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# *Francisella tularensis* Live Vaccine Strain Induces Macrophage Alternative Activation as a Survival Mechanism[1]

**Kari Ann Shirey, Leah E. Cole, Achsah D. Keegan, and Stefanie N. Vogel[2]**

*Francisella tularensis* (*Ft*), the causative agent of tularemia, elicits a potent inflammatory response early in infection, yet persists within host macrophages and can be lethal if left unchecked. We report in this study that *Ft* live vaccine strain (LVS) infection of murine macrophages induced TLR2-dependent expression of alternative activation markers that followed the appearance of classically activated markers. Intraperitoneal infection with *Ft* LVS also resulted in induction of alternatively activated macrophages (AA-M$\phi$). Induction of AA-M$\phi$ by treatment of cells with rIL-4 or by infection with *Ft* LVS promoted replication of intracellular *Ftn*, in contrast to classically activated (IFN-$\gamma$ plus LPS) macrophages that promoted intracellular killing of *Ft* LVS. *Ft* LVS failed to induce alternative activation in IL-4R$\alpha^{-/-}$ or STAT6$^{-/-}$ macrophages and prolonged the classical inflammatory response in these cells, resulting in intracellular killing of *Ft*. Treatment of macrophages with anti-IL-4 and anti-IL-13 Ab blunted *Ft*-induced AA-M$\phi$ differentiation and resulted in increased expression of IL-12 p70 and decreased bacterial replication. In vivo, *Ft*-infected IL-4R$\alpha^{-/-}$ mice exhibited increased survival compared with wild-type mice. Thus, redirection of macrophage differentiation by *Ft* LVS from a classical to an alternative activation state enables the organism to survive at the expense of the host. *The Journal of Immunology,* 2008, 181: 4159–4167.

*F*rancisella tularensis (*Ft*)[3] is an intracellular, Gram-negative coccobacillus that is the etiological agent of tularemia (reviewed in Refs. 1–4). *Ft* has been classified as a category A agent by the Centers for Disease Control and Prevention because it can be spread via an airborne route, is infectious at very low doses, and causes severe disease that may be fatal unless treated rapidly (1). Most basic research studying the pathogenesis of *Ft* has used a live vaccine strain (LVS) (1, 4, 5) because it is attenuated in humans, yet causes a disease resembling human tularemia in mice when administered by some routes of infection (4). Infection of mice with *Ft* LVS results in a pronounced inflammatory response, yet the organism is able to persist within host cells (6, 7).

Macrophages differentiate into functionally distinct immunological populations depending on the cytokine environment. Macrophages exposed to IFN-$\gamma$ and LPS become classically activated macrophages (CA-M$\phi$) and function predominantly in inflammation, tissue damage, killing of intracellular microbes, and increased tumoricidal activity (reviewed in Ref. 8). Conversely, certain cytokines, specifically IL-4 and IL-13, induce an alternatively activated phenotype in macrophages. Alternatively activated macrophages (AA-M$\phi$) are principally associated with allergic and parasitic immune responses, tissue remodeling, angiogenesis, tumor promotion, and humoral immunity (reviewed in Ref. 8), and, more recently, have been associated with Alzheimer's disease (9, 10). Some organisms, such as *Mycobacterium tuberculosis*, induce alternative activation as a means of survival; however, the organism remains in a persistent, nonreplicating state within granulomas (11). A third macrophage differentiation state has been referred to as MII or M$\phi$-II and is induced by costimulation of macrophages through Fc$\gamma$R ligation and TLR signaling (12, 13). Although the MII phenotype was first associated with increased IL-10 and decreased IL-12 production (14), recent evidence suggests that the inhibition of proinflammatory gene expression in MII macrophages extends to inducible NO synthase (iNOS) and other inflammatory gene products regulated by IFN regulatory factor-8 (13). AA-M$\phi$ markers are not expressed by MII macrophages (12). M$\phi$-II cells may play a role in the exacerbation of infectious diseases in which the presence of immune complexes can induce the production of IL-10 from macrophages, allowing for disease progression.

Therefore, we hypothesized that *Ft* may avoid macrophage-mediated killing by altering the macrophage differentiation state from one that is classically activated to one that is alternatively activated. We observed that after an initial, robust proinflammatory response, *Ft* LVS infection induces expression of markers associated with AA-M$\phi$ both in vitro and in vivo. This reprogramming of macrophage differentiation promotes the survival and replication of the bacterium while mitigating the proinflammatory response. The failure of *Ft* LVS infection to induce AA-M$\phi$ in IL-4R$\alpha^{-/-}$ or STAT6$^{-/-}$ macrophages results in a sustained CA-M$\phi$ phenotype and clearance of the bacterium. In vivo, IL-4R$\alpha^{-/-}$ mice exhibited increased survival compared with wild-type (WT) mice. Ab-mediated neutralization of IL-4 and IL-13 also reversed the AA-M$\phi$ phenotype in WT macrophages, blocked production of IL-4 and IL-13 by macrophages, increased IL-12 p70 secretion, and curtailed intracellular replication. Our data support the notion that macrophage differentiation is malleable, allowing for rapid responses to environmental conditions. This study provides new

insights into the innate immune response to *Ft* LVS infection and the mechanism by which *Ft* LVS evades the host innate immune response.

## Materials and Methods

### Reagents

Murine rIL-4 and rIFN-γ were purchased from R&D Systems. *Escherichia coli* K235 LPS was prepared, as previously described (15). Rabbit anti-mouse found in inflammatory zone (FIZZ) 1 Ab was isolated from serum (provided by S. Kunkel, University of Michigan, Ann Arbor, MI) using an ImmunoPure (A) IgG Purification kit (Pierce Endogen), per the manufacturer's protocol. The following Abs were purchased: anti-murine arginase-1 mAb (IgG1; BD Biosciences); polyclonal goat anti-mouse IL-12 p70, anti-IL-4 mAb, and anti-IL-13 mAb (R&D Systems); anti-mouse CD206 (mannose receptor):FITC (IgG2a) (AbD Serotec); anti-murine F4/80 mAb (IgG2b) (Abcam); mouse IgG2a and IgG1, rat IgG2b, goat IgG, and rabbit IgG isotype control Abs (Sigma-Aldrich); and Cy2-conjugated donkey anti-rabbit IgG, Cy2-conjugated donkey anti-mouse IgG, and Cy3-conjugated donkey anti-goat IgG (Jackson ImmunoResearch Laboratories).

Frozen aliquots of *Ft* LVS (ATCC 29684; American Type Culture Collection) were provided by K. Elkins (Food and Drug Administration, Rockville, MD) and stored, as previously described (16).

