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# IL-6 Controls the Innate Immune Response against *Listeria monocytogenes* via Classical IL-6 Signaling

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The cytokine IL-6 plays a protective role in immune responses against bacterial infections. However, the mechanisms of IL-6-mediated protection are only partially understood. IL-6 can signal via the IL-6R complex composed of membrane-bound IL-6R $\alpha$  (mIL-6R $\alpha$ ) and gp130. Owing to the restricted expression of mIL-6R $\alpha$ , classical IL-6 signaling occurs only in a limited number of cells such as hepatocytes and certain leukocyte subsets. IL-6 also interacts with soluble IL-6R $\alpha$  proteins and these IL-6/soluble IL-6R $\alpha$  complexes can subsequently bind to membrane-bound gp130 proteins and induce signaling. Because gp130 is ubiquitously expressed, this IL-6 trans-signaling substantially increases the spectrum of cells responding to IL-6. In this study, we analyze the role of classical IL-6 signaling and IL-6 trans-signaling in the innate immune response of mice against *Listeria monocytogenes* infection. We demonstrate that *L. monocytogenes* infection causes profound systemic IL-6 production and rapid loss of IL-6R $\alpha$  surface expression on neutrophils, inflammatory monocytes, and different lymphocyte subsets. IL-6-deficient mice or mice treated with neutralizing anti-IL-6 mAb displayed impaired control of *L. monocytogenes* infection accompanied by alterations in the expression of inflammatory cytokines and chemokines, as well as in the recruitment of inflammatory cells. In contrast, restricted blockade of IL-6 trans-signaling by application or transgenic expression of a soluble gp130 protein did not restrain the control of infection. In summary, our results demonstrate that IL-6R $\alpha$  surface expression is highly dynamic during the innate response against *L. monocytogenes* and that the protective IL-6 function is dependent on classical IL-6 signaling via mIL-6R $\alpha$ . *The Journal of Immunology*, 2013, 190: 703–711.

The cytokine IL-6 is a central regulator of inflammatory processes. In response to infection but also during chronic inflammation, IL-6 is produced by variety of cells and has pro- as well as anti-inflammatory activities (1, 2). IL-6 is a major inducer of acute phase proteins and is involved in the control of neutrophil and monocyte responses following infection. IL-6 is also a survival factor for lymphocytes and it promotes Ab production by B cells (1–3). More recently, IL-6 has been identified as a decisive cytokine for the differentiation of CD4<sup>+</sup> T cells into Th17 cells, which are considered to be a major proinflammatory T cell population (4, 5). As a consequence, blockade of IL-6 or IL-6 signaling

has a profound impact on innate and acquired immune responses. Impaired IL-6 function causes enhanced susceptibility of mice to infection with various pathogens (6–15) but also results in protection in several mouse models for chronic inflammatory diseases (16–18). Not surprisingly, IL-6 signaling is also an important target of intervention in chronic inflammatory human diseases such as rheumatoid arthritis and Crohn's disease (19).

The IL-6 receptor consists of membrane-bound IL-6R $\alpha$  (mIL-6R $\alpha$ , CD126) and the gp130 chain (CD130). mIL-6R $\alpha$  binds IL-6 but has no intrinsic signaling capacity. Upon IL-6 binding, mIL-6R $\alpha$  interacts with gp130, which subsequently initiates IL-6 signal transduction (2). In contrast to the ubiquitous expression of gp130, expression of mIL-6R $\alpha$  is restricted to hepatocytes and certain subsets of leukocytes (20). Thus, IL-6 can only activate a limited number of cells via classical IL-6R signaling. However, a soluble form of IL-6R $\alpha$  (sIL-6R $\alpha$ ) is constitutively found in the serum, and following inflammation mIL-6R $\alpha$  is actively shed from inflammatory cells, resulting in enhanced local and systemic sIL-6R $\alpha$  concentrations (21, 22). sIL-6R $\alpha$  still binds IL-6, and the IL-6/sIL-6R $\alpha$  complexes can interact with membrane-bound gp130 and thereby provoke signal transduction (23). This process, which is called IL-6 trans-signaling, allows activation of mIL-6R $\alpha$ -negative cells by IL-6 and thereby substantially expands the number of IL-6-responsive cells (2, 3). Under normal conditions, IL-6 levels in human blood are in the range of 1–5 pg/ml. sIL-6R $\alpha$  and soluble gp130 (sgp130) levels, however, are in the range of 50–100 and 100–200 ng/ml, respectively. Secreted IL-6 therefore will bind to sIL-6R $\alpha$  and the complex of IL-6/sIL-6R $\alpha$  will bind to sgp130 and thereby be neutralized. Hence, sIL-6R $\alpha$  and sgp130 constitute a buffer for systemically active IL-6. Only when IL-6 concentrations exceed those of sIL-6R $\alpha$  and sgp130 can IL-6 act systemically (24, 25).

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Abbreviations used in this article: HKL, heat-killed listeria; KO, knockout; mIL-6R $\alpha$ , membrane-bound IL-6R $\alpha$ ; SAA, serum amyloid A; sgp130, soluble gp130; sgp130Fc, soluble gp130 Fc fusion protein; sIL-6R $\alpha$ , soluble IL-6R $\alpha$ ; Tg, transgenic; wt, wild-type.

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IL-6/sIL-6R $\alpha$  complexes also offer a target for therapeutic intervention. Soluble gp130 proteins have been designed that neutralize IL-6/sIL-6R $\alpha$  complexes but do not interfere with signaling via mIL-6R $\alpha$  (24, 25). These proteins have been used to dissect the role of classical IL-6 signaling and IL-6 trans-signaling in different mouse models for infection and inflammatory diseases (22, 26, 27). Because sgp130 proteins show beneficial effects in models of chronic inflammation (16, 28, 29), they might also offer a new option for the treatment of chronic inflammatory diseases in humans (30).

The Gram-positive bacterium *Listeria monocytogenes* can cause severe disease in immune-suppressed individuals, and in pregnant women, infection of the fetus can lead to abortion or to high fatality rates in neonates (31). In mice, infection with *L. monocytogenes* provokes a rapid activation of the innate immune system, which is essential for the restriction of bacteria replication. In particular, the production of proinflammatory cytokines such as TNF- $\alpha$  and IFN- $\gamma$  and the recruitment of inflammatory monocytes to sites of infection are crucial for the early control of *L. monocytogenes* (32, 33). Infection also leads to robust production of IL-6, and IL-6 deficiency or IL-6 neutralization with Abs results in enhanced susceptibility to *L. monocytogenes* (6, 34–41); however, mechanisms of IL-6-mediated protection are so far unclear.

In this study, we characterize the role of different IL-6 signaling pathways for the innate control of *L. monocytogenes*. We demonstrate that following infection, mIL-6R $\alpha$  is rapidly lost from the surface of all leukocytes analyzed. In parallel, high serum levels of IL-6 can be detected, providing the basis for classical IL-6 signaling and IL-6 trans-signaling. Using neutralizing anti-IL-6 mAbs, which block IL-6 classical and trans-signaling, and a sgp130 Fc fusion protein (sgp130Fc), which only blocks IL-6 trans-signaling, as well as IL-6-deficient mice and mice transgenic (Tg) for sgp130Fc, we can show that classical IL-6 signaling but not IL-6 trans-signaling is crucial for the IL-6-mediated control of *L. monocytogenes*.

