Spirochete antigens persist near cartilage after murine Lyme borreliosis therapy

Linda K. Bockenstedt, David G. Gonzalez, Ann M. Haberman, and Alexia A. Belperron

An enigmatic feature of Lyme disease is the slow resolution of musculoskeletal symptoms that can continue after treatment, with some patients developing an inflammatory arthritis that becomes refractory to antibiotic therapy. Using intravital microscopy and the mouse model of Lyme borreliosis, we observed that Borrelia burgdorferi antigens, but not infectious spirochetes, can remain adjacent to cartilage for extended periods after antibiotic treatment. B. burgdorferi was not recovered by culture or xenodiagnosis with ticks after antibiotic treatment of WT mice and all but one of the immunodeficient mice with heightened pathogen burden due to impaired TLR responsiveness. Amorphous GFP* deposits were visualized by intravital microscopy in the entheses of antibiotic-treated mice infected with GFP-expressing spirochetes and on the ear cartilage surface in sites where immunofluorescence staining detected spirochete antigens. Naïve mice were not infected by tissue transplants from antibiotic-treated mice even though transplants contained spirochete DNA. Tissue homogenates from antibiotic-treated mice induced IgG reactive with B. burgdorferi antigens after immunization of naïve mice and stimulated TNF-α production from macrophages in vitro. This is the first direct demonstration that inflammatory B. burgdorferi components can persist near cartilaginous tissue after treatment for Lyme disease. We propose that these deposits could contribute to the development of antibiotic-refractory Lyme arthritis.

Introduction

Lyme disease is an emerging zoonotic infection caused by the Ixodes tick–transmitted spirochete Borrelia burgdorferi (1). Spirochetes deposited in the skin during tick feeding can cause the localized skin rash erythema migrans (EM) or disseminate to cause disease mainly involving other areas of the skin and/or the heart, joints, and nervous system. Although the disease is responsive to antibiotics, up to 25% of patients treated early in the course of infection can experience protracted musculoskeletal symptoms of unclear etiology (2). Patients who present with the late-stage manifestation of arthritis may evolve persistent joint inflammation that no longer responds to antibiotics (3). Although ongoing infection is considered an unlikely explanation for persistent symptoms or disease, it cannot be definitively excluded because B. burgdorferi is difficult to detect by culture except in early infection when EM is present (4). The pathogenic mechanisms underlying the delay in symptom resolution after treatment for early disease and the persistence of objective late signs such as arthritis are incompletely defined.

The mouse model of Lyme borreliosis has provided a useful system for studying B. burgdorferi infection in mammals (5). All laboratory mice are susceptible to infection and develop subacute arthritis and myocarditis similar to that of humans infected with B. burgdorferi. Disease severity is mouse strain dependent, with C3H mice more susceptible than C57BL/6 (B6) mice. Strain differences in disease susceptibility are influenced by both hematopoietic and nonhematopoietic factors and are preserved in SCID mice that lack functional T and B cells (6, 7). Effective pathogen control requires both specific Ab production and phagocyte recognition of B. burgdorferi via TLRs, as demonstrated by the markedly elevated pathogen burdens in mice deficient in B cells, in TLR2, or in the TLR intracellular adaptor molecule myeloid differentiation antigen 88 (MyD88) (8–11). Adaptive immunity is required for resolution of joint inflammation, which usually occurs within 45–60 days of infection even though spirochetes persist in tissues throughout the life span of the mouse (5, 8, 10).

Our laboratory and others have used the C3H mouse model of Lyme borreliosis to evaluate whether viable B. burgdorferi can persist after antibiotic treatment (12–14). Using xenodiagnosis with ticks, we showed that spirochetes could be detected for up to 3 months after treatment with either ceftriaxone or doxycycline for B. burgdorferi infection introduced by tick bite (14). Spirochetes acquired by ticks that were recovered from antibiotic-treated mice lacked genes on plasmids lp25 and lp28-1 associated with infectivity, suggesting that they were attenuated. Nymphs derived from a group of larval ticks that tested positive for B. burgdorferi plasmid DNA after feeding on antibiotic-treated mice could not transmit infection to naive mice. Spirochetes were not cultured from antibiotic-treated mice, even after immunosuppression with cortisone, but low levels of B. burgdorferi DNA were detected by PCR in multiple tissues. A subsequent study examined efficacy of ceftriaxone administered to C3H mice 3 weeks or 4 months after B. burgdorferi infection introduced by needle inoculation (13). Some larval ticks used for xenodiagnosis acquired spirochete DNA from antibiotic-treated mice and, after molting to nymphs, transmitted B. burgdorferi DNA to a small proportion of SCID mice used as blood meal hosts. Rare spirochete-like forms could be detected by immunohistochemistry in connective tissue of the heart and tibiotarsal joints of antibiotic-treated mice, but spirochetes could not be cultured from these tissues. Taken together, these findings suggest that attenuated, noncultivable spirochetes could persist at low levels in mice after antibiotic treatment for disseminated B. burgdorferi infection.