### Macrophage cell culture

Six- to 8-wk-old C57BL/6J, TLR2$^{-/-}$ (B6.129-Tlr2<tm1Kir>/J), and BALB/cByJ mice were purchased from The Jackson Laboratory. Breeding pairs of IL-4Rα$^{-/-}$ and STAT6$^{-/-}$ mice on a BALB/c background were obtained from N. Noben-Trauth (George Washington University Medical Center, Washington, DC) and W. Paul (Laboratory of Immunology, National Institute of Allergy and Infectious Diseases, National Institutes of Health, Bethesda, MD), respectively, and bred within University of Maryland's accredited facility. Peritoneal macrophages isolated from mice 4 days after i.p. injection of sterile 3% thioglycolate were cultured, as described previously (17). Macrophages were plated in 6-well (4 × 10$^6$ cells/well) or 12-well (2 × 10$^6$ cells/well) tissue culture plates (Corning Glass). After overnight incubation, cells were washed with PBS to remove nonadherent cells and were cultured in antibiotic-free medium for 24 h before treatment. All animal experiments were conducted with institutional approval.

Thioglycolate-elicited peritoneal macrophages (99.5% positive for F4/80) were plated in 6-well (4 × 10$^6$) or 12-well (2 × 10$^6$ cells) plates and stimulated with medium alone, rIL-4 (40 ng/ml), or rIFN-γ (20 ng/ml) plus LPS (10 ng/ml) for the indicated times as positive controls for AA-Mϕ and CA-Mϕ, respectively. Macrophages were infected with *Ft* LVS (multiplicity of infection (MOI) = 5; confirmed by colony count) and incubated at 37°C. In initial experiments (e.g., Fig. 1A), macrophages were not washed to remove extracellular bacteria. For all other experiments, cells were washed with PBS 1 h postinfection and placed in RPMI 1640 with gentamicin (50 μg/ml) for 1 h to kill extracellular bacteria. Cells were washed again to remove gentamicin and then lysed at the indicated time for preparation of RNA or measuring intracellular replication of *Ft* LVS. *Ft* LVS in cell lysates were determined by colony count. *Ft* LVS 16S rRNA was also quantified by real-time PCR to confirm the results of colony counts (18).

### Real-time PCR

Total RNA was isolated from macrophage cultures using TRIzol (Invitrogen), according to manufacturer's protocol. Real-time PCR was performed with an ABI 7900HT (Applied Biosystems) sequence detection system and software, as previously described (19). Relative mRNA levels for specific genes are reported as relative gene expression normalized to untreated control samples. The sequences for IL-4, IL-12 p35, IL-12 p40, hypoxanthine phosphoribosyltransferase, iNOS, and 16S *Ft* LVS rRNA have been reported previously (18). New primers were designed using the Primer Express program (Applied Biosystems) in conjuction with GenBank sequences using the Blastn program and include the following: arginase-1, sense (5′-CAGAAGAATGGAAGAGTCAG-3′) and antisense (5′-CAGA TATGCAGGCAGGGAGTCACC-3′); FIZZ1, sense (5′-GGTCCCAGTG CATATGGATGAGACCATAGA-3′) and antisense (5′-CACCTCTTCA CTCGAGGGACAGTTGGCAGC-3′); Ym1, sense (5′-CATGAGCAAG ACTTGCGTGAC-3′) and antisense (5′-GGTCCAAACTTCCATCCTCC A-3′); IL-13, sense (5′-GGAGCTGAGCAACATCACACAA-3′) and antisense (5′-GAATCCAGGGCTACACAGAACC-3′); mannose receptor, sense (5′-GATATGAAGCCATGTACTCCTTACTGG-3′) and antisense

(5′-GGCAGAGGTGCAGTCTGCAT-3′); TGF-β, sense (5′-GCAACATG TGGAACTCTACCAGAA-3′) and antisense (5′-GACGTCAAAAGACA GCCACTCA-3′); GM-CSF, sense (5′-TTGAATGAAGAGGTAGAAGT CGTCTC-3′) and antisense (5′-AATTGCCCCGTAGACCCTG-3′); and FcεRII, sense (5′-CAGCTGGGAGACACTGCAATT-3′) and antisense (5′-ATCTGAACAACCTGGGACTTCTG-3′). All primers were synthesized at the Biopolymer and Genetics Core Facility (University of Maryland).

### Flow cytometry analysis

To preclude the need for scraping cells for flow cytometric studies, macrophages from C57BL/6J, BALB/cByJ, IL-4Rα$^{-/-}$, or STAT6$^{-/-}$ mice were cultured on 6-well low cluster and low adhesion plates (Corning Glass). Washes were conducted in centrifuge tubes, and the cells were replated and treated with medium alone, rIL-4 (40 ng/ml), or IFN-γ (20 ng/ml) plus LPS (10 ng/ml), or infected with *Ft* LVS (MOI = 5) for 24 or 48 h. Cells were harvested for analysis by gentle shaking, washed with PBS, and then fixed with 4% *p*-formaldehyde for FIZZ1 and arginase-1 or with 70% methanol for mannose receptor for 10 min at room temperature. Cells were blocked and permeabilized for 30 min with PBST (PBS, 1% BSA, 1% normal donkey serum, 0.3% Triton X-100) at room temperature. FIZZ1 and IL-12 p70 were detected using polyclonal Abs directed against the proteins, followed by Cy2-conjugated donkey anti-rabbit IgG, or Cy3-conjugated donkey anti-goat IgG, respectively. Arginase-1 was detected by staining the cells with a mAb, followed by a secondary Cy2-conjugated donkey anti-mouse IgG2a Ab, and mannose receptor expression was determined by using a primary FITC-conjugated mAb. Cells were washed in PBST and suspended in PBS for immediate analysis using a FACSCalibur. Analytic gates were set to exclude cellular debris and aggregates. CellQuest software (BD Biosciences) was used to analyze the data.

To assess whether in vivo infection with *Ft* LVS also resulted in AA-Mϕ, C57BL/6 mice were injected i.p. with saline or *Ft* LVS (10,000 CFU) for 3 days. The mice were sacrificed, and peritoneal cells were harvested by lavage. After fixation with 4% *p*-formaldehyde, cells were simultaneously stained for FIZZ1 (an AA-Mϕ marker) and F4/80, a macrophage marker. FIZZ1 was detected as described above, and F4/80 was detected by staining the cells with a mAb, followed by a secondary Cy3-conjugated donkey anti-rat IgG Ab. Cells were washed in PBST and suspended in PBS for FACS analysis. Macrophages were identified as F4/80-positive cells with high forward and side scatter properties. A gate was placed on the F4/80-positive cells to exclude any nonmacrophage cells and cell debris and analyzed for FIZZ1 expression. Experimental and control groups consisted of six animals each.

### Ab neutralization assays

Peritoneal macrophages were treated with medium alone, rIL-4 (40 ng/ml), or rIL-13 (40 ng/ml), or infected with *Ft* LVS. Parallel sets of cells were additionally treated with either isotype control IgG, anti-IL-4 (100 μg), anti-IL-13 (100 μg), or both anti-IL-4 and anti-IL-13 Abs. After 48 h, total RNA was isolated, and FIZZ1, IL-12 p40, and IL-10 mRNA were analyzed by real-time PCR, or protein expression for FIZZ1 and IL-12 p70 determined by FACS analysis. To assess the effect of anti-IL-4 and/or anti-IL-13 Ab treatment on *Ft* LVS survival, macrophages were treated, as described for mRNA or FACS analysis, and, after 48 h, the number of *Ft* LVS in cell lysates was determined by colony counts.