## Materials and Methods

### Abs and reagents

Anti-CD16/CD32 mAb (clone 2.4G2) and anti-IL-6 mAb (MP5-20F3) were purified from hybridoma supernatants according to standard protocols. Fluorochrome-conjugated mAbs against CD4 (RM4-5), CD8 $\alpha$  (53-6.7), CD11b (M1/70), CD19 (eBioID3), CD45 (30-F11), Ly6C (AL-21), Gr-1 (RB6-8C5), F4/80 (BM8), TNF- $\alpha$  (MP6-XT22), IL-6 (MP5-20F3), and IL-6R $\alpha$  (D7715A7) were purchased from BioLegend (San Diego, CA), eBioscience (San Diego, CA), or BD Pharmingen (San Diego, CA).

A soluble fusion protein of the extracellular domains of gp130 and the Fc part of human IgG1 (sgp130Fc) was constructed and purified as described in Jostock et al. (25). As demonstrated before, human sgp130Fc can bind and neutralize mouse IL-6/IL-6R $\alpha$  complexes with similar efficacy as mouse sgp130Fc (25). Purified human IgG was purchased from Dianova (Hamburg, Germany).

Ligand binding capacity and biological activity of sgp130Fc were measured by a Ba/F3-gp130 cell proliferation assay. Ba/F3 cells (murine pre-B cells) stably transfected with human gp130 (Ba/F3-gp130 cells) have been described previously (25). Ba/F3-gp130 cells were seeded with  $5 \times 10^3$  cells per well in 96-well plates. Proliferation was induced by adding 100 ng/ml IL-6 plus 50 ng/ml sIL-6R $\alpha$  and different concentrations of sgp130Fc. Proliferation was measured by the cell titer colorimetric assay (Promega) as described previously (42). The activity assay for the batch of sgp130Fc used for the present study is shown in Supplemental Fig. 1.

### Mice and animal experiments

C57BL/6 mice, IL-6 knockout (IL-6<sup>KO</sup>) mice (6), sgp130Fc<sup>Tg</sup> mice (24), and CCR2<sup>KO</sup> mice (43) were bred at the Central Animal Facilities of the University Medical Center Hamburg-Eppendorf, at the House for Experimental Therapy in Bonn, or at the Victor Hensen Facility of the University of Kiel. IL-6<sup>KO</sup> and sgp130<sup>Tg</sup> mice were on the C57BL/6N background. In experiments with these mouse strains, matched C57BL/

6N mice were used as controls. All other experiments were done with C57BL/6J mice. Animal experiments were conducted according to the German Animal Protection Law.

Mice were i.v. infected via the lateral tail vein with  $2 \times 10^4$  *L. monocytogenes* strain EGD in 200  $\mu$ l PBS. Bacterial inoculi were always controlled by plating serial dilutions on tryptic soy broth agar plates. For determination of bacterial burdens in spleens and livers, mice were killed, organs were homogenized in PBS containing 0.5% Triton X-100, serial dilutions of homogenates were plated on tryptic soy broth agar, and colonies were counted after 48 h incubation at 37°C.

For blocking of IL-6 signaling or trans-signaling, mice were treated with 500  $\mu$ g neutralizing anti-IL-6 mAb (clone MP5-20F3) (44) or 250  $\mu$ g sgp130Fc. Concentrations of anti-IL-6 mAb and sgp130Fc have been shown to be effective in blocking global IL-6 signaling and trans-signaling (26). Control mice received 250  $\mu$ g purified human IgG. Abs and fusion proteins were injected i.p. in a volume of 200  $\mu$ l PBS 1 d prior to *L. monocytogenes* infection. Abs and sgp130Fc proteins were controlled for low endotoxin content using a commercial *Limulus* amoebocyte lysate assay (Lonza, Basel, Switzerland).

### Purification and staining of cells

For characterization of different leukocyte subsets, spleens and livers were cut into small pieces and digested for 45 min with each 0.25 mg/ml collagenase D (Roche, Penzberg, Germany) and collagenase VIII (Sigma-Aldrich, Steinheim, Germany), and 15  $\mu$ g/ml DNase I (Sigma-Aldrich) in RPMI 1640 medium supplemented with glutamine, gentamicin, 2-ME, and 10% heat-inactivated FCS (complete RPMI 1640 medium). The tissue was gently meshed through a 70- $\mu$ m cell strainer and the suspension was centrifuged to pellet the cells. Liver cells were resuspended and centrifuged through a 40/70% Percoll gradient (Biochrom, Berlin, Germany) for 30 min at  $600 \times g$ . Cells were collected from the interface of the gradient and washed in complete RPMI 1640 medium. Erythrocytes were lysed and cells were washed and stored on ice. After collagen digestion, spleen cells were pelleted and erythrocytes were lysed. Cells were washed and stored on ice. For analysis of surface IL-6R $\alpha$  and induction of IL-6, spleen cells were purified without collagen digestion.

For surface staining, cells were washed and incubated for 5 min with rat serum and anti-CD16/CD32 mAb to block unspecific Ab binding. Subsequently, cells were incubated on ice with Abs as indicated. To mark dead cells, 2  $\mu$ g/ml Pacific Orange succinimidyl ester (Invitrogen, Darmstadt, Germany) was added. After 20 min, cells were washed in PBS and fixed with PBS/1% paraformaldehyde. Cells were analyzed using a FACSCanto II and the FACSDiva software (Becton Dickinson, Mountain View, CA).

### Induction of cytokines and intracellular cytokine staining

Cells were cultured in a volume of 1 ml complete RPMI 1640 medium. Cells were stimulated for 6 h with  $10^7$  heat-killed listeria (HKL)/ml. During the final 4 h of culture, 10  $\mu$ g/ml brefeldin A (Sigma-Aldrich) was added. Cultured cells were washed and incubated for 5 min with rat serum and anti-CD16/CD32 mAb to block unspecific Ab binding. Subsequently, cells were stained with Abs against surface proteins, and after 30 min on ice, cells were washed with PBS and fixed for 20 min at room temperature with PBS/4% paraformaldehyde. Cells were washed with PBS/0.2% BSA, permeabilized with PBS/0.1% BSA/0.3% saponin, and incubated in this buffer with rat serum and anti-CD16/CD32 mAbs. After 5 min, fluorochrome-conjugated anti-IL-6 mAb and anti-TNF- $\alpha$  mAb was added. After a further 20 min on ice, cells were washed with PBS and fixed with PBS/1% paraformaldehyde.

For the preparation of HKL, bacteria from an overnight culture were washed twice in PBS and inactivated by incubation for 30 min at 80°C.

