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CX3CR1 Reduces Kidney Fibrosis by Inhibiting Local Proliferation of Profibrotic Macrophages

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A dense network of macrophages and dendritic cells (DC) expressing the chemokine receptor CX3CR1 populates most tissues. We recently reported that CX3CR1 regulates the abundance of CD11c+ DC in the kidney and thereby promotes renal inflammation in glomerulonephritis. Given that chronic inflammation usually causes fibrosis, we hypothesized that CX3CR1 deficiency should attenuate renal fibrosis. However, when we tested this hypothesis using the DC-independent murine fibrosis model of unilateral ureteral obstruction, kidney fibrosis was unexpectedly more severe, despite less intrarenal inflammation. Two-photon imaging and flow cytometry revealed in kidneys of CX3CR1-deficient mice more motile Ly6C/Gr-1+ macrophages. Flow cytometry verified that renal macrophages were more abundant in the absence of CX3CR1 and produced more of the key profibrotic mediator, TGF-β. Macrophages accumulated because of higher intrarenal proliferation, despite reduced monocyte recruitment and higher signs of apoptosis within the kidney. These findings support the theory that tissue macrophage numbers are regulated through local proliferation and identify CX3CR1 as a regulator of such proliferation. Thus, CX3CR1 inhibition should be avoided in DC-independent inflammatory diseases because it may promote fibrosis. The Journal of Immunology, 2015, 194: 1628–1638.
in ischemia/reperfusion injury (44, 49), hypertensive kidney damage (54), and crescentic glomerulonephritis (3, 57). In contrast, the loss of renal phagocytes aggravated renal candidiasis (58) but not pyelonephritis (3). Also, studies on human biopsy material from fibrotic kidneys described upregulation of CX3CR1, supporting a pathogenic role (59). However, in toxin- and obstruction-induced liver fibrosis, CX3CR1 was protective (50).

The role of CX3CR1 in the obstruction-induced standard kidney fibrosis model, UUO, has not been clarified yet. In this study, we have addressed this open question, assuming that the marked loss of renal DCs in CX3CR1 deficiency would attenuate inflammation and fibrosis. However, we found that CX3CR1 was protective in UUO because of a novel inhibitory effect on the other CX3CR1-expressing mononuclear phagocyte type of the kidney, the macrophages.

Materials and Methods

Mice

Eight- to 14-wk-old Cd11cEYFP (60), Cx3cr1GFP/+ and Cx3cr1GFP/GFP mice (61), all on a C57BL/6 background, were bred and kept under specific-pathogen-free condition at the animal facilities of the University of Bonn and Monash Medical Centre. Institutional and Government Review Boards have approved all animal studies.

Induction of UUO

Adult male mice were anesthetized by inhalation of isofluorane. UUO was performed through a midline incision. The left ureter was tied with silk suture at two points and permanently ligated. Sham-operated mice underwent an identical midline incision but without ureteric ligation. At 3, 5, and 7 d postobstruction, mice were sacrificed, and kidneys were collected.

Renal multiphoton microscopy

Intravital multiphoton microscopy of the intact kidney was performed on mice 3 d post-UUO, anesthetized by i.p. injection of 150 mg/kg ketamine hydrochloride and 10 mg/kg xylazine, as described previously (18, 62). The jugular vein was cannulated, and the left kidney was exteriorized through a lateral incision, immersed in bicarbonate buffered saline at 37˚C, and covered with a coverslip. Mice were kept at 37˚C. Time lapse recordings of GFP+ cells in the kidney of Cx3cr1GFP/+ and Cx3cr1GFP/GFP mice were captured using a Leica SP5 multiphoton microscope system (Leica Microsystems, Mannheim, Germany) with a ×20 1.0 NA water immersion objective (18, 63). Fluorophores were excited by a SpectraPhysics MaiTai pulsed infrared laser tuned to 860 nm, with emissions captured by non-descanned detectors with 525/50 nm (GFP), 585/40 nm (tetramethylrhodamine isothiocyanate), and 650/50 nm (Alexa 633) filters.