In this study, we used the elevated pathogen burden in B. burgdorferi–infected Myd88−/− mice and intravital 2-photon microscopy to further investigate spirochetes that may persist after antibiotic therapy. Our results show that infectious spirochetes are rapidly eliminated after institution of antibiotics, but inflammatory B. burgdorferi antigens persist adjacent to cartilage and in the entheses.
We believe this is the first direct demonstration that inflammatory B. burgdorferi components can persist for extended periods after resolution of B. burgdorferi infection, and it raises the possibility that these spirochete remnants contribute to the pathogenesis of antibiotic-refractory Lyme arthritis.

Results

B. burgdorferi DNA can be detected in B6 Myd88<sup>−/−</sup>, but not WT, mice after treatment with doxycycline. To determine whether persistence of B. burgdorferi observed in antibiotic-treated C3H mice could be demonstrated in the disease-resistant B6 background, we infected cohorts of B6 WT and Myd88<sup>−/−</sup> mice with B. burgdorferi strain N40 (BbN40) by tick bite. Infection was confirmed by B. burgdorferi immunoblot (data not shown). At 30 days after infection, mice were treated with doxycycline administered continuously in drinking water to sustain serum drug levels above the mean inhibitory concentration (MIC) for B. burgdorferi during the treatment period (see Methods). Xenodiagnosis with ticks was performed at 1, 4, and 7 weeks after the last day of treatment. At the end of the 15-week experimental period (9 weeks after completion of antibiotics), no spirochetes were detected in doxycycline-treated WT mice. As expected, sham-treated mice showed evidence of spirochetes by all parameters examined, including culture, whereas uninfected controls tested negative.

**Real-time imaging of B. burgdorferi in Myd88<sup>−/−</sup> mice reveals rapid spirochete elimination after antibiotic therapy.** To begin to assess whether DNA in tissue could represent a subpopulation of live spirochetes that remained after antibiotic therapy, we used intravital 2-photon microscopy to image in real time the immediate effects of antibiotics on spirochetes in living mice. Myd88<sup>−/−</sup> mice infected for 21 days with a transformant of B. burgdorferi strain 297 expressing a GFP reporter protein under the control of the flaB promoter (Bb914) (15) were treated with a 5-day course of ceftriaxone or were sham treated. Ceftriaxone was chosen for its bactericidal effects, which could lead to changes in B. burgdorferi more rapidly than doxycycline, a bacteriostatic agent (16). We visualized numerous highly motile spirochetes in the dermis of sham-treated mice (Figure 1A and Supplemental Video 1; supplemental material available online with this article; doi:10.1172/JCI58813DS1); spirochetes were fewer in number and moved more slowly in the calcaneal tendon at all time points analyzed (Figure 1C and Supplemental Video 2). Motility patterns of spirochetes within a field varied, with the majority of spirochetes moving back and forth in tracks along collagen fibers and others.
exhibiting more directional translocation. Less frequently, spirochetes undulated as a traveling wave without positional displacement. We have observed similar patterns of motility in infected WT mice (17). At 24 hours after instituting antibiotics, the population of spirochetes had diminished dramatically in the dermis and tendons (Figure 1, B and D). Remaining spirochetes in the dermis but not the tendon appeared motile, and continuous imaging captured 2 events in which these spirochetes rapidly transformed into spherical objects. In one case, a motile spirochete abruptly stopped and became spherical within the 30-minute observation period (Figure 2 and Supplemental Video 3). In the second case, a spirochete appeared to be tethered at one end and initially wriggling in place (Supplemental Video 5). Continuous imaging revealed that the spirochete flexed once and then rapidly converted into a spherical shape. After imaging at the 1-day time point, we were unable to visualize spirochetes in the skin or tendons of ceftriaxone-treated mice by this technique.

