

Invasion of the Central Nervous System by Intracellular Bacteria

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INTRODUCTION

Invasion of the central nervous system (CNS) is a severe and frequently fatal event during the course of many infectious diseases, and those who survive are often left with permanent neurological dysfunction. One of the key events in the pathogenesis of CNS infections is how microbes interact with and cross the blood-brain or blood-choroid barriers to gain access to the central compartment. Great strides have been made in understanding the mechanisms by which bacterial pathogens with a predominantly extracellular life cycle, such as *Streptococcus pneumoniae*, *Neisseria meningitidis*, and *Haemophilus influenzae*, invade the CNS from the bloodstream (for recent reviews, see references 172, 196, 206, 274, and 419). However the mechanisms used by obligate and facultative intracellular bacterial pathogens, here collectively referred to as intracellular bacteria, to enter the CNS are less well established.

In most scenarios, neuroinvasion occurs in the context of a systemic disease and typically follows bacterial dissemination via the bloodstream. In this respect, intracellular and extracellular bacteria have developed different strategies to solve the biological problem of how to escape elimination by host defenses such as phagocytes and serum proteins. Extracellular bacteria typically produce a polysaccharide capsule that allows them to resist complement-mediated lysis and phagocytosis by leukocytes. As a result, these pathogens typically achieve a high-density bacteremia that is important for CNS invasion (217, 372). Successful entry into the CNS then results in ex-

remely high densities of extracellular bacteria in the cerebrospinal fluid (CSF) (39, 217).

By comparison, the means used by facultative and obligate intracellular bacteria to survive and disseminate within the host are fundamentally different. These differences are reflected in the lower densities of these organisms in the bloodstream and the CSF than of extracellular pathogens. Although this point is intuitively obvious for obligate intracellular bacteria, recent data cited below indicate that even facultative intracellular pathogens, i.e., those that can replicate inside or outside of a eukaryotic cell, cause disease only if they can survive intracellularly. Put another way, if intracellular replication is prevented, the microbes are not pathogenic, suggesting that the intracellular and extracellular pathways are not equal in terms of pathogenicity. This concept is important for understanding how facultative intracellular bacteria enter the CNS, considering that in some infections both extracellular and intracellular organisms are present in the bloodstream to a greater or lesser extent and it is unclear whether one is more neuroinvasive than the other. For example, recent studies of facultative intracellular bacteria including *Listeria monocytogenes*, *Mycobacterium tuberculosis*, and *Salmonella enterica* serovar Typhimurium show that they take up an intracellular location almost from the initial point of invasion of the host (302, 316, 317, 378, 391). Conversely, even obligate intracellular bacteria like *Ehrlichia chaffeensis* have an extracellular portion of their life cycle that could, in theory, contribute to CNS invasion (222). Thus, intracellular bacteria could have more options available to them than do extracellular bacteria for entering the CNS.

Mechanisms used by microbial pathogens to enter the CNS are usually divided according to the cellular route involved and

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TABLE 1. Possible mechanisms used by intracellular bacteria for CNS invasion

Microbe	Possible mechanisms for CNS invasion ^a		
	Phagocyte-facilitated invasion	Direct invasion of endothelial cells	Retrograde transport within neurons
<i>L. monocytogenes</i>	+	+	+
<i>M. tuberculosis</i>	+	?	—
<i>Brucella</i> spp.	+	?	—
<i>Salmonella</i> spp.	+	?	—
<i>R. rickettsii</i>	—	+	—
<i>R. prowazekii</i>	—	+	—
<i>O. tsutsugamushi</i>	+	+	—
<i>E. chaffeensis</i>	+	—	—
<i>C. burnetii</i>	+	—	—
<i>T. whipplei</i>	+	—	—

^a The qualitative scale is as follows: +, likely; ?, questionable; —, unlikely.

whether the organisms breach endothelial cells of blood-brain barrier or specialized epithelial cells of blood-choroid barriers. These routes of entry are commonly referred to as (i) intercellular, i.e., passing between cells, (ii) transcellular, i.e., passing through cells, (iii) leukocyte facilitated, i.e., a Trojan horse-like mechanism, or (iv) nonhematogenous (172, 182, 241, 419). The most common routes of entry for extracellular bacteria are the intercellular and transcellular routes. The transcellular approaches are interesting in that they result in these bacteria temporarily assuming an intracellular location (111, 303, 324). Transcytosis of *Streptococcus pneumoniae* across endothelial cells is an excellent example of this event. Pneumococci bind to the platelet-activating factor receptor on endothelial cells that are activated by proinflammatory cytokines such as tumor necrosis factor alpha (TNF- α) and interleukin-1 α (IL-1 α), in a process dependent on choline binding protein A, a bacterial adhesin (73). Bound bacteria are internalized into vesicles which transport transparent bacterial variants through the endothelial cell or recycle them to the apical surface, whereas most opaque bacterial variants are degraded within the cell (324). Intracellular organisms, on the other hand, typically do not use an intercellular approach, and the transcellular approach is modified by the fact that bacterial replication often takes place within endothelial cells. In addition, at least one bacterium, *L. monocytogenes*, is able to enter the brain via a nonhematogenous route by retrograde transport within cranial nerves.

The purpose of this overview is to elaborate on the mechanisms by which different facultative and obligate intracellular bacteria achieve CNS invasion in the context of human disease. Predominant clinical features reflective of CNS infection are noted, and the salient aspects of the pathological findings at postmortem examination, particularly as they shed light on entry mechanisms, are reviewed. Similarly, we discuss some studies in experimental or natural infection in animals as far as they provide insight into CNS invasion (Table 1). We refer as much as possible to the primary literature describing the original findings so that this paper can be a useful resource to the reader who wants to go into more detail in given aspects. In keeping with the focus of this review, experiments detailing the innate and adaptive host immune responses that follow bacte-

rial entry the CNS are generally not discussed, nor are treatment modalities and prevention strategies discussed. Finally, readers are directed to several recent reviews for a detailed discussion of the blood-brain and blood-choroid barriers in steady state and infection (173, 259, 289, 330).

This review is arranged according to the different microorganisms rather than by the presumed mechanism(s) of entry into the CNS. For most of the organisms considered here, the cellular and molecular mechanisms that enable neuroinvasion are largely unknown or are just now being discovered. Of these, the neuroinvasive mechanisms of *L. monocytogenes* and *Rickettsia rickettsii* are the best studied. This review begins by discussing *L. monocytogenes* because it is a good model organism for illustrating the various modes of CNS entry available to intracellular pathogens. For other bacteria, however, general aspects of neuroinvasion can be inferred from careful histopathological studies of CNS infections and/or from what is known about the cell biology of the organism. Examples of these organisms are *M. tuberculosis*, *Brucella* spp., *Salmonella* spp., *Orientia tsutsugamushi*, *E. chaffeensis*, and *Coxiella burnetii*. Finally, one organism, *Tropheryma whipplei*, which causes a rare, poorly understood disease that includes CNS symptoms but for which little is known about its pathophysiology, is discussed.

FACULTATIVE INTRACELLULAR PATHOGENS

Listeria monocytogenes

L. monocytogenes is a gram-positive, nonsporulating bacillus that is facultatively anaerobic and produces weak beta-hemolysis on blood agar. It is divided into 13 serotypes based on antibody reactivity to somatic and flagellar antigens, but serotypes 1/2a, 1/2b, and 4b account for the majority of human disease (189). It is a facultative intracellular parasite of professional and nonprofessional phagocytes, with a similar intracellular life cycle in both cases (299, 390). Phagocytic cells internalize bacteria through a variety of opsonin-dependent and opsonin-independent mechanisms. In addition, *L. monocytogenes* can induce its own entry into nonprofessional phagocytes by using invasion proteins such as internalin A, internalin B, and P60. Live bacteria delay phagosomal maturation and targeting to the degradative pathway and rapidly lyse the membrane of the acidified phagosome through the action of listeriolysin O (LLO) working in concert with the phospholipases PlcA and PlcB (10, 122, 240, 300, 313, 381). The bacteria then reside freely in the cytoplasm, where they replicate and acquire F-actin-based intracellular motility based on expression of the ActA protein (205). Subsequently, there is invasion of adjacent cells by cell-to-cell spread (326, 381).

Clinical infections with gram-positive rods suggestive of *L. monocytogenes* were first described in the 1890s in humans (cited by Seeliger [349]) and 1911 in rabbits (by Hülphers [174]). Cases of human meningitis caused by gram-positive diphtheroid bacilli later thought to be *L. monocytogenes* were identified in 1915 by Atkinson (19), in 1918 by Dumont and Cotoni (99), and in 1919 by Dick (85). A clear association between *L. monocytogenes* and CNS infection in humans was established in 1945 by Kaplan, who found that meningitis was present in 22 of the first 36 cases reported in the medical

literature (184). Clinical reviews in the 1960s by Gray and by Louria et al. supported this association by identifying meningitis and/or meningoencephalitis in 77 and 78% of patients, respectively (145, 229).

