

Mode of Inoculation of the Lyme Disease Agent *Borrelia burgdorferi* Influences Infection and Immune Responses in Inbred Strains of Mice

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Mice were infected with *Borrelia burgdorferi* by infection via *Ixodes ricinus* and experimental inoculation to determine whether transmission rates of spirochetes and antibody responses are influenced. Mice infected by the natural route were substantially more infective for ticks; two- to sixfold more tick larvae were positive for *B. burgdorferi* than those fed on experimentally inoculated mice. In natural infection, spirochetemia may be greater or spirochetes may be more accessible for transmission. Thus, this form of xenodiagnosis could be used to determine levels of spirochetes in the vertebrate host. Similar levels of antibody were present in all mice; however, those infected by the natural route lacked antibodies to outer surface proteins (Osp) A and B. The small antigen dose given through a tick bite may not have been sufficient to induce rapid OspA or OspB antibodies, thereby allowing the later development of higher levels of spirochetemia.

Since the discovery of *Borrelia burgdorferi* as the cause of Lyme borreliosis, most information on the pathogenesis of the disease and the immune responses to *B. burgdorferi* derives from laboratory models in which animals were experimentally inoculated with viable spirochetes. In mice, hamsters, and white-footed mice (*Peromyscus leucopus*), the first antibodies detected were those to the outer surface proteins (Osp) A and B, to a 39-kDa protein, and to the 41-kDa flagellin (fla) [1–3]. It was subsequently shown that antibodies specific for OspA and OspB are protective and able to prevent *B. burgdorferi*-induced disease in immunodeficient SCID [4, 5] and immunocompetent C3H/HeJ mice [6, 7].

In contrast to results in experimentally inoculated mice, the first antibodies to be detected in patients with Lyme borreliosis recognize the 41-kDa flagellin or the 39-kDa antigen (or both) but not OspA or OspB [8]. The differential immune responses of experimentally inoculated mice versus patients with Lyme disease may be related to the mode of inoculation of spirochetes, as suggested by a study in dogs [9]; only experimentally but not naturally infected dogs expressed antibodies to OspA.

The present study was undertaken to investigate the influence of the route of inoculation of *B. burgdorferi* on infection and antibody response in 3 inbred strains of mice.

Materials and Methods

B. burgdorferi. The European strain ZS7 used in this study was cultivated and quantified as described previously [10].

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Recombinant OspA. Recombinant OspA from strain ZS7 was expressed from the puEX1 expression vector in *Escherichia coli*. The protein was extracted and purified by affinity chromatography using an OspA-specific monoclonal antibody as described previously [5].

Tick colony. The larval and nymphal *I. ricinus* ticks used in this study were derived from a *B. burgdorferi*-free laboratory colony maintained at the Institute of Zoology (Neuchâtel).

Animals. Adult mice of strains AKR/N (H-2^k), C3H/HeJ (H-2^k), and DBA/2 (H-2^d) were bred at the Institute of Zoology. Females 8–10 weeks old were used.

Infection procedures. Unfed *I. ricinus* nymphs were infected using the modified capillary method [11]. Strain ZS7 at a concentration of 1×10^8 cells/mL BSK II medium was used to infect the ticks. Ticks took up $\sim 1.5 \mu\text{L}$ ($\sim 1.5 \times 10^5$ spirochetes). Each experimental group of mice included 2 AKR/N, 2 DBA/2, and 2 C3H/HeJ mice.

For group I, 6 mice were inoculated subcutaneously (sc) in the tail with 0.2 mL of BSK II medium containing 2×10^7 *B. burgdorferi* ZS7 organisms.

For group II, 6 mice were inoculated with a suspension of infected *I. ricinus* nymphs. Immediately after the infection by capillary, 42 infected nymphs were triturated in 1.5 mL of BSK II medium. Spirochetes were counted and adjusted to 3×10^6 spirochetes/mL; 0.2 mL of the suspension ($\sim 6 \times 10^5$ spirochetes/mouse) was inoculated sc in the tail of each mouse.

For group III, 6 mice were exposed to 7 artificially infected *I. ricinus* nymphs. Nymphs were fed in two hollow plastic caps fixed bilaterally on the back of the mouse, and a collar was placed around the neck to prevent grooming. Engorged ticks (4–6) were collected from each mouse.

