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# Cornea derived MSCs therapeutically modulate macrophage immunophenotype and angiogenic function

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# Abstract

Macrophages are crucial drivers of inflammatory corneal neovascularization and thus are potential targets for immunomodulatory therapies. We hypothesized that therapeutic use of cornea derived mesenchymal stromal cells (cMSCs) may alter the function of macrophages. We found that

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#### **Disclosure of Potential Conflicts of Interest**

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cMSCs can modulate the phenotype and angiogenic function of macrophages. In vitro, cMSCs induce apoptosis of macrophages while preferentially promoting a distinct CD14<sup>hi</sup>CD16<sup>hi</sup>CD163<sup>hi</sup>CD206<sup>hi</sup> immunophenotype that has significantly reduced angiogenic effects based on in vitro angiogenesis assays. In vivo, application of cMSCs to murine corneas after injury leads to reduced macrophage infiltration and higher expression of CD206 in macrophages. Macrophages co-cultured ("educated") by cMSCs express significantly higher levels of anti-angiogenic and anti-inflammatory factors compared to control macrophages. In vivo, injured corneas treated with cMSC-educated macrophages demonstrate significantly less neovascularization compared to corneas treated with control macrophages. Knocking down the expression of PEDF in cMSCs significantly abrogates its modulating effects on macrophages, as shown by the reduced rate of apoptosis, decreased expression of sFLT-1/PEDF, and increased expression of VEGF-A in the co-cultured macrophages. Similarly, cMSCs isolated from PEDF knockout mice are less effective compared to wild type cMSCs at inhibiting macrophage infiltration when applied to wild type corneas after injury. Overall, these results demonstrate that cMSCs therapeutically suppress the angiogenic capacity of macrophages and highlight the role of cMSC secreted PEDF in the modulation of macrophage phenotype and function.

# **Graphical abstract**

In the setting of severe corneal injury or inflammation, chemokines and cytokines recruit inflammatory cells which in turn induce corneal angiogenesis via VEGF-A. Cell therapy with corneal derived MSCs could modulate this inflammation in part via PEDF by inducing apoptosis of macrophages and promoting a distinct macrophage immunophenotype (cMSC-educated macrophages - cMEMq) that secrete high levels of sFLT-1, PEDF, and TSG-6.



#### Keywords

Mesenchymal Stromal Cells; Cornea; Macrophages; Inflammation; Angiogenesis; Pigment Epithelial Derived Factor; PEDF

# Introduction

The absence of blood vessels and hematopoietic cells including erythrocytes and leukocytes in the cornea is crucial for optical clarity and maintenance of vision [1]To achieve this, the cornea possesses an extensive array of mechanisms by which immune effector cells and neovascular processes are intricately regulated. These mechanisms include local expression of immunomodulatory factors such as Fas ligand, programmed death-ligand 1 (PD-L1) and TGF- $\beta$ , as well as expression of anti-angiogenic factors such as soluble vascular endothelial growth factor receptor 1 (VEGFR-1; soluble fms-like tyrosine kinase-1), VEGFR3, thrombospondin-1 and pigment epithelium derived factor (PEDF) [2-7].

Corneal neovascularization is often the final consequence of severe corneal infections or inflammation, which alter the balance in favor of angiogenic factors. Under most pathologic conditions that lead to corneal neovascularization, inflammatory cells including neutrophils and macrophages infiltrate the cornea. Macrophages in particular have been shown to play a central role in the development of inflammatory corneal neovascularization by releasing pro-angiogenic and pro-inflammatory cytokines [8-10]. Depleting macrophages largely prevents the development of corneal hemangiogenesis and lymphangiogenesis [11].

Current treatment options for corneal neovascularization include topical anti-inflammatory therapy, mainly corticosteroids, and local administration of anti-VEGF agents [12-14]. While both options can be effective in select situations, they have significant limitations. Corticosteroid-related side effects preclude their long-term use and even their short-term use in many scenarios including infectious and wound healing disorders. Anti-VEGF biological agents, which are used off label for corneal neovascularization, pose a challenge in both administering and maintaining sufficient tissue levels, while also having the potential for adversely affecting the tissue repair processes - given the pleiotropic nature of VEGF signaling [15]. Thus, many patients end up developing corneal neovascularization despite maximal standard therapies, underscoring the need for safer and more effective options in the clinic.