**Figure 2** Image sequences of Bb914 changing from an elongated spirochete to a spherical form. The spirochete image is labeled with an arrow in Figure 1B. (A–D) Images in the xy plane from Supplemental Video 3 depicting changes in morphology of Bb914 over time; arrow (D) shows the spherical form. (E–H) Images from Supplemental Video 4 showing the same spirochete (arrows) imaged in A–D in the yz plane. The blue channel has been suppressed. Scale bars: 30 μm.

*Spirochete antigens can be detected adjacent to ear cartilage in antibiotic-treated Myd88−/− mice.* At the end of the experimental period, all sham-treated mice tested positive for *B. burgdorferi* by both culture of tissues and direct immunofluorescence staining using FITC-conjugated anti-*B. burgdorferi* Ab (DFA) of ear-skin cryosections, which revealed spirochetes throughout the dermis (Table 2 and Figure 3A). Although spirochetes were not cultured from any of the ceftriaxone-treated mice, DFA detected spirochete antigens in the deep dermis adjacent to ear cartilage (Figure 3, B and C); their location was deeper than the level at which intravital imaging had been performed. Residual spirochete antigens also were seen by DFA of ear sections from the BbN40-infected Myd88−/− mice treated with doxycycline described above (Figure 3E). None of the antibiotic-treated tissues contained cultivable spirochetes. Negative cultures suggested that if immunofluorescence staining represented viable spirochetes, the residual organisms were deficient in their ability to multiply when transferred to the culture environment.

**Live imaging reveals antigen deposits but not motile spirochetes adjacent to cartilage of Myd88−/− mice after doxycycline treatment for *B. burgdorferi* infection.** To further assess the significance of DFA detection of spirochete antigens, we conducted a third experiment in which Bb914-infected Myd88−/− mice were evaluated between 2 and 10 weeks after completion of a 1-month course of oral doxycycline (Table 3). Xenodiagnosis with ticks was performed prior to mouse sacrifice, and *B. burgdorferi* DNA was amplified from some ticks that had fed on doxycycline-treated mice. DFA of tick midgut contents, however, did not reveal spirochetes in ticks that acquired blood meals from antibiotic-treated mice, nor were spirochetes cultured from the ticks; only sham-treated mice tested positive for viable *B. burgdorferi* by these modalities (Table 3). As in our previous experiment with BbN40-infected mice, spirochetes could not be cultured from any of the antibiotic-treated mice, but residual spirochete antigens were detected by DFA adjacent to ear cartilage at all time points examined. Using live imaging parameters to include the level of the ear cartilage, we observed that sham-treated mice had not only motile spirochetes in the dermis extending to the cartilage surface, but also large amorphous deposits of nonmotile fluorescent material at the dermal-cartilage interface (Figure 4A and Supplemental Video 6). Similar GFP+ deposits, but not morphologically intact or motile spirochetes, were visualized adjacent to ear cartilage in all doxycycline-treated mice (Figure 4B).

### Table 2

Ceftriaxone eliminates cultivable Bb914 from *Myd88−/−* mice

<table>
<thead>
<tr>
<th>Mouse group</th>
<th>Culture of mouse tissue</th>
<th>DPA of tissues for <em>B. burgdorferi</em></th>
<th>Multiphoton imaging</th>
</tr>
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<tbody>
<tr>
<td></td>
<td>Ear skin</td>
<td>Heart</td>
<td>Tibiotarsal joint</td>
</tr>
<tr>
<td><strong>Sham</strong></td>
<td>3/3</td>
<td>2/3</td>
<td>3/3</td>
</tr>
<tr>
<td>Ceftriaxone</td>
<td>0/6&lt;sup&gt;D&lt;/sup&gt;</td>
<td>0/6</td>
<td>0/6&lt;sup&gt;D&lt;/sup&gt;</td>
</tr>
</tbody>
</table>