Xenodiagnosis and spirochete detection in ticks. To determine the level of spirochetemia, infected mice were exposed to uninfected *I. ricinus* larvae at 14 (xenodiagnosis I, 80 larvae/mouse) and 100 days (xenodiagnosis II, 60 larvae/mouse) after infection. Larvae were put on the head of the mouse, and a collar was placed around the neck to prevent grooming. Mice were kept in separate cages over trays of water, and engorged ticks that had dropped into water were collected daily, placed into vials, and stored at room temperature and 95% humidity.

Ticks were evaluated for *B. burgdorferi* after molting using a direct immunofluorescence antibody test [12].

Western blot and ELISA. Sera derived from infected mice were analyzed by Western blots of spirochetal lysates (strain ZS7) and by ELISA with either soluble *B. burgdorferi* (ZS7) antigen or recombinant OspA as described [5].

Results

Inbred mice of strains AKR/N (H-2^k), C3H/HeJ (H-2^k), and DBA/2 (H-2^d) were infected with *B. burgdorferi* by experimental inoculation with in vitro-propagated spirochetes (group I), experimental inoculation with a tissue suspension of infected ticks (group II), and natural infection via tick bites (group III). The numbers of spirochetes transferred to mice via syringe were $\sim 2 \times 10^7$ for group I and $\sim 6 \times 10^5$ for group II. The numbers of spirochetes transmitted via tick bites can only be estimated and were $\sim 1 \times 10^6$ at most.

At 14 and 100 days after infection, mice were exposed to uninfected *I. ricinus* larvae to assess their capacity to transmit spirochetes. Mice from each of the 3 inbred strains previously infected via tick bites (group III) were shown to infect, on average, about two- to fourfold more *I. ricinus* larvae (42.5% at day 14 and 88% at day 100) than mice infected via syringe with either *B. burgdorferi* organisms (group I, 6% on day 14 and 37.5% at day 100) or a tissue suspension of infected ticks (group II, 10% at day 14 and 33.3% at day 100). In all 3 strains of mice, infection rates of *I. ricinus* larvae were

significantly increased (two- to sixfold) at day 100 compared with those at day 14 independent of the inoculation protocol, reaching 88% (twofold increase) after tick infection (group III) and between 37.5% (sixfold increase, group I) and 33.3% (threefold increase, group II) after experimental inoculation.

Serum samples were taken from mice at days 7, 14, 26, 40, 63, 77, and 110 after infection and analyzed individually for *B. burgdorferi*-specific antibodies by ELISA. All mice, independent of strain or inoculation protocol, expressed similar amounts of specific antibodies as assessed on total cell lysates of *B. burgdorferi* strain ZS7 (figure 1A). However, when assayed on purified recombinant OspA [5] alone, specific antibodies were found only in experimentally inoculated (group I, II) but not in tick-infected mice (group III, figure 1B).

To verify these results, individual sera were analyzed for antibody specificities by Western blot. Figure 2 depicts representative banding patterns obtained with sera from AKR/N, C3H/HeJ, and DBA/2 mice at days 7, 14, 26, and 40 after inoculation. Sera from mice experimentally inoculated with either spirochetes (group I) or lysates derived from infected ticks (group II) contained antibodies to OspA (31 kDa) and OspB (34 kDa) in addition to various other *B. burgdorferi* structures, including flagellin (41-kDa) and antigens with approximate molecular masses of 14, 18, 22, 26, 39, 65, and 70 kDa. In contrast, sera from mice inoculated via infected