Mesenchymal stromal cells (MSCs) have been investigated therapeutically for a wide range of human diseases based on their immunomodulatory, anti-inflammatory, anti-fibrotic and other tissue repair properties; therefore, providing an attractive candidate for many cell based therapies [16-24]. It has been shown that one of major mechanisms of action of MSCs is through their interactions with the immune system, in particular macrophages [25-27]. In this current study, we hypothesized that corneal derived MSCs (cMSCs) can therapeutically modulate the immunophenotype and angiogenic properties of macrophages by promoting a macrophage phenotype that secretes higher levels of anti-inflammatory and anti-angiogenic factors. We specifically characterize and investigate the mechanistic basis of this therapeutic modulation of macrophages by cMSCs.

# **Materials and Methods**

### Mice

All experiments in mice were conducted in compliance with the ARVO Statement for the Use of Animals in Ophthalmic and Vision Research. The protocols were approved by the Animal Care & Use Committee of the University of Illinois at Chicago (UIC; Protocol numbers 15-258 and 14-166). PEDF<sup>-/-</sup> mice on C57BL/6J background were kindly provided by Dr. Paul J Grippo (UIC). C57BL/6J mice were used as wild type.

#### Cell Transplantation

Limbal stem cell deficiency was induced by total corneal epithelial debridement which also creates an inflammatory environment that leads to corneal neovascularization [28, 29]. A 6-0 nylon suture was pre-passed through the lids (in preparation for closing the lids afterwards). Subsequently, 0.2  $\mu$ l thrombin (Ethicon, Johnson and Johnson Inc.) was placed over the debrided cornea. A total of 5000 cells, resuspended in 3  $\mu$ l fibrinogen (Ethicon, Johnson and Johnson Inc.), were transferred over the cornea [30, 31]. After gel formation, the suture was tightened to closed the lids and then tied. Erythromycin ophthalmic ointment was applied. The suture was removed after 3 days.

#### cMSC Culture

Human and mice corneal MSCs were isolated, expanded and characterized as we described before [30] (See Supplementary Figure 1 for PEDF<sup>-/-</sup> cMSCs). Briefly, corneoscleral button from healthy cadaver eyes or freshly enucleated mouse eyes were washed 5 times with PBS containing 2× Antibiotic-Antimycotic and 2× penicillin- streptomycin (both from Thermo Fisher). After removing the central button with a trephine, the limbus was cut into 3 segments and were placed in 2.4 IU of Dispase II (Thermo Fisher) for 1 hour at 37°C. Intact epithelial sheets were removed from stroma. The limbal segments were cut into small pieces and incubated in collagenase Type I (0.5 mg/ml) (Sigma-Aldrich) in DMEM/F12 media (Thermo Fisher) overnight at 37°C or else used directly for explant culture. The digests were filtered through a 70-µm nylon strainer to obtain a single-cell suspension. They were seeded onto 1% gelatin (Sigma-Aldrich)-coated wells of a 6-well tissue culture plate in alpha MEM media supplemented with 10% fetal bovine serum, 1X L-Glutamine, and 1X NEAA (All from Corning, Manassas, VA). Culture media were changed every other day, and cells were sub-cultured by brief digestion with TrypLE Express (Thermo Fisher) when 80% confluent. At least 3 mouse cMSC lines were used from wild type and PEDF-/- mice. For this study, 5 human cMSC lines were isolated from corneoscleral button of cadaveric donors (C-4977, C-4732, C-6179, C-6436, and C-5422) generously provided by Eversight Eye Bank (Chicago IL). Since the research did not involve human subjects and the tissue was from cadaver donors without any identifying or protected health information, IRB approval was not necessary per guidelines of the Office for the Protection of Research Subjects at UIC. Data from different donors were pooled for statistical analysis.

#### MSC Conditioned media

Upon reaching 100% confluency in a T175 flask, the MSCs were washed with 30 ml prewarmed PBS 3 times. The media was then changed to phenol red free alpha MEM media supplemented with 1X L-Glutamine, and 1X NEAA. The conditioned media was collected after 48 hours. The cells were trypsinzed and counted at the same time. The conditioned media was centrifuged with 500 G speed for 15 minutes to remove any cells or debris. The supernatant was transferred to a new tube and was used for experiments or kept at 4° C for up to a week.

#### **MSC** Differentiation

After incubation in appropriate differentiation medium (StemPro<sup>®</sup> Differentiation kits, all from Thermofisher), cMSC differentiation into mesenchymal lineages (osteoblasts and adipocytes) was evaluated using staining with Alizarin Red for calcium and Oil-Red-O for fatty acids, respectively.