### IL-6 and sIL-6R $\alpha$ determination

Serum of naive and infected mice was appropriately diluted and analyzed for IL-6 and sIL-6R $\alpha$  by ELISA according to the manufacturer's protocols (R&D Systems, Wiesbaden, Germany). The addition of sIL-6R $\alpha$  did not interfere with the detection of IL-6 in this ELISA (Supplemental Fig. 2). Spleen and liver tissues of mice were mixed with lysis buffer containing tissue protein extraction reagent (Thermo Scientific, Dreieich, Germany) and protease inhibitors (complete protease inhibitor mixture tablets; Roche). Lysates were stored at  $-80^\circ\text{C}$ . After thawing, lysates were homogenized with a TissueLyser (Qiagen, Hilden, Germany). Lysates were centrifuged, protein content in the supernatant was determined with a BCA protein assay kit (Thermo Scientific), and the concentrations of IL-6 and serum amyloid A (SAA) were determined by ELISA. The murine SAA ELISA was conducted according to the manufacturer's protocol (Tridelta Development, Maynooth, Ireland).

## Real-time RT-PCR analysis

Total RNA of spleen and liver was prepared using TRIzol reagent (Invitrogen, Karlsruhe, Germany) and a TissueLyser (Qiagen) according to the manufacturers' protocols. cDNA was generated with random hexamer primers and Moloney murine leukemia virus reverse transcriptase (Invitrogen). Real-time PCR was performed for 40 cycles (initial denaturation at 95°C for 10 min, denaturation at 95°C for 20 s, primer annealing and elongation at 60°C for 1 min) with 1.5 µl cDNA samples in the presence of 2.5 µl (0.9 µM) specific murine primers and 6.25 µl SYBR Green JumpStart Taq ReadyMix (Sigma-Aldrich) in a StepOnePlus real-time PCR system (Applied Biosystems, Foster City, CA). All samples were run in duplicate and normalized to 18S rRNA to account for small RNA and cDNA variability (45).

The following primers were used: 18S, forward, 5'-CAG GGC CGG TAC AGT GAA AC-3', reverse, 5'-AGA GGA GCG AGC GAC CAA A-3'; CXCL1, forward, 5'-GCA CCC AAA CCG AAG TCA TAG-3', reverse, 5'-CAA GGG AGC TTC AGG GTC AA-3'; CXCL2, forward, 5'-CAC TGC GCC CAG ACA GAA-3', reverse, 5'-CAG GGT CAA GGC AAA CTT TTT G-3'; CCL2, forward, 5'-GGC TCA GCC AGA TGC AGT TAA-3', reverse, 5'-CCT ACT CAT TGG GAT CAT CTT GCT-3'; IL-6, forward, 5'-TGG GAA ATC GTG GAA ATG AGA-3', reverse, 5'-AAG TGC ATC ATC GTT GTT CAT ACA-3'; TNF-α, forward, 5'-AAA TGG CCT CCC TCT CAT CAG T-3', reverse, 5'-GCT TGT ACA AAT TTT GAG AAG-3'; SAA, forward 5'-AGA GGA CAT GAG GAC ACC ATT GCT-3', reverse, 5'-AGG ACG CTC AGT ATT TGT CAG GCA-3'.

## Statistical analysis

All mice per group were independently analyzed. Statistical analysis was performed with GraphPad Prism 4 (GraphPad Software, La Jolla, CA). Quantifiable data were transformed into logarithmic values. Afterward, a Student *t* test or one-way ANOVA and the Bonferroni multiple posttest was performed to define differences among differently treated mice. All results are expressed as arithmetic means ± SD (bar graphs) or median (dot plots). Differences were considered significant with *p* < 0.05.

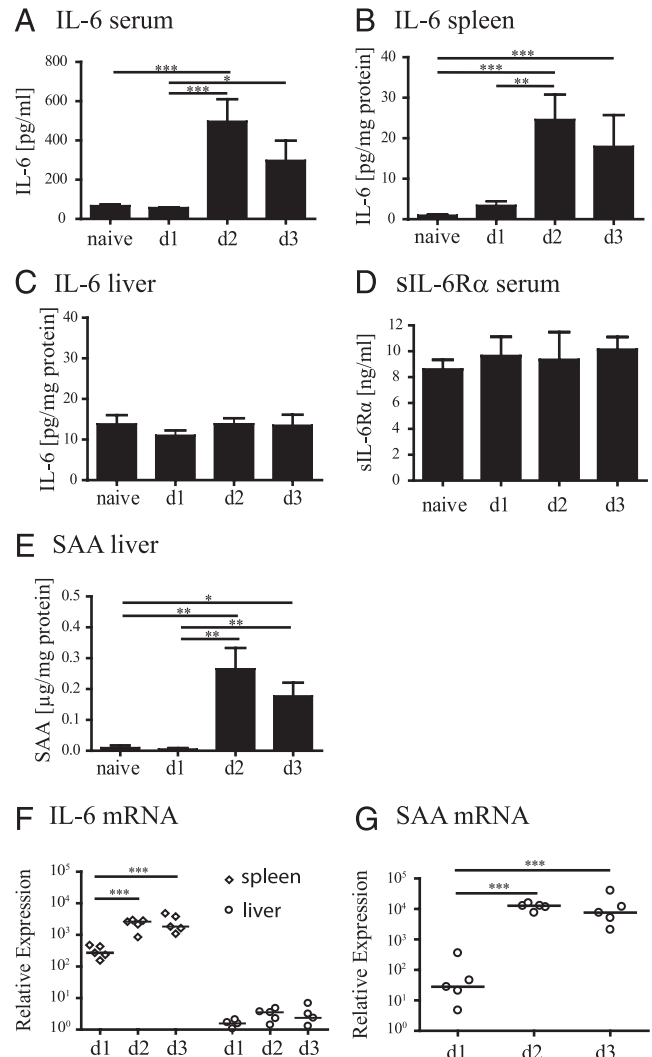
## Results

IL-6 and IL-6Rα expression during early *L. monocytogenes* infection

IL-6 signals via membrane-bound IL-6Rα and gp130 (classical signaling). Alternatively, IL-6 can form complexes with sIL-6Rα, which subsequently interact with gp130 on the cell surface and thereby allow activation of cells lacking surface expression of IL-6Rα. IL-6Rα can be actively shed from the surface of cells under inflammatory conditions. Thus, the mode of IL-6 signaling could switch from classical to trans-signaling following infection. In a first set of experiments, we therefore determined the concentrations of IL-6 and sIL-6Rα in serum and tissues as well as the surface expression level of IL-6Rα on different leukocyte populations following infection of mice with *L. monocytogenes*.

Infection of mice with *L. monocytogenes* caused a profound increase in serum levels of IL-6, which reached a maximum at day 2 postinfection (Fig. 1A). Enhanced IL-6 concentrations were also measured in tissue extracts from spleens but not from liver (Fig. 1B, 1C), although both organs are main target sites for listeria replication. A similar pattern was observed for IL-6 mRNA (Fig. 1F). IL-6 mRNA induction was largely confined to the spleen (100- to 1000-fold induction) and only marginally in the liver (<10-fold induction). Most likely as a consequence of the high serum concentration of IL-6, a strong induction of the mRNA for the IL-6 target gene *saa* followed by enhanced SAA protein concentration was observed in livers of infected mice (Fig. 1E, 1G). In contrast to the strong increase in IL-6, there was only a marginal change in the serum concentrations of sIL-6Rα during the first days of *L. monocytogenes* infection (Fig. 1D).