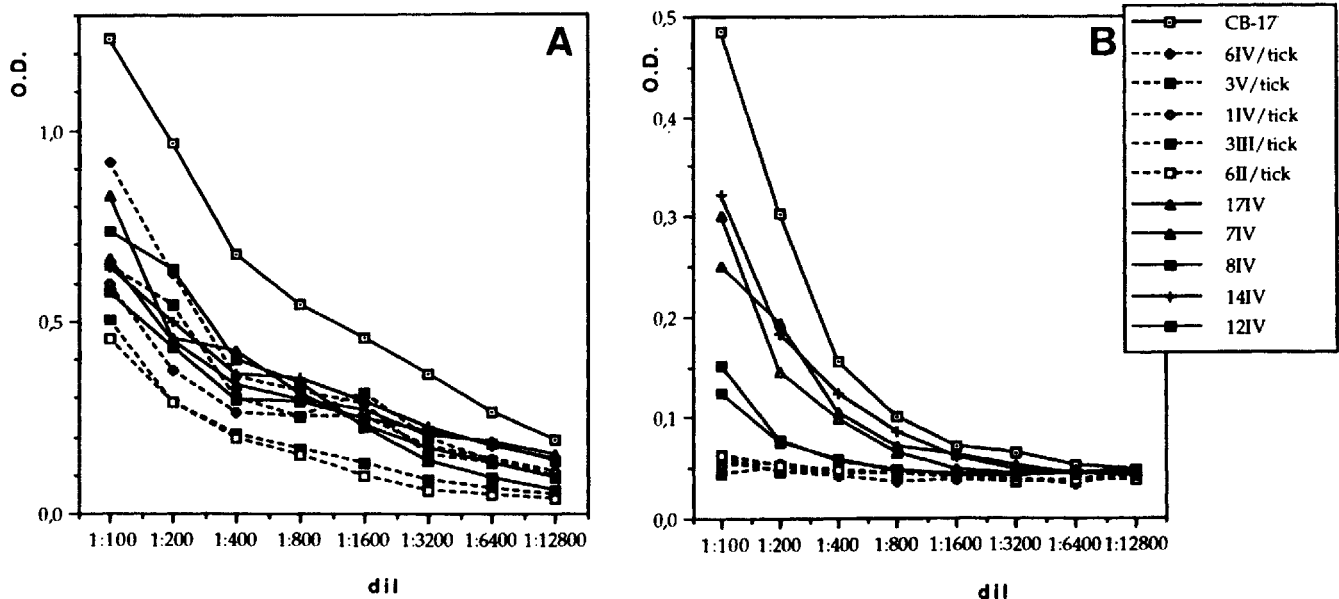


Figure 1. Antibody responses of mice inoculated with 2×10^7 *B. burgdorferi* (—, mouse 14 and 17), tissue lysates derived from experimentally infected ticks (—, 6×10^5 spirochetes/mouse, mouse 7, 8, and 12), or via experimentally infected ticks (---, mouse 1, 3, and 6) by ELISA on whole *B. burgdorferi* proteins (A) and on recombinant (rec) outer surface protein (Osp) A (B). Sera from mouse strains AKR/N (mouse 1, 7, 8, and 14), DBA/2 (mouse 6, 12, and 17), and C3H/HeJ (mouse 3) were tested. Sera were taken 14 (II), 26 (III), 40 (IV), or 63 (V) days after infection. Pooled immune serum derived from 5 CB17 mice was used as control (64 μ L/mL specific immunoglobulin). OD, optical density; dil, dilution.