#### Immunostaining

Corneal whole mount immunostaining was done as described before [32]. Briefly, after enucleation of the eye, the cornea was dissected using a spring scissor. It was then fixed with 4% paraformaldehyde at 4° C overnight. After washing with PBS, it was incubated with 20 µg/ml Proteinase-K (Sigma-Aldrich) for 5 minutes at room temperature followed by 100% methanol for another 30 minutes. It was then incubated with 10% serum and 2% BSA at 4° C overnight. Purified anti-mouse CD31 (Biolegend, cat# 102401) was used as primary antibody and Rhodamine-conjugated donkey anti-rat IgG (Jackson ImmunoResearch, Cat# 712-025-153) as secondary antibody. The slides were visualized and photographed using the Zeiss LSM 710 microscope (Carl Zeiss). To make the quantification more objective, we used VesselJ [33] to quantify the ratio of vascularized area to total corneal area.

# **Tube Formation Assay**

Human umbilical vein endothelial cells (HUVEC, Thermo Fisher) were cultivated in Media 200 fortified with low serum growth supplement (LSGS). At passage 4, the cells were detached and resuspended in Media 200 + 10% LSGS which was used as the base media. The 96-well plate were filled with 75  $\mu$ l Matrigel (Thermo Fisher) and allowed to solidify at 37° C after which HUVECs (2 × 10<sup>4</sup> cells/75  $\mu$ l) were gently seeded on top of the gel. Subsequently, 75  $\mu$ l of conditioned media, complete media or Media 200 was added as the testing condition, positive and negative controls, respectively. After 6 h, network structures were analyzed and photographed at 4× magnification (Leica Microsystems). Total tubule length per image was calculated using Angiogenesis Analyzer plugin of ImageJ (NIH).

#### Enzyme-Linked Immunosorbent Assay (ELISA)

To get macrophage conditioned media, they were washed thoroughly with PBS after 7 days of education and received fresh complete media. After 24 hours, the media was collected and cells were detached and counted. The conditioned media was cleared of cellular debris by centrifugation at 500 g for 15 min. The supernatant was transferred to a new tube and was used for experiments or kept at  $4^{\circ}$  C for up to a week.

Human TSG-6 protein levels in the conditioned media were determined by ELISA as described before [34]. Human PEDF (cat# DY1177-05), sFLT-1 (cat# DVR100B) and VEGF-A (cat# DVE00) proteins were detected with commercially available ELISA kits (all from R&D Systems) according to the manufacturer's protocol. The obtained values were normalized to total cell numbers.

#### siRNA transfection

After reaching 70 to 80% of confluency, cMSCs were transfected with 50 nM PEDF siRNA (Santa Cruz, Cat# sc-40947) or scrambled siRNA (Dharmacon, Cat# D-001205) in TransIT-TKO siRNA transfection reagent (Mirus, Cat# MIR2150). The transfection efficiency was tested with ELISA on the conditioned media.

#### Human Macrophage culture

Macrophages were isolated and cultivated as described before [35, 36]. Briefly, peripheral whole blood buffy coats from healthy donors were purchased from Interstate Blood Bank, INC (Memphis, Tennessee). Peripheral blood mononuclear cells (PBMCs) were isolated from the buffy coats by density-gradient separation. Monocytes were isolated using antihuman CD14 microbeads (Miltenyi Biotech) and autoMACS Pro Separator (Miltenyi) according to manufacturer's protocol. Purity of isolated CD14+ cells was over 95% when checked using an Acuri C6 cytometer (BD Biosciences). Purified CD14+ monocytes were plated into the bottom chamber of 6 well 0.4 um pore size transwell cell culture plates (Costar, Cat# 3450) at a concentration of  $1 \times 10^6$  per well in IMDM media supplemented with 10% human serum blood type AB (Mediatech, Herndon, VA, USA), 1X Non-essential amino acids (NEAA-Lonza), (Thermo Fisher), 1mM sodium pyruvate (Mediatech) and 4 µg/ml recombinant human insulin (Invitrogen). Monocytes were cultured for 7 days at 37° C with 5% CO2 to differentiate to macrophages. On day 7, macrophages were supplemented with fresh media and incubated with cMSCs in the upper chamber of the transwell plate to obtain cMSC educated Macrophages (cMEMqs) at a ratio of 10: 1 of Macrophages:cMSCs, as previously described [36]. After 7 days of co-culture, cMEMqs were collected using Stem Pro Accutase cell detachment (Thermo Fisher) for experiments. For control, macrophages were either cultured in the same media without cMSCs or co-cultured with a nonmesenchymal cell type.