### In vivo infection with Ft LVS

To assess the survival rate of mice infected with *Ft* LVS, BALB/cByJ WT and IL-4Rα$^{-/-}$ mice were inoculated i.p. with either saline or *Ft* LVS (10,000 CFU; four to five mice per treatment in four separate experiments). Survival was monitored for up to 10 days postinfection.

### Cytokine measurements

Cytokine concentrations in cell culture supernatants were measured by ELISA by the Cytokine Core Laboratory (University of Maryland).

### Statistics

Statistical differences between two groups were determined using unpaired, two-tailed Student's *t* test with significance set at $p < 0.05$. For comparisons between three or more groups, analysis was done by one-way ANOVA, followed by a Tukey's multiple comparison test with significance determined at $p < 0.05$. All statistical analyses were performed using Prism GraphPad software.
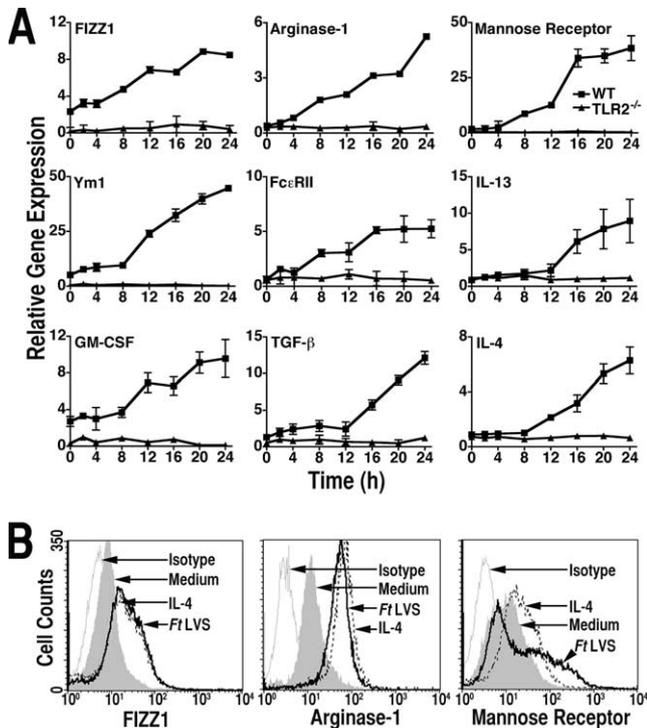
**FIGURE 1.** Induction of AA-Mø by *Ft* LVS. *A*, Peritoneal macrophages from C57BL/6 WT or TLR2$^{-/-}$ mice were infected with *Ft* LVS (MOI = 5). Total RNA was extracted from the macrophage cultures analyzed by real-time PCR for genes associated with AA-Mø. Gene expression is reported as relative gene expression compared with peritoneal macrophages exposed to medium only. All treatments were performed in triplicate, and data are presented as means ± SEM. Data are represented as a single experiment (*n* = 2). *B*, Peritoneal macrophages from C57BL/6 mice were treated, in triplicate, with medium only or rIL-4 (40 ng/ml), or exposed to *Ft* LVS (MOI = 5) for 48 h. Cells were harvested and stained for protein expression of FIZZ1, arginase-1, or mannose receptor by FACS analysis. Data presented are histograms from a single representative experiment (*n* = 4).

## Results

### Ft *LVS infection induces macrophage alternative activation markers*

*Ft* LVS induces a robust inflammatory response in mice and murine macrophages (20–24). Although this array of proinflammatory cytokines and mediators would be expected to control the intracellular bacterial burden, this potent, early inflammatory response fails to control bacterial replication and mice succumb to i.p. or i.v. infection. Therefore, we hypothesized that *Ft* LVS might evade destruction by reprogramming the differentiation state of infected macrophages, leading to a conversion of naive or CA-Mø to an AA-Mø phenotype. To test this hypothesis, homogenous populations of macophages were prepared from thioglycolate-treated WT C57BL/6J and TLR2$^{-/-}$ mice and infected with *Ft* LVS (MOI = 5) for up to 24 h; gene expression for markers associated with AA-Mø was measured by quantitative real-time PCR. *Ft* LVS infection of WT macrophages resulted in an up-regulation of mRNA for all AA-Mø markers examined: FIZZ1 and Ym1 (25–27), arginase-1, the macrophage mannose receptor, and anti-inflammatory cytokines IL-4, IL-13, and TGF-β, as well as GM-CSF, known to induce myeloid cell differentiation (28). In contrast, *Ft* LVS infection failed to stimulate macrophages from TLR2$^{-/-}$ mice to up-regulate mRNA expression of any of these AA-Mø markers (Fig. 1*A*). This finding indicates that TLR2 is

required for AA-Mø gene induction as well as proinflammatory gene expression (20). These findings were confirmed at the protein level: flow cytometric analysis of primary murine macrophages infected with *Ft* LVS revealed a significant (*p* < 0.01) up-regulation of FIZZ1 protein (Fig. 1*B*) that was comparable in magnitude to that observed by stimulation of macrophages with rIL-4, a known inducer of AA-Mø. Increased protein expression in macrophages infected with *Ft* LVS was also observed for arginase-1 and the mannose receptor (Fig. 1*B*).

### Ft *LVS induces alternative activation of macrophages for survival in the host*

CA-Mø and AA-Mø exhibit differential regulation of cytokines, chemokines, and surface receptors. CA-Mø produce high levels of IL-12 p70, TNF-α, and IL-1β, but low levels of IL-4, IL-13, and TGF-β (29, 30). Although our previous reports (18, 20) showed that *Ft* LVS induces a strong proinflammatory response in vivo and in vitro, our current data strongly suggest that after an initial proinflammatory or classical activation response, *Ft* LVS-infected macrophages are directed to differentiate into anti-inflammatory AA-Mø. To determine the proportion of cells positive for either CA-Mø or AA-Mø markers, two-color FACS analysis was performed for concurrent detection of the CA-Mø marker, IL-12 p70, and the AA-Mø marker, FIZZ1. Primary murine macrophages were treated with medium alone, rIFN-γ and LPS (to induce CA-Mø), or rIL-4 only (to induce AA-Mø), or infected with *Ft* LVS for 24 or 48 h. Cells were then fixed and stained concurrently for IL-12 p70 and FIZZ1 and analyzed by flow cytometry. Untreated macrophages expressed essentially no IL-12 p70 or FIZZ1 at 24 and 48 h (Fig. 2, *A* and *B*, *upper left panels*). Consistent with CA-Mø polarization, stimulation of macrophages by rIFN-γ and LPS induced expression of IL-12 p70 (18.6 and 24.3% positive at 24 and 48 h, respectively), but not FIZZ1 (Fig. 2, *A* and *B*, *upper right panels*), whereas alternative activation of macrophages by rIL-4 led to high levels of FIZZ1 (76 and 97.2% positive at 24 and 48 h, respectively), but little IL-12 p70 (Fig. 2, *A* and *B*, *lower left panels*). Macrophages infected with *Ft* LVS for 24 h show a similar phenotype to CA-Mø, with relatively higher expression of IL-12 p70 and lower FIZZ1 expression (Fig. 2*A*, *lower right panel*), but by 48 h postinfection, 82% (*p* < 0.001) of the macrophage population became positive for FIZZ1 (Fig. 2*B*, *lower right panel*), indicating that the majority of macrophages acquire an AA-Mø phenotype late in infection.

