Death from untreated relapsing fever is most common during the first fever episode. With prompt antibiotic treatment, the mortality rate is 2–5% for LBRF and <2% for TBRF. Features associated with a poor prognosis include concurrence with malaria, typhus, or typhoid; pregnancy; stupor or coma on admission; diffuse bleeding; poor liver function; myocarditis; and bronchopneumonia. The mortality rate from the JHR in LBRF, in the absence of adequate monitoring and resuscitation measures, is ~5%. Some patients have survived the crisis or the JHR only to die suddenly either later that day or on the next day. Relapsing fever during pregnancy frequently leads to abortion or stillbirth, but congenital malformations have not been reported. Although it is possible that spirochetes may persist in the CNS or other sequestered sites after bacteremia has resolved, chronic disease or disability from a persistent infection has not been attributed to relapsing fever. Partial immunity against reinfection seems to develop in residents of endemic areas.

PREVENTION

There is no vaccine for either LBRF or TBRF. Reduction of exposure to lice and ticks is the key strategy for prevention. LBRF can be prevented through improved personal hygiene, reduction of crowding, better access to washing facilities, and selected use of pesticides. Infested clothing is an important factor in maintaining body lice. The risk of TBRF can be reduced by construction of houses with concrete or sealed plank floors and without thatched roofs or mud walls. Log cabins pose a particular risk in North America when rodents nest in the roof or beneath the house or porch. Interiors of buildings infested with *Ornithodoros* ticks can be treated with pesticides. If residing in a high-risk environment, individuals should not sleep on the floor, and beds should be moved away from the wall. With an exposure to TBRF, postexposure treatment with doxycycline (200 mg on day 1 followed by 100 mg/d for 4 days) was efficacious in preventing infection in a placebo-controlled trial.

210 Lyme Borreliosis

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DEFINITION

Lyme borreliosis is caused by a spirochete, *Borrelia burgdorferi sensu lato*, that is transmitted by ticks of the *Ixodes ricinus* complex. The infection usually begins with a characteristic expanding skin lesion, erythema migrans (EM; stage 1, localized infection). After several days or weeks, the spirochete may spread to many different sites (stage 2, disseminated infection). Possible manifestations of disseminated infection include secondary annular skin lesions, meningitis, cranial neuritis, radiculoneuritis, peripheral neuritis, carditis, atrioventricular nodal block, or migratory musculoskeletal pain. Months or years later (usually after periods of latent infection), intermittent or persistent arthritis, chronic encephalopathy or polyneuropathy, or acrodermatitis may develop (stage 3, persistent infection). Most patients experience early symptoms of the illness during the summer, but the infection may not become symptomatic until it progresses to stage 2 or 3.

Lyme disease was recognized as a separate entity in 1976 because of a geographic cluster of children in Lyme, Connecticut, who were thought to have juvenile rheumatoid arthritis. It became apparent that Lyme disease was a multisystemic illness that affected primarily the skin, nervous system, heart, and joints. Epidemiologic studies of patients with EM implicated certain *Ixodes* ticks as vectors of the disease. Early in the twentieth century, EM had been described in Europe and attributed to *I. ricinus* tick bites. In 1982, a previously unrecognized spirochete, now called *Borrelia burgdorferi*, was recovered from *Ixodes scapularis* ticks and then from patients with Lyme disease. The entity is now called Lyme disease or Lyme borreliosis.

ETIOLOGIC AGENT

B. burgdorferi, the causative agent of Lyme disease, is a fastidious microaerophilic bacterium. The spirochete's genome is quite small (~1.5 Mb) and consists of a highly unusual linear chromosome of 950 kb as well as 17–21 linear and circular plasmids. The most remarkable aspect of the *B. burgdorferi* genome is that there are sequences for more than 100 known or predicted lipoproteins—a larger number than in any other organism. The spirochete has few proteins with biosynthetic activity and depends on its host for most of its nutritional requirements. It has no sequences for recognizable toxins.

Currently, 13 closely related borrelial species are collectively referred to as *Borrelia burgdorferi sensu lato* (i.e., "*B. burgdorferi* in the general sense"). The human infection Lyme borreliosis is caused primarily by three pathogenic genospecies: *B. burgdorferi sensu stricto* ("*B. burgdorferi* in the strict sense," hereafter referred to simply as *B. burgdorferi*), *Borrelia garinii*, and *Borrelia afzelii*. *B. burgdorferi* is the sole cause of the infection in the United States; all three genospecies are found in Europe, and the latter two species occur in Asia.

Strains of *B. burgdorferi* have been subdivided according to several typing schemes: one based on sequence variation of outer-surface protein C (OspC), a second based on differences in the 16S–23S rRNA intergenic spacer region (RST or IGS), and a third called *multilocus sequence typing*. From these typing systems, it is apparent that strains of *B. burgdorferi* differ in pathogenicity. OspC type A (RST1) strains seem to be particularly virulent and may have played a role in the emergence of Lyme disease in epidemic form in the northeastern United States in the late twentieth century.

EPIDEMIOLOGY

The 13 known genospecies of *B. burgdorferi sensu lato* live in nature in enzootic cycles involving 14 species of ticks that are part of the *I. ricinus* complex. *I. scapularis* (Fig. 475-1) is the principal vector in the eastern United States from Maine to Georgia and in the midwestern states of Wisconsin, Minnesota, and Michigan. *I. pacificus* is the vector in the western states of California and Oregon. The disease is acquired throughout Europe (from Great Britain to Scandinavia to European Russia), where *I. ricinus* is the vector, and in Asian Russia, China, and Japan, where *I. persulcatus* is the vector. These ticks may transmit other agents as well. In the United States, *I. scapularis* also transmits *Babesia microti*; *Anaplasma phagocytophilum*; *Ehrlichia* species Wisconsin; *Borrelia miyamotoi*; and, in rare instances, Powassan encephalitis virus (the deer tick virus) (see "Differential Diagnosis," below). In Europe and Asia, *I. ricinus* and *I. persulcatus* also transmit tick-borne encephalitis virus.

Ticks of the *I. ricinus* complex have larval, nymphal, and adult stages. They require a blood meal at each stage. The risk of infection in a given area depends largely on the density of these ticks as well as their feeding habits and animal hosts, which have evolved differently in different locations. For *I. scapularis* in the northeastern United States, the white-footed mouse and certain other rodents are the preferred hosts of the immature larvae and nymphs. It is critical that both of the tick's immature stages feed on the same host because the life cycle of the spirochete depends on horizontal transmission: in early summer from infected nymphs to mice and in late summer from infected mice to larvae, which then molt to become the infected nymphs that will begin the cycle again the following year. It is the tiny nymphal tick that is primarily responsible for transmission of the disease to humans during the early summer months. White-tailed deer, which are not involved in the life cycle of the spirochete, are the preferred host for the adult stage of *I. scapularis* and seem to be critical to the tick's survival.

Lyme disease is now the most common vector-borne infection in the United States and Europe. Since surveillance was begun by the Centers for Disease Control and Prevention (CDC) in 1982, the number of cases in the United States has increased dramatically. More than 30,000 new cases are now reported each summer, but the actual number of new cases is probably closer to 300,000 annually. In Europe, reported frequencies of the disease are highest in the middle of the continent and in Scandinavia.