CellTracker Orange CMRA (C34551; Molecular Probes) was used as a counterstain to provide visual context. Images were collected with ∼120 μm Z-depth in 6-μm steps, with one Z-stack every 30s for 20–30 min and analyzed using Imaris (version 7.0; Bitplane AG). To determine GFP+ cell motility, individual cells were tracked in vivo and a motile cell defined as one that had moved at least the diameter of the cell body. To determine cell activity, a morphology index was used as previously described (18), relating to the sphericity of cells described by the rate of change in morphology index (ΔMI). Cells with relatively low ΔMI values change morphology less than cells with higher ΔMI values.

Histology and immunohistochemistry

Methyl Carnoy’s fixed paraffin-embedded murine kidneys were sectioned at 1 μm and stained with periodic acid–Schiff’s (PAS) for evaluation of hydropnephrosis and overall injury. For the evaluation of fibrosis, Sirius Red stained sections as well as α-smooth muscle actin (αSMA) immunohistochemistry were evaluated using computer-based morphometric analyses by a pathologist in blinded fashion as described previously (64). In short, for both stainings, the percentage of positively stained area was calculated from 20 cortical fields per section, representing nearly the whole cortex. Indirect immunoperoxidase procedure was performed as described previously (64). HRP coupled to human αSMA (clone 1A4; DakoCytomation, Glostrup, Denmark), which cross-react with murine αSMA was used. Fluorescence microscopy was performed on 5-μm paraffin sections from 4% PFA-fixed kidneys. GFP-expressing cells in Cx3cr1GFP/+ mice were identified by nuclei staining and quantified using the IX71 microscope and the CellR software (Olympus).

Isolation of murine kidney leukocytes

Kidneys were digested with collagenase and DNase-I (both from Sigma-Aldrich) as described previously (7). Briefly, kidneys were digested for 45 min, and tubular fragments from digested kidneys were removed by filtration through a 100-μm nylon mesh.

FIGURE 1. Lack of CX3CR1 increases renal fibrosis in UUO. (A–D) Fibrosis was quantified by Sirius red staining (A and B) and αSMA immunostaining (C and D) of sections of ligated or contralateral kidneys from CX3CR1-competent (Cx3cr1+/+, Cx3cr1GFP/+ [left images or white and light gray bars]) and CX3CR1-deficient (Cx3cr1GFP/GFP [right images or dark gray bars]) mice 7 d after ureter ligation. (A and C) Scale bars, 50 μm. Data are mean ± SEM from three (A and B) and two (C and D) experiments in groups of four to seven mice. *p < 0.05.
Flow cytometry

After treatment with Fc-blocking Abs (2.4G2) cells were stained with fluorochrome-labeled mAbs against F4/80 (CI:A3-1), CD45 (30-F11), CD11c (N418), KI67 (SolA15; isotype control: eBR2a), Gr1 (RB6-8C5), and Annexin V (number 550474; BD Biosciences). For intracellular cytokine staining, single-cell suspensions were incubated in RPMI 1640 medium (10% FCS, penicillin/streptomycin, and glutamine) with 1 μl/ml GolgiPlug and 1 μl/ml GolgiStop (BD Biosciences) for 4 h at 37˚C. After cell surface staining, cells were fixed in Cytofix (BD Biosciences) and permeabilized with PermWash (BD Biosciences). Intracellular staining was performed for 30 min at room temperature using fluorochrome-labeled mAbs against TGFβ (TW7-16B4; isotype control: MOPC-21). For analysis of proliferation, BrdU was administered 3 and 4 d after ligation, and incorporation of BrdU was analyzed on day 5 after ligation by flow cytometry (number 557892; BD Biosciences). Cells were analyzed with a BD Canto II and a BD Fortessa (BD Biosciences), and data were analyzed using Flowjo (Tree Star). For concatenated data presentation, data from all mice within the same group was combined using Flowjo.