Most human infection by *L. monocytogenes* is acquired by consumption of contaminated food. A recent study estimated that 2,500 invasive *L. monocytogenes* infections occur in the United States annually (254). Interestingly, this study concluded that *L. monocytogenes* was responsible for less than 0.1% of all food-borne illnesses but caused 27.6% of deaths attributed to food-borne diseases. *L. monocytogenes* attacks hosts who are immunocompromised by factors such as pregnancy, the extremes of age, solid and hematological malignancies, and liver disease, as well as chronic illnesses not typically thought of as immunosuppressive such as heart disease and diabetes (140, 344). Pregnancy is the single most common predisposing factor and is associated with approximately 35% of cases worldwide (358). Nonperinatal cases of invasive listeriosis in developed nations have an incidence that ranges from 0.1 to 1.1 cases/10⁵ population, 74% of which are in immunocompromised persons, with an overall mortality rate of 36% (358). Regarding CNS infection, active surveillance in the United States showed that *L. monocytogenes* is the second leading cause of bacterial meningitis in patients younger than 1 month or older than 60 years (345). In addition, it is frequently isolated throughout the developed world in community-based studies of bacterial meningitis in adults (101, 114, 175, 360).

L. monocytogenes has a striking predilection for invading the CNS in humans. Reviews published in the past 40 years show that CNS infection is present in 28 to 79% of cases of invasive listeriosis in nonpregnant adults (229, 277, 358) and in 13 to 44% of neonates (119, 251, 278, 348, 377). In contrast to extracellular bacteria that commonly cause CNS infection, i.e., *S. pneumoniae*, *N. meningitidis*, and *H. influenzae*, *L. monocytogenes* infection results in a variety of CNS manifestations including meningitis, meningoencephalitis, rhombencephalitis, cerebritis, and brain abscess. Meningitis and meningoencephalitis are the two most common manifestations in cases not associated with pregnancy (49, 140, 252, 271, 296). The majority of brain abscesses also occur in individuals who are compromised by underlying medical conditions or who are receiving immunosuppressive therapy (77, 102). In contrast, rhombencephalitis, a primary infection of the brain stem, is clearly different since it is found predominantly in noncompromised adults (15, 386). The most common symptoms referable to the CNS are headache and altered sensorium including confusion, lethargy, and coma (44, 140, 215, 271, 296). Seizures and focal neurological findings including cranial nerve palsies are also observed. CSF abnormalities are not diagnostic for *L. monocytogenes* infection and can manifest either a polymorphonuclear or a mononuclear predominance but are reported to show lower total leukocyte and protein concentrations compared with meningitis caused by extracellular bacterial pathogens (44, 215, 271, 296).

Postmortem pathological findings in the brains of humans include focal hemorrhages and areas of necrosis on a gross level, with purulent meningitis, suppurative inflammation with necrosis, microabscesses, and vasculitis with perivascular lymphocytic cuffing and mononuclear cell infiltration of the vessel

walls on histological analysis (15, 146). Bacilli are present in the necrotic parenchymal lesions but are not typically found in the perivascular cuffs. Histological findings in sheep and other ruminants are similar to those in humans, with the exception that humans more typically manifest meningitis or meningoencephalitis whereas brain stem encephalitis is more common in ruminants (51, 72). Histological analysis for sheep shows that focal lesions in the parenchyma usually occur around small vessels. The lesions vary in size from 50 to 500 μm , with decreasing numbers of macrophages but increasing numbers of bacteria and neutrophils as they enlarge that ultimately form a microabscess (72). Such microabscesses are most common in the reticular formation of the midbrain, pons, and medulla but also are found elsewhere. In addition to parenchymal lesions, perivascular meningeal infiltrates are present and consist mostly of mononuclear cells with some neutrophils. The infiltrates are often generalized but can be particularly severe basally and in the cerebral and cerebellar sulci. Immunohistochemical analysis of these lesions shows mixed cellular infiltrates consisting of mononuclear phagocytes and neutrophils with relatively few T and B lymphocytes (211). In parenchymal microabscesses and glial nodules, there is evidence of macrophage activation, as shown by the expression of inducible nitric oxide synthase that colocalizes with bacteria (183). In addition, inducible nitric oxide synthase-negative cells that also express major histocompatibility complex (MHC) class II are present in perivascular lesions and could represent infiltrating dendritic cells or transmigrating monocytes that are not yet fully activated. Ultrastructural analysis by electron microscopy of a specimen from a patient with fatal human meningoencephalitis clearly identified intracellular *L. monocytogenes* free within the cytoplasm of infected macrophages, smooth muscle cells, and endothelial cells (200).

Experimental data indicate that *L. monocytogenes* can use several different mechanisms to invade the CNS (Fig. 1). Infection of the CNS from the bloodstream could occur by (i) direct invasion of endothelial cells of the blood-brain or blood-choroid barriers by blood-borne bacteria or (ii) transportation of bacteria to the CNS within circulating leukocytes in a phagocyte-facilitated (Trojan horse) mechanism. In addition, a neural route exists whereby bacteria reach the CNS from peripheral tissues by intra-axonal transport.

Invasion by blood-borne *L. monocytogenes*. Different lines of evidence support the concept that CNS invasion by blood-borne *L. monocytogenes* is the predominant route of infection in humans; this evidence includes the high concurrence of bacteremia and CNS invasion in cases of invasive disease and the distribution of histological lesions in the brain. The mouse model of experimental *L. monocytogenes* infection has provided key insights into how bacteria enter the CNS in vivo. CNS infection as a manifestation of systemic disease develops in mice infected by a variety of routes including intravenous, intraperitoneal, and subcutaneous injections or by introduction of the bacteria into their drinking water (31, 90, 93, 225, 301). In each of these models, the majority of bacteria can be recovered from the liver and spleen, but the bone marrow also is infected (76). Presumably, overwhelming replication of *L. monocytogenes* in these organs leads to a secondary bacteremia that is composed of cell-free and cell-associated bacteria in an approximately 70:30 ratio (90). CNS invasion follows or coin-