#### Mouse Macrophage Culture

Mouse peritoneal macrophages were isolated as described previously [37]. In brief, mice were euthanized by CO2 asphyxiation and then cleaned with 70% ethanol and mounted onto a Styrofoam block. Outer skin of the peritoneum was cut with a scissor and forceps and gently pulled back to expose the inner skin lining the peritoneal cavity. Using a 22-gauge needle, 10 ml of ice cold IMDM media was injected into the peritoneal cavity with precaution not to puncture any organ. The media, that contains dislodged cells, was aspirated using a 19-gauge needle. The cell suspension was centrifuged and washed twice. Subsequently, they were purified using F4/80 magnetic beads (Miltenyi) according to the manufacturer's protocol. They were plated into 6 well cell culture plates at a concentration of  $1 \times 10^6$  per well in IMDM media supplemented with 10% FBS, 1X NEAA, 1mM sodium pyruvate and 4 µg/ml recombinant human insulin. On day 3, they were supplemented with

fresh media and incubated with mouse cMSCs similar to what was described above for the human macrophage co-culture. On day 10, mouse cMEMqs were collected using Accutase cell detachment for experiments.

## Flow Cytometry

The cells were detached as described above. They were incubated with Fc block (BD Pharmingen, cat#: 564220 for human and Biolegend TruStain fcX<sup>™</sup> [anti-mouse CD16/32], cat# 101319 for mouse) and stained at 4°C for 20 minutes in antibody diluent (PBS with 2% FBS) with cell surface antibodies (Supplementary Table 1). For intracellular staining, fixation buffer (Biolegend, cat# 420801) and intracellular staining perm wash buffer (Biolegend, cat# 421002) were used according to manufacturer's protocol. Compensation was performed using Ultracomp e-beads (Cat# 01-2222-42) ebiosciences, (San Diego, CA). Flow cytometry data was acquired on the MACSQuant analyzer 10 (Miltenyi) and BD LSR Fortessa (BD). Data were analyzed using FlowJo software (TreeStar).

#### Fluorescent-Activated Cell Sorting (FACS)

Seven days after the procedure, mice were euthanized. The cornea was digested in 0.25% trypsin (Thermo Fisher). After filtration through a 70 µl nylon strainer, the cells were stained as described in Flow Cytometry section. Cell sorting was done by a Sony cell sorter (Sony Biotechnology SH800).

#### Apoptosis assay

To measure caspase 3/7 activation, ApoTox-Glo<sup>TM</sup> Triplex Assay (Promega) was used according to the manufacturer's protocol.

#### **Real-time quantitative PCR**

Total RNA was extracted using the miRCURY<sup>tm</sup> RNA isolation kit (Exiqon) according to manufacturer's protocol. After spectrophotometric assessment for quality and concentration (Nanodrop ND-1000; Thermo Scientific), 2 μg of purified RNA was reverse transcribed to single-strand cDNA using High Capacity cDNA Reverse Transcription Kit (Applied Biosystems Inc [ABI]) according to the manufacturer's protocol. TaqMan Universal PCR Master Mix (ABI) or FastStart Universal SYBER GREEN Master (Roche, Germany) was used to make the master-mix. All primers were bought from Integrated DNA technology (IDT; Iowa, USA) (Supplementary Table 2). To detect human sFLT-1, we used 5'-TTTGTTGCAGTGCTCACCTC-3' (Forward) and 5'-GTTGGGAACTGTGGGAAGAAA-3' (Reverse) primer that only detects splice variant 2 of FLT-1 mRNA. Real-time qPCR was performed using QuantStudio 7 Flex Real-Time PCR System (ABI). The mRNA expression levels were normalized to the expression of a housekeeping gene, Glyceraldehyde-3-Phosphate Dehydrogenase (GAPDH). The 2<sup>-CT</sup> method was used to quantify relative mRNA expression.

