Toll-Like Receptor 4 Contributes to Efficient Control of Infection with the Protozoan Parasite *Leishmania major*

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The essential role of Toll-like receptors (TLR) in innate immune responses to bacterial pathogens is increasingly recognized, but very little is known about the role of TLRs in host defense against infections with eukaryotic pathogens. For the present study, we investigated whether TLRs contribute to the innate and acquired immune response to infection with the intracellular protozoan parasite *Leishmania major***. Our results show that TLR4 contributes to the control of parasite growth in both phases of the immune response. We also addressed the mechanism that results in killing or growth of the intracellular parasites. Control of parasite replication correlates with the early induction of inducible nitric oxide synthase in TLR4-competent mice, whereas increased parasite survival in host cells from TLR4-deficient mice correlates with a higher activity of arginase, an enzyme known to promote parasite growth. This is the first study showing that TLR4 contributes to the effective control of** *Leishmania* **infection in vivo.**

The leishmaniases are a spectrum of vector-borne parasitic diseases that are a major international public health problem, affecting the lives of millions of people worldwide (49). Experimental infection of mice with *Leishmania major* is widely used as a model for host resistance or susceptibility. The majority of inbred strains of mice (C57BL/6 and CBA) can control *L. major* infections, and only mice from a few strains (BALB/c) develop a progressive, nonhealing disease (12, 28, 44, 50). The outcomes of *Leishmania* infections are determined by adaptive T–helper (Th) cell responses and their interactions with parasitized host cells, usually macrophages. Th1 responses are associated with healing and parasite killing, whereas Th2 responses are associated with nonhealing diseases and uncontrolled parasite growth (50).

Depending on their activation state, macrophages can either host or kill *Leishmania*. The balance of the activities of inducible nitric oxide synthase (iNOS) and arginase is of crucial importance to parasite fate. These two enzymes are competitively regulated by the cytokines secreted by Th1 and Th2 cells: Th1 cytokines induce iNOS, whereas Th2 cytokines induce arginase (32, 34, 35). The induction of iNOS leads to oxidation of the amino acid L-arginine and the subsequent production of citrulline and nitric oxide (NO). Synthesis of NO has been shown to correlate with the killing of *Leishmania* parasites both in vitro and in vivo (5, 17, 27, 29, 55), and mice lacking iNOS fail to control *L. major* infections in vivo (62). Th2 cytokines, on the other hand, promote an alternative activation of macrophages, resulting in reduced killing ability and in the induction of high arginase levels (14, 16, 34). Arginase cata-

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lyzes the hydrolysis of L-arginine to urea and L-ornithine, and the latter can be used by the parasite for the synthesis of polyamines, which are essential for the growth of *Leishmania* (22).

Innate immunity coordinates the inflammatory response to pathogens, and the contribution of Toll-like receptors (TLRs) to this response is becoming widely recognized. TLRs are triggered by pathogen-associated molecular pattern molecules (PAMPs), which are characteristic of various groups of pathogens. Mammalian cells can express up to 10 different TLRs (48), which share an intracellular domain, called Toll–IL-1R (64), and signal through the myeloid differentiation protein 88 (MyD88) (30). The MyD88 signaling pathway results in the nuclear translocation of NF- κ B and the expression of cytokine genes. *L. major* activates interleukin- 1α (IL- 1α) in macrophages through a MyD88-dependent pathway in vitro (19), and genetically resistant mice lacking the MyD88 adapter protein develop progressive lesions and a polarized Th2 response (36). The PAMPs recognized by some of the TLRs have been identified. For example, TLR2 recognizes peptidoglycan and bacterial lipoproteins on gram-positive bacteria and mycobacteria (2, 6, 56, 60, 66) and is also activated by glycosylphosphatidylinositol anchors and glycoinositolphospholipids from *Trypanosoma cruzi* (9). Enterobacterial lipopolysaccharide (LPS) is the ligand for TLR4, and the activation of this receptor is critical for the response to gram-negative bacteria (21, 41). In contrast to the ample evidence for the recognition of bacterial PAMPs, very little work has been done on the role of TLRs in the host response to infection with eukaryotic parasites. Although MyD88 signaling is important for the healing of *L. major* infections (36), there are no reports about TLR engagement by the intracellular parasite *Leishmania*, and the contribution of TLRs to innate leishmanicidal responses is unknown. Our previous work has suggested a role for TLR4 in the control of *L. major* infections (33). We showed that C57BL/10ScCr mice,

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which carry a homozygous null mutation in the *tlr4* gene (41), were unable to resolve cutaneous lesions and to restrict parasite growth (33). However, these mice carry an additional mutation, resulting in impaired responsiveness to IL-12 (31, 42). Therefore, to evaluate the contribution of TLR4 alone to the host defense against *L. major* infection, we used the progenitor strain of the C57BL/10ScCr mice, C57BL/10ScN, which carries an identical deletion of TLR4 (TLR4 $^{0/0}$) but has intact IL-12 responsiveness (31, 42, 43). We investigated innate and adaptive immune responses to *L. major* infection in TLR4^{0/0} mice and compared these to responses in wild-type mice. The results of our study provide compelling evidence that TLR4 engagement is required for efficient parasite control by innate and adaptive immune responses.