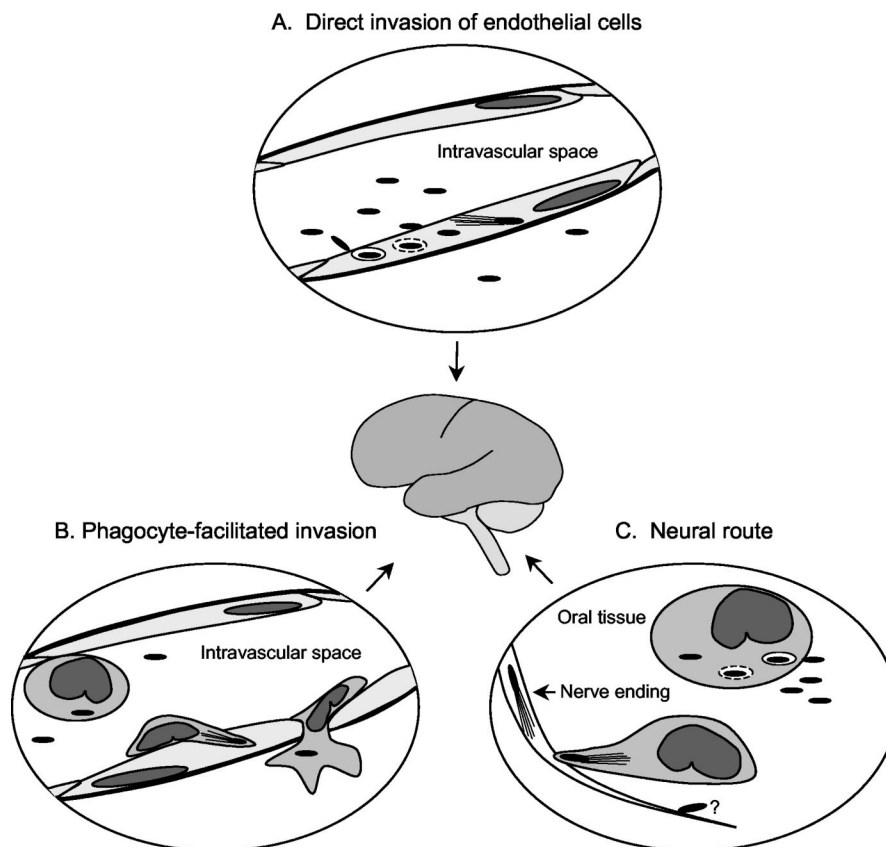


FIG. 1. Routes used by *L. monocytogenes* for neuroinvasion. This cartoon shows a schematic representation of the three different routes used by *L. monocytogenes* to invade the CNS. It should be noted that the intracellular life cycle of *L. monocytogenes*, specifically phagosomal escape and F-actin-based motility, is crucial to each case. (A) Direct invasion of endothelial cells. In this scenario, blood-borne bacteria directly invade endothelial cells of the CNS vasculature, possibly via internalin B or other not yet described mechanisms. Bacteria are internalized into phagosomes, which are lysed largely through the action of LLO, and then escape into the cytoplasm and acquire F-actin-dependent intracellular motility requiring the expression of the ActA protein. The bacteria transcytose the endothelium and then invade surrounding cells including neurons and microglia or escape into the subarachnoid space (not shown). (B) Phagocyte-facilitated invasion. Monocytes that are parasitized in the periphery are recruited to the CNS through as yet unidentified signals. Infected leukocytes adhere to endothelial cells of the CNS vasculature, allowing intercellular spread of bacteria from the phagocyte to the endothelial cell. Alternatively, the leukocyte can transmigrate and deliver bacteria to the CNS parenchyma. (C) Neural route. Bacteria are inoculated into the oral tissues when abrasive food is chewed. Tissue macrophages, or blood phagocytes recruited into the infected tissue, phagocytose the bacteria and become parasitized by intracellular bacteria that facilitate the invasion of cranial nerve neurites by cell-to-cell spread. Bacteria move in a retrograde direction through the nerve axons, eventually reaching the CNS, where they continue to spread intercellularly in the parenchyma. Direct invasion of some nerves *in vitro* has been described, but its role *in vivo* is unclear.

cides with the appearance in the bloodstream of bacteria and parasitized cells and results in histological changes generally similar to those found in humans (31, 226, 301). Data from a gerbil model show that after bacterial infection has been established in the CNS, the majority of bacteria actually reside in the brain parenchyma rather than the CSF (42).

An interesting point in the mouse model is that CNS invasion occurs relatively late in systemic disease, so that animals that succumb early to overwhelming infection die of peripheral organ failure but manifest little in the way of pathological changes in the CNS (18, 72). These experiments suggest that intravenously injected *L. monocytogenes* organisms are not particularly tropic for the brain endothelium in the same way that they are for other cells, in particular hepatocytes (148). Rather, migration of bacteria into the brain from the bloodstream requires a combination of duration and density of bacteremia, neither of which is fully quantified (31). As a complement to

these factors, Autret et al. sought to identify bacterial genes required for brain infection by using *in vivo* signature-tagged transposon mutagenesis (23). These investigators identified 18 hemolytic *L. monocytogenes* mutants attenuated for brain invasion. The transposon was mapped to 10 distinct loci, 5 of which corresponded to putative cell wall components including *GtcA* (five mutants) and *YtgP* (four mutants), whereas the others represented a variety of cellular processes. This method offers great promise for identifying molecular mechanisms for brain invasion *in vivo*.

The location where bacteria enter the CNS in experimental infection, as well as the subsequent lesions produced in response to them, is heavily influenced by how the animals are infected and the ensuing systemic illness. For example, the main inflammatory lesions in the CNS following intravenous and subcutaneous inoculation of mice are in the choroid plexus and the ventricles, suggesting that bacteria enter these sites

first (31, 225, 301). However, additional studies by López et al. concluded that the meningeal vasculature is the most likely site of bacterial invasion, rather than the choroid plexus, based on the sequence of upregulation of intercellular cell adhesion molecule 1 (ICAM-1) and P-selectin on the cerebral endothelium following subcutaneous injection of *L. monocytogenes* (226). By comparison, rhombencephalitis is a prominent feature in mice infected by repeated gastrointestinal inoculation or in gerbils that develop chronic, low-density bacteremia secondary to otitis media (7, 8, 42). In contrast, injection of bacteria into the carotid artery of goats produced lesions in the meninges, the ependyma of the cerebral ventricles, and the aqueduct (17). These manifestations are similar to those caused by extracellular bacteria such as *S. pneumoniae* that produce a high-density bacteremia during clinical infection (419).

Direct invasion of endothelial cells. Pathological analyses demonstrate that *L. monocytogenes* can infect human endothelial cells in vivo. In a fatal case of meningoencephalitis, *L. monocytogenes* organisms were found within vascular endothelial cells, and one bacterium was attached to the luminal surface of a vessel (200). In addition, vasculitis and neutrophilic infiltration of the umbilical vein were identified in a 28-week fetus with congenital listeriosis, suggestive of infection of the endothelium (95). This latter report shows that human umbilical vein endothelial cells (HUVEC), cells that are commonly used in the laboratory but not regarded as in vivo targets, may be infected in naturally occurring disease and elicit neutrophil recruitment as also shown in vitro (91, 212). Nevertheless, widespread vasculitis with readily demonstrable organisms in endothelial cells in the periphery or in the CNS, such as is common in rickettsial diseases, is not common in *L. monocytogenes* infection. In addition, careful histological analysis of CNS lesions in ruminants showed that perivascular lesions typically do not contain bacteria (72, 183). However, it is possible that endothelial cell infection is an early event, the signs of which become less evident as the disease progresses, and as such is not found on postmortem examination.

In vitro data showing that *L. monocytogenes* can invade and replicate within endothelial cells were first obtained by using cultured HUVEC as the target cells (94, 151). Later, bacterial infection was also found in primary human brain microvascular endothelial cells (164, 410) and in simian virus 40 transformed human brain microvascular endothelial cells (150). Endothelial invasion requires intact microfilament and microtubule function, as shown by near complete inhibition of invasion by cytochalasin D and nocodazole, respectively (92, 150, 410). Wild-type *L. monocytogenes* organisms as well as nonpathogenic *Listeria* species bind and enter endothelial cells in vitro, as shown by gentamicin protection assays (91, 150). *L. monocytogenes* possesses many adhesins that may play roles in mediating these interactions (50, 135), but most experimental work to date has focused on the role of proteins of the *L. monocytogenes* *inlAB* operon in binding to and invasion of endothelial cells. Initial findings using an *L. monocytogenes* strain with a transposon-induced mutation in *inlA* suggested a role for internalin A (InlA) in binding to and invading HUVEC under serum-free conditions (94). In apparent contrast, studies using in-frame deletion mutants of *L. monocytogenes* that lack *inlA*, *inlB* or both (Δ *inlAB*) showed that *inlB* played a key role in

both binding and invasion of HUVEC and transformed human brain microvascular cells (150, 286), whereas other studies showed that *inlB* promoted invasion but not adhesion (34, 149). The mechanism for triggering invasion is probably due to the interaction of internalin B (InlB) with the hepatocyte growth factor receptor, Met, triggering phosphorylation of this receptor tyrosine kinase (354). Along these same lines, recent data indicate that gene products of the bacterial *inlGHE* gene cluster suppress InlB-mediated invasion, perhaps by competing for the same receptor (34).

The discrepancies in the experimental evidence showing the importance of InlA versus InlB for endothelial invasion may be caused by several factors. The initial findings supporting a role for InlA could have been compromised by a polar effect of the transposon inhibiting the expression of InlB. Furthermore, different mutants of *L. monocytogenes* were used, as well as different endothelial cells and differential experimental conditions. In particular, the presence or absence of serum has a profound influence on the mechanisms of binding of bacteria to these cells (94). This was demonstrated by experiments conducted in the presence of serum that showed InlA- and InlB-independent invasion of HUVEC and primary brain microvascular endothelial cells (BMEC) (151, 410). It is known that antibodies to surface proteins of *L. monocytogenes* are found in sera from humans following infection with *L. monocytogenes* and also in normal human sera (128, 208). Recent experiments by Hertzog et al. showed that normal human sera, and particularly the immunoglobulin G (IgG) fraction, strongly inhibited *L. monocytogenes* invasion of human brain microvascular endothelial cells, a process dependent on InlB in the absence of serum (164). Given that endothelial cells and extracellular bacteria in the bloodstream are continuously bathed by plasma proteins in vivo, these data question whether InlB-mediated mechanisms play a significant role in the establishment of CNS infection in adult humans. Moreover, although newborn serum also demonstrated inhibitory activity, it was approximately 50-fold less potent than the same concentration of adult serum. Concentrations of >1% were not tested, but this finding is consistent with the extreme susceptibility of the neonatal CNS to invasion by *L. monocytogenes*. The cause of the discrepancy between adult and neonatal serum in the inhibition of binding is not evident because maternal IgG is actively transported across the placenta and into the fetal circulation. At least two possibilities may be considered. First, purified IgG in amounts comparable to a given concentration of serum was not as potent as the serum itself, suggesting that other proteins, e.g., IgA, could also be involved. Second, different IgG subclasses are not transported across the placenta with equal efficiency, and it is possible that the anti-InlB antibodies are of a subtype that does not achieve a sufficiently high concentration in the neonatal circulation (107, 110).

