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 *B. burgdorferi*, 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-positive deposits were visualized by intravital microscopy in the entheses of antibiotic-treated mice infected with GFP-expressing *B. burgdorferi* and on the ear cartilage surface in sites where immunofluorescence staining detected spirochete antigens. Naive 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 naive 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 *E. coli* 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 have 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 *C57BL/6* mice more susceptible than *C3H* mice 3 weeks or 4 months after infection (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 administration 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−/−, 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−/− 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−/− 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−/− 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

<table>
<thead>
<tr>
<th>Mouse group</th>
<th>Culture of mouse tissuea,b</th>
<th>B. burgdorferi DNA in mouse tissuea</th>
<th>Culture of xenodiagnostic ticksa</th>
<th>B. burgdorferi DNA in xenodiagnostic ticksa</th>
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<td>3/3</td>
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<td>0/5</td>
<td>0/5</td>
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</tr>
<tr>
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<td>3/3</td>
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</tr>
<tr>
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<td>12/12</td>
<td>5/12c</td>
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Table 1
Oral doxycycline eliminates cultivable BbN40 from WT and most Myd88−/− mice

Figure 1
Ceftriaxone reduces pathogen burden in Myd88−/− mice within 24 hours of therapy. (A) Image from Supplemental Video 1 of Bb914 in ear skin of sham-treated Myd88−/− mouse at 22 days of infection. Blue, second harmonics of skin collagen. (B) Still image from Supplemental Video 3 of Bb914 in Myd88−/− mouse after 1 day of ceftriaxone. Arrow denotes the spirochete tracked in Figure 2. White, melanin fluorescence. (C) Image of Bb914 aligned with tendon fibers of a sham-treated Myd88−/− mouse. Spirochete length variation may be due to spirochetes in tandem or immediately after replication. See also Supplemental Video 2. (D) Image of Bb914 in the tendon of a Myd88−/− mouse after 1 day of ceftriaxone. These spirochetes did not exhibit detectable movement. Scale bars: 60 μm.
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 Videos 3 and 4). 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. 

**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 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 3B 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). This mate-

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**Table 2**

<table>
<thead>
<tr>
<th>Mouse group</th>
<th>Culture of mouse tissue</th>
<th>Ear skin</th>
<th>DFA of tissues for <em>B. burgdorferi</em></th>
<th>Multiphoton imaging</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sham</td>
<td>0/6[^d^]</td>
<td>3/3</td>
<td>3/3</td>
<td>Tendon 3/3</td>
</tr>
<tr>
<td>Ceftriaxone</td>
<td>3/3/[^a^][^b^]</td>
<td>3/3</td>
<td>2/3</td>
<td>Tendon 3/3</td>
</tr>
</tbody>
</table>

[^a^]Results are reported as the number of mice that had positive cultures for *B. burgdorferi* over the total number of mice examined.  
[^b^]Results are from cultures of urinary bladders, ear skin, and tibiotarsal joints.  
[^c^]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.  
[^d^]Differences between sham- and ceftriaxone-treated mouse groups were statistically significant (*P* = 0.0119, Fisher’s exact test, for all comparisons).  
[^e^]Differences between sham- and ceftriaxone-treated mouse groups were statistically significant (*P* = 0.0476, Fisher’s exact test).
research article

Figure 3
DFA of ear skin cryosections from sham- and antibiotic-treated Myd88–/– mice. Sagittal sections show spirochetes (green) detected with FITC-conjugated anti–B. burgdorferi Ab and DAPI (blue) delineating cell nuclei. Acetone fixation eliminates GFP expressed by Bb914. A–C are representative of sham-treated (A) and ceftriaxone-treated Bb914-infected mice (B and C). D–F are representative of sham-treated (D and F) and doxycycline-treated BbN40-infected mice (E). The epidermis (e), dermis (d), and cartilage (c) are labeled in F; autofluorescence of the cartilage is detected in the red channel. B. burgdorferi is present in the dermis and adjacent to the cartilage in all sham-treated mice. Antibiotic-treated mice only had B. burgdorferi antigens detected at the dermal-cartilage interface, with significant variation in amount from mouse to mouse. Scale bars: 50 μm.