#### **Data Analysis and Statistical Comparisons**

To avoid any observer bias and to increase reproducibility, all the animal surgeries were done by one of the authors who was blinded to the treatment arms. Moreover, all the data

collection and analyses were done in a blinded fashion to minimize inter and intra-observer bias. Eyes with infection were excluded from the study. Corneal infection was defined as corneal edema, hypopyon, and exudate within the first 7 days after the procedure. Results are presented as the mean  $\pm$  SD of three independent experiments. Normality of the data was tested using D'Agostino & Pearson normality test. Based on normality test, Mann-Whitney *U*-test or 2-sided student's *t*-test was performed to determine significance, which was set at p < 0.05. For more than 2 arms comparison, One-way ANOVA with Tukey's post hoc correction was used. All statistics were performed using GraphPad Prism software 7.0 (GraphPad Software) and Microsoft Excel (Microsoft).

# Results

#### cMSCs modulate macrophage immunophenotype

To evaluate the effects of human cMSCs on the macrophages, we used our previously described co-culture method [35, 36]. Macrophages co-cultured with cMSCs demonstrated reduced expression of CD86 and HLA-DR, markers typically associated with inflammatory macrophages, and at the same time showed increased expression of CD163, CD206, CD14 and CD16, markers associated with an inflammation-resolving macrophage immunophenotype (Fig. 1B) [8, 38-40]. Human cMSC conditioned media was found to induce macrophage apoptosis (Fig. 1C).

We further investigated the effect of cMSCs on the macrophage immunophenotype in an *in vivo* corneal injury model. We embedded murine cMSCs in fibrin gels and applied them to the cornea immediately after epithelial injury. Clinical and histologic studies of the corneas after corneal epithelial debridement demonstrated significantly reduced haze with reduced F4/80+ macrophages infiltration in the corneas treated with murine cMSCs compared to those treated with fibrin gels alone (Fig. 2A). To quantify and characterize the infiltrated macrophages, we digested the mice corneas 7 days after treatment and subjected them to flow cytometry. The murine cMSC treated corneas had significantly reduced number of F4/80+ macrophages. However, expression of CD206 was higher in F4/80+ macrophages in the murine cMSC treated corneas compared to fibrin gel treated corneas (P<0.001) (Fig. 2B).

#### cMSCs modulate the angiogenic properties of macrophages

To investigate the effect of cMSCs on the angiogenic function of macrophages, we examined the protein and gene expression of macrophages after 7 days of co-culture with cMSCs – defined as cMSC educated macrophages (cMEMqs). By gene expression, we found the CD14+CD163+ <sup>sorted</sup> population of cMEMqs expressed significantly lower levels of VEGF-A, and significantly higher levels of anti-angiogenic genes (sFLT-1, PEDF and TIMP-1) relative to CD14+CD163+ control sorted macrophages (Fig. 3A). Similarly, cMEMqs secreted more sFLT-1, PEDF, and TSG-6, without any significant change in VEGF-A compared to control macrophages (Fig. 3B).

To investigate the angiogenic function of macrophages *in vivo*, we used a limbal stem cell deficiency model. In this model, the entire corneal epithelium is mechanically debrided

limbus to limbus which results in severe corneal neovascularization. After 7 days of coculture with mouse cMSCs, mouse macrophages were embedded in fibrin gels and applied to the cornea after total epithelial injury. Corneas treated with cMSC educated mouse macrophage (murine cMEMq) had significantly less corneal neovascularization compared to control mouse macrophages treated corneas (Fig. 4A). Likewise, the conditioned media from cMEMqs induced significantly less vascular endothelial tubule formation compared to control macrophage conditioned media (Fig. 4B).

#### cMSC derived PEDF is a key mediator of macrophage modulation

To investigate the mechanism of macrophage modulation by cMSCs, we focused on cMSCsecreted PEDF, given its combined anti-angiogenic and anti-inflammatory effects [1, 41, 42]. We isolated cMSCs from PEDF<sup>-/-</sup> mice and used our *in vivo* inflammatory injury model as described above to investigate the potential role of PEDF. Injured wild type murine corneas treated with PEDF<sup>-/-</sup> murine cMSCs demonstrated more macrophage infiltration with reduced expression of CD206 compared to wild type (PEDF<sup>+/+</sup>) cMSCs treated corneas (Fig. 5A).

By gene expression, mouse macrophages educated with PEDF <sup>-/-</sup> cMSCs expressed more iNOS and less arginase-1 with concomitant reduction in the expression of IL-10, PEDF, and sFLT-1 compared to the control macrophages (Fig. 6A). Similar results were found with human cMSC educated macrophages. In particular, human macrophages were co-cultured with PEDF-knockdown cMSCs and found to secrete more VEGF-A, less PEDF and less sFLT-1 compared to conditioned media from macrophages educated with control (scrambled siRNA) cMSCs (Fig. 6B). The rate of apoptosis in macrophages was likewise significantly reduced when they were exposed to the conditioned media from PEDF-knockdown cMSCs compared to control (Fig. 5B). Overall, these results implicate a mechanistic role for PEDF in the cMSC modulation of macrophages and their angiogenic function.