L. monocytogenes invasion of endothelial cells triggers a robust cellular and has recently been reviewed (390). In brief, the NF- κ B signal transduction pathway is activated, as are phosphatidylinositol-induced lipid mediator generation and diacylglycerol generation (356, 357). In conjunction, there is increased surface expression of the adhesion molecules P- and E-selectin, ICAM-1, and VCAM-1 (91, 193, 212, 410). Infected endothelial cells also produce IL-6 (151) and the chemoattractants IL-8 and monocyte chemoattractant protein 1 (MCP-1) (151,

193). Increased adhesion molecule and chemokine expression leads to increased neutrophil and monocyte adhesion to infected HUVEC and BMEC monolayers (91, 212, 410). A critical role in endothelial cell activation is played by LLO. This is shown by experiments in which purified LLO stimulates endothelial cells while LLO-deficient *L. monocytogenes* mutants and nonhemolytic *Listeria* species are significantly less able to trigger adhesion molecule upregulation and mediator release compared to wild-type *L. monocytogenes* (91, 193). Moreover, transformation of *L. innocua* with the LLO-encoding *hly* gene from *L. monocytogenes* allows these normally nonhemolytic bacteria to produce LLO and also enables them to stimulate endothelial cells in a fashion comparable to that of wild-type *L. monocytogenes* (193). Recent studies on the mechanism by which LLO activates NF- κ B in transfected human embryonic kidney cells (HEK-293) show that LLO activates the I κ B kinase complex to phosphorylate and degrade cytoplasmic I κ B α (192).

Invasion via infected phagocytes. The second mechanism for CNS invasion by *L. monocytogenes* is via infected phagocytes. This is an extension of the Trojan horse model established in studies of neuroinvasion by visna virus in sheep (287) and more recently has been applied to the entry of human immunodeficiency virus type 1 (HIV-1)-infected cells into the CNS of humans (279, 421). The basic concept is that leukocytes are infected in the periphery and then transport intracellular microbes to the CNS and across the blood-brain or blood-choroid barriers. This idea was first applied to a bacterial pathogen when proposed as a means for *Streptococcus suis* to cross the choroid plexus of swine (407). The likelihood that this mechanism applies to *L. monocytogenes*-induced pathogenesis derives in part from the facts that these bacteria are so well adapted to the intracellular environment that intracellular parasitism is absolutely required for pathogenesis. The latter point is demonstrated by the finding that *L. monocytogenes* mutants that lack *hly*, and thus do not escape the phagosome and replicate intracellularly, are also more than 1,000-fold less virulent in vivo (122). Moreover, intracellular transport of *L. monocytogenes* from the gut to the lymph nodes by dendritic cells or other phagocytes appears to be involved in the early in vivo dissemination of bacteria following gastrointestinal infection (239, 302). This demonstrates that *L. monocytogenes* also utilizes host phagocytes for trafficking at another crucial step in its pathogenesis.

The notion that parasitized phagocytes are present in the CNS in human disease is supported by the presence of F-actin-coated bacteria within macrophages from a case of human rhombencephalitis (200). Similarly, *L. monocytogenes*-infected phagocytes were identified in the peripheral blood (90) and the ventricular system of the brain (301) in experimentally infected mice. Differential quantification of cell-free and cell-associated bacteria in whole blood revealed that approximately 30% of bacterial CFU present were isolated with leukocytes (90). In these experiments, more than 90% of cells associated with bacteria were mononuclear cells, suggesting that neutrophils did not participate in dissemination of intracellular bacteria. Further characterization of these cells showed that nearly all of them were CD11b⁺, supporting their identification as monocytes (D. A. Drevets, P. J. M. Leenen, T. Nikolic, M. J. Dillon, J. E. Schawang, and C. Sunderkötter, Program Abstr. 43rd

Intersci. Conf. Antimicrob. Agents Chemother., abstr. B-1048, 2003). Moreover, the viability and virulence of cell-associated bacteria was established by showing that infected peripheral blood leukocytes initiated disseminated infection, including CNS infection, when transferred into recipient mice. In addition, bacteria were capable of spreading directly from parasitized peripheral blood leukocytes to endothelial cells in vitro (90). Taken together, these data show that *L. monocytogenes*-infected cells are present in the peripheral circulation and in the CNS of experimentally infected mice and that cell-associated bacteria retain virulence. Nevertheless, it was not clear whether these cells played a key role in the initiation of CNS infection or were merely present but irrelevant in terms of pathogenesis.

A recent study tested whether leukocyte-associated bacteria could establish CNS infection in the absence of extracellular bacteria. This was accomplished by treating *L. monocytogenes*-infected mice with a continuous infusion of gentamicin delivered by surgically implanted osmotic pumps (93). Gentamicin is an antibiotic that is rapidly bactericidal against extracellular *L. monocytogenes* but crosses cell membranes poorly and is not concentrated in the cytoplasm of cells (166). This antibiotic is commonly and effectively used in vitro to kill extracellular *L. monocytogenes* (see, for example, references 94, 148, and 300), but because intracellular replication and cell-to-cell spread continue unabated, its role in the treatment of human infections is limited to use in combination with other antimicrobials (166). Mice with gentamicin-secreting pumps had serum and plasma gentamicin concentrations that were 16- to 256-fold greater than the MIC for the bacterial strains used for infection (93). Gentamicin-treated animals had 17-fold fewer CFU bacteria in the bloodstream at the time of sacrifice, but essentially all bacteria were cell associated, compared with only 26% in untreated mice. Nevertheless, the quantity of bacteria in the brain was similar in gentamicin-treated and untreated animals. Additionally, brain infection developed in gentamicin-treated mice infected by oral inoculation. Single-cell analysis of blood leukocytes from mice infected with an *actA-gfpuv-plcB*-transfected *L. monocytogenes* in which there is a 200 to 500-fold increase of *gfpuv* expression when bacteria escape from phagosomes to the cytosol (120) confirmed that intracellular bacteria were in fact parasitizing circulating phagocytes. Taken together, these data demonstrate that *L. monocytogenes* parasitizes circulating phagocytes in vivo and indicate that CNS infection can be established by infected leukocytes in the absence of viable cell-free bacteria.

It is not known whether *L. monocytogenes*-infected phagocytes are recruited to the CNS by specific or nonspecific mechanisms or whether infected phagocytes have altered migratory characteristics. For the endothelial counterpart in the CNS, it has been shown that it is activated for leukocyte adhesion prior to bacterial invasion as evidenced by upregulation of ICAM-1 on large veins, arterioles, subarachnoid venules, and capillaries and de novo P-selectin expression on subarachnoid vessels and to a lesser extent on parenchymal vessels and in the choroid plexus (226). The mechanism for this endothelial activation involves LLO-dependent triggering of NF- κ B nuclear translocation in cerebral vessels (193). In vitro studies show that once infected phagocytes adhere to endothelium, bacteria can invade endothelial cells by cell-to-cell spread in an *hly*- and

actA-dependent process (90, 94, 150). After invading endothelial cells, bacteria presumably continue to spread deeper into the CNS parenchyma, thus causing the characteristic meningoencephalitis of *L. monocytogenes*. The finding that $\Delta plcB2$ mutants of *L. monocytogenes* which are deficient in the phosphatidylcholine-specific phospholipase C, but not $\Delta inlAB$ mutants have reduced virulence following intracranial injection suggests that cell-to-cell spread rather than direct invasion occurs (341). As an alternative to adhering to and infecting the endothelium, infected phagocytes could transmigrate and enter the brain tissue. In this case, bacteria contained within phagocytes could spread to cells such as neurons and microglia (89).