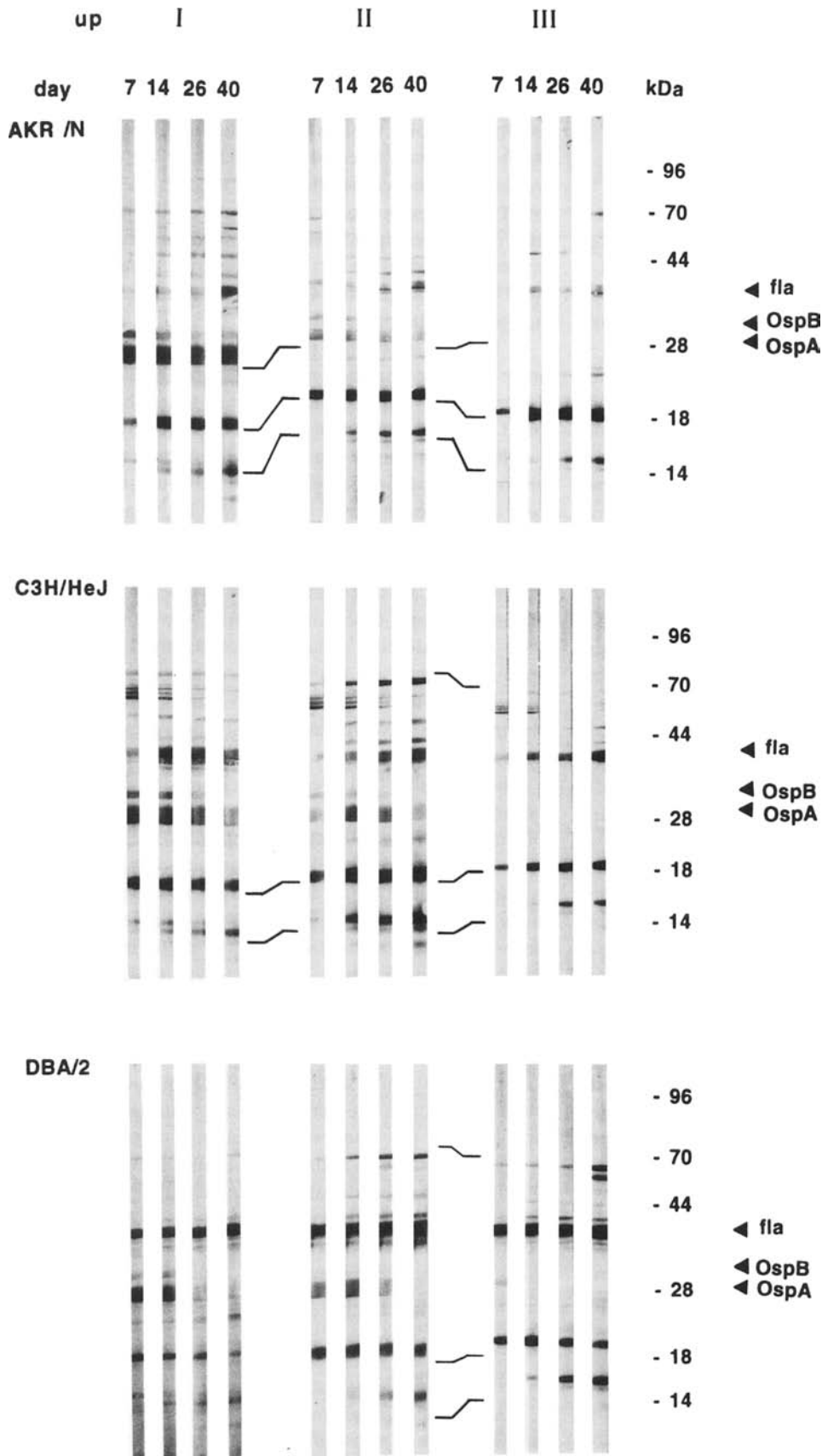


Figure 2. Western blot analysis of sera taken from individual mice of strains AKR/N, C3H/HeJ, and DBA/2 at indicated days after infection. Mice were inoculated with 2×10^7 *B. burgdorferi* spirochetes in BSK II (I), tissue lysates from experimentally infected ticks (6×10^5 /mouse, II), or via experimentally infected ticks (III). Representative data from 1 mouse of each group are shown. Osp, outer surface protein; fla, flagellin.

ticks (group III) contained no or few antibodies to OspA and OspB but had a banding pattern similar to that of the sera from groups I and II. Similar results were obtained with sera taken at days 63, 77, and 110 (data not shown).

Discussion

This investigation showed that the rate of transmission of spirochetes to noninfected tick larvae is much higher in mice infected via tick bites than in those experimentally inoculated by needle injection and that antibodies to OspA and OspB are generated only in response to experimentally but not to tick-delivered organisms.

At first glance, the finding that naturally infected mice in turn infect more ticks than mice experimentally inoculated with cultured spirochetes is surprising, since the number of spirochetes transferred by needle inoculation was at least two to three orders of magnitude higher. This discrepancy may be related to the site of infection, the immune reaction of the host, the number of spirochetes transferred (antigenic load), or the phenotypic changes of the spirochetes within the vector or host or both.

It is possible that skin areas chosen by ticks for blood meals are more heavily infested with spirochetes whereas others are relatively free of the organism. In that case, more naive ticks would become infected when feeding on a recipient that was itself infected by tick bites. At present, detailed information on that point is lacking and requires more intensive studies. It is more likely that the discrepancy of transmission rates observed in naturally versus experimentally infected mice is due to qualitative differences in their immune responses. In fact, sera from all mice infected by any of the three protocols contained similar quantities of spirochete-specific antibodies independent of the greatly differing numbers of spirochetes transferred ($<10^6$ by tick bites, 6×10^5 – 2×10^7 by needle infection); however, antibodies to OspA or OspB were observed only in mice experimentally inoculated by needle injection and not in those infected via tick bites. The finding that the transmission of *B. burgdorferi* organisms to ticks correlated inversely with the presence of antibodies to OspA and OspB in mice therefore suggests that these antibodies are responsible for the elimination of spirochetes and consequently for the reduction in the rate of transmission. This assumption is also supported by previous studies showing that OspA- and OspB-specific antibodies protect SCID [4, 5] and C3H/HeJ mice [6, 7] against infection and development of arthritis after experimental and tick-mediated infection with *B. burgdorferi*, when these antibodies are present at the time of inoculation [13, 14].