# Discussion

In this study, we focused on the modulatory effects of cMSC on macrophages and show that cMSC-educated macrophages have significantly reduced angiogenic effects both *in vitro* and *in vivo*. cMSCs specifically induce markers that are associated with the anti-inflammatory and pro-resolving function of the macrophages. Mechanistically, we found cMSC secreted PEDF as an important mediator of its macrophage modulation.

It is well accepted that the therapeutic effects of MSCs, most commonly derived from the bone marrow, are mediated largely through their immunomodulatory and anti-inflammatory properties [43-47]. MSCs have been shown to modulate both innate and adaptive immune mechanisms [48-51]. They have also been implicated in angiogenesis and wound repair [52-54]. Macrophages, in particular, have been studied as one of the major mediators of MSCs' immunomodulatory and regenerative effects [25-27, 55-60]. The current study provides the first direct evidence that MSCs derived from the cornea modulate the immunophenotype and angiogenic function of macrophages in a distinct manner. An important finding of our studies is that cMSCs induce apoptosis of macrophages *in vitro* as a

potential mechanism of suppressing the monocytic inflammatory response. Likewise, cMSC treated corneas *in vivo* demonstrated significantly reduced macrophage recruitment.

Macrophages exhibit extraordinary phenotypic plasticity during different stages of inflammation and repair, where they dynamically orchestrate an immune and later a reparative response specific for that tissue [61]. Traditionally, macrophages have been classified into subtypes, such as classic versus alternatively activated, or M1 and M2, mostly defined by in vitro cytokine stimulation and subsequent marker/cytokine expression profile [8]. This study provides new insights into the distinct phenotype of macrophages "educated" by cMSCs. Our results demonstrated that cMSCs promote markers of M2 (CD163, CD206) while suppressing those associated with M1 phenotype (CD86, HLA-DR). However, it may be simplistic to consider macrophage plasticity just as a simple binary phenomenon [62]. A satisfactory paradigm should take into account the specific function of the macrophages, along with spatial and temporal attributes of their environment, including the stage of inflammation and their tissue of residence [61]. Consistent with their dynamic nature, macrophages could be both anti-angiogenic and/or pro-angiogenic, depending on the stage of inflammation, playing a critical and complex role in regulating angiogenesis at sites of tissue injury [63-65]. We have shown that cMSCs promote macrophages with a distinct phenotype favorable for the terminal stages of corneal tissue repair and remodeling. Specifically, cMSC-educated macrophages demonstrated high expression of CD163, CD206, CD14 and CD16 consistent with a profile of macrophages that help resolve inflammation in part by phagocytizing apoptotic cells, including neutrophils [39, 66, 67].

In the cornea, macrophages have been shown to play a central role in the development of inflammatory corneal neovascularization and depleting macrophages largely prevents the development of hemangiogenesis and lymphangiogenesis [9, 11]. In our study, cMSCs, when used therapeutically, not only reduced macrophage infiltration after injury *in vivo* but also in co-culture experiments *in vitro*, enhanced the macrophages' secretion of anti-angiogenic factors (sFLT-1 and PEDF), consistent with the role of these macrophages during the latter phases of corneal tissue repair and remodeling [64, 65, 68]. It is well-known that macrophages have remarkable plasticity based on the spatial and temporal attributes of their environment, including the stage of inflammation and their tissue of residence [62]. Therefore, the observed phenotype of cMEMqs are not likely to be end stage and can potentially change with time.

Our findings also implicate cMSC derived PEDF as an important mediator of their modulating effects on macrophages. During skin wound healing, PEDF is typically found in the remodeling phase where regression of neo-vessels takes place, a phenomenon that is important not only in the cornea but also in vascularized tissues such as the skin [69]. PEDF selectively inhibits the formation and promotes the regression of neo-vessels but has no detrimental effect on mature vascular structures [41]. The inhibitory effect of PEDF on VEGF-induced angiogenesis is mediated in part via  $\gamma$ -secretase-dependent cleavage of the C terminus of VEGF receptor (VEGFR)-1, which in turn inhibits VEGFR-2-induced angiogenesis [70]. PEDF has likewise been recognized to exert anti-inflammatory effects in part by inhibiting macrophage recruitment and activation [42, 71]. Our results demonstrate that cMSC derived PEDF is not only involved in the induction of macrophage apoptosis, but

also contributes to the development of anti-inflammatory macrophages which, in the terminal stage of repair, function by removing immune cells, granulation tissue and unnecessary neovasculature.

