MCP-1 Deficiency Delays Regression of Pathologic Retinal Neovascularization in a Model of Ischemic Retinopathy

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Purpose. The present study investigates whether retinal neovascularization (NV) and apoptosis are altered in MCP-1−/− mice in the OIR model.

Methods. Postnatal day (P) 7 MCP-1−/− and C57BL/6 (B6) mice were exposed to 75% oxygen for 5 days and then recovered in room air. Immunostaining was performed to localize macrophages/microglia within retinal whole mounts and cross-sections. Retinopathy was qualitatively assessed in FITC-dextran-perfused retinas, and preretinal NV was quantified on P17, P21, and P24. TUNEL analysis was used to compare apoptosis between B6 and MCP-1−/− mice.

Results. MCP-1−/− and B6 mice revealed normal vascular development in room air controls and similar vaso-obliteration in oxygen-exposed mice on P12. MCP-1−/− mice exhibited significantly reduced vascular tuft-associated F4/80+ cells compared with B6 mice. FITC-dextran-perfused retinas exhibited prominent neovascular tufts on P17, and quantification of preretinal nuclei revealed no significant differences between MCP-1−/− and B6 mice. In contrast, on P21 and P24, MCP-1−/− mice exhibited significant increases in preretinal neovascular nuclei compared with B6 controls. These increases in NV in the MCP-1−/− mice were associated with a significant reduction in vascular tuft apoptosis.

Conclusions. The results demonstrate that the absence of MCP-1 does not alter normal retinal vascular development. Furthermore, MCP-1−/− mice exhibit a similar neovascular response on P17. However, the reduction in tuft-associated macrophages/microglia in the MCP-1−/− mice correlates with reduced vascular tuft apoptosis and delayed regression of retinal NV. These findings suggest that macrophages/microglia may contribute to tuft regression through their proapoptotic properties. (Invest Ophthalmol Vis Sci. 2008;49:4195–4202) DOI:10.1167/iovs.07-1491

Pathologic angiogenesis is a key component of many diverse diseases, including ischemic retinopathies.1,2 A balance between proangiogenic and antiangiogenic factors determines whether endothelial cells (ECs) will proliferate, migrate, and incorporate into pathologic vessels or undergo apoptosis, causing the nascent vessels to regress.3 Our laboratory and others4–6 have reported that Fas-ligand (FasL) can play a counterbalancing role to the proangiogenic growth factors in the eye. Pigment epithelial-derived factor (PEDF) and thrombospondin-1, potent endogenous inhibitors of angiogenesis, induce EC apoptosis through the Fas/Fasl pathway.7 Activated macrophages can induce apoptosis in target cells, including ECs, through cell surface Fasl, tumor necrosis factor-alpha (TNF-α), and TNF-related apoptosis-inducing ligand (TRAIL).8–13 Activated resident and infiltrating macrophages are dynamic cell populations that have the potential to play multiple roles in disease pathogenesis and development.14 In addition to their proapoptotic properties during postnatal disease, macrophages are required for EC apoptosis during the regression of ocular capillaries in normal eye development, as demonstrated in the seminal studies by Lang et al.15,16 In contrast, macrophages are a rich source of angiogenic cytokines and have been linked to choroidal, tumoral, and inflammatory angiogenesis.17–20 These studies suggest that macrophages/microglia have the potential to play dual roles during pathologic angiogenesis.

Several laboratories have reported the immunolocalization of microglia and macrophages to preretinal neovascular tufts in the mouse model of oxygen-induced retinopathy (OIR).21–24 As shown in our previous study, the F4/80+ population peaks on postnatal day (P) 17 to P21 in the OIR model, which is associated with a significant increase in the chemokine monocye chemoattractant protein-1 (MCP-1, CCL2) on P17. Proliferation of resident microglia was not observed, suggesting the increase in F4/80+ cells is secondary to infiltrating monocytes/macrophages.22 To further characterize the role that macrophages/microglia play in retinal neovascularization (NV), vascular tuft regression, or both, we used MCP-1−/− deficient mice (MCP-1−/−) in the OIR model. The mouse OIR model reproducibly develops preretinal NV but is also characterized by the predictable regression of neovascular tufts through apoptosis.5,24,25 In several models of inflammation and injury, MCP-1−/− mice consistently exhibit a reduction in infiltrating macrophages, with an associated alteration in tissue injury.26–30 Therefore, using MCP-1−/− mice in the model of OIR should help clarify the role of infiltrating macrophages in this model of retinal NV.

