

Monocyte chemoattractant protein 1 and fractalkine play opposite roles in angiogenesis *via* recruitment of different macrophage subtypes

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Abstract

• **AIM:** To explore the interaction between macrophages and chemokines [monocyte chemoattractant protein 1 (MCP-1/CCL2) and fractalkine/CX3CL1] and the effects of their interaction on neovascularization.

• **METHODS:** Human peripheral blood mononuclear cells, donated by healthy volunteers, were separated and cultured in RPMI-1640 medium containing 10% fetal bovine serum, then induced into macrophages by stimulation with 30 µg/L granulocyte macrophage-colony stimulating factor (GM-CSF). The expression of CCR2 and/or CX3CR1 in the macrophages was examined using flow cytometry. Macrophages were then stimulated with recombinant human CCL2 (rh-CCL2) or recombinant human CX3CL1 (rh-CX3CL1). The expression of angiogenesis-related genes, including *VEGF-A*, *THBS-1* and *ADAMTS-1* were examined using real-time quantitative polymerase chain reaction (PCR). Supernatants from stimulated macrophages were used in an assay of human retinal endothelial cell (HREC) proliferation. Finally, stimulated macrophages were co-cultured with HREC in a migration assay.

• **RESULTS:** The expression rate of CCR2 in macrophages stimulated by GM-CSF was 42%±1.9%. The expression rate of CX3CR1 was 71%±3.3%. Compared with vehicle-treated groups, gene expression of *VEGF-A* in the macrophages was greater in 150 mg/L CCL2-treated groups ($P<0.05$), while expression of *THBS-1* and *ADAMTS-1* was significantly lower ($P<0.05$). By contrast, compared with vehicle-treated groups, expression of *VEGF-A* in 150 mg/L CX3CL1-treated groups was significantly lower ($P<0.05$), while

expression of *THBS-1* and *ADAMTS-1* was greater ($P<0.05$). Supernatants from CCL2 treated macrophages promoted proliferation of HREC ($P<0.05$), while supernatants from CX3CL1-treated macrophages inhibited the proliferation of HREC ($P<0.05$). HREC migration increased when co-cultured with CCL2-treated macrophages, but decreased with CX3CL1-treated macrophages ($P<0.05$).

• **CONCLUSION:** CCL2 and CX3CL1 exert different effects in regulation of macrophage in expression of angiogenesis-related factors, including *VEGF-A*, *THBS-1* and *ADAMTS-1*. Our findings suggest that CCL2 and CX3CL1 may be candidate proteins for further exploration of novel targets for treatment of ocular neovascularization.

• **KEYWORDS:** chemokine; macrophage; proliferation; migration; angiogenesis

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INTRODUCTION

Angiogenesis, formation of new blood vessels from existing vessels, is precisely controlled by angiogenic and angiostatic cytokines^[1]. It is involved in various pathological processes, including inflammation, wound healing, and tumor growth^[2-4]. When pathological angiogenesis occurs in the eye, as in neovascular age-related macular degeneration (nAMD) or proliferative diabetic retinopathy (PDR), vision may be heavily impaired and may progress to blindness^[5-6]. In an aging population with rising living standards, the morbidity rate of nAMD and PDR is increasing, leading to decreased quality of life and increasing health care burdens^[7]. Therefore, ophthalmologists are paying increasing attention to prevention of vision loss by inhibiting ocular neovascularization.

Many studies suggest that macrophages migrate to lesion sites and modulate angiogenesis *via* the action of a number of cytokines^[8]. The recruited macrophages or resident macrophages differentiate into subtypes depending on the

specific microenvironment^[9]. The first step is migration to lesion sites after binding to chemokines^[10]. Chemokines are members of a superfamily of small signaling molecules that regulate migration of target cells^[11]. Monocyte chemoattractant protein 1 (MCP-1/CCL2), a member of the C-C motif-chemokine subfamily, exhibits chemotactic potential for monocytes/macrophages *via* binding to its cognate receptors CCR2 or CCR4. In addition, CX3C chemokine ligand 1 (CX3CL1/fractalkine), a member of C-X-C motif-chemokine subfamily, can also recruit mononuclear cells (including macrophages) to lesion sites by binding to CX3CR1^[12-15]. Several studies suggest that CCL2 and CX3CL1 are involved in development of pathological neovascularization, including monocytes/macrophages recruitment, and mediation of cytokine secretion by target cells^[16-17].