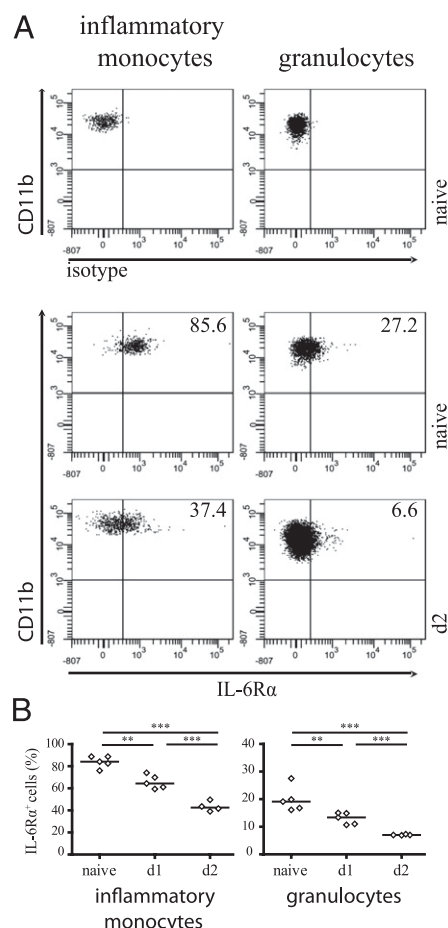
In a next set of experiments, we determined the expression of IL-6Rα on innate immune cells following *L. monocytogenes* infection. As shown before (46, 47), infection of mice with *L. monocytogenes* caused a rapid accumulation of neutrophils and



**FIGURE 1.** IL-6 and sIL-6Rα during the early phase of *L. monocytogenes* infection. C57BL/6 mice were i.v. infected with  $2 \times 10^4$  *L. monocytogenes*. On days 1, 2, and 3 postinfection, mice were killed and IL-6 concentrations were determined in serum (**A**) as well as spleen (**B**) and liver extracts (**C**). IL-6Rα concentrations were measured in the serum (**D**) and SAA in liver extracts (**E**). mRNA levels for IL-6 were determined in the spleen and liver (**F**) and for SAA in the liver (**G**) by quantitative RT-PCR. mRNA levels were normalized to 18S RNA and are given as fold change compared with the mean values of naive mice. Bars in (**A**)–(**E**) represent means ± SD from five to seven individually analyzed mice per time point. Graphs in (**F**) and (**G**) give values for individual mice and the median. Figures are representative of at least two independent experiments. \**p* < 0.05, \*\**p* < 0.01, \*\*\**p* < 0.001.

inflammatory monocytes in spleen and liver (Supplemental Figs. 3, 4). Flow cytometric analysis revealed IL-6Rα surface expression on inflammatory monocytes as well as on CD4<sup>+</sup> and CD8<sup>+</sup> T cells of naive mice. In contrast, neutrophils and B cells displayed only relatively low mIL-6Rα expression (Figs. 2, 3). Closer examination of CD4<sup>+</sup> and CD8<sup>+</sup> T cells showed that in particular naive CD4<sup>+</sup> T cells (CD44<sup>low</sup>CD62L<sup>+</sup>) were IL-6Rα<sup>+</sup>. Activated and memory CD4<sup>+</sup> T cells (CD44<sup>high</sup>CD62L<sup>+</sup> or CD44<sup>high</sup>CD62L<sup>+</sup>) contained both IL-6Rα<sup>+</sup> and IL-6Rα<sup>−</sup> populations. Following infection, we observed a significant loss of IL-6Rα surface expression on all cells analyzed (Figs. 2, 3). Loss of IL-6Rα expression was particular strong on CD4<sup>+</sup> T cells and included the naive CD44<sup>low</sup>CD62L<sup>+</sup> CD4<sup>+</sup> T cell population (Fig. 3). These results indicate that following *L. monocytogenes* infection, all analyzed cell populations





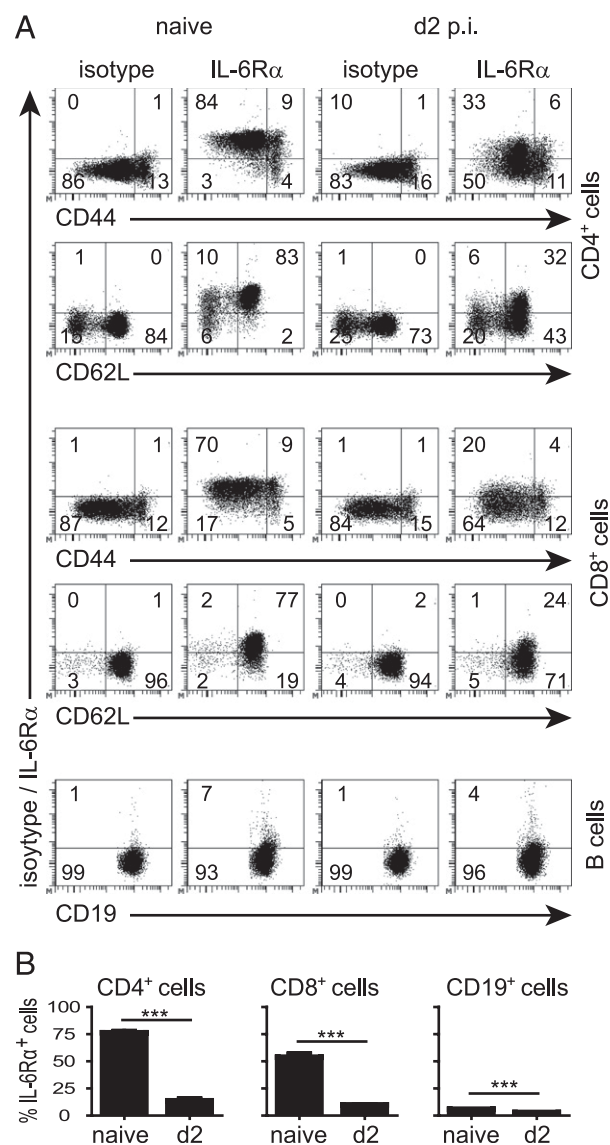
**FIGURE 2.** IL-6Rα on inflammatory monocytes and granulocytes during *L. monocytogenes* infection. C57BL/6 mice were i.v. infected with  $2 \times 10^4$  *L. monocytogenes*. On the indicated time points postinfection, spleen cells were analyzed for the expression of IL-6Rα. (A) Representative results for the surface expression of IL-6Rα on gated cell populations. Numbers give percentage values of mIL-6Rα<sup>+</sup> cells among the gated cell population. (B) Combined results from groups of individually analyzed mice. Results are representative of two independent experiments. Bars in (B) represent median values. \*\**p* < 0.01, \*\*\**p* < 0.001.

rapidly lose the ability to respond to IL-6 via classical IL-6 signaling and become dependent on IL-6 trans-signaling.

Loss of surface IL-6Rα expression could be due to proteolytic shedding (22). However, it is also possible that the receptor is internalized in response to IL-6 signaling. Finally, IL-6 could interfere with the binding of the anti-IL-6Rα mAb to mIL-6Rα. To distinguish between these processes, IL-6<sup>KO</sup> mice were infected with *L. monocytogenes* and the expression of mIL-6Rα was analyzed. On all leukocyte subsets including CD4<sup>+</sup> T cells, CD8<sup>+</sup> T cells, granulocytes, and inflammatory monocytes, we observed a similar loss of mIL-6Rα expression in wild-type (wt) and IL-6<sup>KO</sup> mice (Fig. 4 and data shown). Therefore, loss of mIL-6Rα expression is most likely due to shedding and is not a consequence of receptor internalization or epitope masking.