MATERIALS AND METHODS

Mice. C57BL/10ScSn (wild-type), C57BL/10ScN (TLR4^{0/0}), and BALB/c mice were bred under specific-pathogen-free conditions in the animal facilities at the Max-Planck-Institut für Immunbiologie, Freiburg, Germany. C57BL/10ScN mice have a homozygous deletion of 74,723 bp at the *tlr4* locus which removes all three exons (41, 42). Six- to 10-week-old animals of both sexes were used for this study.

Parasites and infection. *L. major* LV39 (MRHO/SU/59/P strain) was maintained in a virulent state by monthly passaging in mice (25). For infections, $2 \times$ 10⁶ stationary-phase parasites were injected subcutaneously into the hind foot pads of mice. The lesions were measured by determining the increase in the foot pad thickness compared to the uninfected contralateral foot pad with a dial gauge caliper (Kröplin Schnelltaster, Schlüchtern, Germany).

To determine whether *L. major* promastigotes express LPS-like activity, the parasites were analyzed by a *Limulus* test for endotoxin activity. The results showed that the concentration of *L. major* promastigotes used for infection (2 \times 10^6 per mouse) did not cause detectable LPS activity (<1 pg). The parasite growth medium also contained no detectable endotoxin activity.

Macrophages. Bone marrow was obtained by flushing the femurs of naïve 5- to 6-week-old mice. Bone marrow precursor cells were cultured as previously described (33) in hydrophobic Teflon bags (Biofolie 25, Heraeus, Hanau, Germany) in Dulbecco's modified Eagle's medium containing 10% heat-inactivated fetal calf serum, 5% horse serum, and the supernatant of L929 fibroblasts at a final concentration of 15% (vol/vol) as a source of colony-stimulating factors, which drive cell proliferation toward a pure population of bone-marrow-derived macrophages (BMM ϕ). After 9 to 10 days of culturing, macrophages were harvested and 5×10^5 cells ml⁻¹ were plated and stimulated in the presence or absence of 20 U of gamma interferon (IFN- γ) ml⁻¹, 200 U of tumor necrosis factor alpha (TNF- α) ml⁻¹, 20 U of IL-4 ml⁻¹, 1 µg of LPS ml⁻¹, and 25 × 10⁵ *L. major* parasites ml^{-1} .

Determination of arginase activity. Arginase activity was measured in macrophage lysates as previously described (34). Briefly, cells were lysed with 100 µl of 0.1% Triton X-100. After 30 min on a shaker, 100 μ l of 25 mM Tris-HCl was added. To 100 μ l of this lysate, 10 μ l of 10 mM MnCl₂ was added, and the enzyme was activated by heating for 10 min at 56°C. Arginine hydrolysis was conducted by incubating the lysate with 100 μ l of 0.5 M L-arginine (pH 9.7) at 37°C for 15 to 20 min. The reaction was stopped with 800 μ l of H₂SO₄ (96%)- H_3PO_4 – (85%)- H_2O (1/3/7 [vol/vol/vol]). The urea concentration was measured at 540 nm after the addition of 40 μ l of α -isonitrosopropiophenone (dissolved in 100% ethanol) followed by heating at 95°C for 30 min. One unit of enzyme activity was defined as the amount of enzyme that catalyzed the formation of 1 umol of urea per min.

Determination of in vitro killing and survival of *L. major* **parasites.** Mature BMM ϕ (5 \times 10⁵ ml⁻¹) were plated and infected 4 h later with 25 \times 10⁵ *L. major* parasites ml⁻¹ in the presence of IFN- γ (20 U ml⁻¹) and TNF- α (200 U ml⁻¹) (Th1 conditions) or IL-4 (20 U ml⁻¹) (Th2 conditions). After 48 h, the macrophages were washed and lysed (33), and a limiting dilution assay was performed (25, 58) to determine the number of viable parasites.