Transfer of bone marrow monocytes

Single-cell suspensions from the bone marrows were prepared by flushing the bones with sterile and cold PBS. Bone marrow monocytes were enriched by excluding CD19+CD3+CD11c+ cells. Bone marrow cells (1–5×10⁶) were transferred into mice.

Detection of cytokines

The protein levels of cytokines in the supernatant of mechanically homogenized kidneys were determined by Luminex assay. The growth factors M-CSF and GM-CSF were determined by using ELISA (R&D Systems). All other levels were determined by a flow cytometry–based bead assay (eBioscience).

RNA isolation and RT-PCR

Whole-tissue RNA from kidneys was extracted using NucleoSpin RNA II kit (Macherey-Nagel), according to the manufacturer’s instructions. RNA was reverse transcribed into cDNA using the High-Capacity Reverse Transcription Kit (Applied Biosystems). Quantitative PCR was performed for 40 cycles using SYBR Green (Applied Biosystems). Quant iTect Primer assays (Qiagen) were used to detect mouse gene expression of Stat1 (QTO0159915), Irf5 (QTO0252623), Retina (QTO0254359), and Mrc1 (QTO013012). All samples were run in duplicates and normalized to Hprt primers (Invitrogen), forward primer (5’-GTCCCAGCGTCGTGATTAGC-3’) and reverse primer (5’-GGGCCACAATGTGATGGCCTCC-3’).

Statistical analysis

Results are expressed as mean ± SEM. Comparisons were drawn using parametric T-Test and nonparametric Mann–Whitney test (Prism 5; Graphpad Software, San Diego, CA). Significant p values were expressed as follows: *p < 0.05; **p < 0.01; ***p < 0.001.

Results

CX3CR1 deficiency enhances renal fibrosis

The chemokine receptor CX3CR1 has been shown to promote renal inflammation in several models (3, 49, 54, 57–59, 65). Given that chronic kidney inflammation often leads to fibrosis, we hy...
pothesized that CX3CR1 deficiency should attenuate renal fibrosis. To test this hypothesis, we induced one of the standard kidney fibrosis models, UUO in mice expressing or lacking CX3CR1. We used Cx3cr1GFP/GFP mice that homozygously express a GFP reporter under the control of the CX3CR1 promoter and are consequently CX3CR1 deficient. As controls, we used Cx3cr1GFP/+ mice, which express the GFP reporter heterozygously and CX3CR1-competent wild-type mice (Cx3cr1+/+ mice). Surprisingly, kidney fibrosis in CX3CR1-deficient mice was notably aggravated compared with control mice 7 d after UUO, as evidenced by higher Sirius Red collagen staining in the ligated kidneys of Cx3cr1GFP/GFP mice (Fig. 1A, 1B). Also, αSMA staining for profibrotic myofibroblasts was stronger in ligated kidneys of CX3CR1-deficient mice (Fig. 1C, 1D). These findings revealed an unexpected antifibrotic role of CX3CR1 in renal fibrosis.

CX3CR1 alters the abundance, morphology, and migration of renal mononuclear phagocytes

To define the mechanisms underpinning the protective role of CX3CR1, we focused on the cells that express CX3CR1 in the kidney, namely mononuclear phagocytes, including DCs and macrophages. As reported before (3), CX3CR1-deficient (Cx3cr1GFP/GFP) mice normally possess far less GFP+ renal phagocytes than CX3CR1-competent (Cx3cr1GFP/+ ) mice (Fig. 2A, 2B). Also, on day 7, after ureter ligation, fewer phagocytes were seen, albeit only 10% less (Fig. 2A, 2B), suggesting the appearance of new phagocytes that partially compensated for their deficiency under homeostatic conditions. Of note, the message for the Th1 mediator Stat1 was partially compensated for their deficiency under homeostatic conditions. Of note, the message for the Th1 mediator Stat1 was partially compensated for their deficiency under homeostatic conditions. Of note, the message for the Th1 mediator Stat1 was partially compensated for their deficiency under homeostatic conditions. Of note, the message for the Th1 mediator Stat1 was partially compensated for their deficiency under homeostatic conditions. Of note, the message for the Th1 mediator Stat1 was partially compensated for their deficiency under homeostatic conditions.