#### *Inflammatory monocytes are not an essential source of IL-6 and sIL-6Rα*

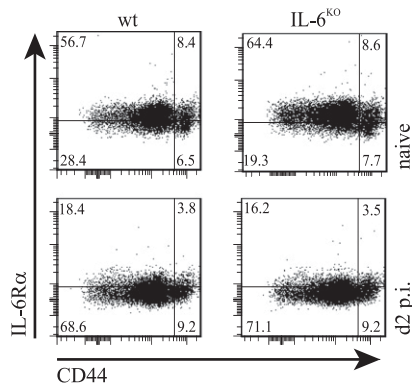
Different cell types have been identified as sources of IL-6 during inflammation and infection (1). Because we observed a strong accumulation of granulocytes and inflammatory monocytes, we tested whether these cells were potential sources of IL-6 during early *L. monocytogenes* infection. Upon stimulation of spleen cells with HKL, a large fraction of inflammatory monocytes were in-



**FIGURE 3.** IL-6Rα on T and B lymphocytes during *L. monocytogenes* infection. C57BL/6 mice were i.v. infected with  $2 \times 10^4$  *L. monocytogenes*. On day 2 postinfection, spleen cells were analyzed for the expression of IL-6Rα. (A) Graphs show representative results for surface expression profiles for IL-6Rα or isotype control and CD44 or CD62L for CD4<sup>+</sup> and CD8<sup>+</sup> T cells, and for IL-6Rα or isotype control on CD19<sup>+</sup> B cells. (B) Combined results from groups of individually analyzed mice. Bars represent means  $\pm$  SD for three or four individually analyzed mice per time point and are representative of three independent experiments. \*\*\**p* < 0.001.

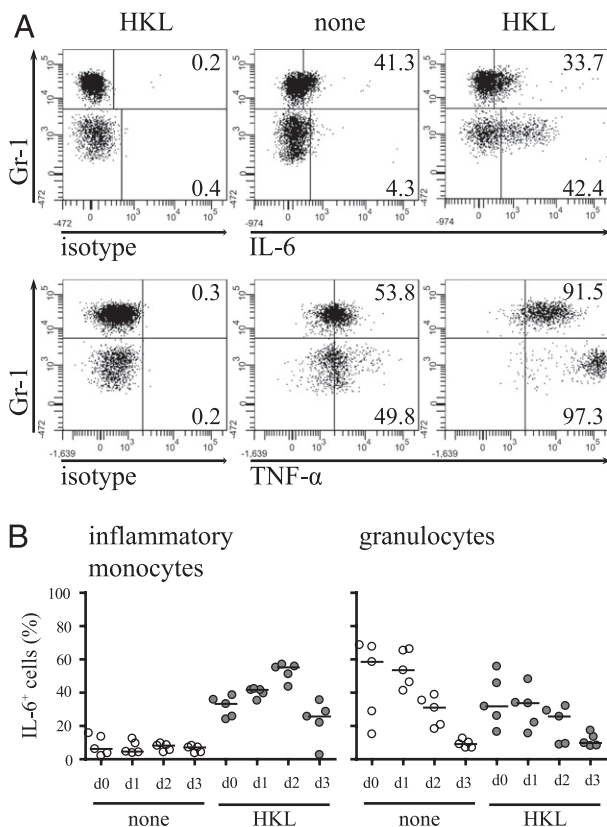
deed able to produce IL-6 as determined by intracellular cytokine staining (Fig. 5). Frequencies of inflammatory monocytes producing IL-6 in response to HKL increased until day 2 of infection and then declined again. In these cells, IL-6 production was accompanied by a strong expression of TNF-α. Without stimulation, neutrophils showed already low expression levels of IL-6, but expression did not increase after addition of HKL. Frequencies of granulocytes producing IL-6 declined postinfection.

Because inflammatory monocytes expressed IL-6Rα on their surface and were able to secrete IL-6 upon stimulation with listeria, we determined the impact of this cell population on serum levels of IL-6 and IL-6Rα during the early phase of *L. monocytogenes* infection. Serbina and Pamer (48) recently demonstrated that following *L. monocytogenes* infection, mobilization of



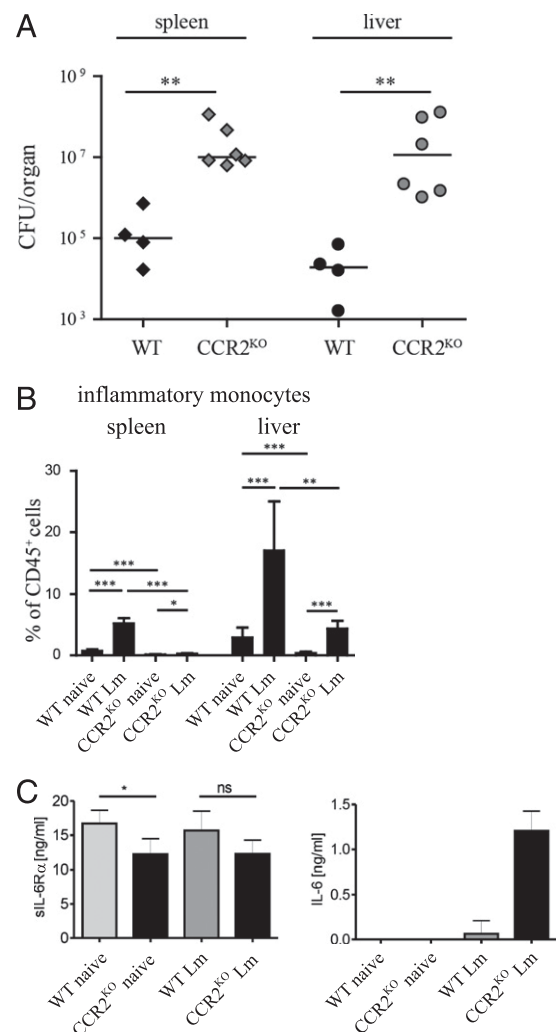
**FIGURE 4.** Loss of IL-6Rα surface expression occurs in the absence of IL-6. C57BL/6 mice and IL-6<sup>KO</sup> mice were i.v. infected with  $2 \times 10^4$  *L. monocytogenes*. On day 2 postinfection, spleen cells were analyzed for the expression of IL-6Rα. Graphs show representative results for surface expression profiles of IL-6Rα and CD44 on CD4<sup>+</sup> T cells. Results are representative of two independent experiments with three mice per group.

inflammatory monocytes from the bone marrow and accumulation of these cells in infected tissues were strictly dependent on the chemokine receptor CCR2. Therefore, wt and CCR2<sup>KO</sup> mice were infected with *L. monocytogenes* and analyzed 3 d postinfection. Determination of *L. monocytogenes* burden in spleen and livers revealed 100-fold higher titers in CCR2<sup>KO</sup> mice (Fig. 6A).