RT-PCR analysis. For reverse transcription (RT)-PCR, total RNAs were extracted from mouse tissues and macrophages by use of TriReagent (Sigma) according to the manufacturer's instructions. Total RNAs were treated with DNase I (Ambion) and then reverse transcribed with an oligo(dT) primer (Promega) and Omniscript reverse transcriptase (Qiagen). cDNA samples were standardized, based on PCR amplification of the housekeeping gene β -actin. TLR4, arginase 1, and iNOS were amplified by PCR as described previously (1, 63, 65)

and were analyzed by densitometry using a UVP bioimaging system and Labworks analysis software.

Determination of parasite load. The number of living *L. major* parasites in infected tissues was determined with the parasite limiting dilution assay (25, 58). Briefly, serial dilutions of the footpad homogenate were distributed in replicate wells, and the plates were incubated at 26°C. After 10 to 14 days, the results were read microscopically and the number of viable parasites in the tissue was determined as previously described (25).

Cytokine measurements. Cells $(5 \times 10^6 \text{ ml}^{-1})$ from the lymph nodes draining the lesions of individual mice were restimulated with 1×10^6 *L. major* promastigotes ml^{-1} . Forty-eight hours later, the culture supernatants were harvested and cytokines and chemokines were measured by an enzyme-linked immunosorbent assay (ELISA) performed according to the suppliers' protocols. Detection limits were 20 pg ml⁻¹ for IL-10, 7.8 pg ml⁻¹ for IL-13, 2 pg ml⁻¹ for IL-6, 1 U ml^{-1} for IFN- γ , and 4 pg ml⁻¹ for monocyte chemoattractant protein 1 (MCP-1).

Flow cytometric analysis. For flow cytometric analysis, lymphoid cells were stimulated as described above. Six days later, the cells were harvested and a Ficoll gradient was created. The detection of intracellular cytokine was performed as described previously (40). Briefly, 1.5×10^6 cells were stimulated for 4 hours with 50 ng of phorbol 12-myristate 13-acetate (Sigma) and 500 ng of ionomycin (Calbiochem), or as a control, in the presence of complete medium alone, with 10 μ g of brefeldin A (Sigma) added during the last 2 hours. Before surface labeling was done with an anti-CD4 antibody (clone H129.19 or RM4-5; PharMingen), cells were preincubated with 1μ g of the rat anti-mouse monoclonal antibody CD32/CD16 (FcyII/III receptor) (PharMingen) to reduce nonspecific binding. Cells were washed, fixed with 2% formaldehyde (Sigma), and permeabilized with 0.5% saponin (Sigma) before anti-cytokine antibodies or isotype controls were added (anti-IL-4, clone BVD4-1D11; anti-IFN- γ , clone XMG1.2; anti-IL-10, clone JES5-16E3; and appropriately labeled rat immunoglobulin G_1 [PharMingen]). Detection of the intracellular cytokines was done with an EPICS XL instrument (Beckman Coulter), and data were analyzed with Beckman Coulter Expo32 software.

Statistical analyses. Experimental results were analyzed with GraphPad PRISM, version 2.0 (San Diego, Calif.), using a two-tailed Mann-Whitney test, and differences were considered statistically significant at P values of ≤ 0.05 .

RESULTS

Contribution of TLR4 to the control of parasite growth in vivo. To determine whether the presence of TLR4 contributes to the control of *L. major* parasites in vivo, we infected wildtype and TLR40/0 mice with *L. major* promastigotes and assessed the parasite loads at the local sites of infection. The results presented in Fig. 1A show that 24 h after infection, there was a significantly higher number $(P < 0.05)$ of parasites present in the lesions of TLR4^{0/0} mice than in those of wildtype mice. Interestingly, iNOS mRNA expression was detected in the footpads of the wild-type mice at 1 day postinfection but was not detectable at the sites of infection of TLR4^{0/0} mice (Fig. 1B). Arginase 1 mRNA was detectable in both groups of mice (Fig. 1B). TLR4 mRNA was detected in the footpads of naïve and infected wild-type mice (Fig. 1B), indicating that TLR4-expressing cells are present at the local sites of infection. However, the phenotype of the TLR4-expressing cells in vivo remains to be determined.

Thus, our results demonstrate that TLR4 is required for more efficient parasite control during the innate immune response to *L. major* infection.