In vivo infection showed a similar trend of AA-Mø induction in peritoneal macrophages. Mice were injected i.p. with saline (mock) or *Ft* LVS (10,000 CFU) and sacrificed on day 3 postinfection. Peritoneal cells were harvested and fixed, as described above. The cells were stained for FIZZ1 (an AA-Mø marker) and F4/80 (a macrophage marker) and analyzed by two-color FACS. Mock-infected mice showed low expression of FIZZ1, but high expression of F4/80 (Fig. 3, *top panel*). Consistent with the *lower right panels* of Fig. 2, *A* and *B*, macrophages from mice infected with *Ft* LVS were alternatively activated, as evidenced by the significant (*p* < 0.001) increase in FIZZ1 expression (Fig. 3, *bottom panel*).

We next tested the hypothesis that *Ft* LVS survives within macrophages by inducing an AA-Mø phenotype. Primary murine macrophages were treated with medium only, rIFN-γ and LPS, or rIL-4 for 48 h. Cells were then infected with *Ft* LVS (MOI = 5) for 1 h, followed by 1 h of gentamicin treatment to kill extracellular bacteria. After gentamicin treatment, cells were either immediately lysed or incubated for an additional 24 h in fresh medium and then lysed. Intracellular *Ft* LVS in cell lysates was quantified by colony counts. At 1 h postinfection, all infected cells contained
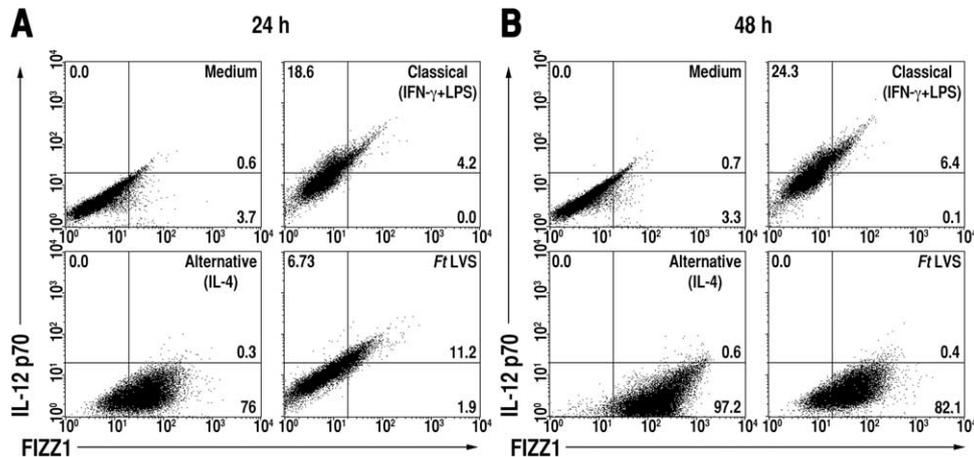
**FIGURE 2.** *Ft* LVS redirects macrophage differentiation from CA-Mø to AA-Mø over time. Peritoneal macrophages from C57BL/6 mice were treated with medium only, IFN-γ (20 ng/ml) plus LPS (10 ng/ml), or rIL-4 (40 ng/ml), or infected with *Ft* LVS (MOI = 5) for either 24 (*A*) or 48 (*B*) h. Cells were harvested and simultaneously stained for FIZZ1 and IL-12 p70. Protein expression was determined by FACS analysis. The numbers in the quadrants indicate the percentage of cells within that quadrant and have been rounded to the nearest one-tenth of a percent. All treatments were performed in triplicate, and data shown are from a single representative experiment (*n* = 3).

approximately the same number of *Ft* LVS (Fig. 4*A*), indicating that there were no differences in bacterial uptake among the three treatment groups. However, by 24 h postinfection (Fig. 4*A*),

CA-M$\phi$ exhibited significantly lower bacterial loads when compared with AA-M$\phi$ induced by either rIL-4 or infection only. Comparison of the relative gene expression for *Ft* LVS 16S rRNA, a surrogate marker for colony counts (18), yielded similar results (Fig. 4*B*). Most importantly, similar numbers of bacterial CFU were recovered from the *Ft*-infected medium- or IL-4-pretreated macrophages, supporting the hypothesis that *Ft*
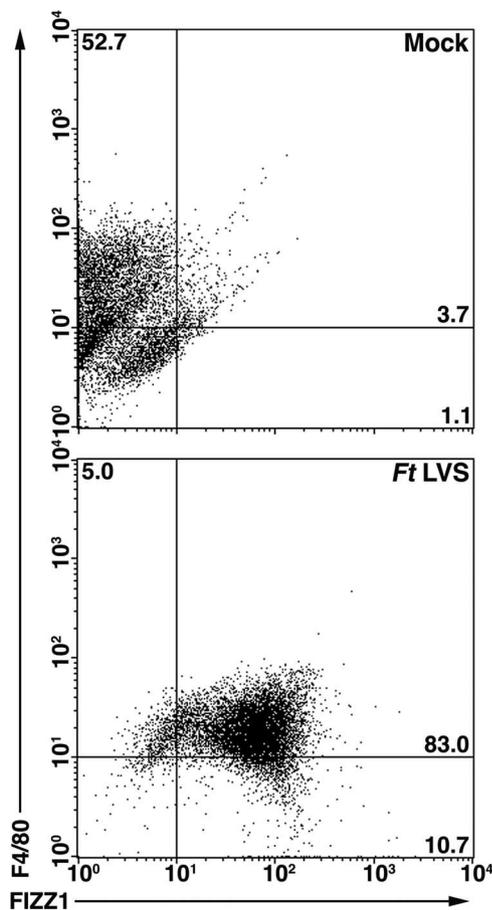