Vascular endothelial growth factor A (VEGF-A), one of the most potent proangiogenic factors, promotes vascular endothelium migration and proliferation^[18]. Thrombospondin 1 (*THBS-1*) and *ADAMTS-1* (a disintegrin and metalloproteinase with a thrombospondin motif), are principal anti-angiogenic factors. These factors suppress neovascularization by counteracting VEGF-A, thereby inhibiting endothelial cell migration and survival^[19]. In our study, we examined the effects of recombinant human CCL2 protein (rh-CCL2) and recombinant human CX3CL1 protein (rh-CX3CL1) on macrophages *in vitro*.

MATERIALS AND METHODS

Samples Preparation Human peripheral blood was obtained from healthy volunteers. Informed consent was obtained from all donors. The study was performed in accordance with ethical standards of the Declaration of Helsinki and the Ethics Committees of Soochow University, Suzhou, China, who approved all the protocols.

Reagents and Antibodies Human retinal endothelial cells (HREC) were purchased from Ya Ji Biologic Technologies (Shanghai, China). The RNeasy Mini Kit and RNase-Free DNase Set were purchased from QIAGEN Technical Services (Hilden, Germany). PrimeScript RT Master Mix and DRR041A SYBR Premix Ex Taq (Perfect Real Time) were purchased from TAKARA bio (Dalian, China). RPMI-1640, DMEM cell culture medium, fetal bovine serum (FBS) and phosphate-buffered saline (PBS) were obtained from ThermoFisher Scientific (Pittsburgh, USA). Recombinant human granulocyte macrophage-colony stimulating factor (GM-CSF) protein was purchased from R&D Systems (Minneapolis, USA). Ficoll-Paque PLUS was purchased from GE Healthcare Life Sciences (Pittsburgh, USA). Recombinant Human CCL2 protein and recombinant human CX3CL1 protein were purchased from BioLegend (San Diego, USA). Mouse anti-human CD68 PE-conjugated antibody (IC20401P) was purchased from R&D Systems (Minneapolis, USA). Rabbit anti-human CCR2 antibody (aa20-100) and rabbit anti-human CX3CR1 antibody

(aa175-189) were provided by LifeSpan BioSciences (Seattle, USA). Primers were synthesized by GENEWIZ (Suzhou, China). The Cell Counting Kit-8 (CCK-8) was obtained from Dojindo Molecular Technologies (Shanghai, China). Transwell plates with 8.0 μ m pore polycarbonate membrane insert were purchased from Corning Life Science (New York, USA).

Cells Isolation and Culture Totally 100 mL human peripheral blood was mixed with equal volumes of Hank's balanced salt solution. Peripheral blood mononuclear cells (PBMC) were then isolated by density gradient centrifugation. After rinsing with PBS twice, cells were resuspended in RPMI-1640 medium supplemented with 10% FBS (mass fraction). About 1×10^5 cells per well were seeded in a 6-well plate. Non-adherent cells were removed 2h after seeding, and culture medium was replaced with fresh medium the following day. Cultured cells were observed under phase-contrast microscope. Totally 30 μ g/L of GM-CSF was added to the medium for one week, and the medium was replaced every three days. Macrophages were harvested after induction by GM-CSF.