Materials and Methods

Animals

Breeding pairs of C57BL/6 (B6) mice and MCP-1−/− mice (on a B6 background) were purchased from The Jackson Laboratory (Bar Harbor, ME), provided food and water ad libitum, and kept on a 12-hour light/12-hour dark cycle. Mice were housed and bred in the Oregon Health & Science University animal care facilities and were treated in compliance with the National Institutes of Health guidelines and the guidelines outlined in the ARVO Statement for the Use of Animals in Ophthalmic and Vision Research. Using the mouse model of OIR established by Smith et al.33 MCP-1−/− and B6 control pups, along with nursing females, were exposed to 75% oxygen for 5 days beginning on
P7 and then allowed to recover in room air on P12. Room air control
litters of MCP-1−/− and B6 mice were maintained in conditions iden-
tical to those for the hyperoxia-exposed animals. Hyperoxia-exposed
(O2 [O2 refers to hyperoxia exposed between P7 and P12]) and room
air control pups were killed by CO2 euthanization or cervical dislo-
cation on P12, P14, P17, P21, and P24. One eye was carefully enucle-
ated from each mouse and was placed in 10% neutral-buffered formalin
overnight and routinely processed for paraffin embedding. These eyes
were sectioned at 5-μm intervals, mounted on slides (SuperFrost Plus;
Fisher Scientific, Pittsburgh, PA), and stored at room temperature until
used for immunohistologic and TUNEL analysis. The retina of the
contralateral eye was dissected for mRNA isolation and analysis, and
subsets of mice were used for retinal flat mount studies.

Immunohistochemistry
To assess microglia and macrophages, retinal whole mounts and cross-
sections were immunolabeled as previously described. Briefly, for
whole mount analysis, the eyes were enucleated, and the lenses and
retina were dissected out and fixed in 4% paraformaldehyde (PFA). The
lenses were removed, and the retinas were incubated with a rat
anti–mouse F4/80 antigen antibody (Serotec, Raleigh, NC) for 56 to 48
hours at 4°C. Retinas were incubated with biotinylated rabbit anti–rat
IgG (Vector Laboratories, Burlingame, CA). ABC-AP complex (Vector)
was applied to retinas and visualized (Fast Red; BioGenex Laboratories,
San Ramon, CA). Retinas were incised radially, and the vitreous was
removed and flat mounted with antifade (SlowFade; Molecular Probes,
Eugene, OR).

Retinal tissue sections were routinely deparaffinized and rehy-
drated before antigen retrieval by proteolytic digestion with 0.1%
trypsin (Sigma, St. Louis, MO) for 5 minutes at room temperature. After
a 1-hour blocking step in 2% goat serum, sections were incubated
overnight at 4°C with F4/80 antibody in a humidifying chamber.
Sections were incubated with an FITC-conjugated goat anti-rat IgG
antibody (Serotec), counterstained with DAPI, and mounted with ant-
tifade (SlowFade; Molecular Probes).

Retinal whole mounts and cross-sections were visualized by light
and fluorescence microscopy and were photographed with a digital
camera (DC500; Leica Microsystems, Bensheim, Germany). To quanti-
tate the F4/80−/− labeled cells of B6 and MCP-1−/− mice, vascular tufts
from P17O2 retinas were photographed in a masked fashion (2–4
classes (2–4 images/section) using four representative sections 60 μm apart (n = 8
eyes). F4/80−/− cells were then counted in a masked fashion from the
digital images (B6, n = 53; MCP-1−/−, n = 53). Additional tissue
sections were used for evaluation of intraretinal blood vessels after
immunolocalization with a type IV collagen antibody (Collaborative
Biomedical Products, Bedford, MA), as previously described.