To assess the contribution of TLR4 to the *L. major*-specific response, we determined the parasite load at 5 days postinfection, at a time when the healing phenotype of the TLR4 competent mice is unequivocally established (28 days) (33) and at a time when TLR4-competent mice had completely resolved their lesions (11 weeks). The lesions of *L. major*-infected TLR40/0 mice contained significantly larger parasite loads 5,

FIG. 1. Parasite loads and protein expression in mice. (A) Parasite loads in the footpads of TLR4-competent and TLR4-deficient mice 24 h after *L. major* infection. Groups of wild-type $(n = 4)$; filled circles) and TLR4-deficient ($n = 4$; open circles) mice were infected with 2 \times 106 *L. major* promastigotes. At 1 day postinfection, the numbers of viable parasites in the infected footpads were determined. Data show the results of one representative experiment of three independent experiments. Each symbol represents one mouse, and the horizontal black bars represent the means for four individual mice. $*, P < 0.05$. (B) Expression of iNOS, arginase, and TLR4 mRNAs in the foot pads of *L. major*-infected TLR4-competent and TLR4-deficient mice. The footpads of wild-type $(+/+)$ and TLR4-deficient $(0/0)$ mice were infected with $2 \times 10^6 L$. *major* parasites. After 24 h, gene expression was analyzed in the footpads of naïve and infected mice by semiquantitative RT-PCR. Samples were standardized by densitometric comparison of the amplification of the house keeping gene β -actin.

28, and 78 days after infection (Fig. 2A). We must note that TLR40/0 mice harbored 13.4- and 3.8-fold more parasites than the TLR4-competent mice 1 and 5 days after infection, respectively. Interestingly, 5 days after infection, the parasite loads had increased 49.2-fold for wild-type mice, whereas for TLR40/0 mice the increase was 14.2-fold. These differences in initial parasite control are likely due to more efficient immediate defense systems in TLR4-competent mice. While the parasite loads in the footpads of TLR4^{0/0} mice were still increasing at 28 days postinfection, they had started to decrease in wild-type mice (Fig. 2A). At 11 weeks postinfection, even though TLR4^{0/0} mice had resolved their lesions, they still harbored significant amounts of parasites, whereas no parasites were detectable in the lesions of TLR4-competent mice. These data are in agreement with our previous findings (33) and confirm that TLR4-competent C57BL/10ScSn mice are able to control parasite growth and to resolve lesions. In contrast, TLR40/0 mice not only had significantly larger parasite loads throughout the course of infection, but they were also less efficient at resolving the lesions and still had significantly larger cutaneous lesions and parasite loads after 2 months of infection, at a time when the lesions of the control mice had healed (Fig. 2B).

These results show that TLR4^{0/0} mice cannot efficiently control parasite replication and resolve cutaneous lesions. Therefore, the activation of TLR4 is not only crucial for the innate immune response, but it is also essential for a successful adaptive immune response to *L. major* parasites.

Influence of TLR4 on parasite-specific cytokine and chemokine production. The levels of *L. major*-specific cytokines and chemokines secreted by cells in the draining lymph nodes of wild-type and $TLR4^{0/0}$ mice were determined at 4 weeks postinfection. The results presented in Fig. 3 show that lymph node cells from $TLR4^{0/0}$ mice produced higher levels of IL-10 in response to antigenic restimulation than did cells from infected wild-type mice. Levels of IL-6 and IL-13 secreted by lymph node cells from *L. major*-infected TLR4^{0/0} mice were also higher, but IL-4 was not detectable. The levels of *L. major*-specific IFN- γ were higher in the supernatants of lymph node cells cultured from TLR40/0 mice than in those from wild-type mice (Fig. 3). Lymph node cells from *L. major*-infected TLR40/0 mice secreted more MCP-1 than did those from wild-type mice (Fig. 3). Thus, even though the differences were not statistically significant $(P > 0.05)$ 4 weeks after infection, lymph node cells from *L. major*-infected TLR4^{0/0} mice secreted higher levels of Th1 and Th2 cytokines than did those from infected wild-type mice. Similar results were obtained by determining the frequency of cytokine-expressing CD4⁺ T cells by intracellular cytokine staining. Four weeks after infection, there was a slightly higher frequency of IFN- γ -, IL-4-, and IL-10-expressing $CD4^+$ T cells in the lymph nodes draining the lesions of TLR40/0 mice than in those of control mice, and the mean fluorescence intensities were also slightly higher (*P* 0.05) (Table 1) in response to antigenic restimulation in vitro. The cytokine levels detected directly ex vivo were below the detection limit (1%).