Flow Cytometry Macrophages derived from isolated monocytes were verified by flow cytometry. Adherent cells were pooled and cell pellets were obtained by centrifugation at 1600 rpm at 4 °C for 5min. Cells were then resuspended in PBS at a density of 1×10^6 cells/mL. Totally 100 μ L of PBS containing approximately 1×10^5 cells were then pipetted into each polypropylene tube. The samples were divided into CD68+CCR2 and CD68+CX3CR1 groups. Group CD68+CCR2 was stained with PE-conjugated mouse anti-human CD68 antibody (1:100) and rabbit anti-human CCR2 antibody (1:100); Group CD68+CX3CR1 was stained with PE-conjugated mouse anti-human CD68 antibody (1:100) and rabbit anti-human CX3CR1 antibody (1:100). The samples were then incubated in the dark for 30min. Following PBS washing and two centrifugations, the samples were incubated with appropriate secondary antibodies in the dark for 30min. The samples stained with non-immunized mouse IgG mAb and rabbit IgG were used as isotype controls. All samples were detected using flow cytometry. The data was then analyzed using FlowJo version 10 software (USA).

Intervention on Macrophages with CCL2 or CX3CL1 Monocyte-derived macrophages were seeded on 6-well plates at a density of 5×10^5 cells/well. Cells were pre-cultured in RPMI-1640 supplemented with 10% FBS containing GM-CSF. Medium was replaced with serum-free medium for 6h followed by replacement with complete medium supplemented with 10% FBS. rh-CCL2 or rh-CX3CL1 in various concentrations was added to appropriate groups and the cells were incubated for 12h at 37 °C in a humidified incubator with 5% CO₂. The cells were then pooled for subsequent assays.

Quantitative Reverse Transcription Polymerase Chain Reaction The mRNAs of angiogenesis-related cytokines

expressed by macrophages were examined by quantitative reverse transcription polymerase chain reaction (qRT-PCR). Total mRNA of the cells treated with rh-CCL2 or rh-CX3CL1 was extracted with the RNeasy mini Kit according to manufacturer's instructions. The resultant RNA preparations were further treated with ribonuclease-free deoxyribonuclease (DNase) I (Life Technologies, Gaithersburg, MD, USA) to remove genomic DNA. Next, cDNAs from the RNA preparations were generated by reverse transcription. *VEGF-A*, *THBS-1* and *ADAMTS-1* were amplified with SYBR Premix Ex Taq on a CFX96 Touch™ Real-Time PCR Detection System (Bio-Rad Laboratories, USA). mRNA expression levels were measured according to the Ct value produced by each gene. Relative expression levels normalized to housekeeping gene β -actin were evaluated using the $2^{-\Delta\Delta Ct}$ method, where Ct represents the threshold cycle, and β -actin was used as a reference gene. Primers used are listed in Table 1.

Proliferation Assay In order to confirm whether the cytokines released by macrophages regulate the proliferation of HREC, we carried out a proliferation assay. Cultured macrophages were treated with rh-CCL2 or rh-CX3CL1 for 24h. Then the medium was replaced with fresh medium for another 24h. Supernatants were then collected. Totally 5×10^3 HREC in 100 μ L DMEM were seeded in 96-well plates and divided into control, rh-CCL2 and rh-CX3CL1 groups. Each group was replicated in five wells. Six hours later, when the HREC were attached to the plate, the medium was removed and an equivalent volume of the collected supernatants from macrophages culture medium were added to appropriate groups. After 12 to 48h stimulation of HREC with the supernatant, the absorbance was measured at 6, 12, 24 and 48h. The inhibition rate (IR) for the proliferation of cells in different groups was compared to control groups. The data was analyzed using GraphPad Prism 6.

Migration Assay To evaluate the effect of intervened macrophages on the migration of HREC, a modified Boyden chamber assay was performed as described previously^[20]. Briefly, 2.5×10^4 macrophages in 500 μ L DMEM were seeded in the lower chambers and were treated with rh-CCL2 or rh-CX3CL1. After 24h incubation, the medium was replaced and fresh DMEM was added for another 24h. The 1×10^4 HREC in 100 μ L DMEM were seeded in upper chambers. After co-culture for 24h, migrated HREC were fixed prior to staining with 0.5% crystal violet solution. Non-invading cells were removed by a cotton swab. The infiltrated cells were counted under phase contrast microscopy.