<sup>A</sup>Results are reported as the number of mice that had positive cultures for *B. burgdorferi* over the total number of mice examined. <sup>B</sup>Results are from cultures of urinary bladders, ear skin, and tibiotarsal joints. <sup>C</sup>The mice testing positive for *B. burgdorferi* by multiphoton imaging were evaluated at day 1 after ceftriaxone therapy, and tissues harvested for culture were collected within 1 hour after administration of a third dose of ceftriaxone. <sup>D</sup>Differences between sham- and ceftriaxone-treated mouse groups were statistically significant (*P* = 0.019, Fisher’s exact test, for all comparisons). <sup>E</sup>Differences between sham- and ceftriaxone-treated mouse groups were statistically significant (*P* = 0.0476, Fisher’s exact test).
Spirochete antigens can be detected in joints of antibiotic-treated C3H Myd88\(^{-/-}\) mice. Because B. burgdorferi DNA was detected in the joints of antibiotic-treated mice, we postulated that this site could also contain deposits of B. burgdorferi antigens. Subsequent to the above studies, we developed a technique to image the articular surfaces and adjacent entheses of the patellofemoral joint. Strikingly, was only found at imaging depths corresponding to the regions viewed sagitally on ear-skin cryosections that tested positive by DFA (Figure 3). As further evidence that infectious spirochetes had been eliminated by antibiotic treatment, only transplanted ear skin from sham-treated mice introduced infection into naive Myd88\(^{-/-}\) mice, as documented by IgG seroconversion to multiple B. burgdorferi antigens and culture (Supplemental Figure 1A). In contrast, similar transplants from antibiotic-treated mice induced reactivity to only 1 or 2 B. burgdorferi antigens, and spirochetes were not cultured from recipient mice (Supplemental Figure 1, A and B). This pattern of reactivity was reproduced when mice were immunized with homogenates of sham-treated mouse ears (Supplemental Figure 1C).

Discussion
This study demonstrates that antibiotics (ceftiraxone or doxycycline) eliminate infectious B. burgdorferi from mice with disseminated infection, even when the host is immunocompromised and initially has a high pathogen burden. Through use of Myd88\(^{-/-}\) mice and intravital microscopy, we have shown that ceftiraxone rapidly (within 24 hours) reduces pathogen burden in the skin, a preferential site of B. burgdorferi infection, and that both ceftiraxone and doxycycline eliminate infectious spirochetes as assessed by real-time qPCR from mouse tissues.

Table 3

<table>
<thead>
<tr>
<th>Mouse group</th>
<th>Culture of mouse tissue(^{A,B})</th>
<th>ospA PCR from mouse tissues(^{A})</th>
<th>DFA of mouse tissues(^{A})</th>
<th>Culture of xenodiagnostic ticks(^{A,C})</th>
<th>ospA PCR from xenodiagnostic ticks(^{A})</th>
<th>DFA of tick midguts(^{A})</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sham</td>
<td>4/4</td>
<td>4/4</td>
<td>4/4</td>
<td>4/4</td>
<td>5/5</td>
<td>5/5</td>
</tr>
<tr>
<td>Doxycycline</td>
<td>0/5(^{b})</td>
<td>3/8(^{b})</td>
<td>9/9(^{b})</td>
<td>0/7</td>
<td>7/8</td>
<td>0/7</td>
</tr>
</tbody>
</table>

\(^{A}\)Results are reported as the number of mice that had positive cultures for B. burgdorferi over the number of mice examined. \(^{B}\)Results are from cultures of urinary bladders, ear skin, and one tibiotarsal joint. One mouse from each group died during tick feeding; ticks were retrieved from the sham-treated mouse for analysis. \(^{C}\)Ticks placed on mice 2 weeks after sham or doxycycline treatment were not analyzed by culture, PCR, or DFA for B. burgdorferi. \(^{b}\)Results from doxycycline-treated mice were significantly different from those of sham-treated mice (Fisher’s exact test, \(P = 0.0014\) for comparison of mouse tissue culture; \(P = 0.0256\) for comparison of B. burgdorferi DNA in mouse tissues; \(P = 0.0013\) for comparison of DFA of tick midguts). \(^{5}\)Ear skin of doxycycline-treated mice exhibited immunofluorescence staining adjacent to ear cartilage, but not in the dermal connective tissue.
time imaging, culture, and xenodiagnosis with ticks. Our studies also provide what we believe is the first direct evidence that spirochete remnants can persist adjacent to ear cartilage and within joint entheses. Notably, these remnants contain immunogenic material that can induce IgG responses to B. burgdorferi antigens in naive recipients and stimulate macrophages to produce TNF-α in vitro.