Parasite survival and arginase activity in BMM ϕ from **TLR40/0 and wild-type mice.** We showed in Fig. 1 and 2 that TLR40/0 mice harbor larger parasite loads in their infected foot pads than do wild-type mice. To assess whether the macro-

phages from TLR40/0 mice provide a more permissive environment for parasite growth, we determined whether the absence of TLR4 influences the survival of parasites in BMM ϕ in vitro. The results presented in Fig. 4 show that *L. major* growth is strongly enhanced in BMM ϕ activated with Th2 cytokines (alternatively activated BMM ϕ) from TLR4 $^{0/0}$ mice. Thus, TLR4-competent macrophages provide a less permissive environment for *Leishmania*, controlling the growth of intracellular parasites more efficiently than TLR4^{0/0} macrophages. In the absence of exogenous cytokines, macrophages from TLR40/0 mice supported parasite growth more efficiently (1.5- to 3.5 fold more parasites survived) (data not illustrated) than did those from wild-type mice. Furthermore, these results confirm that the activation of macrophages with type 1 cytokines results in parasite killing, whereas activation with type 2 cytokines promotes parasite growth. We conclude from the differential survival of L . *major* parasites in wild-type and $TLR4^{0/0}$ mac-

rophages that TLR4 contributes to an efficient host defense against *Leishmania* parasites.

Since iNOS and arginase activity have been associated with parasite killing and parasite growth, respectively, we determined their relative expression levels in *L. major*-infected macrophages from wild-type and TLR4^{0/0} mice. The activation of *L. major*infected macrophages by IL-4 induced increased arginase 1 mRNA expression, whereas activation with type 1 cytokines (IFN- γ plus TNF- α) upregulated iNOS mRNA expression (Fig. 4). The weak expression of arginase 1 in macrophages activated with type 1 cytokines is due to the autocrine production of IL-10 (35). Similar results were obtained by analyzing iNOS mRNA expression with an RNase protection assay and by measuring the levels of nitric oxide in the culture supernatants (data not shown). Control stimulation with LPS upregulated both arginase 1 and iNOS mRNAs in macrophages from wildtype mice, but not in those from TLR4^{0/0} mice (data not shown).

FIG. 3. Cytokine and chemokine production by lymph node cells from *L. major*-infected wild-type and TLR4^{0/0} mice. Groups of wildtype (filled circles) and $TLR4^{0/0}$ (open circles) mice were infected with 2×10^6 *L. major* promastigotes in one hind footpad. At 4 weeks postinfection, 5×10^6 popliteal lymph node cells were restimulated with 10^6 *L. major* parasites. Supernatants were harvested after 48 h and tested for their cytokine contents by ELISA. Each symbol represents one mouse, and the horizontal black bars represent the means for four individual mice. $**$, $P > 0.05$.

Thus, the differential survival of *Leishmania* parasites in macrophages (Fig. 4) correlates with the induction of iNOS and arginase mRNAs in these cells.

In vitro arginase activity in *L. major***-infected macrophages from TLR40/0 and wild-type mice.** Since increased arginase levels have been shown to promote the multiplication of *Leishmania* parasites (22) and since TLR4^{0/0} macrophages are more permissive for *Leishmania* growth (Fig. 4), we determined the enzymatic activity of arginase in macrophages from wild-type and $TLR4^{0/0}$ mice. The results presented in Fig. 5A show that the stimulation of BMM ϕ in vitro with IL-4 alone induced similar arginase activities in wild-type and TLR40/0 mice. However, the simultaneous infection of these macrophages with *L.*

FIG. 4. Parasite survival in macrophages from $TLR4^{0/0}$ and wildtype mice. BMM ϕ (5 \times 10⁵ ml⁻¹) from naïve wild-type (black bars) and TLR4^{0/0} (open bars) mice were infected with 25×10^5 *L. major* parasites ml⁻¹ in the presence of IFN- γ (20 U ml⁻¹) and TNF- α (200 U ml⁻¹) (Th1 conditions) or IL-4 (20 U ml⁻¹) (Th2 conditions). After 48 h of incubation, macrophages were lysed and a limiting dilution assay was performed to determine the number of viable parasites. Data show the results of one representative experiment of four independent experiments. In addition, gene expression was analyzed by semiquantitative RT-PCR. Samples were standardized by densitometric comparison with the amplification of the housekeeping gene β -actin. Data show the results of one representative experiment of two independent experiments.

major parasites clearly creates synergism with IL-4 in vitro, resulting in the induction of 3.8-fold more arginase activity in macrophages from TLR4^{0/0} mice than in those from wild-type mice. The arginase activity in parasitized macrophages from TLR40/0 mice in vitro was comparable to that induced in macrophages from the nonhealer BALB/c strain (Fig. 5A). The simultaneous activation of *L. major*-infected macrophages with IL-10 and IL-4 enhanced the arginase activity even further (for TLR4 $^{0/0}$ mice, 1,443 mU/10⁶ cells; for the wild type, 622 mU/ 106 cells; data not illustrated). *L. major* parasites themselves constitutively express arginase activity (8); however, it is unlikely that the parasite arginase activity contributed to the values shown in Fig. 5A, since the arginase activity of 2.5×10^6 *L. major* promastigotes ml^{-1} was below the limit of detection (data not shown).