Statistical Analysis The data in this study were assessed using the Shapiro-Wilk test according to the normal distribution. Variables were expressed as mean \pm SD. The equality of variances was assessed by Levene's test. Various expression levels in each group were determined by one-way ANOVA,

Table 1 The sequence of primers used in quantitative RT-PCR

Genes	Sequence	Production length (bp)
<i>VEGF-A</i>	F: AATGCAGACCAAGAAAGATAGAGC	149
	R: GAGGCTCCAGGGCATTAGAC	
<i>THBS-1</i>	F: AGACTCCGCATCGCAAAGG	157
	R: TCACCACGTTGTTGTCAAGGG	
<i>ADAMTS-1</i>	F: TTCCACGGCAGTGGTCTAAAG	100
	R: CCACCAGGCTAACTGAATTACG	
β -actin	F: GCCGTCTCCCTCCATCGTG	102
	R: TCTCTTGCTCTGGGCCTCGTC	

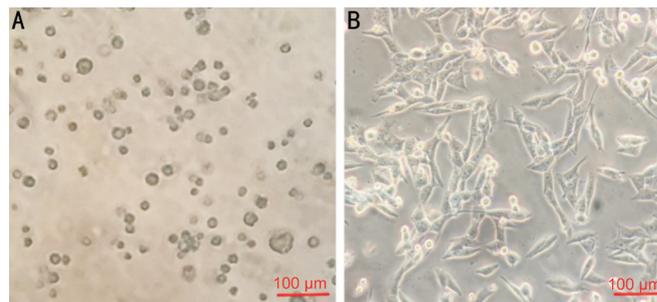


Figure 1 Macrophages differentiated from PBMC Cells were isolated from human peripheral blood of healthy individuals by density gradient centrifugation. Cells were cultured with RPMI-1640 supplemented with 10% FBS, 30 μ g/L of GM-CSF was added to induce the cells different to macrophages. A: The vitality of cells declined gradually over 2wk, and numerous cells were dead; B: When PBMC treated with GM-CSF, the vitality of cells was enhanced markedly.

and for multiple comparisons Dunnett's test was used for intergroup comparisons. $P < 0.05$ were considered statistically significant. Statistical analyses were performed with GraphPad Prism 6.0 (GraphPad Software, Inc., USA).

RESULTS

Induction of Macrophages by GM-CSF The cells isolated from peripheral blood completely adhered to the plates 2h after plating. The shapes of these cells were ovoid, pyramidal, or fusiform. Primary cultured cells survived for approximately 2wk without stimulation by GM-CSF. Nevertheless, we detected numerous dead cells floating in the medium. The viability of cells was enhanced markedly after addition of GM-CSF. Cells grew stably with GM-CSF stimulation and the mortality was markedly reduced (Figure 1). Induced macrophages were passaged every 3d and the cells were used for assays within five passages.

CCR2 and CX3CR1 Expression on Macrophages To examine the induction rate of macrophages from PBMC, cells were stained with CD68 (a macrophage marker) and were detected using flow cytometry. Cells treated with GM-CSF expressed CD68 (91% \pm 4.4%), indicating that the cells were macrophages. To further verify whether CCR2 and CX3CR1 receptors for CCL2 and CX3CL1 were expressed