Retinal Fluorescein Angiography
Qualitative assessment of retinal vasculature was performed on hyper-
oxia-exposed and room air control mice at P12, P17, and P21.31 Mice
were given a general anesthetic cocktail, injected subcutaneously,
that contained ketamine, xylazine, and acepromazine. Mice were then
perfused through the left ventricle with a solution of 1 mL PBS and 50
mg high molecular-weight (2 × 105), FITC-conjugated dextran (Sigma).
Animals were killed by cervical dislocation, eyes were enu-
cleated, and lenses and retinas were dissected away and fixed in 4% PFA
for 3 hours at 4°C. The lens was removed, and the retina was incised radially. After the vitreous was removed, the retina was flat
mounted with antifade (SlowFade; Molecular Probes). Retinal vessels
were visualized by fluorescence microscopy and photographed with a
digital camera (DC500; Leica).

Neovascular Nuclei Quantification
NV was quantified by counting the vascular nuclei that extended
anterior to the internal limiting membrane in hematoxylin and eosin-
stained sections. Retinal vascular nuclei were counted in a masked
fashion and averaged, avoiding hyaloid vessel nuclei near the optic disc
and lens (P17O2 and P21O2, n = 1 eye each from 8 to 10 animals, 15
sections per eye; P24O2, n = 1 eye each from 4 animals, 15 sections
per eye). Data are expressed as mean ± SEM.

Apoptosis Analysis
TUNEL assay was performed on retinal cross-sections from hyperoxia-
exposed MCP-1−/− and B6 mice, comparing the degree of apoptosis
occurring in the neovascular tufts on both P17O2 (n = 8 eyes, 10
sections per eye) and P21O2 (n = 6 eyes, 10 sections per eye). Sections
were labeled with the use of a detection kit (Apoptag Peroxidase In
Situ Apoptosis Detection Kit; Intergen, Purchase, NY) according to the
manufacturer’s instructions. After labeling the exposed 3′-OH ends of
DNA fragments, apoptotic cells were visualized with DAB substrate
and counterstained with methyl green. While taking care to avoid the
optic nerve, representative sections were assessed for TUNEL cells
located beyond the inner-limiting membrane and were quantified in a
masked fashion. Data are expressed as mean ± SEM.

To further characterize the apoptotic response, a subset of sections
were immunolabeled with an activated caspase-3 antibody. Briefly, after
deparaffinization and rehydration, sections were boiled in 1 mM EDTA
for antigen retrieval. After a blocking step, sections were incubated
overnight at 4°C with cleaved caspase-3 antibody (Cell Signaling, Dan-
ers, MA). Retinal sections were then incubated with biotinylated goat
anti-rabbit IgG (Vector). ABC-HRP complex (Vector) was applied,
visualized with DAB substrate, and counterstained with methyl green.

Quantitative RT-PCR
Retinas were dissected on P14, P17, and P21 from room air and
hyperoxia-exposed B6 mice and pooled (n = 4) for RNA analysis. For
comparison on the peak day of NV, P17 hyperoxia-exposed MCP-1−/−
retinas were also dissected and pooled for RNA analysis. RNA was
extracted using Qiagen columns (RNeasy; Qiagen, Valencia, CA) ac-
cording to manufacturer’s instructions. Total RNA was purified by
DNase treatment to remove potential genomic DNA contamination.
cDNA was then synthesized using oligo(dT)-primed M-MLV reverse
transcriptase (Promega, Madison, WI) for 2 hours at 37°C. Relative
mRNA expression of F4/80 normalized to β-actin was quantified in
triplicate by real-time PCR using a thermocycler (Chromo4; MJ Re-
search, Watertown, MA) and supermix (IQ SYBR Green; Bio-Rad, Her-
cules, CA). Annealing temperatures and plate-read temperatures were
predetermined using control mRNA. Melting curves were determined
after product formation, and samples were run on a gel to confirm
product size. A standard curve was generated and amplified simulta-
eously, allowing for determination of relative concentrations of un-
known samples. Concentrations are reported in relative fluorescent
units. Mouse-specific primer sets were synthesized by Integrated DNA
Technologies, Inc. (Coralville, IA) for the following F4/80 sequences
sense, 5′-GGCTGCTGGTTGATACAGGAGA-3′; antisense, 5′-GGGTGTT-
GACGACAGCTGAATGA-3′. A primer pair for the constitutively ex-
pressed β-actin gene was included in each assay as an internal loading
control, as follows: sense, 5′-ATGCCAACACAGTGCTGTCT-3′; anti-
sense, 5′-AAGCATTGGCGTGGAC-3′.