The finding that the humoral immune response to tick-delivered spirochetes did not include recognition of OspA and OspB is analogous to the situation in patients with Lyme disease, who develop these antibodies, if at all, only during later stages of the infection [8]. It is therefore possible that

the immune response of the host is influenced by inoculation route. This hypothesis is substantiated by previous studies demonstrating that experimentally infected dogs [9] and inbred [1, 3, 13] and wild mice (Kurtenbach K, personal communication) express antibodies to OspA and OspB, whereas naturally infected animals do not. The host immune system may indeed be affected by the tick itself, which has been shown before to secrete immunomodulatory constituents during feeding [15]. However, such a mechanism is hard to reconcile with the fact that the suppression seen affects antibodies to OspA and OspB but not to other spirochetal antigens, including flagellin.

It is more likely that the inability of naturally infected mice to produce antibodies to OspA and OspB is due to suboptimal doses of spirochetes delivered during the tick's blood meal. This assumption is strongly supported by recent studies showing that mice experimentally inoculated with different numbers of spirochetes (10^1 – 10^8) express similar quantities of antibodies to *B. burgdorferi* but that those with specificities for OspA and OspB are seen only in animals receiving $>10^4$ organisms (unpublished data). These results not only indicate that the number of spirochetes transmitted by ticks during a blood meal is rather low but also suggest that the qualitative differences of humoral immune responses observed are due to differential immunogenicity of individual antigens, such as OspA, OspB, and flagellin, in the inoculum. Experiments to test this hypothesis are underway.

The possibility that phenotypic changes of the spirochetes occurring in the tick or mouse environment (or both) as demonstrated previously for certain isolates [16] may contribute to the suppression of antibody responses to OspA and OspB cannot be formally excluded; however, this seems less likely, since similar phenotypic patterns were obtained with the in vitro-propagated strain ZS7 and its reisolates from either the vector or the murine host. Compared with the first xenodiagnosis at 14 days after infection, the percentage of infected ticks increased from two- to sixfold in all experimental groups of mice at day 100 after infection. These data suggest that spirochetes may survive or multiply in mice even if protective antibodies are generated early during disease. It is possible that the kinetics of appearance of anti-OspA and anti-OspB antibodies and/or their serum level only controls the development of disease, as demonstrated previously [4, 13], but are insufficient to eradicate the spirochetes from the host. Further studies will be directed toward the development of immunization protocols that allow the establishment of sterile immunity against *B. burgdorferi* infection in this animal model.

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DNA Fragment Length Polymorphism Analysis of *Mycobacterium tuberculosis* Isolates by Arbitrarily Primed Polymerase Chain Reaction

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Strain identification of *Mycobacterium tuberculosis* would prove whether transmission had occurred between individuals. A method to characterize strains of *M. tuberculosis* has been developed utilizing polymerase chain reaction (PCR). Purified chromosomal DNA of cultured clinical samples of *M. tuberculosis* were subjected to PCR using short (10-12 nucleotide) oligonucleotide primers. PCR products visualized after agarose gel electrophoresis and ethidium bromide staining demonstrated that different strains of *M. tuberculosis* give different banding patterns. This technique was used to confirm the relationship between cases of tuberculosis in several clusters, prove the lack of relationship between 2 isolates with the same antibiotic-resistance pattern, confirm a suspected mislabeling event, and suggest the source of infection in a case of tuberculous meningitis. This method is rapid and simple and does not require radioactive probes.

Tuberculosis control in developed countries is based mainly on the interruption of the disease transmission by case finding and treatment. Strategies for case finding gener-

ally include case tracing from an index patient with the assumption of airborne spread to individuals in close contact. Direct proof of transmission by strain identification of the organisms is rarely done since all strains of *Mycobacterium tuberculosis* are serologically homogeneous [1]. Until recently, strain identification could be done only by phage typing, which is time-consuming and technically demanding. Analysis of the resistance profile is of limited value since the number of drugs is finite and an identical resistance profile does not prove that the strains are identical.

Restriction fragment length polymorphism (RFLP) can differentiate strains of *M. tuberculosis*. RFLP could be analyzed after ethidium bromide staining [2] or after Southern

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