**FIGURE 3.** In vivo *Ft* LVS infection results in AA-M$\phi$. C57BL/6 mice were administered saline or *Ft* LVS (10,000 CFU) i.p. Three days later, the mice were sacrificed and peritoneal macrophages were harvested and simultaneously stained for FIZZ1 and F4/80. Protein expression was determined by FACS analysis. The numbers in the quadrants indicate the percentage of cells within that quadrant and have been rounded to the nearest one-tenth of a percent. Six mice were used for each treatment, and data shown are from a single representative mouse.
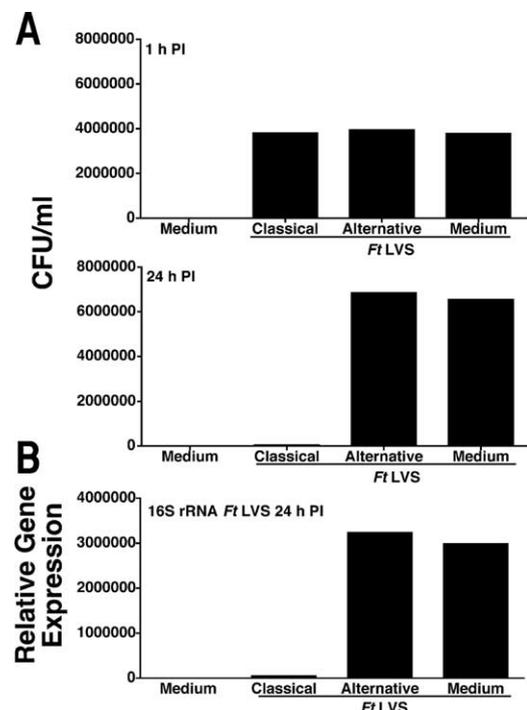


**FIGURE 4.** *Ft* LVS induction of AA-M$\phi$ results in intracellular survival. Peritoneal macrophages from C57BL/6 mice were treated with medium only, rIFN-γ (20 ng/ml) plus LPS (10 ng/ml), or rIL-4 (40 ng/ml) for 48 h. The cells were then infected with *Ft* LVS (MOI = 5) for either 1 or 24 h. Intracellular *Ft* LVS was determined by (*A*) colony counts from the plated lysates or (*B*) real-time PCR amplification of the *Ft* LVS 16S rRNA in the same samples as analyzed in *A*. All treatments were performed in triplicate, and data are presented as means ± SEM (the SEMs are too small to be seen). Data are derived from a single representative experiment (*n* = 4).

LVS induces macrophages to become alternatively activated for its own survival.

### Failure of Ft LVS to induce AA-Mφ results in prolonged inflammatory response

IL-4 and IL-13 use the same signaling pathways to induce AA-Mφ (31–33). Both cytokines signal through the IL-4Rα chain and induce the recruitment of the transcription factor STAT6 that translocates to the nucleus, where it activates transcription of STAT6-dependent genes. To assess whether Ft LVS is dependent upon this signaling pathway to induce AA-Mφ, thioglycollate-elicited macrophages derived from WT BALB/cByJ, IL-4Rα$^{-/-}$, or STAT6$^{-/-}$ mice were stimulated with medium alone or rIL-4 alone, or infected with Ft LVS (MOI = 5) for 24, 48, or 72 h, and AA-Mφ and CA-Mφ gene expression and protein production were measured. As expected, FIZZ1 gene expression increased upon stimulation of WT macrophages with either rIL-4 or Ft LVS (Fig. 5A, top panels). In contrast, macrophages deficient in either IL-4Rα or STAT6 failed to induce FIZZ1 mRNA in response to IL-4 stimulation; this was also observed upon infection of macrophages with Ft LVS. These results support the hypothesis that both IL-4Rα- and STAT6-dependent signaling are required for differentiation of AA-Mφ directly induced by Ft LVS infection. These data were further supported by flow cytometric analyses for detection of FIZZ1 protein in WT, IL-4Rα$^{-/-}$, and STAT6$^{-/-}$ primary macrophages (Fig. 5B, top graphs). In contrast to the failure of Ft LVS-infected IL-4Rα$^{-/-}$ and STAT6$^{-/-}$ macrophages to develop AA-Mφ markers, Ft LVS infection of IL-4Rα$^{-/-}$ and STAT6$^{-/-}$ macrophages prolonged expression of steady-state levels of IL-12 p35 and IL-12 p40 mRNA that resulted in increased levels of IL-12 p70 protein (Fig. 5B, bottom graphs). Similar enhancements in proinflammatory gene expression were observed for TNF-α and IL-1β (data not shown). These data indicate that the failure to induce STAT6 signaling via IL-4Rα leads to the prolongation of the CA-Mφ phenotype. Consistent with these observations, infected WT cells that expressed AA-Mφ markers were highly permissive for Ft LVS replication, whereas cells deficient in IL-4Rα or STAT6 had significantly lower bacterial loads, as measured by relative levels of Ft-specific 16S rRNA (Fig. 5C).

### Neutralization of IL-4 and IL-13 blocks Ft LVS-induced AA-Mφ activation and intracellular Ft replication

Because macrophages deficient in IL-4Rα or STAT6 failed to become alternatively activated, we hypothesized that neutralization of the cytokines required for AA-Mφ activation would be expected to yield similar results. To test this hypothesis, WT macrophages were treated without or with medium only, an isotype control IgG, anti-IL-4, and/or anti-IL-13 Abs. Treatment of cells with either anti-IL-4 or anti-IL-13 Abs, but not medium or the isotype control Ab, significantly ($p < 0.001$) inhibited expression of FIZZ1 in cells treated with rIL-4 or rIL-13, or infected with Ft LVS, and the presence of both neutralizing Abs completed inhibited FIZZ1 mRNA (data not shown) and protein expression in infected cells (Fig. 6A). Conversely, macrophages stimulated with rIL-4 or rIL-13, or infected with Ft LVS, subsequent to Ab treatment, exhibited a compensatory increase in the expression of IL-12 p40 mRNA (data not shown) and IL-12 p70 protein (Fig. 6A). This observation further supports the data obtained in macrophages derived from IL-4Rα$^{-/-}$ and STAT6$^{-/-}$ mice and indicates that macrophage-derived IL-4 and IL-13 drive AA-Mφ differentiation in Ft LVS-infected cells, and conversely, neutralization of IL-4 and IL-13 produced by the macrophages results in a CA-Mφ phenotype. Ab neutralization of either IL-4 or IL-13 resulted in significantly lower bacterial counts in comparison with cells treated with medium only
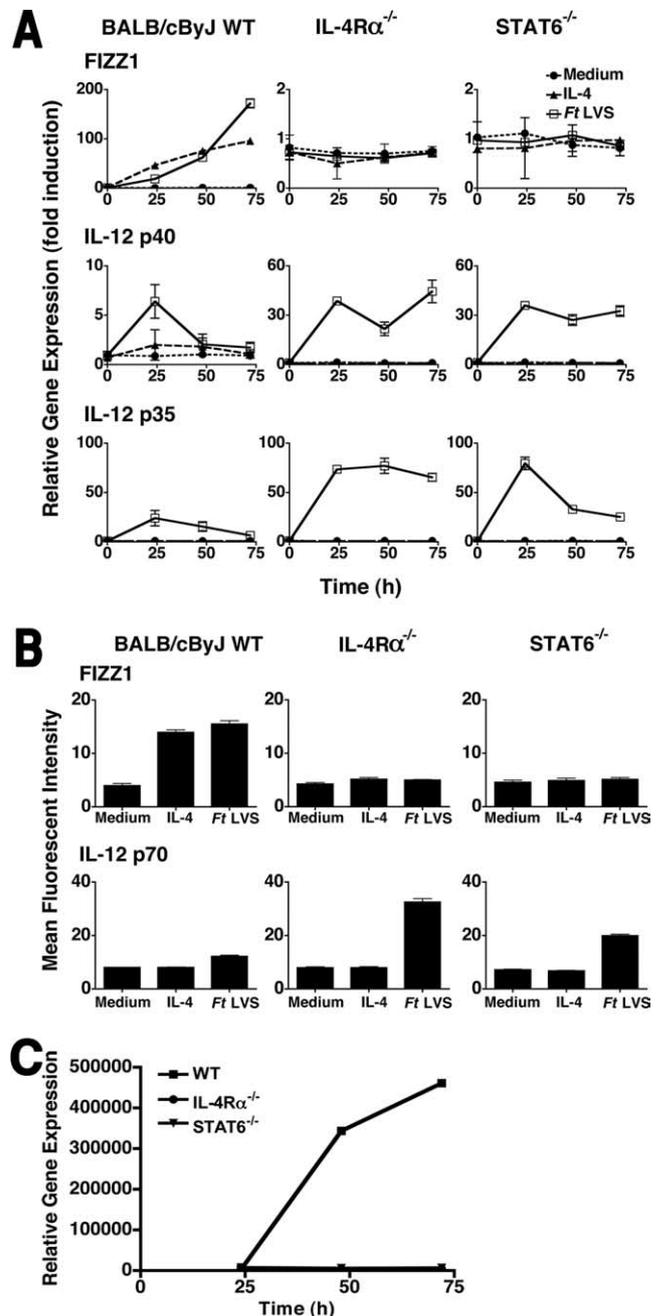