Previous studies examining antibiotic efficacy in murine Lyme borreliosis used the BbN40 strain (12–14). Here, we found that antibiotics were effective against 2 strains of B. burgdorferi (BbN40 and Bb914) that represent genotypes that differ in their capacity for dissemination and pathogenicity (18, 19). Improved efficacy of doxycycline in WT mice as compared with that found in previous studies may be due to its administration in drinking water, which optimizes the time interval for which serum drug levels remain above the MIC for B. burgdorferi (20). With the exception of a single doxycycline-treated Myd88−− mouse, cultivable spirochetes were eliminated from all antibiotic-treated Myd88−− mice infected with either B. burgdorferi strain. Presence of infection in 1 of a total of 21 Myd88−− mice was easily demonstrated by several modalities, including culture of mouse tissues and the ticks used for xenodiagnosis. This contrasts with our earlier report, in which intact organisms were documented only by xenodiagnosis in a subset of doxycycline-treated animals (14). Administration of doxycycline by gavage twice daily results in periods in which serum drug levels fall below the MIC for BbN40, which may allow for the persistence of attenuated, noncultivable organisms (21). The failure of antibiotics to eliminate B. burgdorferi from a single Myd88−− mouse may have been due to a drinking pattern that led to inconsistent doxycycline levels.

The significance of B. burgdorferi DNA in xenodiagnostical ticks and in mouse tissues after antibiotic therapy is unclear. The nymphs used in the study were from a specific pathogen-free colony derived from larvae that fed on uninfected laboratory mice; the colony was routinely tested for B. burgdorferi by PCR. In humans, spirochete DNA has been detected by PCR for up to 9 months after treatment for Lyme arthritis, but its presence does not correlate with relapse or duration of arthritis (22). Microbial DNA can persist in mammalian tissues for extended periods (years) when sequestered in cellular debris even though the microbe itself is no longer viable (23, 24). Some B. burgdorferi DNA could remain intact if it is sequestered in cellular debris such as the GFP deposits. Alternatively, spirochete DNA could represent a minor subpopulation of B. burgdorferi that is not killed by the antibiotic treatment.

It has been proposed that B. burgdorferi persists in the human host by transforming into cysts (25) because in vitro studies show that it can alter its morphology under stress conditions such as nutrient deprivation (26). Our real-time imaging captured spirochetes changing into spherical forms, but indirect evidence indicates that they are not bacterial cysts. The image sequences bear substantial similarity to in vitro observations of spirochete ingestion by phagocytes in both morphologic changes (Supplemental Video 6) and the time course for entry of ingested B. burgdorferi into phagolysosomes (∼20 minutes [ref. 10] vs. 25 minutes in Supplemental Videos 3 and 5). Formation of true bacterial cysts and endospores involves programmatic structural changes that occur over hours to days, not minutes, as we observed (27).

For other bacterial species, phenotypic variants known as “persisters” have been identified based on growth patterns that differ from the bulk of the population and their tolerance to antibiotics (28–30). The study of persisters of biologic relevance is difficult because they are genetically identical to other cells in the population and have a temporary phenotype induced by environmental stress such as antibiotics. In the context of biofilms, this type of persister is thought to contribute to infection relapse when the antibiotic is removed, as these revert to the growth characteristics and antibiotic susceptibility of the original population (28). If the spirochete DNA detected in mouse tissues after antibiotics represents a subpopulation of live spirochetes, this population differs in several ways from persisters studied in other bacterial species. Residual spirochetes, if present, did not revert to cultivable microorganisms once conditions become more favorable to replication, as would be expected to occur when tissues were placed in culture medium. Reversion of a dormant population to an active phenotype was not detected by real-time imaging; motile spirochetes were not seen in tissues of immunodeficient BALB/c mice at any time point after completion of antibiotics. It is unlikely that spirochetes lost GFP expression because the gfp expression cassette in Bb914 is inserted into cp26, a plasmid ubiquitously present in spirochetes and required for viability (15, 31). Although a mutation that would reduce expression could occur in either gfp or its flaB promoter, it is improbable that spirochetes with such a mutation would also be the only remaining population selectively resistant to antibiotics. If residual amorphous debris noted in the enthesis contained morphologically altered but viable spirochetes, as has been described for an rpoS mutant in the tick midgut (32), such forms did not resume the usual spirochete morphology and replicate when placed in culture medium. Importantly, both xenodiagnosis and tissue transplants demonstrated that any spirochetes remaining after antibiotics could not establish infection in the tick or a new mammalian host, even when the recipient mammal was immunocompromised. This is consistent with our previous study.