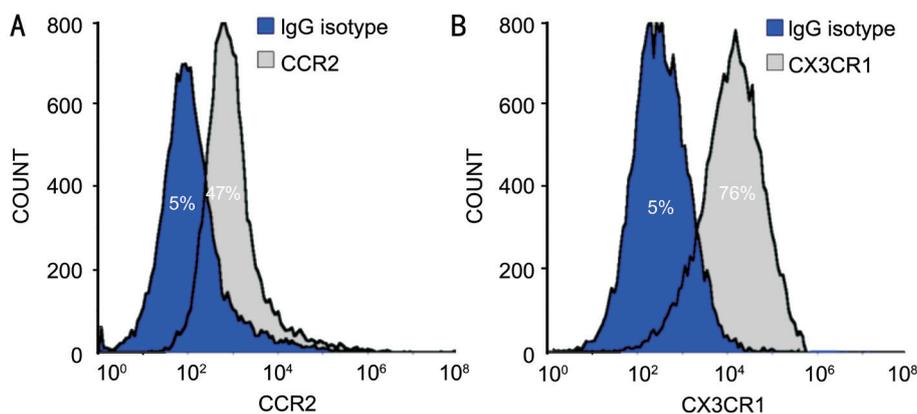


Figure 2 CCR2 and CX3CR1 expression on macrophages Cultured cells were collected following centrifugation and resuspension to $1 \times 10^6/\text{mL}$. Each group was stained with IgG isotype and rabbit anti-human CCR2 antibody (1:100) or anti-CX3CR1 antibody (1:100). A: Positive expression of CCR2 on GM-CSF treated cells was $42\% \pm 1.9\%$; B: Positive expression of CX3CR1 on GM-CSF treated cells was $71\% \pm 3.3\%$. Positive expression represent the value of experimental group minus the value of the control group.

on macrophages respectively, anti-CCR2 and anti-CX3CR1 antibodies were employed. Both CCR2 and CX3CR1 were co-expressed with CD68 on the membrane of cultured cells. CX3CR1 expression was higher than CCR2 expression on macrophages. The mean expression of CCR2 was $42\% \pm 1.9\%$ while the mean expression of CX3CR1 was $71\% \pm 3.3\%$ (Figure 2). These results suggest that PBMC could be induced into macrophages by GM-CSF. Additionally, both the relative cognate receptors for CCL2 and CX3CL1 were expressed on macrophages. Expression of these receptors is crucial for macrophages to engage in further biological functions by binding to relevant chemokines.

Angiogenesis-related Cytokine Expression in Various Macrophage Subtypes mRNA levels of *VEGF-A* were higher when stimulated with rh-CCL2, while mRNA levels of *THBS-1* and *ADAMTS-1* were significantly lower ($F=20.85$, $P=0.0001$; $F=9.35$, $P=0.0036$; $F=4.632$, $P=0.0323$ respectively). Following stimulation with 150 mg/L rh-CCL2, *VEGF-A* expression was higher than that of the control group ($t=5.809$, $P=0.0004$). By contrast, expression of *THBS-1* and *ADAMTS-1* was lower ($t=5.268$, $P=0.0008$; $t=2.522$, $P=0.0357$). Stimulation with rh-CX3CL1 markedly reduced *VEGF-A* mRNA expression, while it markedly increased mRNA expression of *THBS-1* and *ADAMTS-1* ($F=12.89$, $P=0.0010$; $F=4.379$, $P=0.0292$; $F=5.021$, $P=0.0260$ respectively). When stimulated with rh-CX3CL1, at a concentration of 150 mg/L, mRNA expression levels of *VEGF-A* were lower ($t=4.100$, $P=0.0034$) while expression of *THBS-1* and *ADAMTS-1* were higher ($t=2.720$, $P=0.0199$; $t=2.456$, $P=0.0396$; Figure 3).

Divergent Macrophage Populations Modulate HREC Proliferation in Opposite Directions In order to validate whether macrophages treated with various chemokines influence proliferation of HREC differently, we carried out a CCK-8 assay using supernatants pooled from cultured macrophages stimulated with rh-CCL2 or rh-CX3CL1. CCL2-

treated supernatants did indeed promote proliferation of HREC in a time-dependent manner ($IR=5.99\% \pm 1.42\%$, $P<0.05$). On the contrary, compared to control groups, proliferation was markedly inhibited when HREC were incubated with the supernatant of the CX3CL1-treated group for 48h ($IR=-6.93\% \pm 1.74\%$, $P<0.05$; Figure 4). This data suggests that both CCL2 and CX3CL1 regulate the expression of cytokines by macrophages. However, the effects of CCL2 and CX3CL1 on macrophages are opposite, depending on their cellular cytokine profile.