**FIGURE 5.** Failure to induce AA-Mφ results in prolonged inflammatory response by the host. A, Peritoneal macrophages from WT BALB/cByJ, IL-4Rα$^{-/-}$, or STAT6$^{-/-}$ mice were treated with medium only or rIL-4 (40 ng/ml), or exposed to Ft LVS (MOI = 5) for 0–72 h. Total RNA was extracted and analyzed by real-time PCR for genes associated with AA-Mφ and CA-Mφ. Gene expression is reported as relative gene expression normalized to untreated control samples. All treatments were performed in triplicate, and data are presented as means ± SEM. B, Peritoneal macrophages from WT BALB/cByJ, IL-4Rα$^{-/-}$, or STAT6$^{-/-}$ mice were treated with medium only or rIL-4 (40 ng/ml), or exposed to Ft LVS (MOI = 5) for 48 h. Cells were harvested and stained for expression of either FIZZ1 or IL-12 p70 protein and subjected to FACS analysis. All treatments were performed in triplicate, and data are presented as means ± SEM. Data are derived from a single representative experiment (n = 3). C, Real-time PCR for Ft LVS 16S rRNA was conducted using the same samples as A. Gene expression is reported as relative gene expression.

or with the isotype control Ab; however, neutralization of both cytokines resulted in nearly complete inhibition of bacterial replication (Fig. 6B). Finally, Ab-mediated neutralization of IL-4 and
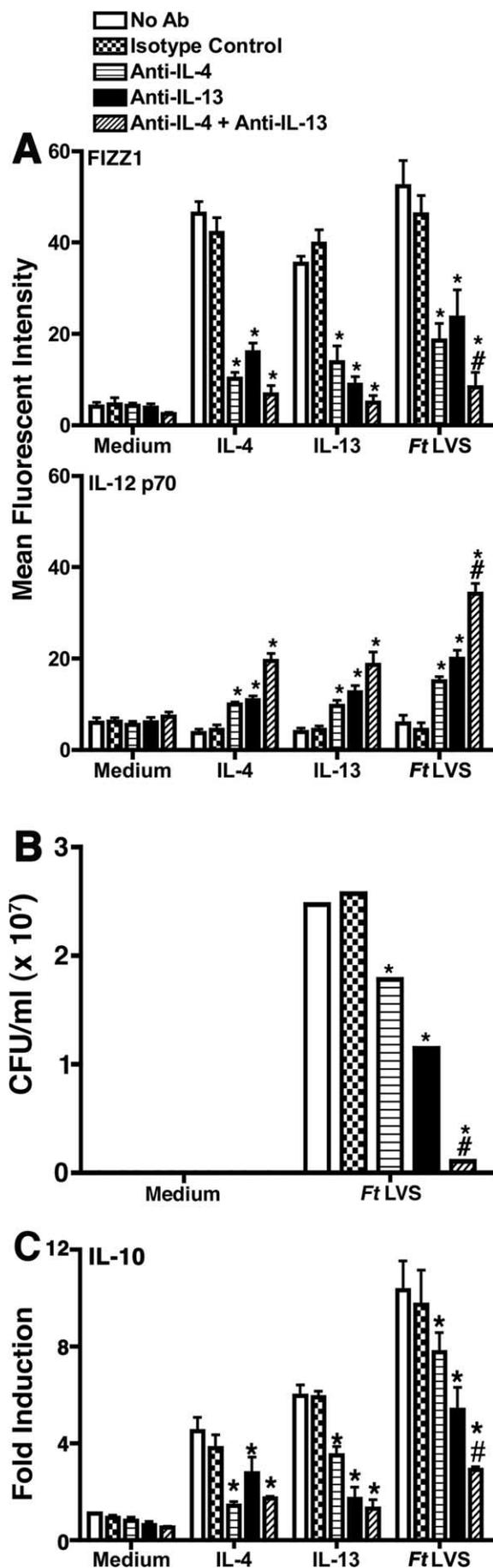
IL-13 led to a diminished capacity of *Ft* LVS-infected macrophages to produce IL-10 (Fig. 6*C*), thus mitigating the potential effects of this anti-inflammatory cytokine on induction proinflammatory cytokines such as IL-12.

### Ft *LVS infection of macrophages alters the cytokine milieu to favor development of the AA-Mø phenotype*

*Ft* LVS infection (MOI = 5) strongly induced both IL-4 and IL-13, with maximum production detected 72 h postinfection (Fig. 7*A*). *Ft* LVS also induced the production of both IL-1$\beta$ and IFN-$\gamma$; however, production of these cytokines peaked at 48 h, with a slight decrease by 72 h (Fig. 7*A*). These observations are consistent with the hypothesis that *Ft* LVS infection results in a cytokine milieu that favors the development of an AA-Mø phenotype.