As noted earlier, we only studied cMSCs in the context of their therapeutic effects as a cell therapy and did not specifically examine the role of cMSCs that already exist in the cornea during inflammation. We also focused specifically on cornea-derived MSCs and did not study MSCs from other sources. However, it has been shown that bone marrow derived MSCs are also able to modulate macrophage immunophenotype and function [26, 27, 35, 36, 47, 49, 55, 58, 72, 73]. Further studies are needed to determine if MSCs from other sources can similarly modulate the angiogenic function of macrophages.

Overall, this study advances our understanding of the mechanisms by which cMSCs modulate macrophages and specifically adds PEDF to the list of important mediators (such as TSG-6, stanniocalcin 1, and prostaglandin E2) which MSCs secrete to modulate macrophages [26, 34, 74-77]. It will be interesting to further examine the role of PEDF in mediating the therapeutic effects of MSCs from other source including the bone marrow.

# Conclusion

Our study reveals that corneal derived MSCs can therapeutically modulate the immunophenotype and the angiogenic function of the macrophages in part through PEDF. These results highlight the potential of tissue specific MSCs for targeted immunomodulatory and anti-angiogenic cell-based therapies.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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# **Significance Statement**

Macrophages play a central role in pathologic corneal neovascularization. We found that mesenchymal stromal cells from the cornea (cMSCs) therapeutically modulate the inflammatory and angiogenic functions of macrophages. In particular, cMSCs induce macrophage apoptosis and preferentially promote macrophages that secrete high levels of anti-angiogenic factors. We further demonstrate that cMSC modulation of macrophages significantly depends on cMSC secreted pigment epithelium-derived factor.



#### Figure 1. cMSCs modulate macrophage immunophenotype

A: Control Mq and cMEMqs were expanded as described in the methods. On day +14, human control Mq or cMEMqs were analyzed by flow cytometry. Cell surface expression of CD86, HLA-DR, CD206 (Mannose receptor), CD163 (scavenger receptor), CD14 and CD16 were determined in control Mq and cMEMqs. Histograms represent MFI (Mean fluorescent intensity) in unstained Mq, control Mq and cMEMqs. **B:** The mean from independent biological replicated are presented as the bar charts. Educating macrophages with cMSCs reduced the expression of CD86 and HLA-DR, and increased the expression of CD163, CD206, CD14 and CD16 (*n*=4, 2-sided t-test \* *P*<0.001). The values shown are mean  $\pm$  SD (error bars). **C:** Conditioned media from human cMSCs induced more apoptosis in macrophages compared to unconditioned media (*n*=5, Mann-Whitney *U*-test: *P*=0.0079) Boxes show the interquartile (25%–75%) range, whiskers encompass the range (minimummaximum), and horizontal lines represent the mean. MFI: median fluorescein intensity, Cas: Caspase, <sup>CM</sup>: conditioned media, Ctrl: control, Mq: macrophage, cMEMq: cMSC-educated macrophage.



# Figure 2. cMSCs modulate macrophage phenotype and recruitment into the corneal stroma after injury

A: Representative immunofluorescence staining after topical application of murine cMSCs to the injured murine cornea, demonstrating significantly reduced infiltration of F4/80+ macrophages, and increased CD206+ cells in the stroma compared to control treated corneas. Scale bar, 50  $\mu$ m. **B**: To quantify, corneas were digested and subjected to flow cytometry which demonstrated significantly reduced the percentage of F4/80+ cells to total cells in murine cMSCs treated corneas compared to the control. Among the F4/80+ gated cells, there was the higher expression of CD206 in the cMSCs treated corneas compared to control treated corneas compared to control treated corneas compared to control treated corneas compared to the cMSCs treated corneas compared to control treated corneas (*n*=5, 2-sided t-test \* *P*<0.001).