These findings led us to examine in more detail the renal GFP+ phagocytes in UUO using multiphoton microscopy, which allows analyzing their morphology and motility over time. In both healthy CX3CR1-competent and CX3CR1-deficient mice, GFP+ cells with dendritic protrusions were present within the interstitium (see Supplemental Videos 1, 2).

After UUO, GFP+ cells tended to form clusters around damaged tubules (see Supplemental Videos 3, 4). In addition to GFP+ cells with dendritic morphology, the kidneys of CX3CR1-deficient mice with UUO contained larger cells (Fig. 3A) devoid of dendrites (Fig. 3B), which were more motile (Fig. 3C). Furthermore, the phagocytes in CX3CR1-deficient mice contained a subset that underwent little shape change over the time of analysis (Fig. 3D). Although these data did not reach statistical significance, they prompted us to hypothesize that GFP+ phagocytes with monocyte/macrophage properties might be involved in the increased fibrosis in CX3CR1-deficient mice with UUO.

CX3CR1 differentially regulates the numbers of renal DCs and Gr1+ macrophages in UUO

We hypothesized that higher numbers of renal macrophages might explain the profibrotic effect of CX3CR1 deficiency in UUO (Fig. 1) because it has been previously reported that renal macrophages but not DCs mediate fibrosis in this model (16, 18, 66). However, Cx3cr1GFP/GFP mice possessed similar or lower numbers of GFP+ renal phagocytes under homeostatic conditions, in experimental glomerulonephritis and in kidney infection compared with CX3CR1-competent controls (3, 58), consistent with our present findings in UUO (Fig. 2A, 2B). We therefore discriminated between GFP+ kidney macrophages and DCs by costaining with F4/80 and CD11c. We found that CX3CR1 deficiency reduced renal CD11c+ DC numbers in the ligated and contralateral kidney (Fig. 4A, upper gate in the dot plots) consistent with the morphological changes shown in Fig. 3. Next, we focused on CD11c+ GFP+ cells (Fig. 4A, lower gate in the dot plots). We found that >90% of these cells expressed F4/80 (Fig. 4B) identifying these GFP+/CX3CR1-expressing cells as macrophages. Furthermore, these data also exclude considerable contaminations of other cells than monocytes/macrophages and DCs in the microscopy data (Figs. 2, 3). Because the microscopy findings above (Fig. 3) hinted at the appearance of monocytes/macrophages, we focused on the analysis of macrophages expressing the Gr1 marker, which designates mononuclear phagocytes recently recruited from the circulation (67, 68) and which is related to helper macrophages expressing Ly6C (4), a component of Gr1. Gr1+ macrophages were considerably more numerous already on day 3 after UUO (Fig. 4B, 4C). In the absence of CX3CR1, these cells were more abundant than in CX3CR1-competent mice, whereas the abundance of kidney-resident Gr1− macrophages was unchanged (Fig. 4C). On day 7, fewer recruited macrophages were seen, whereas numbers of Gr1− macrophages had further increased (Fig. 4C). Thus, the lack of CX3CR1 reduced the number of CD11c+ DCs, but increased the numbers of CD11c− Gr1+ macrophages in UUO. This shift from DCs to macrophages explains why the total number of GFP+ phagocytes was only somewhat reduced in CX3CR1-deficient mice (Fig. 2), and the relative increase of macrophages might explain the increased kidney fibrosis in the mice.

