our studies demonstrated that in primary mouse MCs,  $\sigma$ R1 is localized to both sites.  $\sigma$ R1 to the ER was consistent with findings by Hyashi and Su,<sup>29</sup> who localized  $\sigma$ R1 to the ER of oligodendrocytes. They did not

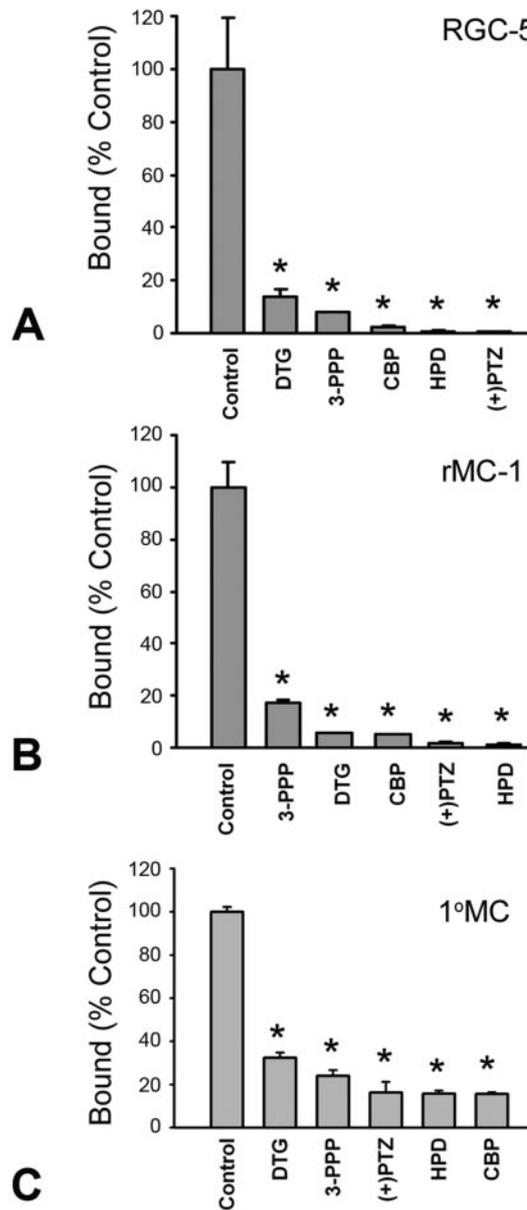


**FIGURE 3.** Saturation kinetics of  $\sigma$ R1 binding in membrane preparations. The two retinal cell lines (RGC-5 and rMC-1) and primary mouse MCs were cultured as described, membranes were prepared, and the binding of [ $^3$ H](+)-pentazocine over a broad concentration range was measured in (A) RGC-5 cells, (B) rMC-1 cells, and (C) primary mouse MCs. Values are the mean  $\pm$  SE for two independent experiments performed in triplicate. *Inset*: Scatchard plot.

report localization to the nuclear membrane; however, they did not use antibody markers, so it is unknown whether oligodendrocytes might also place  $\sigma$ R1 on the nuclear membrane.

Previous studies of  $\sigma$ R in retina have demonstrated  $\sigma$ R1 binding activity; however, these studies used whole retina from large models (bovine)<sup>21,22</sup> and did not attempt to study the  $\sigma$ R1 binding activities of individual retinal cell types. If we

are to postulate a role for  $\sigma$ R in mediating neuroprotection in retinal cells, it is essential to be able to quantify the  $\sigma$ R1 binding activity in the isolated cells. We were interested in  $\sigma$ R1 binding activity in ganglion cells and in Müller cells. Ganglion cells die in several retinal diseases,<sup>44-46</sup> and we predict that activation of the  $\sigma$ R1 may be beneficial to their survival; Müller cells play a key role in neuronal survival<sup>30</sup> and may activate  $\sigma$ R1 during stressful episodes, providing protection for adjacent neurons. In the present studies, we exploited the retinal ganglion cell line RGC-5 to study  $\sigma$ R1 binding activity in neurons. Although isolation of ganglion cells from mouse retina is

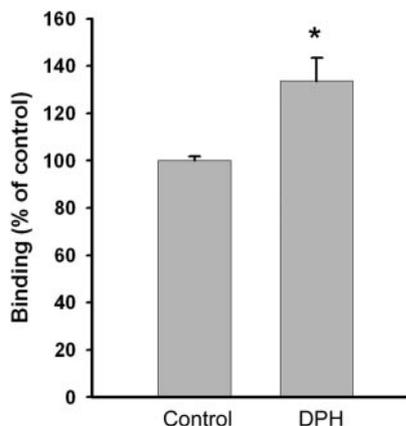


**FIGURE 4.** Inhibition of [ $^3$ H](+)-pentazocine binding by  $\sigma$ R1 ligands. RGC-5, rMC-1, and primary mouse MCs were cultured as described, membranes were prepared, and the inhibition of the binding of [ $^3$ H](+)-pentazocine was measured using several known ligands for  $\sigma$ R1. (A) RGC-5 cells, (B) rMC-1 cells, and (C) primary mouse MCs. Data are shown as relative percentage of control binding (100%) measured in the absence of ligands. Values are the mean  $\pm$  SE for two independent experiments performed in triplicate. 3-PPP, (+)-3-(3-hydroxyphenyl)-N-(1-propyl)piperidine; CBP, carbetapentane; DTG, 1,3-di-(2-tolyl)guanidine; HPD, haloperidol; (+)PTZ, (+)-pentazocine. \*Significantly different from control;  $P < 0.01$ .