**FIGURE 5.** Induction of IL-6 in inflammatory monocytes and granulocytes. C57BL/6 mice were i.v. infected with  $2 \times 10^4$  *L. monocytogenes*. On the indicated time points postinfection, spleen cells were cultured for 6 h with or without HKL. Production of IL-6 was determined by intracellular IL-6 staining. (A) Representative results for gated inflammatory monocytes (Gr-1<sup>low</sup>) and granulocytes (Gr-1<sup>high</sup>). Numbers give the percentage values for cytokine-positive granulocytes and inflammatory monocytes, respectively. (B) Combined results from individually analyzed mice. Results are representative of two independent experiments. Bars represent median values.

Compared to infected wt mice, infected CCR2<sup>KO</sup> mice completely failed to recruit inflammatory monocytes to the spleen and had significantly reduced frequencies of these cells in the liver (Fig. 6B). In contrast, CCR2<sup>KO</sup> mice displayed high frequencies of granulocytes in spleen and liver following infection (data not shown). Determination of IL-6 revealed increased serum IL-6 levels in infected CCR2<sup>KO</sup> mice when compared with infected wt mice (Fig. 6C). Serum concentrations of sIL-6Rα were reduced in naive and infected CCR2<sup>KO</sup> mice when compared with wt controls (Fig. 6C), but reduction did not reach a significant level in infected animals (naive, wt  $16.73 \pm 0.86$  ng/ml versus CCR2<sup>KO</sup>  $12.21 \pm 1.13$  ng/ml,  $p = 0.018$ ; infected, wt  $15.68 \pm 1.28$  ng/ml versus CCR2<sup>KO</sup>  $12.25 \pm 1.57$  ng/ml,  $p = 0.054$ ). In summary, our results indicate that although inflammatory monocytes are fundamental in control of *L. monocytogenes* they are not an essential source for IL-6 and they only partially contribute to the generation of sIL-6Rα.

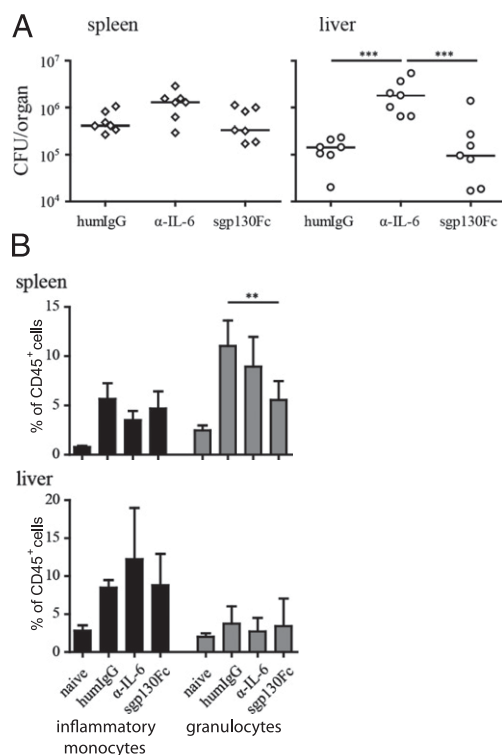


**FIGURE 6.** Inflammatory monocytes are not an essential source of IL-6 and sIL-6Rα. C57BL/6 wt and CCR2<sup>KO</sup> mice were i.v. infected with  $2 \times 10^4$  *L. monocytogenes*. On day 3 postinfection, spleen and liver were analyzed. (A) *Listeria* titers in spleen and liver of infected animal. (B) Accumulation of inflammatory monocytes in spleen and liver. (C) Serum concentration of sIL-6Rα and IL-6. Individual values and the median are shown in (A). Bars in (B) and (C) give the means  $\pm$  SD for four to five individually analyzed mice per group and time point. Figures are representative of two independent experiments. \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

### Blocking of conventional IL-6 signaling but not IL-6 trans-signaling caused impaired control of *L. monocytogenes* infection

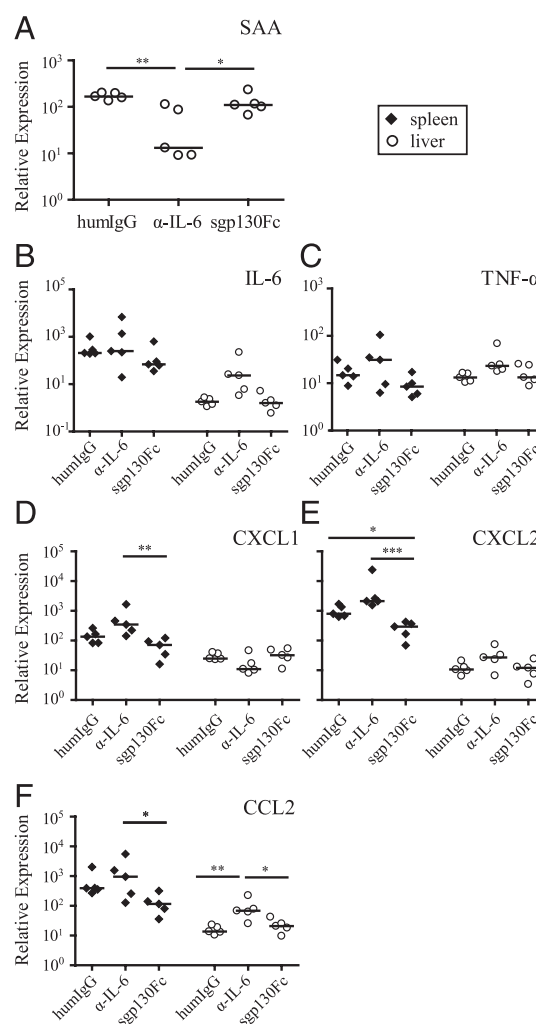
Our results demonstrated the presence of IL-6 and sIL-6R $\alpha$  during the early phase of *L. monocytogenes* infection. Thus, IL-6 could interact with target cells both via classical and trans-signaling. To determine the impact of classical and trans-signaling on the control of *L. monocytogenes*, mice were treated with a neutralizing anti-IL-6 mAb, which blocks all IL-6 signaling (44), or with a sgp130Fc fusion protein to selectively block IL-6 trans-signaling (26–29). Treatment of mice with a neutralizing anti-IL-6 mAb caused an increase in the listeria burden in the spleen and particularly in the liver of infected mice (Fig. 7A). In contrast, mice treated with sgp130Fc showed listeria titers in both organs, which were comparable to infected control animals. Thus, classical IL-6 signaling but not IL-6 trans-signaling was important for the control of *L. monocytogenes* infection.

Following infection, we observed pronounced accumulation of inflammatory monocytes in spleen and liver, which was not further affected by the treatment with anti-IL-6 mAb or sgp130Fc (Fig. 7B). At day 3 postinfection, there was still an enlarged neutrophil population in spleens of infected mice. Pretreatment of mice with sgp130Fc but not with anti-IL-6 mAb resulted in a reduction of this neutrophil population. In the liver, neither treatment with anti-IL-6 mAb or with sgp130Fc caused an alteration of neutrophil frequencies.