Statistical Analysis
Results are expressed as the mean ± SEM. Statistical significance was
determined using the Student’s t-test for comparison between 2 groups
or one-way ANOVA for multiple-group comparison (Prism; GraphPad,
San Diego, CA). P < 0.05 was considered statistically significant in all
statistical analyses used.

RESULTS
Reduced Retinal F4/80+ Cells in MCP-1–Deficient Mice
We examined retinal flat mounts and cross-sections using the
macrophage/microglia marker F4/80 to determine whether
MCP-1−/− mice exhibited reduced macrophage infiltration. Whole mount immunostaining for F4/80 antigen in B6 room air control mice demonstrated increased F4/80+ cells throughout the retina on P17 (Figs. 1A, 1D). In comparison, the P17O2 B6 retinas showed a marked increase in F4/80+ cells throughout the retina (Figs. 1B, 1E). However, the increase in F4/80+ cells was substantially reduced in the P17O2 MCP-1−/− mice, as observed in the retinal flat mounts (Figs. 1C, 1F). This observation was confirmed by evaluating retinal cross-sections with immunohistochemistry. As previously described, the F4/80 antibody labels microglia in the inner plexiform layer and the outer plexiform layer in B6 room air control retinas on P17 (Fig. 1G, arrows).21 In contrast to the room air controls, B6 mice exhibited numerous F4/80+ cells surrounding the neovascular tufts on P17 in the oxygen-injured retinas, as reported previously (Fig. 1H, arrow).22 An increase in mRNA expression of retinal F4/80 correlated with increased F4/80+ cells observed in B6 oxygen-exposed retinas compared with room air controls (Fig. 1I). The F4/80 PCR results correlate with our previous quantification of retinal F4/80+ cells in the OIR model.22 Similar to the flat mount results, the retinal cross-sections showed that the MCP-1−/− mice exhibited reduced numbers of F4/80+ cells in the retina (Fig. 1J, arrow). F4/80+ cells were also quantified in retinal cross-sections, and there was a significant reduction in the MCP-1−/− mice compared with B6 mice on P17O2 (Fig. 1K). Real-time PCR also revealed a reduction in F4/80 expression in MCP-1−/− mice compared with B6 controls on P17O2 (Fig. 1L). Despite the reduction in F4/80+ cells, the microglia still appeared to have migrated from the plexiform layers to the anterior retina in the hyperoxia-exposed MCP-1−/− mice, similar to the B6 mice (Figs. 1H, 1J).

**Figure 1.** Reduced F4/80+ macrophages/microglia in MCP-1−/− mice. F4/80-labeled cells (red) were localized in retinal whole mounts from B6 room air control mice (A, D), along with B6 hyperoxia-exposed mice (B, E), and from MCP-1−/− mice (C, F). Compared with P17 room air control B6 mice (A, D), an increase in F4/80-positive cells was observed in hyperoxia-exposed B6 mice (B, E). Examination of the hyperoxia-exposed MCP-1−/− mice (C, F) demonstrated a reduced number of F4/80+ cells compared with hyperoxia-exposed B6 mice (B, E) on P17. Immunolabeling of F4/80+ macrophages/microglia (green) in retinal cross-sections revealed cells localized in the inner plexiform layer on P17 in B6 room air controls (G, arrow). However, in P17 hyperoxia-exposed retinal cross-sections, F4/80+ cells were located within the neovascular tufts in the B6 mice (H, arrow), whereas fewer macrophages/microglia were associated with tufts in the MCP-1−/− mice (I, arrow). Original magnifications: (A–C) ×25, (D–I) ×400. Quantitative real-time PCR for F4/80 expression normalized to β-actin in B6 OIR time course (J) and in MCP-1−/− mice compared with B6 mice on P17O2 (L). Quantification of F4/80+ cells (K) in retinal cross-sections. *P < 0.002; †P < 0.05.