Figure 4
Amorphous GFP deposits on the dermal-cartilage interface of Bb914-infected mice. (A) Multiphoton image of typical GFP+ spirochete (white arrow and inset) and amorphous GFP deposits (yellow arrow) on the surface of ear cartilage (red) of Bb914-infected mice (see also Supplemental Video 6). (B) Multiphoton image of a doxycycline-treated mouse ear showing GFP deposits (yellow arrows), but not typical spirochete forms (see also inset). (C) Multiphoton image of the dermal-cartilage interface of an uninfected mouse. The green channel was enhanced to show absence of fluorescent green deposits. Images were acquired using a ×20 objective and optically zoomed to a scanned field of view of 300 × 300 μm in A and C and 150 × 150 μm in B. The objects delineated by arrows in A and B are enlarged 3-fold and 1.7-fold, respectively, in the corresponding insets. Scale bars: 30 μm.
in which immunosuppression with corticosteroids did not result in positive xenodiagnoses or recrudescence of *B. burgdorferi* infection in WT mice after antibiotic treatment (14).

A recent study in the primate model of Lyme borreliosis reported detection of spirochetes after antibiotic therapy, a conclusion that should be interpreted with caution (33). Subtherapeutic antibiotic levels were documented in experimental animals and measures of infection in the control sham-treated monkeys were not sufficiently robust to distinguish between inadequate treatment and the persistence of a subpopulation with antibiotic tolerance. Furthermore, monkeys were infected by needle inoculation of $3 \times 10^8$ stationary phase spirochetes, several orders of magnitude greater than the spirochete inoculum size introduced by an *Ixodes* nymph, which is estimated to be in the hundreds (34). Spirochetes are known to lose plasmids and pathogenicity with expansion and serial passaging in vitro (35), and stationary phase cultures used in that study likely comprised a genetically and phenotypically heterogeneous population of organisms. Although phenotypic variants that exhibit antibiotic tolerance may have been introduced into the primates in large numbers and propagated during the 4-month infection period prior to introduction of antibiotics, the study falls short of showing that they can evolve from tick-transmitted spirochetes or low-dose needle inocula in an immunocompetent host.

The finding of fluorescent deposits containing *B. burgdorferi* antigens in the knee enthesis after antibiotic treatment has important implications for the pathogenesis of human Lyme arthritis. These deposits have inflammatory potential and can induce TNF-α production from C3H/HeJ macrophages, which are LPS unresponsive (36). Such debris could contribute to prolonged inflammatory responses in the joint after infectious spirochetes have been eradicated. This possibility cannot be addressed in WT or *Myd88*−/− mice because histopathologic evidence of arthritis resolves even though spirochetes persist in the joint for the life span of the infected mouse (10, 14, 37). In humans, treatment-resistant Lyme arthritis has been attributed to localized autoimmunity because *B. burgdorferi* DNA can no longer be detected in synovial fluid or synovectomy tissue (22, 38–40), and immune responses that crossreact with self antigens have been found (41–44). Infection-induced autoimmunity, however, does not explain why this form of arthritis resolves over time, usually within 4 years (3, 45). Recently, a human TLR1 polymorphism (1805GG) that reduces TLR1/2 responsiveness has been
correlated with greater inflammation after *B. burgdorferi* infection and the development of antibiotic-refractory Lyme arthritis (46). In mice, defects in TLR2 responsiveness lead to elevation in pathogen burden similar to that seen with MyD88 deficiency (9–11). It is possible that the TLR1 polymorphism permits a higher pathogen burden within ligamentous attachments of the joint (regions in humans that have not been sampled for DNA analysis) and that inflammatory spirochete debris remains after antibiotic treatment. Release of foreign antigens could occur as a consequence of biomechanical stress at the entheses and contribute to inflammation at this site and in adjacent synovium. Residual cellular debris that contains spirochete antigens could also perpetuate self responses that might arise during the course of infection, and these would be expected to subside once foreign antigens are eventually cleared.