CA-M$\phi$ produce iNOS that enables the cell to kill many intracellular parasites through the production of NO. AA-M$\phi$ counteract iNOS activity by producing arginase-1 that competes with iNOS for the same substrate, arginine, thereby allowing survival of intracellular microbes (34, 35). Arginase-1 and iNOS mRNA were also measured concurrently in *Ft*-infected macrophages. These two gene products were reciprocally regulated, as follows: arginase-1 mRNA expression was detectable by 24 h and increased to its peak at 48–72 h, whereas iNOS mRNA decreased from its peak at 24 h to background levels by 72 h (Fig. 7*B*). This correlates well with the kinetics of IL-4 and IL-13 production in *Ft*-infected macrophages (Fig. 7*A*) that facilitate differentiation of AA-M$\phi$, thus enabling the survival and replication of *Ft* LVS (Fig. 4).

### Failure to develop AA-Mø improves survival in vivo

Although AA-M$\phi$ induction is beneficial to *Ft*, we hypothesized that the failure to induce AA-M$\phi$ in vivo may protect against lethal infection with *Ft* by allowing CA-M$\phi$ to control *Ft* replication. To test this hypothesis, BALB/cByJ WT and IL-4R$\alpha^{-/-}$ mice were injected i.p. with saline (mock) or *Ft* LVS (10,000 CFU), and survival was monitored for 10 days. All of the mock-infected WT and IL-4R$\alpha^{-/-}$ mice survived; however, 5% of *Ft*-infected WT mice died on day 3, with 100% mortality by day 7. In contrast, 10% of IL-4R$\alpha^{-/-}$ mice challenged with *Ft* LVS died at day 4 postinfection, and 33% of the mice survived through day 10 ($p < 0.001$). These data suggest that the prevention of AA-M$\phi$ induction during *Ft* LVS infection increases survival.

**FIGURE 6.** Neutralization of IL-4 and IL-13 during *Ft* LVS infection results in prolonged macrophage classical activation. *A*, Peritoneal macrophages were cultured in medium, treated with rIL-4 (40 ng/ml) or rIL-13 (40 ng/ml), or infected with *Ft* LVS (MOI = 5). Cultures were also treated with medium only or an isotype-matched IgG control Ab or neutralizing anti-IL-4 (100 $\mu$g), anti-IL-13 (100 $\mu$g), or both Abs for 48 h. Cells were harvested and stained for expression of FIZZ1 and IL-12 p70 protein and subjected to FACS analysis. All treatments were performed in triplicate, and data are presented as means ± SEM. Data are represented as a single representative experiment (*n* = 3). *B*, Peritoneal macrophages were cultured in medium alone, or infected with *Ft* LVS. Cultures were treated with either medium only; an isotype control IgG; or neutralizing anti-IL-4, anti-IL-13, or both for 48 h. Intracellular *Ft* LVS was determined by colony counts from plated lysates. *C*, Peritoneal macrophages were treated the same as in *A*. Total RNA was extracted and analyzed by real-time PCR for IL-10. Gene expression is reported as fold induction normalized to untreated control samples. All treatments were performed in triplicate, and data are presented as means ± SEM. Statistical significance for comparison between one neutralizing Ab and control samples is represented as ∗, whereas statistical significance for treatment with both neutralizing Abs is represented as #.
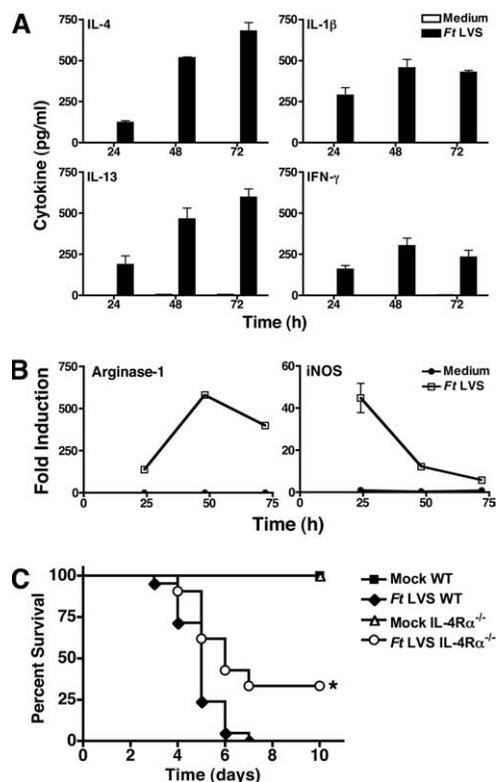
**FIGURE 7.** Production of IL-4 and IL-13 promotes the AA-Mφ phenotype while attenuating the CA-Mφ phenotype during *Ft* LVS infection. Peritoneal macrophages were treated with medium alone or infected with *Ft* LVS (MOI = 5). *A*, Cell supernatants were harvested at the indicated times and analyzed by ELISA for production of IL-4, IL-13, IL-1β, and IFN-γ. *B*, Total RNA was extracted and analyzed by real-time PCR for arginase-1 and iNOS mRNA. Gene expression is reported as fold induction normalized to untreated control samples. All treatments were performed in triplicate, and data are presented as means ± SEM of four individual experiments. *C*, BALB/cByJ WT and IL-4Rα$^{-/-}$ mice were i.p. inoculated with either saline or *Ft* LVS (10,000 CFU). Survival was monitored for up to 10 days. Data are pooled from four separate experiments with four to five mice/treatment/experiment. Statistical significance for comparison between *Ft* LVS-infected WT and *Ft* LVS-infected IL-4Rα$^{-/-}$ mice is represented as ∗ ($p < 0.001$).

## Discussion

Macrophage activation is central for controlling the replication of intracellular pathogens such as *Francisella*. However, despite the fact that *Ft* LVS initially induces a robust inflammatory response, the mice nonetheless die, usually 4–5 days after i.p. infection. Our observations suggest that *Ft* may avoid macrophage-mediated killing by directly altering the macrophage differentiation state from one that is classically activated and highly microbicidal to one that is alternatively activated, and therefore, permissive to microbial survival and replication at the expense of the host. *Ft* LVS infection of murine macrophages induced TLR2-dependent expression of AA-Mφ markers (e.g., FIZZ1, Arg-1, Ym1, SR) that followed expression of CA-Mφ markers (e.g., IL-12, iNOS).

The innate immune response to *Ft* is complex and involves multiple pattern recognition receptors. Despite the fact that every single host gene that we have examined to date requires TLR2 signaling for its expression (e.g., TNF-α, IL-1β, IFN-γ, IFN-β, iNOS, IL-6, KC, MCP-1, RANTES, and others (18, 20), in addition to those shown in Fig. 1), we have more recently observed that expression of a subset of these TLR2-dependent genes additionally requires the organism to escape from the phagosome, implying that

cytosolic sensing and signaling are also necessary. Using an *Ft* LVS mutant that is unable to escape from the phagosome, in conjunction with IFN-β-deficient macrophages, we further demonstrated that those genes whose induction required both TLR2 and cytosolic sensing were also IFN-β dependent. In addition to this uncharacterized cytosolic sensor, activation of the inflammasome, as has been reported for the response to *Francisella novicida* (36), is also necessary for secretion of IL-1β in response to *Ft* LVS, despite the fact that induction of IL-1β mRNA is solely TLR2 dependent (37). In preliminary experiments, we have observed that TLR2 signaling by synthetic triacylated lipopeptide is also not sufficient for induction of all AA-Mφ markers (e.g., FIZZ-1, arginase-1, and mannose; data not shown). Experiments are ongoing to identify subsets of genes that encode AA-Mφ markers that can be induced by *Ft* infection solely through TLR2 vs those that require additional signaling pathways for their expression.