### Delayed Regression of Neovascular Tufts in MCP-1−/− Mice

Fluorescein-perfused retinal flat mounts and type IV collagen-immunostained retinal sections revealed similar retinal vascular development on P12 in room air control B6 and MCP-1−/− mice (data not shown). Hyperoxia-exposed B6 and MCP-1−/− mice exhibited central vaso-oblation (asterisks) and a delay in development of the deep vascular network on P12O2, as previously described (Figs. 2A, 2B).23 In addition, the superficial and deep networks exhibited a similar degree of vaso-oblation in type IV collagen–immunostained retinal sections in B6 and MCP-1−/− mice on P12O2 (data not shown). Fluorescein angiography also revealed a similar pattern of avascular regions (asterisks) on P17O2 in B6 and MCP-1−/− mice (Figs. 2C, 2D). Higher magnification of retinal flat mounts on P17O2 demonstrated a similar neovascular response in B6 and MCP-1−/− mice (Figs. 2E, 2F). On P21O2, B6 mice experienced vascular tortuosity but decreased NV, as shown in fluorescein-perfused flat mounts (Fig. 2G), whereas MCP-1−/− mice continued to exhibit numerous neovascular tufts (Figs. 2H, arrows).

The extent of retinal NV was quantified in cross-sections by counting the average number of vascular nuclei extending beyond the inner limiting membrane (Fig. 3). On P17O2, the peak day of NV, hyperoxia-exposed B6 and MCP-1−/− mice exhibited similar degrees of preretinal NV (arrows), with 24.7 ± 5.3 and 24.6 ± 1.3 preretinal nuclei per cross-section, respectively (Figs. 3A, 3B). By P21O2, the average number of preretinal nuclei per section had decreased substantially in the hyperoxia-exposed B6 mice to 6.7 ± 1.1 (Fig. 3C). Such vascular regression is typical for this model.24,25 In contrast, MCP-
1−/− mice continued to have a significant number of preretinal neovascular nuclei, 30.1 ± 6.1, on P21O2 (Fig. 3D). Despite continued vascular regression in hyperoxia-exposed B6 and MCP-1−/− mice on P24O2, a difference persisted in the number of preretinal vascular tuft nuclei per cross-section (1.3 ± 0.28 vs. 9.9 ± 2.2, respectively; Figs. 3E, 3F). The numeric differences in preretinal neovascular nuclei are also depicted in graphical form (Fig. 3G). These observations indicate that the absence of the macrophage chemokine MCP-1 was not associated with reduced retinal NV on P17O2 but rather with delayed regression of retinal NV on P21O2 and P24O2 in the MCP−/− mice compared with the B6 mice.

Reduced Vascular Tuft Apoptosis in MCP-1−Deficient Mice

To determine whether the absence of MCP-1 alters the retina’s apoptotic response during vascular tuft regression, TUNEL staining was performed to label apoptotic cells. The apoptotic cells located exclusively within the neovascular tufts were quantified and normalized as a percentage of neovascular nuclei. Compared with the relatively high percentage of TUNEL+ cells (arrows) seen in the hyperoxia-exposed P17 B6 mice (9.9% ± 1.1%; Fig. 4A), a significant decrease in preretinal apoptotic cells (arrows) was observed in hyperoxia-exposed MCP-1−/− mice on P17 (1.6% ± 0.5%; Fig. 4B). Percentage differences for P17O2 preretinal TUNEL+ cells are represented in graphical form (Fig. 4E). The reduction of tuft apoptosis on P17O2 in the MCP−/− mice correlated with the reduced number of infiltrating macrophages observed in the MCP−/− mice. In contrast to P17O2, the percentage of TUNEL+ cells was not significantly different on P21O2 between the B6 (Fig. 4C) and the MCP−/− (Fig. 4D) mice (15.1% ± 4.3% and 8.6% ± 2.8%, respectively; Fig. 4E). However, the relative increase in tuft apoptosis on P21O2 compared with P17O2 (8.6% ± 2.8% vs. 1.6% ± 0.5%) in the MCP−/− mice correlated with the gradual regression of the NV observed between P21O2 and P24O2.
Activated caspase-3 was also localized in P17O2 sections. Qualitative assessment of the cleaved caspase-3 immunostaining demonstrated numerous positive cells localized to the neovascular tufts of B6 mice (Fig. 4F), whereas the neovascular tufts in the MCP-1−/− mice exhibited only an occasional caspase-3–positive cell (Fig. 4G).