Divergent Macrophage Populations Modulate the Migration of HREC The migration of endothelium is an initial step for angiogenesis. We performed a migration assay to characterize macrophages stimulated by various cytokines. When co-cultured with CX3CL1-treated macrophages, the infiltration number of HREC crossing through the upper chamber decreased. Positive-stained migrated cell numbers in the CX3CL1 group were 119 ± 63 and 236 ± 34 in control group ($P<0.05$). The number of migrated HREC increased (410 ± 100) when incubated with CCL2-treated macrophages compared to control group ($P<0.05$; Figure 5). This suggests that different macrophage subtypes, induced by different chemokines, have the ability to mediate HREC behavior of migration by secreting different cytokines.

DISCUSSION

Chemokines and their cognate receptors are involved in a number of physiological and pathological processes via their chemotactic properties: embryonic development, immunity, wound healing, inflammation, tumor growth and angiogenesis^[21-23]. In the process of neovascularization, CCL2 directly promotes vascular endothelial migration. CCL2 has the ability to induce macrophages to release angiogenic factors such as VEGF-A, IFN- γ and IL-6, resulting in enhanced angiogenesis^[24-26]. At early stages of neovascularization, with stimulation by CCL2, macrophages

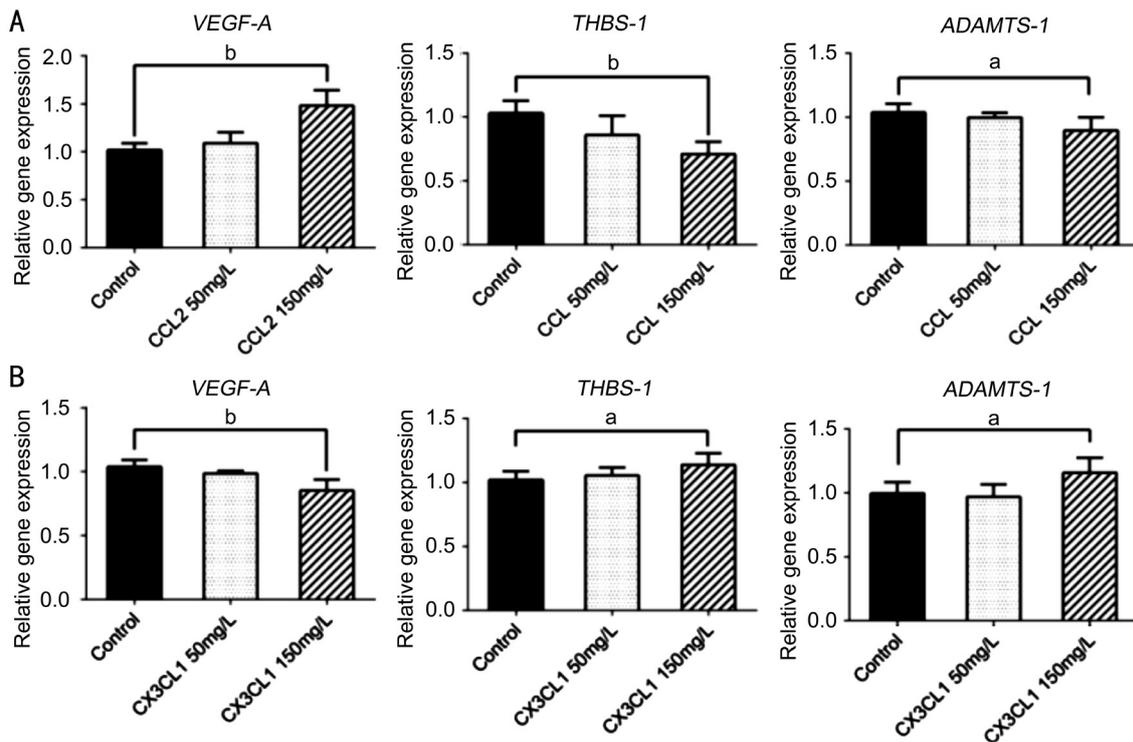


Figure 3 Angiogenesis-related cytokines expression in different subtype macrophages A: *VEGF-A* mRNA level was upregulated, while *THBS-1* and *ADAMTS-1* levels were downregulated when 150 mg/L rh-CCL2 added to macrophages; B: *VEGF-A* mRNA level was downregulated, while *THBS-1* and *ADAMTS-1* levels were upregulated when 150 mg/L rh-CX3CL1 added to macrophages. Levels of mRNA expression were analyzed by the Ct value produced by each gene and the relative expression level compare to housekeeping gene was calculated by standard $2^{-\Delta\Delta Ct}$ method. Data presented as mean±SD, ^a*P*<0.05, ^b*P*<0.01.