CX3CR1 reduces TGF-β production by macrophages

Next, we examined whether CX3CR1 also modulates the functionality of macrophages. To this end, we analyzed the production of TGF-β, the key profibrotic molecule in the kidney (65, 69). Most of the TGF-β–producing leukocytes indeed expressed F4/80 (Supplemental Fig. 1). Importantly, the frequencies (Supplemental Fig. 1) and numbers (Fig. 4D) of TGF-β–producing Gr1+ and Gr1− macrophages were increased in ligated kidneys of CX3CR1-deficient mice. M1/M2 differentiation markers such as Arg1, Mrc1, Irf5, Stat6, and Retnla1 were little altered in CX3CR1

FIGURE 3. In vivo multiphoton microscopy of Cx3cr1GFP/+ and Cx3cr1GFP/GFP mice. (A) Analysis shows the proportion of cells with a diameter larger than 8 μm (Cx3cr1GFP/+ [open squares]; Cx3cr1GFP/GFP [gray circles]). (B and C) In the absence of CX3CR1, there is a trend (p = 0.09) toward a higher proportion of GFP+ cells without protrusions (B) and higher number of motile cells (C) (Cx3cr1GFP/+ [open squares]; Cx3cr1GFP/GFP [gray circles]). (D) Change in morphology index (ΔMI). Analysis of cellular shape change over time shows that CX3CR1-deficient mice tend to have more cells that change shape < 30 min. The x-axis in (D) represents groups of cells that are changing their shape more over time (Cx3cr1GFP/+ [open squares]; Cx3cr1GFP/GFP [gray circles]). Videos can be seen in the Supplemental Videos 1–4; data (A–D) represent the mean ± SEM of five to six mice per group.
deficiency (Fig. 5), indicating that the functional changes of macrophages could not be explained by effects on M1/M2 differentiation (70, 71). These data demonstrate that CX3CR1 not only negatively regulates the numbers of profibrotic macrophages but also their fibrogenic potential.

CX3CR1 regulates neither recruitment nor differentiation of kidney macrophages

The inhibitory effects of CX3CR1 on renal macrophages could theoretically be due to reduced recruitment of macrophage precursors such as monocytes, reduced intrarenal differentiation into macrophages, reduced proliferation, increased survival or be due to a combination of these factors. Therefore, we examined all of these potential mechanisms.

To establish a system to analyze macrophage precursor recruitment to the kidney, we first adoptively transferred bone marrow–derived monocytes expressing a YFP reporter for CD11c, which allows convenient discrimination between DCs and macrophages, into CD45.1-congenic mice with or without UUO. Only a few monocytes reached the kidneys of healthy mice and the contralateral kidney of mice with UUO (Fig. 6A), suggesting that renal phagocytes possess a low turnover rate. In contrast, ligated kidneys contained clearly detectable monocytes 2 d after transfer, and some of them were YFP+ (Fig. 6A), indicating differentiation into DCs.

To investigate whether CX3CR1 is required for the entry of these precursors into the kidney, we transferred monocytes from CX3CR1-deficient mice into CX3CR1-competent wild-type recipients 1 d after ureter ligation (Fig. 6B, 6C). After an additional 2 d, less CX3CR1-deficient phagocytes had entered the ligated kidneys than did CX3CR1-competent phagocytes (Fig. 6C). Notably, no difference was seen in the blood (Supplemental Fig. 2), excluding an effect of CX3CR1 on the abundance of circulating macrophage precursors. Thus, CX3CR1 promoted, rather than inhibited, phagocyte precursor recruitment. Furthermore, the levels of the chemokine MIP-1α that can attract macrophage precursors were decreased in ligated and also in contralateral kidneys of CX3CR1-deficient mice, and there were trends toward reduced levels of RANTES, MCP-1, and MCP-3 (Fig. 6D). This is consistent with reduced macrophage precursor recruitment and may be explained by the reduction in chemokine-producing DCs in CX3CR1-deficient mice (3).
The chemokine receptor CX3CR1 is widely expressed by DCs and macrophages (74), but its functional role is still not fully clear. In this paper, we report a previously undescribed antifibrotic effect of CX3CR1: it reduced the inflammation-induced proliferation of TGF-β-producing renal macrophages, which are critically important in experimental kidney fibrosis (15–18). This finding was unexpected, given the proinflammatory and profibrotic properties of CX3CR1 not only in diseases of the kidney (3, 49, 54, 57–59, 65), but also of the lung (75), liver (50), intestine (28), skin (55), arteriosclerotic vessels (76), and the CNS (33). CX3CR1 has been shown to contribute to the inflammatory recruitment of monocytes from the circulation into all these organs (44). We found that CX3CR1 had a similar effect in UUO so that fewer monocytes lacking this receptor entered the inflamed kidneys. The lower levels of monocyte-attracting chemokines and of inflammatory mediators that we detected in kidneys of CX3CR1-deficient mice may be responsible for the reduced kidney recruitment of these cells. It is likely that these cells originated from monocytes, which have been shown to give rise to tissue macrophages under inflammatory conditions (5). The reduction of monocyte-attracting chemokines likely resulted from the paucity of DCs in kidneys of CX3CR1-deficient animals because DCs are the principal producers of chemokines and inflammatory mediators in the kidney (77–79). However, these findings were hard to reconcile with the accumulation of macrophages in CX3CR1-deficient mice, implying that other mechanisms must exist that overcompensated for the reduced attraction of macrophage precursors.