Invasion by the neural route. The third route of CNS invasion, infection via neural transport, was explored several decades ago by Asahi et al. These investigators injected *L. monocytogenes* into the oral membranes of mice and goats or fed the animals with abrasive food that had been soaked in bacterial cultures (18). Using these techniques, 47 (72%) of 65 injected mice and 13 (65%) of 20 mice fed dried boiled fish developed signs of brain stem encephalitis 6 days after infection. The remaining animals either died of systemic disease prior to day 6 (some of these also had CNS lesions at autopsy) or remained asymptomatic. Examination of the brain and cranial nerves revealed mononuclear infiltrations along the length of the trigeminal nerve from the distal end in the lips to the proximal terminus in the medulla, in addition to neutrophilic perivascular cuffing throughout the brain stem. Similarly, injection of bacteria into the lips of goats produced neurological disease in 4 of 11 animals and feeding them with bean shells soaked in bacteria produced symptoms in 2 of 8 animals. Histological analysis also showed lesions similar to those in the mice, in particular a mononuclear infiltrate in the trigeminal nerve from the maxillary and mandibular branches to the medulla, usually with focal meningitis. These pathological findings were very similar to those for naturally occurring cases of *L. monocytogenes* encephalitis in sheep and goats. These authors concluded that transport of *L. monocytogenes* from the oral cavity to the brain stem via the trigeminal nerve was the likely mechanism by which CNS disease was established in ruminants.

Some of these conclusions were reconfirmed in a subsequent study by Cordy and Osebold, who found encephalitis in 33% of mice infected by oral scarification (72). They also found that 80% of mice that died early, with a median time to death of 4.9 days, did die of systemic disease without histological encephalitis. In contrast, mice developing encephalitis had a mean time to death of 9.3 days, but fewer than 10% of these mice showed evidence of neuronal involvement. Although the authors did find infected trigeminal nuclei and roots in 4 of 15 field cases of sheep, they suggested that mononuclear infiltrates in the nerve sheaths, including the trigeminal nerve, could have resulted from centrifugal spread of inflammation from the surrounding meninges. Subsequent studies by others using electron microscopy identified intra-axonal bacteria in cases of spontaneous *L. monocytogenes* encephalitis in sheep (58, 59, 283). This finding was reproduced by experimental inoculation of the *L. monocytogenes* into the dental pulp of sheep, in which case the animals developed brain stem encephalitis on the same side as the dental infection (28). In addition, recent studies of mice confirmed a direct relationship between

the site of neuronal infection and the resulting CNS disease. In these experiments, rhombencephalitis developed ipsilateral to the site of *L. monocytogenes* injection into the facial muscle and the cut end of the facial nerve, whereas transverse myelitis followed injection into the triceps surae muscle and the cut sciatic nerve (12). Myelitis could be prevented by surgical disruption of the sciatic nerve proximal to the site of bacterial injection. Furthermore, bacteria could be identified in axons and within mononuclear cells in the peripheral nerve fascicles. Taken together, these data clearly indicate that bacterial conveyance along a neural route can ultimately result in brain stem encephalitis or transverse myelitis.

In vitro studies showed that invasion of neurons by cell-free *L. monocytogenes* is a rare event (290, 291), and until recently it was unclear how *L. monocytogenes* entered neurons from infected tissue. In vitro studies by Dons et al. compared the infectivity of different types of neurons and showed that this characteristic differs significantly among them (88). They found that cultured rat dorsal root ganglion neurons could be invaded efficiently by internalin-dependent mechanisms whereas hippocampal neurons were resistant to this process. In addition, bacteria were found to move in antegrade as well as retrograde directions. Dramsi et al. also found that rat spinal chord neurons were resistant to direct bacterial invasion (89). However, they showed that these neurons were efficiently invaded via cell-to-cell spread by bacteria from infected phagocytes. Jin et al. subsequently tested the mechanism of cell-to-cell spread of bacteria from phagocytes to neurons in vivo. These investigators compared the bacterial loads in the trigeminal ganglion and brain of *RAG-1*-deficient mice injected in the snout with wild-type *L. monocytogenes* or with $\Delta plcB2$ mutants, bacteria which are impaired in phagosomal escape and cell-to-cell spread (182). The results showed that mice infected with $\Delta plcB2$ mutants had a significantly longer survival time and lower bacterial counts in the trigeminal ganglion and brain than did mice infected with wild-type bacteria, even though equivalent bacterial loads had been injected into the snouts of the two groups. In addition, bacteria were identified within MHC class II-expressing cells, presumably tissue histiocytes, in the snout by immunofluorescence microscopy. The authors interpreted these data to indicate that phagocytes, not neurons, are the first cells infected in the snout injection model and that intercellular spread between the infected phagocytes and neurons was important for initiating CNS infection. This interpretation was supported by a second study in which macrophage colony-stimulating factor-deficient *op/op* mice were infected in the snout and then the bacterial loads in the trigeminal ganglion and in peripheral organs were compared with those in *op/+* controls (181). The results showed that *op/op* mice had fewer bacteria in the brain despite being more susceptible to systemic infection. The finding that *op/op* mice developed CNS infection at all could be explained by systemic spread of bacteria or by transfer via macrophage colony-stimulating factor-independent macrophages present in the tissues (273).

Taken together, these data provide a clear picture of certain events in the neuronal route of CNS invasion. Bacteria inoculated into the oral tissues are initially taken up by tissue macrophages and/or by mononuclear phagocytes that are rapidly recruited from the circulation. Some phagocytes are par-

asitized, and bacteria from them invade, presumably from cell-to-cell spreading, the distal ends of nerves that form branches of the trigeminal ganglion. Bacteria move in a centripetal fashion through the trigeminal ganglion into the brain stem. It is not clear if this mobility requires the *L. monocytogenes* actin-polymerizing machinery, since inert beads also can be transported within axons (403). Once bacteria arrive in the brain stem, encephalitis is established by continued cell-to-cell spread. This mechanism of infection is clearly important in ruminants, which may chew material coarse enough to abrade the oral mucosa and in which *L. monocytogenes*-contaminated silage has been linked to epidemics of encephalitis (230, 389). However, the relevance of the neuronal route in humans is less clear because most of the foodstuffs linked to human diseases, e.g., salads (22, 334), meat products (57a, 347), jellied pork tongue (49, 141), milk (119), and cheeses (179, 224), do not have the abrasive qualities required to inoculate submucosal tissues with sufficient numbers of bacteria to cause disease. Whether gingivitis or dental caries could be a contributing factor that enables this mechanism to operate in humans is not known.

Brucella Species

Brucellae are small, nonmotile, and non-spore-forming gram-negative coccobacilli. They are generally considered to be facultative intracellular parasites of professional and non-professional phagocytes. However, recent data indicate that *Brucella* spp. are better adapted for intracellular survival than in the extracellular environment and, as such, should be thought of as facultative extracellular pathogens rather than the converse (266). The intracellular niche of virulent brucellae is within membrane-bound endosomes that differ somewhat between professional and nonprofessional phagocytes. In phagocytic cells, the *Brucella*-containing phagosomes acidify but do not fuse with lysosomes, so that the bacteria reside in and replicate in specialized endosomes recently called "brucellosomes" (14, 207, 325). By comparison, in nonprofessional phagocytes, early endosomes that contain *Brucella* enter the autophagocytic pathway and bacteria replicate within autophagosome-like compartments in the endoplasmic reticulum (294, 295). Recent data show that the *VirB* operon, a type IV secretion pathway that is induced on phagosomal acidification, plays a key role in intracellular parasitism and is essential for pathogenicity (280, 359). It is hypothesized that this system secretes effector molecules that alter the host cell endosomal pathways to promote the formation of an endoplasmic reticulum-derived compartment that is conducive to the replication of Brucellae rather than to fusion with lysosomes in a degradative pathway (57, 258).

There are six nomen species in the genus *Brucella*; however, human disease is caused predominantly by *B. abortus*, *B. melitensis*, and *B. suis* (70). In addition, two cases of human neurobrucellosis were reported recently that were caused by a *Brucella* sp. closely related to *B. pinnipediae*, a bacterium found in marine mammals (365). The history of brucellosis is interesting in that bone lesions consistent with *Brucella* infection have been identified in 17.4% of adult skeletal remains from Herculaneum, a Roman town covered by the Mount Vesuvius eruption in 79 AD (52). In more modern times, systemic in-

fection caused by *Brucella* spp. was referred to as Malta fever or undulant fever, and its etiologic agent was isolated from humans and identified in 1886 as *Micrococcus melitensis* (46). A wide variety of acute and chronic CNS infections caused by *Brucella* spp. were recognized soon thereafter (78).

Pathogenic brucellae are found worldwide, and the incidence of clinical disease in humans varies greatly. It has been nearly eradicated from some countries, but in areas of endemic infection such as Peru, Kuwait, and Saudi Arabia, there are more than 200 cases/100,000 population (69). Human infection typically is acquired by ingestion of unpasteurized milk or cheese or through occupational exposure to infected animals, in particular sheep, goats, swine, camels, and cattle (16, 47, 69, 103, 233, 269). There is no obvious predilection for individuals with underlying diseases (45). The frequency with which *Brucella* attacks the CNS is relatively low, from 1.3 to 7% of total cases in recent series, and neurobrucellosis is found much less commonly in children than in adults (45, 157, 233, 253). The mortality rate of neurobrucellosis in the postantibiotic era is 0 to 5.5%, but permanent neurological deficits, particularly deafness, are common (27, 253, 270).