**FIGURE 7.** Role of classical IL-6 signaling and IL-6 trans-signaling in the innate immune response against *L. monocytogenes*. C57BL/6 mice were treated i.p. with anti-IL-6 mAb, sgp130Fc or control IgG (humlgG). One day later, mice were i.v. infected with  $2 \times 10^4$  *L. monocytogenes*. After a further 3 d, mice were analyzed. (A) Listeria burden in spleen and liver. (B) Accumulation of inflammatory monocytes and neutrophils in spleen and liver. Values of individual mice and the median are shown in (A). Bars in (B) give the means  $\pm$  SD of seven individually analyzed mice per group. Results in (A) are representative of four independent experiments and in (B) of two independent experiments. \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

To determine changes in the degree of inflammation following different treatments, the mRNA levels of SAA, IL-6, TNF- $\alpha$ , and of different chemokines were measured. *L. monocytogenes* infection caused a strong induction of SAA mRNA in livers of mice, which was reduced by anti-IL-6 mAb but not by sgp130Fc treatment (Fig. 8A). The mRNA for IL-6 was enhanced in spleens of infected mice but was not further affected by anti-IL-6 mAb or sgp130Fc treatment (Fig. 8B). There was limited induction of IL-6 mRNA in the livers of mice. Treatment with anti-IL-6 mAb caused enhanced IL-6 mRNA expression, but the difference did not reach a significant level. The mRNA for TNF- $\alpha$  was increased in spleens and livers of infected mice, but neither anti-IL-6 mAb nor sgp130Fc caused a change in expression (Fig. 8C). We detected enhanced levels of the neutrophil-attracting chemokines CXCL1 and CXCL2 and of the monocyte-attracting chemokine CCL2 (Fig. 8D–F). All three chemokines were more strongly induced in the spleen than in the liver. For all three chemokines, we observed



**FIGURE 8.** Role of classical IL-6 signaling and IL-6 trans-signaling for induction of cytokines and chemokines during the early phase of infection with *L. monocytogenes*. C57BL/6 mice were treated i.p. with anti-IL-6 mAb, sgp130Fc, or control IgG. One day later, mice were i.v. infected with  $2 \times 10^4$  *L. monocytogenes*. After a further 3 d, mRNA expression of SAA (A), IL-6 (B), TNF- $\alpha$  (C), and of the chemokines CXCL1 (D), CXCL2 (E), and CCL2 (F) was determined. mRNA levels were normalized to 18S RNA and are given as fold change compared with the mean values of naive mice. Graphs give the values for individual mice and the median. Results are representative of four independent experiments. \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

the strongest expression levels in spleens of anti-IL-6 mAb-treated mice, which were significantly higher than the expression levels in spleens of sgp130Fc-treated mice.

#### IL-6 deficiency but not Tg expression of sgp130Fc resulted in impaired control of *L. monocytogenes*

To confirm results from studies with anti-IL-6 mAb and sgp130Fc treatment, IL-6<sup>KO</sup> mice and sgp130Fc<sup>Tg</sup> mice were infected with *L. monocytogenes*. sgp130Fc<sup>Tg</sup> mice express a gp130Fc fusion protein under the control of the liver-specific phosphoenolpyruvate carboxykinase promoter, resulting in constitutively high serum levels of sgp130Fc protein and selective blockade of IL-6 trans-signaling (24). Deficiency in IL-6 caused enhanced listeria titers in spleen and liver (Fig. 9A). In contrast, Tg expression of sgp130Fc did not impair the control of *L. monocytogenes* infection. Listeria titers were even slightly lower than titers of infected wt mice. Similar to mice treated with sgp130Fc, sgp130Fc<sup>Tg</sup> mice demonstrated reduced accumulation of granulocytes in spleens of infected mice (Fig. 9B). However, sgp130Fc<sup>Tg</sup> mice in addition displayed diminished frequencies of inflammatory monocytes in spleen and liver.

## Discussion

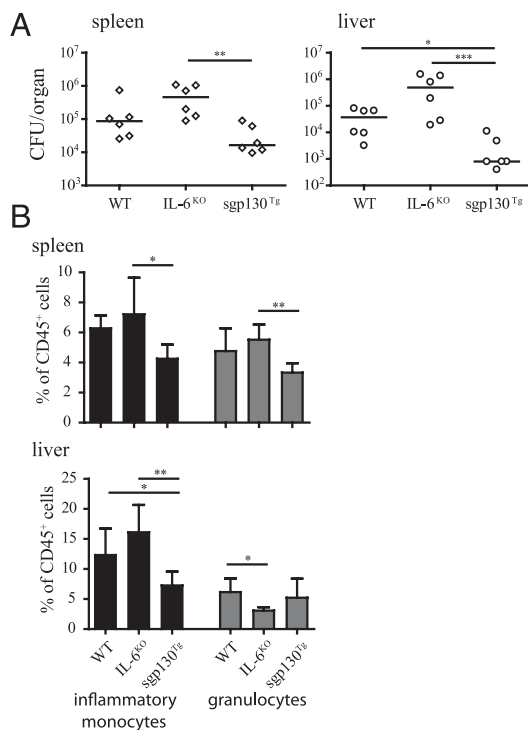
Following *L. monocytogenes* infection, we measured high serum levels of IL-6 at days 2 and 3 postinfection, which is in agreement with published results (39–41). Inflammatory monocytes are essential for the protective innate immune response against *L. monocytogenes* (33, 48), and we could demonstrate that these cells produced IL-6 as well as TNF- $\alpha$  following stimulation with HKL. However, despite their fundamental role in *L. monocytogenes*

control and their IL-6 response to listeria Ag, inflammatory monocytes were not an essential source for systemic IL-6. Infection of CCR2<sup>KO</sup> mice, which failed to mobilize inflammatory monocytes from the bone marrow (48), resulted in even enhanced IL-6 serum concentration when compared with infected control mice. Alternative IL-6 sources could include other myeloid cells such as tissue-resident macrophages, dendritic cells, and granulocytes or T and B cells, but also nonhematopoietic cells such as endothelial or epithelial cells (49). Owing to the more severe infection, some of these cells might respond even stronger in CCR2<sup>KO</sup> mice.

In naive mice, we detected mIL-6R $\alpha$  expression on T cells and inflammatory monocytes. In particular, CD4<sup>+</sup> T cells expressed high levels of mIL-6R $\alpha$ . Granulocytes expressed only low mIL-6R $\alpha$  levels and B cells were largely negative. Naive T cells (CD62L<sup>high</sup>CD44<sup>low</sup>) were uniformly mIL-6R $\alpha$ <sup>+</sup>, and effector as well as memory T cells (CD62L<sup>low</sup>CD44<sup>high</sup> or CD62L<sup>high</sup>CD44<sup>high</sup>) contained mIL-6R $\alpha$ <sup>+</sup> and mIL-6R $\alpha$ <sup>−</sup> subpopulations. Following *L. monocytogenes* infection, all analyzed leukocyte subsets lost mIL-6R $\alpha$  expression. On T cells, reduction of mIL-6R $\alpha$  was observed on naive as well as preactivated subpopulations. A similar reduction of mIL-6R $\alpha$  expression was also observed in infected IL-6<sup>KO</sup> mice, which excludes IL-6-induced receptor internalization or interference of IL-6 with anti-IL-6R $\alpha$  mAb binding to IL-6R $\alpha$  as a cause for low IL-6R $\alpha$  surface staining (50). Shedding of mIL-6R $\alpha$  by ADAM10 and ADAM17 has been identified as a general mechanism of sIL-6R $\alpha$  generation in mice (51), and it is very likely that shedding is responsible for the loss of IL-6R $\alpha$  surface expression following *L. monocytogenes* infection. TCR triggering has been described as a signal for IL-6R $\alpha$  shedding (52, 53); however, the global loss of IL-6R $\alpha$  on naive and effector cells argues against a specific TCR signal at this early stage of infection. It is more likely that loss of IL-6R $\alpha$  expression is due to shedding induced by inflammatory mediators such as proinflammatory cytokines, chemokines, or acute phase proteins (52, 53). Our present work aims at defining these mediators in our infection model.