Induction of AA-Mφ by treatment of macrophages with rIL-4 or rIL-13, or by infection with *Ft* LVS promoted replication of intracellular *Ft*, in contrast to CA-Mφ derived by IFN-γ plus LPS treatment. In vivo, infection of mice with *Ft* LVS also led to induction of AA-Mφ. Multiple groups have shown the importance of IFNs in the control of *Francisella* infection (36–39). IFN-γ$^{-/-}$ mice have been shown to be highly susceptible to otherwise sublethal doses of *Ft* (40). Cole et al. (37) showed that even though endogenous IFN-β contributed to control intracellular *Ft* replication within the first 24 h of infection, *Ft* was able to overcome this inflammatory signal and replicate over time. Anthony et al. (41) first showed that pretreatment of cells with IL-4 or GM-CSF permitted *Ft* replication. Our data expand and clarify these previous findings by showing that the cytokine treatments impact the antimicrobial state of the macrophages through the induction of either classical or alternative activation. We have further shown that infection of the macrophages by *Ft* only is sufficient to elicit an AA-Mφ phenotype in the absence of exogenous IL-4. The concept of macrophage plasticity has long since been recognized, and Gratchev et al. (42) showed that the cytokine milieu facilitates the macrophages to change phenotypes. Our data clearly demonstrate that *Ft* LVS-infected macrophages change phenotypes, as evidenced by a switch from a CA-Mφ-producing and -secreting IL-12 p70 to AA-Mφs that produce multiple markers associated with alternative activation, e.g., FIZZ1, arginase-1, etc. (Fig. 2).

IL-4 and IL-13 drive many immune and anti-inflammatory responses that are associated with a Th2 response (31, 42). The IL-4Rα chain is essential for signaling by both cytokines. *Ft* LVS failed to induce alternative activation in IL-4Rα$^{-/-}$ or STAT6$^{-/-}$ macrophages and prolonged expression of CA-Mφ activation markers, indicating that IL-4- and/or IL-13-mediated signaling underlie AA-Mφ differentiation in the *Ft*-infected cells. Neutralization of either IL-4 or IL-13 blunted AA-Mφ activation during *Ft* LVS infection, but prolonged CA-Mφ activation. Because these same events were seen in both the RAW 267.4 cell line and macrophages derived from Rag2$^{-/-}$ mice (data not shown), our neutralization data strongly support the conclusion that the macrophages are, in fact, the source of the IL-4 and IL-13 and that both cytokines contribute to AA-Mφ differentiation in infected macrophages. Leiby et al. (39) previously reported that neutralization of IL-4 with a mAb enhanced survival of mice i.p. infected with *Ft* LVS compared with control mice. Although they reported that "the differences were not statistically significant," the potential contribution of the compensatory activity of IL-13 induced during infection, which also drives AA-Mφ activation (Fig. 6), was not examined in their studies. Our data expand upon their findings by showing that neutralization of both IL-4 and IL-13 is required to

limit intracellular replication of *Ft* maximally, and that mice deficient in IL-4Rα, which fail to differentiate AA-Mφ, have significantly higher survival rates ($p < 0.001$) compared with WT mice (Fig. 7*C*).

Apart from the fact that *Ft* LVS infection also greatly increases intracellular replication (Fig. 4), up-regulation of mannose receptor expression (Fig. 2) may also contribute to the dissemination of this pathogen. Schulert and Allen (43) reported that the mannose receptor mediates uptake of *Ft*, and more efficiently so in the presence of complement. Thus, increased uptake of *Ft* via mannose receptors may facilitate spread from cell to cell for survival as well as replication. IL-4 production and secretion by infected cells might be envisioned to act on adjacent cells to increase mannose receptor expression, thus allowing more efficient uptake of *Ft* once it escapes a neighboring cell. This mechanism might also apply to the observed increase in scavenger receptors in infected macrophages that have also been shown to enhance uptake of this pathogen (44).

In addition to wound repair and regulation of the immune response (8), AA-Mφ provide host resistance to certain parasites (45, 46). AA-Mφ have developed multiple functions for resistance, such as acidic mammalian chitinase that induces eotaxin and MCP-1 during infection with *Heligosomoides polygyrus*, and in *Schistosoma mansoni* egg granulomas (47–49). FIZZ2, which is expressed exclusively in the gastrointestinal tract and induced during infection with *Trichuris muris*, interferes with parasite chemotaxis toward host tissues (50, 51). In addition to its role in angiogenesis, Ym1 acts as a chemotactic factor for eosinophils (52), which play an essential role in antihelminth responses (53). The outcome of parasitic infections is also dependent on both the parasite as well as the susceptibility of the host. C57BL/6 mice are resistant to infection with *Trypanosoma congolense* because they have a predominantly CA-Mφ-type environment during the early stages of infection that allows NO, generated by iNOS, to control growth of the parasite; however, AA-Mφs develop later in infection and contribute to wound repair. BALB/c mice infected with *T. congolense* have a simultaneous mix of CA-Mφ and AA-Mφ during the early stages of infection, resulting in uncontrolled parasite growth (54). Our data show *Ft* LVS is able to strongly induce AA-Mφ in both C57BL/6 as well as BALB/c backgrounds. Both strains are susceptible to *Ft* and typically succumb to infection within 4–7 days (Fig. 7*C*) (55). However, the inability of IL-4Rα$^{-/-}$ mice to develop AA-Mφ correlates with increased survival, presumably due to prolonged activation of the CA-Mφ phenotype and control of intracellular replication.

Collectively, our data support a novel paradigm for the host response to the intracellular bacterium, *Ft* LVS: initially, the host interaction with *Ft* LVS results in a brisk up-regulation of proinflammatory genes that is generally associated with development of CA-Mφ. With time, however, the organism ultimately redirects macrophage differentiation to that of an AA-Mφ phenotype by inducing the production of IL-4 and IL-13 that ultimately facilitates intracellular survival of the organism with a concomitant mitigation of the proinflammatory response. Thus, the balance of environmental signals induced during *Ft* infection of macrophages results in a constantly evolving host immune response that can be either beneficial or detrimental to the host and pathogen alike. This organism clearly takes advantage of the plasticity of macrophages to differentiate for its own survival and suggest new approaches for therapeutic intervention.

## Acknowledgments

## Disclosures

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