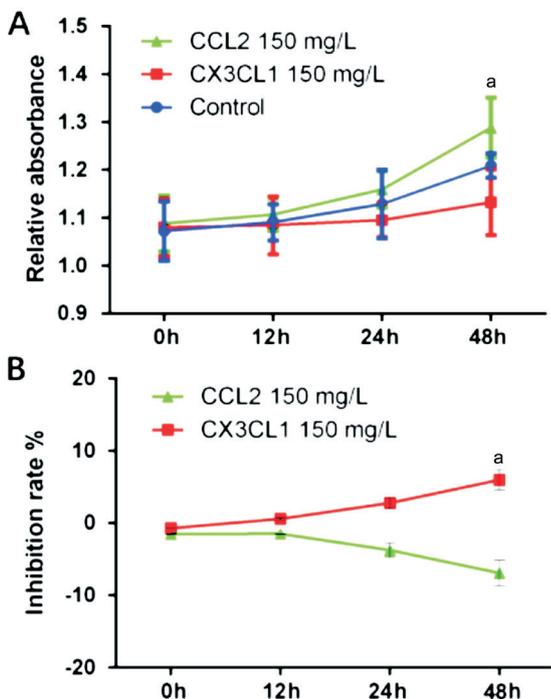


Figure 4 The macrophages treated by CCL2 or CX3CL1 modulate the proliferation of HREC The macrophages treated with rh-CCL2 or rh-CX3CL1 for 24h, the medium was replaced and cultured for another 24h, the supernatant of each group was collected then added to cultured HREC. The proliferation of HREC was detected using CCK-8 assay. A: OD values of different groups; B: The inhibitory rate of CCL2- and CX3CL1-treated group. Data presented as mean±SD, ^a*P*<0.05.

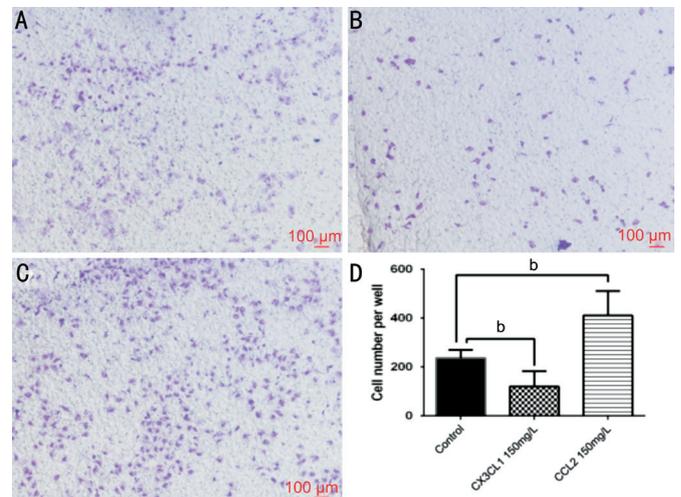


Figure 5 Cytokines released by CX3CL1- or CCL2-treated macrophages has different effects on the migration of HREC HREC 1×10^4 were seeded in the upper chamber, the lower chamber were treated macrophages seeded at 2.5×10^4 per well. A: The migrated HREC of control group; B: The migrated HREC when co-cultured with rh-CX3CL1 treated macrophages; C: The migrated HREC when co-cultured with rh-CCL2-treated macrophages; D: The statically result of each group. Data presented as mean±SD, ^b*P*<0.01.

secrete a variety of pro-angiogenic cytokines. In the case of ocular neovascularization, there is evidence that macrophages are involved in modulating vascular formation. Wallace *et al*^[27] and Tsutsumi-Miyahara *et al*^[28] reported that macrophages are

implicated in the development of choroidal neovascularization. With the systemic deletion of macrophages, choroidal neovascularization was suppressed in an experimental choroidal neovascularization model^[29].