These results demonstrate that in the absence of TLR4, alternatively activated *L. major*-infected macrophages express

TABLE 1. Intracellular cytokine staining in lymph node cells from *L. major*-infected wild-type and TLR4^{0/0} mice^a

Mouse strain	$%$ CD4 ⁺ cells	$%$ CD4 ⁺ IFN- γ ⁺ cells	MFI^b	$%$ CD4 ⁺ IL-4 ⁺ cells	MFI^b	$\%$ CD4 ⁺ IL-10 ⁺ cells	MFI ^b
Wild-type TLR4 ^{0/0}	50.3 ± 2.0	6.1 ± 1.2	5.4 ± 0.2	1.3 ± 0.1	4.5 ± 0.2	2.3 ± 0.7	4.0 ± 0.2
	52.1 ± 2.8	7.7 ± 3.3	6.2 ± 1.5	1.9 ± 0.7	5.6 ± 1.0	2.8 ± 1.1	4.6 ± 0.2

a Groups of wild-type and TLR4^{0/0} mice were infected with 2×10^6 *L. major* promastigotes in one hind footpad. At 4 weeks postinfection, 5×10^6 popliteal lymph node cells were stimulated with 10⁶ *L. major* parasites. Six days later, a Ficoll gradient was performed and the cells were restimulated with PMA and ionomycin in the presence of brefeldin-A, as described in Materials and Methods. The frequencies of IFN- γ -, IL-4-, and IL-10-expressing CD4⁺ T cells were determined by flow cytometry, and the values represent averages \pm standard deviations for four mice. The results of one of two similar experiments are presented. *b* MFI, mean fluorescence intensity.

FIG. 5. (A) Differential induction of arginase activity in *L. major*infected macrophages from TLR4-competent or TLR4-deficient mice. BMM ϕ (5 \times 10⁵ ml⁻¹) from naïve wild-type (black bars), TLR4^{0/0} (white bars), and BALB/c (hatched bars) mice were cultured in the presence and/or absence of 25×10^5 *L. major* parasites ml⁻¹ and IL-4 (20 U ml^{-1}) . After 48 h, the arginase activity in macrophage lysates was measured. Data show the results of one representative experiment of five independent experiments. (B) MCP production by *L. major*infected macrophages from TLR4-competent or TLR4-deficient mice. BMM ϕ (5 \times 10⁵ ml⁻¹) from naïve wild-type (black bars) and TLR4^{0/0} (white bars) mice were cultured in the presence or absence of 25×10^5 L. *major* parasites ml^{-1} and IL-4 (20 U ml⁻¹). Supernatants were harvested after 48 h and tested for their MCP-1 contents by ELISA. Data show the results of one representative experiment of three independent experiments.

more arginase activity in vitro, which is likely to promote the polyamine synthesis required for parasite growth. Thus, the activation of parasitized macrophages with type 2 cytokines not only induces more arginase activity in the absence of TLR4, but also results in enhanced parasite growth.

Induction of cytokine and chemokine production in *L. major***-infected macrophages.** To identify the cytokines and chemokines produced as a consequence of parasite interactions with macrophages, we tested the culture supernatants for the presence of IL-1 α , IL-6, IL-10, IL-12 p70, and MCP-1. The infection of macrophages from wild-type and $TLR4^{0/0}$ mice

with *L. major* for 48 h did not result in detectable cytokine production (data not shown). The stimulation of infected macrophages with type 1 cytokines only induced IL-6 production (for wild-type mice, 171 ± 36 pg ml⁻¹; for TLR4^{0/0} mice, 215 \pm 69 pg ml⁻¹), whereas stimulation with type 2 cytokines did not induce any detectable cytokine production (data not shown). These results confirm that *L. major* infection results in a poor activation of macrophages, a feature that helps the parasites to escape immune defense mechanisms. The chemokine MCP-1 was constitutively produced by naïve macrophages, and infection with *L. major* resulted in slightly increased levels. IL-4 formed synergism with the parasites to induce an increase in MCP-1 production, which was further enhanced in the absence of TLR4 (Fig. 5B).