**DISCUSSION**

Previous studies by our laboratory and others22–24 have shown that macrophages/microglia colocalize with neovascular tufts in the mouse model of OIR and exhibit an associated increase in retinal MCP-1 expression. A recent study using in vivo immunostaining also confirms an intimate association between F4/80-labeled cells and retinal vessels in this model.22 In vivo studies using transgenic models of MCP-1 overexpression confirm the important role this chemokine plays in the recruitment of monocytes and macrophages to sites of tissue expression.33,34 Subsequently, MCP-1−/− mice were generated by targeted gene disruption to confirm the essential role of this chemokine in several models of inflammation and ischemia.26–30 Similarly, in the present study, MCP-1−/− mice exhibited reduced recruitment of F4/80+ cells, allowing us to examine the contribution of infiltrating macrophages to the pathogenesis of OIR. However, because microglia migrate to the retinal plexiform layers between P0 and P5 in response to macrophage colony-stimulating factor (MCSF), this F4/80+ cell population was already present in the retina on P7, having migrated from its normal array in the plexiform layers to the anterior retina in response to oxygen-induced injury in B6 and MCP-1−/− mice.21,35 Hence, the neovascular tufts in the MCP-1−/− mice showed a significant reduction in tuft-associated F4/80+ cells but were not devoid of this cell population.

A balance between angiogenic and antiangiogenic factors determines whether vascular remodeling (regression) rather than vascular stabilization occurs after the initiation of retinal...

[FIGURE 4. Decreased neovascular tuft apoptosis in MCP-1−/− mice. Representative TUNEL-stained sections from B6 (A, C) and MCP-1−/− (B, D) mice on P17O2 and P21O2. Apoptotic cells were observed within the neovascular tufts on P17O2 in B6 mice (A, arrows) and, to a lesser extent, in MCP-1−/− (B, arrows) mice. On P21O2, B6 mice (C, arrows) had numbers of apoptotic cells similar to those for MCP-1−/− mice (D, arrows). TUNEL+ cells localized exclusively within the neovascular tufts were counted on P17O2 and P21O2 and were reported as a percentage of TUNEL+ cells per neovascular nuclei (E). There was a significant difference between B6 and MCP-1−/− on P17O2 (*P < 0.0001). Representative cleaved caspase-3–stained sections from B6 (F, arrows) and MCP-1−/− (G, arrows) mice on P17O2. Original magnification, ×400.]
NV. VEGF and other angiogenic cytokines (antiapoptotic) have been well characterized for their contribution to NV in the mouse model of OIR. In contrast, Fas/Fasl, endostatin, and thrombospondin-1 are known to serve as negative regulators of angiogenesis through the induction of EC apoptosis in the OIR model. Reduced vascular tuft apoptosis on P17 in the MCP-1−/− mice correlated with reduced numbers of vascular tuft F4/80+ cells, suggesting that macrophages/microglia may play a role in tuft regression through the induction of EC apoptosis. Similarly, in a model of choroidal neovascularization (CNV) induced by subretinal injection of basement membrane matrix (Matrigel; BD Biosciences, Franklin Lakes, NJ), MCP-1−/− mice develop more CNV in conjunction with reduced macrophages in the subretinal lesion. In addition, after laser injury to the retina, Fasl is downregulated in ocular macrophages of old mice, with an associated enhancement of CNV, compared with young mice. A recent study using an experimental model of retinal detachment also confirmed the ability of activated macrophages/microglia to induce apoptosis in target cells in the eye through Fas/Fasl, whereas a reduction in apoptosis was observed in MCP-1−/− mice. We have also observed that TRAIL-deficient mice exhibit increased retinal NV on P17 and delayed regression of the vascular tufts on P21 in the OIR model (Hubert KE, et al. IOVS 2008;49:ARVO EAbstract 3295). TRAIL expression was also colocalized with a subset of tuft-associated F4/80+ cells in control B6 mice, suggesting that macrophages/microglia may induce EC apoptosis and contribute to vascular tuft regression by TRAIL. In addition to the classic death receptor pathways, nitric oxide and hypochlorous acid are two macrophage products that have been characterized as mediators of EC apoptosis. Ocular macrophages are also required for EC apoptosis during regression of the ocular capillaries in normal eye development, which has now been shown to be mediated by WNT/b. Taken together, these studies support the concept that macrophages/microglia are capable of inducing endothelial apoptosis through a variety of proapoptotic pathways in the normal and diseased eye.