Discussion
This study demonstrates that antibiotics (ceftriaxone 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 ceftriaxone rapidly (within 24 hours) reduces pathogen burden in the skin, a preferential site of B. burgdorferi infection, and that both ceftriaxone and doxycycline eliminate infectious spirochetes as assessed by real-time PCR. Both tissues used as immunogens resulted in IgG seroconversion to several B. burgdorferi proteins (Figure 6A), whereas tissues from uninfected mice induced only weak reactivity to a protein band of approximately 41 kDa protein. In addition, IgG from infected mice bound antigens present in patella homogenates of sham- and antibiotic-treated mice (Figure 6B). These antigens had molecular weights similar to those detected on B. burgdorferi immunoblots (Figure 6B). Homogenates of patellar tissue from both sham- and ceftriaxone-treated mice induced TNF-α production from C3H/HeJ macrophages in vitro (Figure 7). Ear skin of sham-treated mice contained insufficient amounts of spirochetal antigens to stimulate production of TNF-α from macrophages in vitro (data not shown).

Table 3
Doxycycline eliminates cultivable Bb914 from MyD88–/– mice

<table>
<thead>
<tr>
<th>Mouse group</th>
<th>Culture of mouse tissueα,β</th>
<th>ospA PCR from mouse tissuesα</th>
<th>DFA of mouse tissuesα</th>
<th>Culture of xenodiagnostic ticksα,δ,ε</th>
<th>ospA PCR from xenodiagnostic ticksα,δ,ε</th>
<th>DFA of tick midgutsα,δ,ε</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>
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</tr>
<tr>
<td>Doxycycline</td>
<td>0/9β</td>
<td>3/8β</td>
<td>9/9β</td>
<td>0/7</td>
<td>7/8</td>
<td>0/7β</td>
</tr>
</tbody>
</table>

αResults are reported as the number of mice that had positive cultures for B. burgdorferi over the number of mice examined. βResults 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. γTicks placed on mice 2 weeks after sham or doxycycline treatment were not analyzed by culture, PCR, or DFA for B. burgdorferi. δResults 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). εEar 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 enthese. 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 xenodiagnostic 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 Myd88−/− 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.
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 entheses 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 enthesis 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 (49–51). Indeed, our real-time imaging has demonstrated that B. burgdorferi is highly vulnerable to antibiotics in vivo. Although no single method for detecting viable spirochetes is absolute, the development of antibiotic-resistant Lyme arthritis.

**Methods**

*Mice.* Breeding pairs of B6 Myd88−/−mice, produced by Shizua 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/He mice (Jackson Laboratories) to the F6 gen-

**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 7.5 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%.

**Sporochetes.** 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 Stoenner Kelley II medium at 33°C, and then enumerated using a Petroff-Hauser counting chamber. Mice were infected either by infestation with 5 B. burgdorferi–infected Ixodes nympha or by intradermal inoculation with 106 spirochetes in 100 μl BSK II medium into the interscapular region of the back. The MIC and mean bactericidal concentrations (MBC) of 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 infestation 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 xenodiagnoses. 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 primer pairs: ospA forward 5′-GGTCAACAACACATTGAAGT3′, ospA reverse 5′-GTCAGTGTCAATATTGTC3′; flaB forward 5′-TGTCGGCTACCTGATTGAA3′, flaB reverse 5′-TGCAAAACCGTCTAACATT3′. 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′-GCCCAGTGTCACCTGTCC3′ and tick β-actin reverse 5′-GAGTGGCCAGATCATGTTCGACC3′ (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
using the following primer pair, yielding a 128-bp product: 16S rRNA forward 5′-TTCGTCGCTGTCGCTGAAAAGA-3′ and 16S rRNA reverse 5′-AACCTTGAGGCTCTTCACATTA-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′-GGCCGCTCGTGATGGCTTTGGGACCA-3′ and β-tubulin reverse 5′-CAGGCCTGTCAAATGTGGCAACCAGATCGGT-3′.

**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 containing 5% sucrose). Control mice were sham treated with 0.9% normal saline s.c. or 5% sucrose in drinking water, respectively. For the 4-months–infected C3H mice, ceftriaxone was administered at the above dose twice daily for 5 days, then daily until sacrifice after 18 days of therapy. Serum doxycycline levels were first optimized in a cohort of mice administered the drug in drinking water. Blood samples were obtained from mice at days 1, 3, and 4 after institution of antibiotics, and drug levels were assessed in triplicate 10-μl serum samples using an agar plate–based bacterial growth inhibition assay with *Staphylococcus aureus* (25923; ATCC) (14). Mice administered oral doxycycline had mean serum β levels of 2 ± 0 μg/ml at days 1 and 3 and 1.62 ± 0.25 at day 4. Water bottles containing freshly reconstituted doxycycline were changed every 4 days during the period of antibiotic treatment.

**Culture of *B. burgdorferi* from mouse tissues.** Blood, ear skin, and urinary bladders were cultured in BSK II medium as described (5). Cultures were incubated at 33°C and analyzed at 2 weeks for motile spirochetes. Cultures with no growth were held for an additional 15 weeks before they were considered negative.