Persistent musculoskeletal symptoms and/or objective findings (e.g., arthritis) after antibiotic therapy for human Lyme disease have been attributed to host genetic factors, irreversible infection-induced tissue damage, protracted resolution of inflammation, and infection-induced fibromyalgia or autoimmunity (3, 47, 48). Ongoing infection is considered unlikely given the lack of improvement in these complaints with prolonged antibiotic therapy for human Lyme disease (1, 47, 48). Engorged nymphs used for xenodiagnosis were retrieved and housed at 22°C in environmental chambers for up to 8 days prior to analysis. *I. scapularis* nymphs were used to introduce infection into mice (14). Uninfected nymphs were provided by Durland Fish (Yale School of Public Health, New Haven, Connecticut, USA). Nymphs infected as described with BbN40 at the larval stage were used to introduce infection into mice (14). Uninfected nymphs were used for xenodiagnosis as described previously (14). From 2 to 5 nymphs were placed on the ears of mice to introduce infection and for xenodiagnosis. Engorged nymphs used for xenodiagnosis were retrieved and housed at 22°C in environmental chambers for up to 8 days prior to analysis.

**PCR of *B. burgdorferi* DNA.** DNA was extracted from engorged ticks using either the Allprep DNA/RNA Mini Kit (QIAGEN) or the Isoquick DNA Isolation Kit (Orca Research) according to the manufacturers’ protocols. *B. burgdorferi* ospA and flaB genes were amplified using the following primers: ospA forward 5'-GGTCAACACACTTGAAGTT-3', ospA reverse 5'-GCAGTGCCAATATTCT-3'; flaB forward 5'-TGGCCGGTACCT-GATTGAA-3', flaB reverse 5'TGCAATTACCGCTGTAACTTT-3'. Robocycler (Stratagene) settings were 35 cycles of 94°C denaturation for 1 minute, 55°C annealing for 1 minute, and 72°C extension for 1 minute followed by a final extension set of 72°C for 10 minutes, yielding product sizes of 332 bp and 255 bp for ospA and flaB targets, respectively. *I. scapularis* β-actin was amplified as a control for DNA amplification from ticks, using the following primer pairs: tick β-actin forward 5'-GCGATGTGACCTGTCGCC-3' and tick β-actin reverse 5'-GATGGACCCAGATCATGGTGACCC-3' (14). The 400-bp β-actin product was amplified under the same parameters as for ospA except that the annealing temperature was 59°C and the cycle number was 30. PCR was performed on DNA extracted as described (14) from mouse ear skin, urinary bladders, and knee or tibiotarsal joints. ospA was amplified as described above; the 16S rRNA gene was amplified.

**Methods.** Mice. Breeding pairs of B6 Myd88−/− mice, produced by Shizuka Akira (52), were obtained from Joseph Craft (Yale University). Age- and sex-matched B6 WT mice were purchased from the Jackson Laboratories. B6 Myd88−/− mice were backcrossed with C3H/HeN mice (Jackson Laboratories) to the F6 generation. Female C3H/He mice (6 to 8 weeks old; Jackson Laboratories) were used as a source of macrophages. Myd88−/− mice were maintained as a colony in specific pathogen–free housing with autoclaved food, water, and bedding. Sulfamethoxazole-trimethoprim (0.25 mg/ml; Sulfaftrim, Sigma-Aldrich) was added to drinking water to reduce opportunistic infection. Sulfatrim has no effect on *B. burgdorferi* infection or disease and was provided to Myd88−/− mice throughout all experiments and to WT mice in experiments comparing WT with Myd88−/− mice. Mice were killed by carbon dioxide asphyxiation.