Despite endogenous proangiogenic properties of MCP-1, we did not observe a reduction in NV on P17 in the MCP-1−/− mice compared with B6 mice. In addition, the unaltered NV observed on P17 in MCP-1−/− mice suggested that infiltrating macrophages are not a major contributor to NV in this model. These observations are consistent with Müller cell-derived VEGF playing the major role in stimulating angiogenesis in this model, as previously described. However, with reduced numbers of infiltrating macrophages, the shift toward tuft regression that normally occurs in B6 mice between P17 and P21 is delayed in MCP-1−/− mice. Similarly, in a preliminary study in B6 mice, macrophages were systemically depleted with liposomal-clodronate, resulting in a reduction of neovascular tuft-associated F4/80+ cells with an associated twofold increase in neovascular nuclei on P210, supporting the concept that infiltrating macrophages contribute to tuft regression (Davies MH, et al. IOVS 2006;47:ARVO EAbstract 3223). In contrast to the MCP-1−/− mice, mice deficient in the intracellular apoptosis inhibitor Bcl-2 exhibit significantly less NV than B6 control mice in the OIR model. In vivo studies using capillary tube assays revealed a requirement of only 22% apoptotic ECs for a significant decrease in microvascular density. Hence, it appears that the B6 mice had a level of EC apoptosis that allowed for tuft regression between P17 and P21 in contrast to the delayed regression we observed in the MCP-1−/− mice, whereas Bcl-2−/− mice likely had a level of apoptosis that did not allow for a significant level of NV, despite the presence of VEGF. However, the vascular tufts eventually regressed in the MCP-1−/− mice, suggesting that additional apoptotic mechanisms lead to tuft regression after P21. We did observe an increase in the percentage of tuft cells undergoing apoptosis in the MCP-1−/− mice on P21. Growth factor withdrawal secondary to waning levels of retinal VEGF likely contributed to the eventual regression of the vascular tufts after P21 in the MCP-1−/− mice. Retinal Fas was also up-regulated on P21 in the OIR model, whereas EC up-regulated this death receptor under irregular flow conditions (vascular tufts). Thus, the tuft ECs in the nascent capillaries were vulnerable to autoregulatory Fasl-induced cell death from neighboring ECs. Finally, the resident microglia likely also contributed to eventual regression of the neovascular tufts.

Macrophages have been shown to exhibit distinct functions during injury and repair, with early-arriving macrophages promoting tissue injury and late-arriving macrophages promoting tissue recovery through the induction of apoptosis. The concept of macrophages/microglia promoting vascular tuft regression through apoptosis does not exclude a proangiogenic function for these cells during an earlier phase in the OIR model. In fact, two recent studies have explored the contribution of microglia to the development of the retinal vasculature and the retinal response to oxygen-induced injury during the phase of hyperoxia exposure. These studies suggest that microglia play a survival-promoting role for the nascent vessels in the immature retina. In models of laser-induced CNV, macrophages are localized to sites of tissue injury before the onset of NV, and systemic macrophage depletion results in reduced pathologic NV. Hence, in an inflammatory model with an infiltration of macrophages and neutrophils before NV, the macrophages appear to play a significant role in promoting angiogenesis. However, in the OIR model, macrophages/microglia are localized to the neovascular tufts on P17 and P21, peaking well after the onset of NV. We have also previously reported that neutrophil infiltration is not observed in the mouse model of OIR, supporting the notion that this model is not a proinflammatory model. Alternatively, in a rat model of ischemia-induced NV, depletion of monocytes/macrophages leads to the suppression of retinal NV, supporting an angiogenic role for infiltrating macrophages in the context of this model and species. Macrophages have also been linked to tumor angiogenesis, specifically when they are localized to regions of tumor hypoxia, where they release proangiogenic cytokines. Even in the same tumor, macrophages can display distinct and alternative functions, depending on the local microenvironment. Hence, macrophages/microglia have the potential to play many different roles in the processes of inflammation, angiogenesis, vascular regression, and neoplasia, depending on the temporal, spatial, and species biological context.

In summary, we demonstrate that the absence of MCP-1 expression in the mouse model of OIR results in reduced infiltration of macrophages, reduced vascular tuft apoptosis, and delayed regression of retinal NV. This suggests that the normal vascular regression observed in this model could potentially be mediated by the proapoptotic properties of macrophages/microglia. Further characterization of macrophage/microglia factors that induce retinal EC apoptosis may lead to new pharmacologic approaches in treating pathologic retinal NV, such as using intravitreal injection of soluble Fasl or TRAIL.

References


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