**DFA of ear skin.** Ears were flash frozen in OCT and stored at –80°C for group analysis. Cryosections of 6 μm were fixed in acetone and allowed to air dry. After blocking with PBS containing 5% goat serum for 30 minutes or more, sections were stained with FITC-conjugated goat polyclonal Ab to *B. burgdorferi* (1:30 dilution; KPL Inc.) and DAPI (1 μg/ml final concentration; Sigma-Aldrich). Slides were rinsed twice in PBS, then air dried, and coverslips were mounted with Fluoromount-G. Sections were analyzed using an Olympus BX40 widefield immunofluorescence microscope equipped with digital imaging.

**Intravital microscopy.** Imaging was performed in real time using an upright multiphoton laser scanning microscope. Differences between antibiotic- and sham-treated mice were so marked that it was not possible to blind the imaging personnel to the mouse groups. Images were acquired using an Olympus BX61WI fluorescence microscope with a ×20, 0.95NA water immersion Olympus objective and dedicated single-beam LaVision TriM scanning laser (LaVision Biotech) that was controlled by Imrspector software. The microscope was outfitted with a Chameleon Vision II Ti:Sapphire Laser (Coherent) with pulse precompensation. Emission wavelengths of 390–480 nm (blue), 500–550 nm (green, GFP), and 565–665 nm (orange-red) were collected with an array of 3 photomultiplier tubes (Hamamatsu). Mice were anesthetized with an i.p. injection of ketamine (100 mg/kg) and xylazine (10 mg/kg) prior to shaving and Nair application. After prepping, the anesthetized mouse was placed on a custom-designed stereotaxic restraint platform for skin imaging. A plane of deep anesthesia was maintained using a mixture of isoflurane gas and oxygen delivered via a nosecone. For calcaneal tendon imaging, the mouse was immobilized on a stereotactic platform with tweezers to clamp the hip and the spine, and traction was applied to the foot to position the area to be imaged. In some cases, a small incision was made in the skin overlying the tendon to optimize imaging. Patellar enthesis images were acquired after mouse euthanasia. Patellae were harvested, and the articular surface of the patella and adjacent rectus femoris and patellar ligaments were imaged immediately. Image stacks of 1 to 60 optical sections with 1- to 2-μm z-spacing were acquired every 1 to 25 seconds for 1 to 60 minutes with the laser tuned to a wavelength of 940 nm. Each xy plane spanned 300 μm in each dimension with an xy resolution of 0.59 μm per pixel. Velocity software (Improvision) was used to create QuickTime-formatted videos of image sequences. All videos are displayed as 2D maximum intensity projections, unless otherwise stated.

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

**Immunization of mice.** *Myd88*−/− mice were immunized by i.p injection with homogenized cryopreserved ear tissue or 80 μg fresh patellar homogenates mixed 1:1 with alum in a total volume of 200 μl. For patellae immunizations, mice were boosted 2 weeks later with 40 μg antigen mixed 1:1 in alum. Sera were analyzed by immunoblot 2 weeks after the last immunization.

**Assay for TNF-α production.** Patellae harvested aseptically from sham- and ceftriaxone-treated *B. burgdorferi*-infected and uninfected C3H *Myd88*−/− mice (females only) were rinsed twice in tissue homogenization buffer (PBS with 100 U/ml penicillin, 100 μg/ml streptomycin, and 50 μg/ml gentamicin) and then homogenized in 300 μl of buffer on ice. Homogenates were then centrifuged at 828 g for 5 minutes to remove particulate debris. The protein concentration in the homogenate supernatants was determined using a modified Bradford assay (Bio-Rad). Splenocytes harvested from female C3H/HeJ mice were resuspended after rbc lysis at 5 × 10⁹/ml in Clicks’s medium supplemented with 10% FBS, 100 U/ml penicillin, 100 μg/ml streptomycin, 2 mM l-glutamine, and 20 μM 2-mercaptoethanol. Splenocytes were aliquoted at 0.5 ml total volume into 24-well plates and incubated for 2 hours at 37°C, after which nonadherent cells were removed by rinsing twice with fresh medium. Adherent macrophages were stimulated with the indicated amounts of protein for 24 hours, after which TNF-α production was assessed by cytokine-specific ELISA according to the manufacturer’s recommendations (eBioscience).

**B. burgdorferi immunoblot.** 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, after which immunoblots were performed using the indicated dilution of sera and 1:1,000 dilution alkaline phosphatase-conjugated horse anti-mouse IgG (H&L chain–specific) (Vector) as described (10). Bound Ab were visualized by enzymatic reaction with BCIP/NBT phosphatase substrate (KPL).

**Statistics.** 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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