**Spirochetes.** BbN40 or a GFP-expressing stable transfectant of Bb297 (Bb914) was used (15). Bb914 was provided by Melissa Caimano (University of Connecticut Health Center, Farmington, Connecticut, USA) and exhibits infectivity for mice and ticks similar to that of WT strain 297. Frozen aliquots of low-passage spirochetes were thawed, expanded to log phase in Barbour Steenberner Kelley II medium at 33°C, and then enumerated using a Petroff-Haussser counting chamber. Mice were infected either by inoculation with 10⁶ spirochetes in 100 μl BSK II medium into the interscapular region of the back. The MIC and mean bactericidal concentrations (MBC) for ceftriaxone and doxycycline for each spirochete strain were determined by serial dilution of antibiotics in liquid cultures of *B. burgdorferi* as described (13). For BbN40, the MIC and MBC for doxycycline were 0.5 μg/ml and 0.25 μg/ml, respectively; the MIC and MBC for ceftriaxone were 0.065 μg/ml and 0.25 μg/ml, respectively. For Bb914, the MIC and MBC for doxycycline were 0.5 μg/ml and 1.0 μg/ml, respectively; the MIC and MBC for ceftriaxone were 0.008 μg/ml and 0.25 μg/ml, respectively.

**Tick infection of mice.** Laboratory-reared *Ixodes scapularis* nymphs were provided by Durland Fish (Yale School of Public Health, New Haven, Connecticut, USA). Nymphs infected as described with BbN40 at the larval stage were used to introduce infection into mice (14). Uninfected nymphs were used for xenodiagnosis as described previously (14). From 2 to 5 nymphs were placed on the ears of mice to introduce infection and for xenodiagnosis. Engorged nymphs used for xenodiagnosis were retrieved and housed at 22°C in environmental chambers for up to 8 days prior to analysis.

**Figure 7**

Patellae homogenates from sham- and ceftriaxone-treated mice induce TNF-α production from C3H/HeJ macrophages. (A) Macrophages were stimulated with 10 μg/ml of patellae homogenates from the indicated mice. Results represent the mean ± SEM of 4 or more samples from each group. The lower detection limit of the assay was 75 pg/ml. *P = 0.0179, Kruskal-Wallis test; P < 0.05 for sham versus uninfected and ceftriaxone versus uninfected groups; P > 0.05 for sham versus ceftriaxone groups, Dunn’s multiple comparisons test. (B) Dose response of macrophages to the indicated patellae homogenates. Values of duplicate samples differed by less than 10%.
using the following primer pair, yielding a 128-bp product: 16S rRNA forward 5′-TGCTAAGTTGATGCTGCAAAGAA-3′ and 16S rRNA reverse 5′-AATTCTTCTGGCTTCCAACTTATAA-3′. Parameters for 16S rRNA amplification were identical to those set for the ospA gene except that the annealing temperature was 60°C. A 300-bp target of mouse β-tubulin was amplified as for the ospA gene using the following primer pairs: β-tubulin forward 5′-GGGCCCTCTTGTGATGGCTTGGCACAAC-3′ and β-tubulin reverse 5′-CAGCTGTCAATGTGGCACACGATCGGT3′.

Antibiotic treatment of mice. Mice were treated with ceftriaxone (16 mg/kg) administered s.c. twice daily for 5 days or a 30-day course of doxycycline administered continuously in water bottles (0.2% doxycycline in drinking water) (14). Mice were boosted 2 weeks later with 40 μg protein for 1:1 in alumin. Sera were analyzed by immunoblot 2 weeks after the last immunization.

DFA of ear skin. Cryopreserved ear tissue pieces of 3 mm2 were transferred subdermally through a small incision in the back skin of Myd88–/– mice anesthetized with ketamine and xylazine. The incision was sealed using surgical glue.

B. burgdorferi immunoblots. Proteins in B. burgdorferi lysates or patellar homogenates were separated by SDS-PAGE and transferred to nitrocellulose membranes for immunoblot analysis. Membranes were blocked with 3% BSA, and immunoblots were performed using the indicated dilution of sera and 1:1,000 dilution alkaline phosphatase-conjugated goat anti-mouse IgG (H+L chain–specific) (Vector) as described (10). Bound Ab were visualized by enzymatic reaction with BCIP/NBT phosphate substrate (KPL).

Statistical analysis. Fisher’s exact test was used to assess differences between antibiotic- and sham-treated mouse groups. Differences between TNF-α responses of cells stimulated with uninfected mouse tissue versus tissue that had been infected with B. burgdorferi (sham- or antibiotic-treated) were analyzed using the Kruskal-Wallis test with Dunn’s multiple comparisons test. P values of less than 0.05 were considered significant.

Study approval. Experimental protocols involving mice were approved by the Animal Care and Use Committee at Yale University School of Medicine. All mice were cared for according to Yale Animal Care and Use guidelines.

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