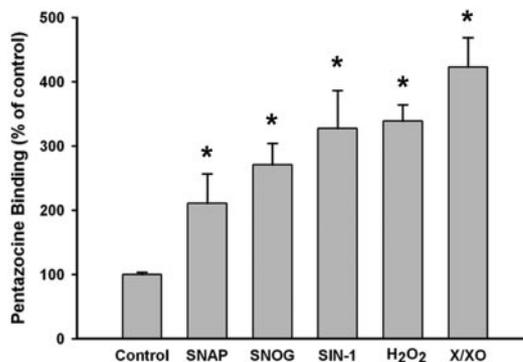
possible,<sup>35</sup> these cells are terminally differentiated, making it difficult to obtain sufficient numbers of these neurons to carry out  $\sigma$ R1 binding assays. For studies of  $\sigma$ R1 in Müller cells, we used the rMC-1 cell line and complemented the data using primary MCs isolated from mouse. Given the plethora of genetic defects that cause retinal abnormalities in commercially available strains of mice, the ability to study  $\sigma$ R1 expression and activity in primary mouse MC cultures afforded an excellent opportunity to dissect  $\sigma$ R1 activity and expression in Müller cells of various mouse models of retinal disease.

Our studies of  $\sigma$ R1 binding characteristics in Müller cells showed that the affinity of  $\sigma$ R1 in primary mouse MCs was comparable to that of two retinal cell lines, RGC-5 and rMC-1. In all three cases,  $\sigma$ R1 bound its ligand, (+)-pentazocine, with great affinity. The density of receptors, as indicated by the  $B_{max}$  value, was similar between the primary mouse MCs and the two retinal cell lines. The ability of various ligands to inhibit the binding of (+)-pentazocine differed slightly among the cell types. The two cell lines seemed more sensitive to the inhibitory effects of CBP, HPD, and (+)-pentazocine, whereas the primary mouse MCs seemed slightly less sensitive to these inhibitors. Nonetheless, the binding in these primary mouse MCs was inhibited by 75% to 80% over a wide range of ligands tested. Finally, the known allosteric effects of phenytoin to stimulate binding of 3-PPP were borne out in these studies and provide strong evidence that the binding studied in these retinal cells was indeed mediated by  $\sigma$ R1.

The value of the present work is that it forms a scaffold on which to study the regulation of  $\sigma$ R1 activity in isolated retinal cells. We now have substantial baseline data for two retinal cell lines and primary mouse MCs about the dissociation constant, receptor density, and ligand-specificity characteristics of these receptors. Thus, we are poised to compare these benchmarks under conditions that are known to represent disease states in the retina. For example, in diabetic retinopathy, a variety of factors such as hyperglycemia, oxidative stress, increased levels of NO, and inflammation are thought to figure prominently in the eventual compromised functions of several retina cell types.<sup>44</sup> It will now be possible to evaluate the effects of these factors on the  $\sigma$ R1 binding characteristics in isolated retinal cells. Although we did not explore this comprehensively in the present study, we performed preliminary experiments using NO donors and ROS donors to determine the effects on  $\sigma$ R1 binding. Our studies showed that oxidative stress led to in-



**FIGURE 5.** Allosteric effects of phenytoin on the binding of [<sup>3</sup>H]-3-PPP. The rMC-1 cell line was cultured and membranes were prepared as described. The allosteric effect on [<sup>3</sup>H]-3-PPP binding activity in the absence (Control) or presence of phenytoin (DPH) was measured after 90-minute incubation, as described. Values are mean  $\pm$  SE for two experiments performed in triplicate. \*Significantly different from control;  $P < 0.01$ .



**FIGURE 6.** Regulation of  $\sigma$ R1 binding activity by NO and ROS donors. rMC-1 cells were cultured as described and were treated for 6 hours with NO (3-nitroso-N-acetylpenicillamine [SNAP, 250  $\mu$ M], S-nitroso-glutathione [SNOG, 250  $\mu$ M], or 3-morpholinosydnonimine [SIN-1, 100  $\mu$ M]), ROS (hydrogen peroxide [H<sub>2</sub>O<sub>2</sub>, 0.00025%, wt/wt] solution), or xanthine/xanthine oxidase (X/XO, 25  $\mu$ M/10 mU/mL), after which the binding activity of  $\sigma$ R1 was assayed. Values are mean  $\pm$  SE for two experiments performed in triplicate. \*Significantly different from control;  $P < 0.01$ .

creased  $\sigma$ R1 binding activity. Future studies will investigate this phenomenon comprehensively, determining whether gene and protein expression are altered under conditions of oxidative stress and characterizing the kinetic parameters associated with this increased activity. In addition to characterizing the binding activity under conditions of oxidative stress, we will use the subcellular localization information obtained in this study for future experiments designed to determine whether oxidative stress induces  $\sigma$ R1 to translocate from its typical location at the nuclear and ER membranes to other cellular sites, such as the plasma membrane.

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