Signs and symptoms referable to CNS involvement typically include headache with or without meningeal irritation and can occur early in the course of brucellosis or more than 1 year after the onset of systemic symptoms (27, 253, 270, 276, 353). Neurobrucellosis can result in a multitude of nervous system manifestations that can only be summarized here. The most common presentation is as a typical meningitis or meningoencephalitis that has an acute onset and can occur either as the only site of infection or in the context of systemic disease (45, 253, 353). Patients with acute infection can have cranial nerve palsies that usually resolve completely with antibiotics, whereas those with chronic CNS infection often have permanent neurological deficits. The various chronic manifestations are perhaps best divided into those presenting with peripheral neuropathy or radiculopathy and those presenting with more diffuse CNS involvement that includes myelitis with cranial nerve involvement and a syndrome of parenchymatous dysfunction (253, 353). Symptoms of the peripheral neuropathy and radiculopathy include back pain, areflexia, and paraparesis with involvement of the proximal nerve radicals. In patients with diffuse CNS involvement, myelitis is evidenced by back pain, spastic paraparesis, and demyelination and can also occur with cerebellar dysfunction. The syndrome of parenchymatous dysfunction can occur at any location in the CNS but most commonly affects the cerebellum, spinal cord, and cerebral white matter. Meningovascular complications, in particular mycotic aneurysms, ischemic strokes, and subarachnoid hemorrhage, are relatively common (45, 118, 159, 253).

Examination of the CSF typically reveals an elevated protein concentration, a depressed glucose concentration, and a moderate leukocytosis composed mainly of lymphocytes (27, 253, 270, 353). The exception is the cerebellar syndrome, in which the protein concentration is elevated but there is no leukocytosis. Cultures of CNS tissue and fluid are frequently sterile; however, bacteria are occasionally recovered from the CSF and from the brain granuloma. Pathological changes of the meninges are almost always found. They are typified by diffuse involvement manifested by thickening due to acute and chronic inflammatory cell infiltration as well as connective cell prolifer-

eration that is sometimes more prominent in the basal areas (118, 276). Granulomatous inflammation of the meninges and of the brain parenchyma is also found and can be accompanied by central necrosis (276, 365). In the spinal cord, there is chronic inflammatory cell infiltration into the parenchyma and the overlying arachnoid membrane as well as areas of demyelination. Vascular inflammation is present and ranges in character from minimal and mononuclear to massive and neutrophilic; it can lead to significant adventitial thickening, necrosis, and the formation of aneurysms. Furthermore, there is inflammation of the perineurium of nerve roots in the peripheral nervous system that probably accounts for the radiculopathies (253, 276).

The precise mechanisms by which brucellae enter the CNS are not known. The extreme degree to which brucellae are adapted to the intracellular environment suggests that phagocyte-facilitated infection is a likely mechanism. Consistent with this, it has been shown that *B. suis* prevents apoptosis of infected and uninfected human monocytes in vitro and that monocytes and lymphocytes taken from *Brucella*-infected patients are resistant to spontaneous apoptosis and to apoptosis triggered by anti-Fas monoclonal antibody (152, 382). This would prolong the life span of infected cells and in theory would give more opportunity for them to enter the CNS. In addition, brucellae can invade nonprofessional phagocytes, and so it is likely that endothelial cell invasion could be another possible route for CNS infection; however, this has not been studied. In terms of potential mechanisms for entry into non-professional phagocytes, a two-component regulatory system, BvrR-BvrS, that is important for invasion and intracellular parasitism has been identified in *B. abortus* (366). Transposon-induced inactivation of *bvrR-bvrS* inhibits the invasion of mouse peritoneal macrophages and HeLa cells, and mutants that are internalized do not inhibit phagolysosome fusion. Moreover, BvrR-BvrS mutants are avirulent in mice. Further studies showed that this system regulates the expression of *B. abortus* outer membrane proteins, in particular Omp3a (formerly known as Omp25) and Omp3b (155). Omp3a/Omp25 probably plays an important role since mutants of *B. abortus*, *B. melitensis*, and *B. ovis* that lack this protein are less pathogenic in mice, cattle, and goats (104–106). Internalization of *Brucella* spp. by cells occurs by a zipper-like mechanism involving as yet unidentified cellular receptors, in which sialic acid-mediated recognition by a bacterial lectin could play an important role (79). Internalization of bound bacteria requires recruitment of F-actin and, in nonprofessional phagocytes, activation of GTPases of the Rho subfamily (154, 325).

Mycobacterium tuberculosis

M. tuberculosis is a member of the *M. tuberculosis* complex along with *M. africanum*, *M. bovis*, and *M. microti* (121). The characteristic staining quality of these organisms is their ability to resist decolorization of carbol fuchsin by acid-alcohol in the Ziehl-Neelsen process due to the high lipid content of the cell wall. *M. tuberculosis* is an aerobic, nonmotile, non-spore-forming bacillus that also stains weakly gram positive. The main cellular reservoirs of these organisms in the host are tissue macrophages, although the organisms also exist extracellularly. There are numerous macrophage receptors capable of binding

M. tuberculosis that can participate in its internalization, including CD11b, CD14, the macrophage mannose receptor, scavenger receptor A, Toll-like receptor 4, and CD44 (reviewed in references 36, 108, 219, 292, and 392). Intracellular parasitism is accomplished by a variety of mechanisms that include altered trafficking of bacteria during endocytosis, interference with host Ca²⁺ signaling pathways, and induction of maturational arrest of the phagosome (reviewed in references 82, 293, and 332). The result is that bacteria reside in a membrane-bound compartment which has an elevated pH and does not contain mature lysosomal enzymes.

Tuberculosis is a disease whose history and impact on humanity are impossible to underestimate (331). Morphological and molecular evidence of *M. tuberculosis* infection has been found in the fossilized remains of Pleistocene era bison that lived approximately 18,000 years ago in North America (329) and in human mummies from ancient Egypt and pre-Columbian Peru (209, 336, 420). Green published the first description of tuberculous meningitis as a specific entity in 1836 (147), approximately half a century prior to Koch's discovery in 1882 that *M. tuberculosis* was the causative agent of human tuberculosis (204). In the vast majority of human cases, tuberculosis is acquired through inhalation of bacillus-containing droplets into the lungs (reviewed in reference 26). Bacilli are then transported to draining lymph nodes, and the ensuing inflammatory response eventually leads to the formation of the characteristic granuloma and to activation of a type 1 T-helper cell-mediated immune response. During the early stage of infection prior to the development of a containing response, there is a low-titer bacteremia in which bacteria disseminate to distant sites. Active tuberculosis develops when containment fails so that bacillary replication and caseous necrosis occur in preexisting granulomas (332). Pulmonary involvement is the predominant form of clinical tuberculosis and usually accounts for 80 to 90% of cases, depending on the era in which the data were collected. Extrapulmonary tuberculosis, i.e., disease outside the lungs, can be found in nearly every organ system. In the pre-AIDS era, extrapulmonary tuberculosis was found in 10 to 20% of cases of active tuberculosis. Meningitis was the predominant feature in 5 to 8% of extrapulmonary cases, so that the overall estimation of CNS infection was approximately 1% of active cases (9, 115, 195a). The presence of HIV-1-induced immunodeficiency, however, is a powerful cofactor for the development of disseminated tuberculosis, and therefore the proportion of extrapulmonary cases has increased (for reviews, see references 156 and 352). Nevertheless, whether HIV-1 coinfection also increases the occurrence of meningitis is not completely clear. In a study of 2,205 patients diagnosed with tuberculosis, Berenguer et al. identified meningitis in 10% of HIV-1-coinfected patients compared with 2% of non-HIV-1-coinfected individuals (32). Supporting this finding of an increased prevalence of meningitis, an autopsy study from Kenya found meningeal involvement in tuberculosis in 26% of HIV-infected adults (305). However, other studies have reported that HIV-1-infected and uninfected patients have similar rates of CNS tuberculosis (352, 374). Current mathematical models suggest that tuberculosis caused 8.3 million new cases worldwide in 2000 (71). Hence, although the exact incidence of tuberculous meningitis is unknown, these data imply

that *M. tuberculosis* is the most common cause of CNS infection by an intracellular bacterium.