DISCUSSION

The results of our study demonstrate unequivocally that TLR4 plays a role in the host defense against parasitic protozoa. We show that after infection with *L. major*, TLR4 contributes to both innate and adaptive immune responses: TLR40/0 mice are clearly less efficient at controlling parasite growth at the local site of infection in the early (days 1 and 5) as well as the late (weeks 4 and 11) phases of infection. The more efficient parasite control of TLR4-competent mice 1 day after infection correlated with the expression of iNOS at the local site of infection. TLR4^{0/0} mice not only had larger parasite burdens, but they were also less efficient in the resolution of cutaneous lesions. Indeed, about 2 months after infection, the cutaneous lesions were healed in the TLR4-competent mice, whereas TLR4^{0/0} mice still displayed pronounced foot pad swelling. In agreement with the increased parasite growth in the footpads of TLR4^{0/0} mice, alternatively activated macrophages were more permissive for parasite growth in the absence of TLR4 in vitro. This increased parasite proliferation was associated with more arginase activity; indeed, our results show that IL-4 formed synergism with *L. major* in the induction of enhanced arginase activity in TLR40/0 macrophages, indicating that TLR4 signaling contributes to the regulation of L-arginine metabolism. The growth of *Leishmania* parasites within the macrophage environment is dependent on their ability to ensure the provision of nutrients. Arginase catalyzes the hydrolysis of L-arginine into urea and L-ornithine, and the latter is used by parasites to generate polyamines, which are essential for their proliferation. A correlation between intracellular parasite growth and arginase induction was also observed when infected alternatively activated macrophages from nonhealer and healer strains of mice were compared in vitro: parasite growth and arginase levels were significantly higher in macrophages from susceptible BALB/c mice than in those from resistant C57BL/6 mice (22). In a *Schistosoma mansoni* infection model, the inducible expression and activation of arginase 1 in vivo directly correlated with a type 2 cytokine response, and arginase activity was functionally related to pathology in the granulomas induced by schistosome eggs (20). Several studies with mice have indicated that the balance between arginase and iNOS is an important mechanism for controlling macrophage function (16, 32, 34). Indeed, the expression of iNOS or arginase in vivo has been suggested to be a better predictor of the pathology induced by schistosome eggs

than the Th1 or Th2 cytokine levels (20). There is ample evidence that the iNOS pathway is crucial for the killing of *Leishmania* parasites (5, 17, 27, 29, 55, 62). Here we showed that increased parasite killing in wild-type mice correlates with iNOS expression in *L. major*-infected macrophages and in the footpads of TLR4-competent mice infected for 24 h. These data confirm that TLR4 signaling contributes to the induction of the iNOS pathway (52, 59). Interestingly, the engagement of TLR4 by LPS results in the induction of IFN- α/β (54), and the early expression of iNOS is dependent on IFN- α/β (11). Since IFN- α/β has a protective role in experimental leishmaniasis (4), it is tempting to speculate that the beneficial effect of TLR4 in the control of *L. major* infection is in part associated with TLR4-mediated induction of early IFN- α/β .

Of the cytokines that were elevated in the absence of TLR4 after in vivo infection, the antigen-specific production of IL-10 is of particular interest. The severity of visceral leishmaniasis is strongly associated with increased IL-10 levels (24, 50), while CD4⁺ T cells from *L. major*-infected nonhealer mice express high levels of IL-10 mRNA (45) and IL-10-deficient mice from a nonhealer background can control *L. major* infections (3, 23). Since IL-10 acts primarily on activated macrophages to decrease the secretion of proinflammatory cytokines and to prevent parasite killing (23), it is likely that the increased levels of IL-10 in *L. major*-infected TLR4^{0/0} mice contribute to the observed reduction in host defense. Here we demonstrated that IL-10 forms synergism with IL-4 and *L. major* to increase arginase 1 activity in vitro. This is in agreement with previous work showing that arginase 1 expression is induced by IL-4 and IL-13 (35) and that IL-10 strongly forms synergism with Th2 cytokines for the induction of arginase in vitro (34). In addition, IL-10 alone induces low levels of this enzyme (32). Global gene expression analysis has shown that IL-10 influences not only arginase 1, but also arginase 2, induction in macrophages (26) and that IL-10 also causes an upregulation of IL-4R α expression (26). Thus, a functional consequence of the exposure of macrophages to IL-10 is an enhanced sensitivity to IL-4 and IL-13, with both cytokines promoting the alternative activation of macrophages, which express higher levels of arginase and thus promote the growth of intracellular pathogens. By the same criteria, the increased levels of IL-10 in *L. major*-infected TLR40/0 mice are likely to enhance macrophage responsiveness to Th2 cytokines, thereby enhancing the permissiveness of these host cells for parasite growth. Thus, our results demonstrating that alternatively activated parasitized macrophages from TLR4-deficient mice express increased arginase activity in vitro (Fig. 5) and are more permissive for parasite growth in vitro (Fig. 2) suggest that TLR4-mediated signaling modulates the effector function of parasitized macrophages in favor of the pathway resulting in parasite killing. However, increased arginase activity in alternatively activated, parasitized macrophages in response to IL-4 in vitro does not necessarily reflect the complex mechanisms underlying the increased parasite survival rate in the absence of TLR4 in vivo.