Despite the profound loss of mIL-6R $\alpha$  on all analyzed leukocyte subsets, we observed only a marginal increase in serum concentration of sIL-6R $\alpha$  following *L. monocytogenes* infection. This result would argue that leukocytes are not a major source of sIL-6R $\alpha$  or that concentrations of sIL-6R $\alpha$  are tightly controlled and excessive sIL-6R $\alpha$  is rapidly eliminated from the circulation. Interestingly, naive and infected CCR2<sup>KO</sup> mice showed a reduction of sIL-6R $\alpha$  concentrations. Thus, inflammatory monocytes or other CCR2-dependent cells represent a source for sIL-6R $\alpha$  already under homeostatic conditions. Recently, it was demonstrated that IL-6R $\alpha$  deficiency in hepatocytes (AlbCre<sup>+</sup>Il6ra<sup>fl/fl</sup> mice) results in a 30% reduction of sIL-6R $\alpha$  serum levels, and deficiency in granulocytes and macrophages (LysMCre<sup>+</sup>Il6ra<sup>fl/fl</sup> mice) results in a 60% reduction of these levels (54). Because lysozyme M is most likely also expressed in inflammatory monocytes, our results are in accordance with this conclusion and add inflammatory monocytes to the list of cells involved in constitutive sIL-6R $\alpha$  production.

Differential blockade of global IL-6 signaling by neutralizing mAb and IL-6 trans-signaling by sgp130Fc indicated that classical signaling is sufficient for early control of *L. monocytogenes* infection. Infection of IL-6<sup>KO</sup> and sgp130Fc<sup>Tg</sup> mice confirmed this observation. Compared to wt mice or mice treated with sgp130Fc proteins, sgp130Fc<sup>Tg</sup> mice demonstrated even better control of *L. monocytogenes* infection. It is currently unclear why sgp130Fc<sup>Tg</sup> mice show this improved control. The constant Tg expression of sgp130Fc and the constant inhibition of IL-6 trans-signaling could



**FIGURE 9.** Role of classical IL-6 signaling and IL-6 trans-signaling in the immune response against *L. monocytogenes*. C57BL/6 wt, IL-6<sup>KO</sup>, and sgp130Fc<sup>Tg</sup> mice were i.v. infected with  $2 \times 10^4$  *L. monocytogenes*. After a further 3 d, mice were analyzed. (A) Listeria burden in spleen and liver. (B) Accumulation of inflammatory monocytes and neutrophils in spleen and liver. Values of individual mice and the median are shown in (A). Bars in (B) give the means  $\pm$  SD of six individually analyzed mice per group. \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .



cause alterations in development and cellular composition of the immune system, which might be responsible for the enhanced control of infection. In mice with a global blockade of IL-6, the increase in listeria titers was more pronounced in the liver, and thus IL-6 could be either particularly important for the innate response in the liver (e.g., through the hepatic induction of acute phase proteins) or there might be compensatory mechanisms in the spleen. Listeria infection induced a strong upregulation of the mRNA for several proinflammatory cytokines and chemokines. We observed a slight increase in chemokine mRNA expression following anti-IL-6 mAb treatment; however, it is unclear whether this was due to defective IL-6 signaling or due to a higher bacterial load. In contrast, treatment of mice with sgp130Fc did not change the mRNA expression levels of any of the analyzed factors. Infection with *L. monocytogenes* induced accumulation of neutrophil granulocytes in spleen and liver. Interestingly, sgp130Fc treatment caused a reduction of neutrophil accumulation in the spleen. Recent studies from Shi et al. (46) and Carr et al. (47) analyzed the role of neutrophils for the control of *L. monocytogenes*. In those studies, depletion of neutrophils had either no consequences (46) or only affected control of infection when relatively high doses of bacteria were applied (47). Under the latter conditions, depletion of neutrophils mainly impaired control of infection in the liver (47). Thus, it is not surprising that reduction in neutrophil numbers in the spleen was not accompanied by a more severe infection.

It has been demonstrated that IL-6 trans-signaling is responsible for a switch from a granulocyte to a monocyte/macrophage response (55). We observed reduction of monocyte frequencies in sgp130Fc<sup>Tg</sup> mice, but there was no change in monocyte frequencies in sgp130Fc- or anti-IL-6-treated animals. Thus, blockade of classical IL-6 signaling or IL-6 trans-signaling does not generally result in a reduced recruitment of monocytes to sites of *L. monocytogenes* infection. The failure to observe profound changes in the monocyte responses is most likely due to the infection model. Control of *L. monocytogenes* infection is highly dependent on recruitment of monocytes to the site of infection. Chemokines such as CCR2 ligands have been shown to release monocytes from the bone marrow (48). *L. monocytogenes* or material from these bacteria can induce production of CCR2 ligands in infected sites but also within the bone marrow, and this increase in local chemokines might be sufficient for releasing monocytes into the circulation (56). Alternatively, CCR2 ligands and chemokines in general appear to be dispensable for the accumulation of monocytes in the infected spleen and liver (48, 57). Thus, monocyte recruitment following *L. monocytogenes* infection could be largely independent of IL-6 and IL-6-mediated chemokine production at the sites of infection.

In summary, our results indicate that the protective function of IL-6 occurs in the absence of IL-6 trans-signaling, which also implies that protection is mainly due to mIL-6Rα<sup>+</sup> cells. The identity of these cells is currently unclear. Although recruitment of inflammatory monocytes is not impaired by IL-6 deficiency, these cells might be less functional without IL-6 stimulation. IL-6 could also operate in the activation and differentiation of NK cells, of unconventional "innate" T cells, or of conventional T cells. Finally, IL-6 might induce protective mechanisms in hepatocytes, for example, via the induction of the acute phase proteins. The response of these different cell types is currently under investigation by our group. Our results also suggest that IL-6 trans-signaling is not essential for protection against certain bacterial infections. Owing to this residual antibacterial immune response, blockade of IL-6 trans-signaling could therefore be of advantage to a global blockade of IL-6 for the treatment of chronic inflammatory diseases.

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## Disclosures

Dr. Rose-John is an inventor on patents describing the function of sgp130Fc. He is also a shareholder of the CONARIS Research Institute (Kiel, Germany), which is commercially developing sgp130Fc proteins as therapeutics for inflammatory diseases. The other authors have no financial conflicts of interest.

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