CNS infection by *M. tuberculosis* occurs in individuals of any age. Large autopsy series of patients dying of tuberculosis identified CNS involvement in 19.3 to 42.2% of pediatric cases compared with only 2.9 to 5.9% of adult cases (21, 234, 321). Most cases in children occurred between the ages of 6 months and 4 years, whereas adult cases clustered in patients aged 20 to 50 years. In addition to HIV-1 coinfection, underlying conditions associated with tuberculous meningitis include recent measles infection and malnutrition in children (416) and alcoholism, malignancies, and immunosuppressive medications in adults (202, 281, 393). In terms of prevention of CNS infection, *M. tuberculosis* is exceptional in that it is the only intracellular bacterium for which a vaccine, i.e., BCG, is available that has a protective effect against meningitis (327). Nevertheless, the salutary effect of BCG vaccination against tuberculous meningitis is found only in children, not in adults, and some authors suggest that it merely delays the age of onset of meningitis.

M. tuberculosis infection of the CNS has several manifestations, including involvement of the meninges, space-occupying lesions in the brain parenchyma, and focal disease of the spinal chord and its osseous structures (for reviews, see references 123, 265, 369, and 380). Meningitis is the most common form of CNS disease and is discussed in depth here. The typical presenting signs and symptoms include fever, headache, and altered mental status, ranging from lethargy to coma (136, 188, 281, 298, 393). Meningismus, seizures, and focal neurological signs, in particular cranial nerve palsies, are also common. Spinal fluid analysis usually reveals an elevated protein concentration, hypoglycorrhachia, and a moderately (typically <500 cells/mm³) increased leukocyte count with lymphocyte predominance. Nevertheless, patients with normal CSF or CSF that shows a neutrophil predominance are occasionally found; most of these individuals have impaired host defenses (188, 214). Interestingly, most series comparing tuberculous meningitis in patients with and without HIV-1 infection do not find striking differences between the two groups of patients in the clinical features of the disease or in the CSF analysis (32, 188, 190, 346, 383). In contrast, HIV-1-coinfected individuals do show more frequent abnormalities such as space-occupying lesions or hydrocephalus on brain-imaging studies. Analysis of cytokines and chemokines in the CSF of patients with tuberculous meningitis shows elevated levels of IL-1 β , TNF- α , IL-8, MCP-1, and macrophage inflammatory protein-1 α (MIP-1 α), in amounts that are usually below those found in patients with bacterial meningitis but remain elevated for a much longer period after treatment (4, 245, 246, 417). Acid-fast bacteria have been recovered from the CSF in as many as 91% of centrifuged specimens (370), but recovery is usually much lower than that due to low concentrations of bacteria in the CSF. The mortality rate of tuberculous meningitis is substantial, and survivors often have permanent neurological deficits. Untreated patients usually die within 3 weeks of the onset, but effective antituberculosis treatment had lowered the mortality rate to 10 to 20% in the pre-AIDS era (265). By comparison, more recent studies report mortality rates ranging from 40 to 70% (137, 188, 298). A correlation of higher mortality with HIV-1 coinfection is sometimes present (190, 383).

The pathological features of tuberculous meningitis have

been extensively reviewed and offer the best understanding to date of how the CNS becomes infected (20, 33, 74). A thick, gelatinous exudate covers the surface of the brain and is most abundant on the basilar surface in the optic chiasm and within the ventricles. Dilatation of the ventricles is common, due in part to obstruction of the normal CSF flow by the protein-rich exudate. The fibrinous exudate contains a variety of cells including erythrocytes, neutrophils, and macrophages in early lesions, whereas lymphocytes are more common in older lesions. In the gelatinous exudate, there are focal areas of caseous necrosis with surrounding epithelioid cells and giant cells. The inflammatory process in the meninges typically extends through the perivascular spaces into the cortex, resulting in foci of encephalitis properly described as a diffuse meningoencephalitis. The critical lesions in terms of pathogenesis are called "Rich foci" and were first described in the early 1930s (318). These are small nodules, about 1 mm in diameter, which follow a vascular distribution. They are found most frequently in the meninges but also are present in the ependyma and the brain parenchyma. Interestingly, Rich foci are not found more frequently in the basilar areas of the brain than in other locations, and the localization of the exudate to this area is thought to be due to the normal pattern of CSF flow (234, 318). Microscopically, there is an accumulation of inflammatory cells in the subendothelial region of the intima of the blood vessel, with formation of a nodule in the adventitia. A detailed study of blood vessels in tuberculous meningitis by Winkelmann and Moore suggested that the earliest lesions begin in the subendothelium with an infiltration of lymphocytes, monocytes, and plasma cells (411). As the lesion progresses, there is cellular proliferation with displacement of the intact endothelium with or without tubercle formation. In later stages, the vessel is completely involved with a contiguous inflammatory process that began within it or was the result of direct extension of inflammation from the subarachnoid space into the vessel. This process can lead to complete occlusion of the vessel.

Early investigators confirmed that there was blood-borne dissemination of *M. tuberculosis*. Data first reported by Villemin in 1868, and subsequently confirmed by others, showed that injection of blood from experimentally infected animals, or from human patients, transmitted tuberculous infection to recipient animals (66). However, until the pioneering work of Rich and McCordock in 1933, it was thought that tuberculous meningitis was the "direct and immediate result" of hematogenous spread of bacilli during acute miliary tuberculosis (234, 318). These investigators showed that intravenous and intra-arterial injection of large amounts of the virulent H37RV strain of *M. tuberculosis* into rabbits and guinea pigs did not produce meningitis or meningeal lesions despite the presence of widespread visceral miliary disease (reviewed in reference 21). In addition, autopsy studies of adult humans showed that meningitis also occurred in the absence of miliary involvement elsewhere and that miliary tuberculosis did not always lead to meningitis or even to frequent tuberculous lesions in the CNS vasculature (318). These observations led to the hypothesis that hematogenous dissemination of bacilli from primary foci caused vascular lesions in the meninges and elsewhere but this event was temporally separated from bacillary invasion of the CSF and meningitis. The latter events happened when a vascular lesion ruptured and subsequently introduced bacteria

into the CSF. This hypothesis was supported by the identification of CNS vascular foci (Rich foci) in 87 to 94% of patients with tuberculous meningitis (21, 234, 318), as well as in individuals dying of tuberculosis who did not have histological evidence of meningitis (234). Subsequent experiments showed that direct inoculation of acid-fast bacteria into the CSF could replicate many of the pathological findings of natural cases, and that TNF- α is a key proinflammatory cytokine once bacteria enter the CSF (318, 385). Recently, this sequence of events has been demonstrated in experiments by Bolin et al., using swine infected intravenously with *M. bovis* (43). In these experiments, three of six pigs developed meningitis 17 to 38 days after infection. Postmortem examination of the brains identified granulomatous lesions, some of which also contained acid-fast bacteria, in the walls of meningeal blood vessels. These data substantiate the central hypothesis that Rich foci develop soon after hematogenous dissemination and offer a new animal model for in vivo studies of the pathogenesis of tuberculous meningitis.

The mechanisms used by *M. tuberculosis* to disseminate and enter blood vessels to establish Rich foci is not completely understood. In theory, hematogenous dissemination of cell-free bacteria and dissemination of bacteria within phagocytes are both possible. With regard to the CNS invasion by cell-free bacteria, data from human autopsies and animal experiments cited above suggest that the vast majority of cell-free bacteria that enter the bloodstream, either from erosion of caseous foci into the vascular or lymphatic systems or by experimental injection, are removed by visceral organs. Direct interactions between mycobacteria and endothelial cells are not well characterized. By comparison, recent data indicate that trafficking of intracellular bacteria by phagocytes could play a key role. For example, experimental data show that alveolar macrophages and newly recruited monocytes are important in transporting mycobacteria from the alveolar spaces to the draining lymph nodes (37, 41, 378) and that dendritic cells also may participate in this process (126, 180, 375). In addition, data from intratracheally infected mice show that mycobacteria enter microfold cells (M-cells), specialized antigen-sampling cells that overlie the mucosa-associated lymphoid tissues that line the bronchial epithelium. The bacteria are then transported to draining lymph nodes by macrophage colony-stimulating factor-dependent cells, i.e., monocytes and most macrophages (378). Once mononuclear phagocytes are infected, a number of physiological and phenotypic changes occur that could alter their cellular trafficking. For instance, infected monocytes and macrophages show increased expression of surface ICAM-1 (227), increased migratory activity across epithelial and endothelial bilayers (37), and activation of intracellular signaling pathways and secretion of proinflammatory cytokines (29).