The detection of increased levels of MCP-1 in the draining lymph nodes of TLR4^{0/0} mice at a time when the acquired immune response is unequivocally established suggests a negative correlation with parasite killing. MCP-1 influences innate immunity through its effects on the recruitment of monocytes and adaptive immunity by its effects on T-helper cells, specifically by controlling the polarization of Th2 cells (18, 61). Indeed, MCP-1-deficient mice from a nonhealer background are resistant to *L. major* infections and unable to mount a Th2 response (18). Our results support these observations, as higher levels of MCP-1 in TLR $4^{0,0}$ mice correlate with a more pronounced type 2 response, less efficient control of parasite replication, and lesion resolution. Although a pathogenic role for MCP-1 has been shown previously (10, 15) and although MCP-1 has been implicated in Th2 polarization, beneficial effects of MCP-1 in cutaneous leishmaniasis have also been reported. Infection with *L. major* has been shown to induce a rapid transient induction of MCP-1 1 day after infection in mice from healer strains (61), and MCP-1 has been reported to form synergism with IFN- γ to promote parasite killing in vitro (46). Mice that are from a genetically resistant background but are unable to express the chemokine receptor CCR2, the ligand for MCP-1, are susceptible to *L. major* infection (51), while increased levels of MCP-1 are found in the lesions of patients with self-healing disease (47).

Our data show that TLR4 signaling helps parasitized macrophages to kill intracellular *L. major* more efficiently and to control arginase activity, an effect that might be ascribed to the balance of Th1 and Th2 responses. Indeed, signaling through TLRs has been associated with the induction of Th1 responses and a lack of signaling has been associated with the induction of Th2 responses (39, 53). Although we showed that TLR4 contributes to the efficient control of *L. major* parasites during both innate and adaptive immune responses, we cannot exclude the possibility that the natural mutation in the *tlr4* locus (37, 42, 43) leads to altered homeostatic responses affecting early parasite survival. Indeed, it cannot be excluded that mechanisms of early defense, such as complement activation (49) , are altered in TLR $4^{0/0}$ mice compared to wild-type mice. We have obtained evidence that cell recruitment to the site of parasite inoculation is different for TLR4-competent and TLR40/0 mice 10 hours after infection (P. Kropf, N. Freudenberg, C. Kalis, M. Modolell, S. Herath, C. Galanos, M. Freundenberg, and I. Müller, submitted for publication). This initial difference in cell recruitment could contribute to the more efficient invasion of host cells 1 day after infection. Even though the parasite load is consistently higher in the absence of TLR4, it is interesting that the dynamics of parasite replication between days 1 and 5 are different. These differences in parasite growth could be due to (i) more efficient invasion of the host cells, (ii) more efficient growth of the parasites, or (iii) more efficient killing of the parasites.

In addition, in macrophages from TLR4^{0/0} mice, there may be more competition for polyamines, which is essential for their growth due to the higher parasite number in these cells at day 1. Furthermore, it is also possible that cooperation between different TLRs or between TLR4 and parasite-specific receptors (7, 13) is required, and we are currently addressing these issues. Regardless of the underlying mechanism, we have unequivocal evidence that the increased parasite load in the TLR40/0 mice used in the present study was indeed due to the absence of TLR4, as the insertion of a TLR4 transgene in these mice enabled them to control parasite replication as efficiently as the wild-type mice (Kropf et al., submitted).

Based on observations in bacteria, it can be hypothesized that TLRs recognize PAMPs associated with *Leishmania* parasites and that these interactions are essential for the implementation and maintenance of active innate and adaptive immune responses to infection. However, at least in vitro, *L. major* parasites do not have a direct effect on the activation of TLR4 in a dual luciferase reporter system (H. P. Price and D. F. Smith, unpublished data). Although at a first glance these data seem to indicate that TLR4 does not interact with parasite molecules in vitro, they may also indicate that in vivo multiple innate immune recognition receptors interact during pathogen recognition (7, 13) and that in vitro transfection systems do not necessarily reflect the complexity of host-parasite interactions in vivo. Because TLR4 not only interacts with pathogen-associated PAMPs, but can also recognize endogenous ligands such as heat shock proteins (38) and components of the extracellular matrix (57), we cannot yet conclude whether TLR4 activation after *L. major* infection is due to the recognition of parasite PAMPs or an interaction with endogenous ligands. TLRs may represent novel targets for the activation of the adaptive immune response and TLR agonists may be useful for the prevention and treatment of leishmaniasis.

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