#### Figure 3. cMSCs modulate the angiogenic properties of macrophages

A: CD14+CD163+ population of human cMEMqs were isolated by fluorescence-activated cell sorting and subjected to quantitative RT-PCR demonstrating lower expression of VEGF-A and higher expression of PEDF, sFLT-1, and TIMP-1 compared to CD14+CD163+ sorted control macrophages (n=5, P<0.001 for all comparison). **B:** The conditioned media from cMEMqs contains more sFLT-1 (1296 ± 715.3 vs. 276.6 ± 168.0 pg/ml), PEDF (17741 ± 8546 vs. 4956 ± 3851 pg/ml) and TSG-6 (10357 ± 2190 vs. 2757 ± 823.7 pg/ml) compared to the conditioned media from control untreated macrophages (n=5). The values shown are mean ± SD (error bars). 2-sided *t*-test: \* P<0.0001. Mq: macrophage, cMEMq: cMSC-educated macrophage.



**Figure 4. cMSC-educated macrophages demonstrate reduced angiogenic properties A:** Murine macrophages were co-cultured with murine cMSCs as described in methods (cMEmMq). Applying cMEmMqs in fibrin gels to the cornea of wild-type mice after injury resulted in significantly less neovascularization compared to control macrophages on day ten (n=5, 2-sided *t*-test: \* P < 0.0001). Scale bar, 500 µm. **B:** Conditioned media from cMEMqs inhibited tubule formation by human umbilical vein endothelial cells (HUVEC cells) compared to conditioned media from control untreated macrophages (n=5, 2-sided *t*-test: \* P< 0.0001). Mq: macrophage, cMEmMq: cMSC-educated mouse macrophage, NV: neovascularization, CM: conditioned media.



# Figure 5. cMSC derived PEDF is a key mediator of macrophage modulation

A: Injured wild-type corneas treated either with cMSCs obtained from the PEDF<sup>-/-</sup> mice (placed in fibrin gels) or from wild-type PEDF <sup>+/+</sup> mice were digested and subjected to flow cytometry. Injured wild-type corneas treated with cMSC<sup>PEDF-/-</sup> had more infiltration of F4/80+ macrophages with lower expression of CD206 compared to the those treated with cMSC<sup>PEDF+/+</sup> (*n*=5; \**P*=0.03). **B:** knocking down PEDF by siRNA decreased PEDF protein by almost 90% compared to scrambled siRNA (n=9, 2-sided t-test: \*P<0.0001). Conditioned media from PEDF knockdown cMSCs induced less apoptosis in macrophages compared to conditioned media from control cMSC (*n*=5, Mann-Whitney *U*-test: *P*=0.007). The values shown in bar graphs are mean ± SD (error bars). Boxes show the interquartile (25%–75%) range, whiskers encompass the range (minimum–maximum), and horizontal lines represent the mean. 2-sided *t*-test used for all comparison. MFI: median fluorescein intensity, SC: scrambled, Mq: macrophage, Cas: Caspase.



Figure 6. cMSC derived PEDF is a key mediator of macrophage angiogenic function A: RT-qPCR shows mouse macrophages educated with wild type PEDF<sup>+/+</sup> cMSCs, expressed more arginase-1, CD206, IL-10, PEDF, and sFLT-1; and less iNOS mRNA compared to control untreated macrophages. Conversely, macrophages educated with PEDF  $^{-/-}$  cMSCs, expressed less arginase-1, CD206, IL-10, PEDF, and sFLT-1 with more iNOS (*n*=5; *P*<0.001 for all comparison). **B:** Macrophages educated with PEDF knockdown cMSCs secreted more VEGF-A (448.3 ± 104.1 vs. 229.4 ± 125.1 pg/ml; *P*= 0.03) and less PEDF (7162.1 ± 2918.0 vs. 22373 ± 9683.0 pg/ml; *P*= 0.02) and sFLT-1 (249.1 ± 98.54 vs. 1172 ± 473.4 pg/ml; *P*= 0.008) compared to those trained with control cMSCs. The mean TSG-6 expression although decreased was not statistically significant (1160 ± 1309.0 vs. 3035 ± 3032.0 pg/ml) (*P*= 0.32). The values shown in bar graphs are mean ± SD (error bars). 2-sided *t*-test used for all comparison. M<sup>PEDF</sup> EmMq: cMSC<sup>PEDF</sup>-educated mouse macrophage, M<sup>Sc siRNA</sup> EMq: cMSC<sup>Scrambled siRNA</sup> educated macrophage, M<sup>PEDF siRNA</sup> EMq: cMSC<sup>PEDF siRNA</sup> educated macrophage.