One such mechanism might be a negative effect of CX3CR1 on the survival of tissue macrophages. However, previous studies in disease models of liver, CNS, gut, and arteriosclerotic vessels (74) demonstrated that CX3CR1 signaling increased rather than reduced the life span of macrophages. In contrast to these findings, one recent study reported increased macrophage numbers and liver fibrosis in CX3CR1-deficient mice after surgical bile duct ligation, a model that resembles UUO by being induced by mechanical tissue damage (50). The discrepancy between macrophage accumulation and decreased macrophage life span in that study was suggested to have resulted from the release of proinflammatory mediators from CX3CR1-deficient dying macrophages that attracted further profibrotic macrophages. Although we did observe more dying macrophages after UUO in CX3CR1-deficient mice, we found neither increased monocyte-attracting chemokines and inflammatory cytokines, nor did we find a stronger influx of inflammatory monocytes, arguing against inflammation resulting from increased macrophage death.
from premature macrophage death in the kidney as the underlying reason for macrophage accumulation.

Another hypothetic mechanism is that CX3CR1 might cause immigrating monocytes to differentiate preferentially into macrophages rather than DCs. DC numbers are severely reduced in CX3CR1-deficient mice (3), which would be consistent with altered differentiation. DCs do not play a major role in UUO, whereas macrophages are critical (18, 49, 54). This is not unexpected given that DCs primarily regulate adaptive immune responses, explaining why they affected fibrosis in models that involve adaptive immune cells, such as ischemia/reperfusion injury, crescentic glomerulonephritis and hypertensive nephropathy (3, 49, 54, 57). By contrast, a model of mechanic tissue injury like UUO is more likely mediated by cells of the innate immune system. The frequency of donor YFP*CD45.2*F4/80*CD11c* DCs (upper region) and of YFP*CD45.2*F4/80*CD11c* macrophages (lower region) in the kidney was determined 3 d after ureter ligation (bone marrow cells were transferred 1 d after ligation). (B) Sorted CD19*CD3*CD11c* bone marrow cells from CX3CR1GFP/+ mice were transferred into wild-type mice 1 d after ligation. Three days after ureter ligation GFP*F4/80*CD11c* DCs (upper region) and GFP*F4/80*CD11c* macrophages (lower region) were detected by flow cytometry. (C) Sorted CD19*CD3*CD11c* bone marrow cells from CX3CR1GFP/+ or CX3CR1GFP/GFP mice were transferred into wild-type mice. Three days after ureter ligation, the number of F4/80*GFP*CD11c* macrophages (left panel) and of GFP*F4/80*CD11c* DCs (right panel) were determined by flow cytometry. The group designated without transfer was used to control for specific detection of transferred cells. (D) Kidneys from CX3CR1GFP/+ (white squares) or CX3CR1GFP/GFP (gray circles) mice were homogenized and levels of the chemokines MCP-1 (top left), MCP-3 (top right), MIP1α (bottom left), and RANTES (bottom right) were determined 3 d after ureter ligation. Control animals did not undergo unilateral ureteral ligation. Data represent three (A-C) and two (D) experiments in groups of five to six mice. Results in (D) are mean ± SEM. *p < 0.05, **p < 0.01.
system, like macrophages. However, adoptive transfer experiments revealed that monocyes expressing or lacking Cx3CR1 gave rise to similar ratios between macrophages and DCs, arguing against an effect of Cx3CR1-signaling on the monocyes’ decision to differentiate into either DCs or macrophages. This conclusion was corroborated by unchanged levels of the growth factors that usually govern this decision, M-CSF and GM-CSF.