It is well established that macrophages participate in various pathological processes, including myocardial ischemia, tumor angiogenesis, as well as ocular neovascularization^[30]. Macrophages are characterized by their heterogenic phenotypes. Particular micro-environments induce macrophages differentiation into distinctive subtypes. In addition to pro-angiogenic macrophages, there are anti-angiogenic macrophages that release angiostatic cytokines to suppress angiogenesis. Hence, when recruited by different types of chemokines such as CCL2, CCL5 and CX3CL1, macrophages release a series of angiogenic or angiostatic factors such as *VEGF-A*, *THBS-1* and *ADAMTS-1* to tip the balance of cytokine expression to either pro-angiogenesis or anti-angiogenesis^[31-32]. It would contribute our understanding of the mechanisms of pathological neovascularization if we knew more about the pro-angiogenic or anti-angiogenic functions of infiltrated polarized macrophages.

In our previous study, we showed that high CCR2 and low CX3CR1 expressing macrophages promoted corneal neovascularization by secreting VEGF and bFGF, while high CX3CR1 and low CCR2 expressing suppressed corneal neovascularization by secreting *THBS-1* and *ADAMTS-1*. This suggested that CCL2/CCR2 and CX3CL1/CX3CR1 signaling are involved in corneal neovascularization, mediated by macrophages^[33]. There is substantial evidence that the character of functions exhibited by macrophages depends on their microenvironment that polarizes macrophages into specific subtypes^[9,34-35]. In present study, we explored the effects of CCL2 and CX3CL1 on macrophage secreting functions by stimulating them with the functional recombinant proteins of CCL2 and CX3CL1, respectively. We found that mRNA expression of VEGF-A was increased when stimulated with rh-CCL2. In contrast, expression of *THBS-1* and *ADAMTS-1* decreased. VEGF-A is a pivotal angiogenic factor characterized by its strong pro-angiogenic function while *THBS-1* and *ADAMTS-1* are angiostatic factors that suppress neovascularization. The effects of CCL2 upregulating *VEGF-A* expression and simultaneously downregulating expression of *THBS-1* and *ADAMTS-1* drive the balance toward a pro-angiogenic direction. This demonstrates that CCL2 has an angiogenic effect by interacting with macrophages.

The expression of *VEGF-A* was reduced while expression of *THBS-1* and *ADAMTS-1* was increased in macrophages when the cells stimulated with rh-CX3CL1. This suggests that macrophages interacting with CX3CL1 secrete angiostatic cytokines and may have anti-angiogenic properties. The finding agrees with a previous study^[36]. Zheng *et al*^[37] also reported

that the highly-expressed CX3CR1 subtype of macrophages has an inhibitory effect on pathological angiogenesis.

To further investigate whether macrophages treated by CCL2 and CX3CL1 influence HREC function, we performed proliferation and migration assays. We showed that CCL2-stimulated macrophages promote proliferation and migration of HREC. In contrast, the potency of proliferation and migration was suppressed when co-cultured with CX3CL1-stimulated macrophages. This suggests that macrophages stimulated with various chemokines secrete a distinct profile of cytokines, giving rise to opposite effects on angiogenesis. These results are also consistent with reports that define macrophages as M1 and M2 subtypes^[38]. Further investigation is needed to determine whether CCL2 and CX3CL1 induce macrophages to polarize into M1 or M2 subtypes. Our data suggest that CCL2 and CX3CL1 have opposite effects on angiogenesis.

In conclusion, we found both CCL2 and CX3CL1 alter the secretion profile of cytokines in macrophages toward pro-angiogenic or anti-angiogenic directions, and that these effects influence biological functions in HREC. These data may provide new insights into the mechanisms of recruited macrophages on angiogenesis, and may provide potential therapeutic targets for ocular neovascularization.

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