In vivo multiphoton microscopic analysis revealed morphologic alterations of Cx3CR1-deficient renal phagocytes. Although there were slight changes in the morphology of Cx3CR1+ kidney DCs, such as more yet shorter dendrites (our unpublished observations), it is unlikely that these alterations impacted fibrogenesis given such as more yet shorter dendrites (our unpublished observations), it is unlikely that these alterations impacted fibrogenesis given that both Gr1– and Gr1 + macrophages produced TGF-

In summary, we report a novel inhibitory effect of CX 3CR1 on the monocytes’ decision to differentiate into either DCs or macrophages. This conclusion was corroborated by unchanged levels of the growth factors that usually govern this decision, M-CSF and GM-CSF.

Traditionally, tissue macrophages were thought to arise from circulating monocytes recruited in response to locally produced chemokines (11). Recent findings have revealed that some tissue macrophages are derived from embryonic precursors (22). A sudden demand because of inflammatory conditions is met by the circulating monocytes recruited in response to locally produced chemokines (11). Recent findings have revealed that some tissue macrophages are derived from embryonic precursors (22). A sudden demand because of inflammatory conditions is met by the circulating monocytes, including those that had just recently been recruited to the kidney, showed stronger proliferation in the absence of Cx3CR1. These findings are consistent with the concept that local proliferation of tissue macrophages regulates their numbers and extend this concept by showing that also recently recruited macrophages are locally regulated in the inflamed kidney. It remains to be seen whether this mechanism also contributes to fibrosis in other organs.

CX3CR1 not only affected the abundance, but also the functional state of renal macrophages. In its absence, more macrophages produced TGF-β, the key profibrotic mediator in the kidney (69). A previous study showed that CCR2-dependent macrophages produced TGF-β in UUO and that the lack of these cells significantly reduced renal fibrosis (16). Also, we found that both Gr1– and Gr1 + macrophages produced TGF-β and that production was increased in the absence of Cx3CR1. TGFβ production suggested acquisition of a profibrotic functional state change. This state could not be explained by M2 differentiation, nor was it associated with higher renal inflammation. Recently, the classical M1/M2 paradigm was refined by the discovery of an activation-independent functional state that does not fall into the M1 or M2 definition (71). Our findings are consistent with this notion by demonstrating that TGF-β–producing macrophages cannot be categorized as classical M1 and M2 macrophages. Future studies are required to clarify whether these profibrotic macrophages belong to a distinct functional state.

In summary, we report a novel inhibitory effect of Cx3CR1 on local macrophage proliferation and on fibrogenesis in the kidney.
in the present study, so that CX3CR1 inhibition in such diseases may turn out profibrotic.

In cregenic GN, the previously described proinflammatory influence of CX3CR1 on DCs prevailed (3) because macrophages are downstream of DCs in the pathomechanism of this disease and because DCs outnumber macrophages in the kidney. However, in DC-independent renal diseases like UUO, the inhibitory effect of CX3CR1 on macrophages becomes manifest, as we demonstrated in the present study, so that CX3CR1 inhibition in such diseases aggravates fibrosis. This may apply also to organs in which CX3CR1 does not regulate DCs, like the liver or the lung (3). In this case, therapeutic CX3CR1 inhibition in liver and lung disease may turn out profibrotic.

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Disclosures
The authors have no financial conflicts of interest.

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