Salmonella Species

Salmonella spp. are gram-negative, facultative anaerobic, motile, non-lactose-fermenting, non-spore-forming bacilli measuring 2 to 3 by 0.4 to 0.6 μm in size. All *Salmonella* isolates are currently classified as *S. enterica*, which is subdivided into more than 2,300 serovars according to somatic O, surface Vi, and flagellar H antigens. For historical reasons and

convenience, serovar names are often used as species names. Salmonellae are facultative intracellular bacteria that can invade and replicate within many different cells in vitro. Nevertheless, in vivo data show that intracellular survival of bacteria within macrophages is essential for virulence in mice (117) and that the preferred in vivo niche for *Salmonella* is within CD11b⁺ macrophages in the liver and the spleen rather than in the extracellular milieu (247, 319, 335). Intracellular *Salmonella* organisms replicate within a spacious phagosome, a modified, acidified endosome (6). This is due largely to the actions of virulence genes of the *Salmonella* pathogenicity island 2 type III secretion systems and the PhoP-PhoQ two-component regulatory system (reviewed in reference 168).

The genus is named after the U.S. veterinary surgeon Daniel E. Salmon, who first isolated *S. enterica* serovar Choleraesuis from porcine intestine in 1885. Ghon (134) and Cole (68) were the first to recognize human *Salmonella* meningitis in the first decade of the 20th century. *S. enterica* serovars Typhi and Paratyphi cause typhoid and paratyphoid fever, respectively, and colonize only humans. Therefore, they are acquired either by close contact with a person who is colonized or infected with these organisms or, more commonly, by ingestion of food or water contaminated by the excreta of such a person (262). It is estimated that there are at least 16 million new cases of typhoid fever globally each year, with 600,000 deaths (177). While it was a scourge of European and U.S. urban areas in the 19th century, most of the burden of typhoid fever is now in the developing world. The annual incidence of typhoid fever in an urban area of Kalkaji, New Delhi, India, has recently been reported as 980 per 100,000 (364). Incidence peaks in 5- to 12-year-old children, but children aged 1 to 5 years are also commonly infected; although less frequently infected, children younger than 1 year experience more severe disease (364). In the developed world, typhoid fever is largely a disease of travelers returning from areas of endemic infection, with occasional point source epidemics (3).

In contrast to *S. enterica* serovars Typhi and Paratyphi, nontyphoidal *Salmonella* (NTS) is a prime source of common gastroenteritis. It is widely dispersed in nature, including the gastrointestinal tracts of domesticated and wild mammals, reptiles, birds, and insects. In humans, NTS infections are most often associated with consumption of inadequately prepared food products of animal origin, including meat, poultry, eggs, and dairy products, although a variety of other foods have been implicated. In the United States, the annual incidence has doubled in the last 20 years. There are an estimated 1.4 million cases yearly, resulting in an estimated 16,430 hospitalizations and 582 deaths (254). Globally, by far the most common group with NTS illnesses are children younger 5 years (142). Host conditions associated with an increased risk of salmonellosis include gastric hypoacidity, extremes of age, alteration of the endogenous bowel flora, diabetes mellitus, malignancy, rheumatological disorders, reticuloendothelial blockade (sickle cell disease, malaria, or bartonellosis), and therapeutic immunosuppression (167). In addition, patients with HIV infection have an estimated 20- to 100-fold increased risk of salmonellosis and are more likely to have severe invasive disease (262).

Enteric fever (typhoid or paratyphoid fever) is a severe systemic illness characterized by abdominal symptoms and fever. After an incubation period of 5 to 21 days, diarrhea of several

days' duration may develop and typically resolves before the onset of fever. Nonspecific symptoms such as chills, diaphoresis, headache, anorexia, cough, weakness, sore throat, dizziness, and muscle pains are frequently present before the onset of fever. In addition to headache, CNS manifestations of enteric fever in the form of neuropsychiatric manifestations from encephalitis, including confusion and psychosis, occur in 5 to 10% of patients. *S. enterica* serovar Typhi meningitis has been reported but is very rare (67, 220). Mortality was 15% in the preantibiotic era but is now less than 1% in the United States (262). Gastroenteritis caused by NTS is not clinically distinctive. After an incubation period of 6 to 72 h (depending on the inoculum and host status), patients typically present with acute onset of fever, diarrhea, and abdominal cramping (167). There is a wide spectrum of severity of illness. Approximately 5% of patients with gastroenteritis develop bacteremia, and <1% have focal infections such as osteomyelitis, soft tissue infection, urinary tract infection or, endocarditis (262).

CNS infection is present in approximately 0.1 to 0.9% of cases (167). CNS infection by NTS is primarily a disease of children, particularly neonates and infants (218, 418). Only about 10 cases of *Salmonella* meningitis have been reported in adults (186, 220). In a recent review of 144 cases of NTS bacteremia in children, the median age was 10.5 months, and only 3 (2.1%) developed meningitis (418). In general, presenting symptoms and signs of *Salmonella* meningitis are fever (73%), meningismus (51%), vomiting or anorexia (43%), seizures (43%), lethargy or coma (34%), and irritability (12%) (the percentages are frequencies) (67). CSF findings include a neutrophilic pleocytosis, elevated protein concentration, hypoglycorrhachia, and, in 75%, a positive Gram stain (67). In a review of 13 infants with *Salmonella* meningitis, serious complications were common. In this series, 69% of infants had seizures, 38% developed hydrocephalus, whereas subdural effusions and empyema were present in 31 and 23%, respectively. Overall mortality in the last 40 years has ranged from 0 to 59% (13, 171, 191, 218, 231, 256). Neurological complications, including mental retardation, cerebral palsy, and visual and hearing impairment, have been found in 13 to 62% of survivors. In addition to typical meningitis, focal intracranial infections with NTS can occur with or without meningitis. Principal among these are subdural empyema and brain abscess (178, 213, 232, 236, 328, 337, 404).

It is unknown whether salmonellae invade the CNS primarily as cell-free or intracellular bacteria. With regard to direct interactions of bacteria with endothelial cells, *in vitro* data showed that smooth, wild-type *S. enterica* serovar Minnesota was poorly adherent to bovine pulmonary endothelial cells compared with a rough mutant and that phagocytosis of the latter bacterium by the endothelial cells was induced by opsonization with C1q (333). However, interactions between NTS and endothelial cells remain a poorly studied phenomenon. Trafficking of *Salmonella*-infected phagocytes is a likely but as yet unproven mechanism for CNS infection. In terms of trafficking of infected phagocytes from the blood into other anatomic spaces, monocytes bearing *Salmonella* antigens show increased binding to synovial endothelium and transendothelial migration, and *Salmonella* antigens have been identified within synovial cells in patients with reactive arthritis (143, 201). However, this does not necessarily indicate that infected

cells may go elsewhere, in particular into the CNS. Nevertheless, recent descriptions of the mechanisms by which leukocytes transport *Salmonella* across the gastrointestinal epithelium do provide a useful example for how phagocytes can transport bacteria across an epithelial barrier. Vazquez-Torres showed that CD18⁺ phagocytes transported *S. enterica* serovar Typhimurium mutants that lacked invasion genes across the gut epithelium in a process that bypassed both epithelial cells and M-cells (391). More recent studies showed that CD11b⁺ CD8α⁻ dendritic cells are rapidly recruited to the gut epithelium in the presence of intraluminal *Salmonella* (316, 317). These dendritic cells emerge between the epithelial cells by interacting with the tight-junction proteins claudin-1 and zonula occludens-1 and then extend dendrites into the lumen without disrupting the integrity of the epithelial lining (317). Thus, dendritic cells that sample luminal contents directly can internalize bacteria and then transport them to lymphoid tissues (169, 316, 317).

OBLIGATE INTRACELLULAR PATHOGENS

Rickettsiaceae

Human pathogens of the family *Rickettsiaceae* are obligate intracellular parasites of professional and nonprofessional phagocytes that invade the CNS as part of a systemic infection. They are for the most part arthropod borne and have a limited capacity to survive outside a host. The notable exception is *Coxiella burnetii*, an organism that does not have an arthropod intermediate host. It is efficiently transmitted via aerosol and survives well in the environment outside the host. This family of pathogens also varies with regard to their cellular targets and intracellular niches. The *Rickettsiaceae* can be divided into three categories according to their targets during natural infection: (i) organisms that parasitize vascular endothelial cells, (ii) organisms that parasitize phagocytes, and (iii) organisms that parasitize both endothelial cells and phagocytes (Table 2). The first category includes members of the *R. rickettsii* group, *R. rickettsii* and *R. conorii*, and the typhus group, *R. prowazekii* and *R. typhi* (350). The organisms that principally target phagocytes are *E. chaffeensis*, *Anaplasma (Ehrlichia) phagocytophila*, and *C. burnetii*. By comparison, only *Orientia tsutsugamushi* is routinely found in endothelial cells as well as in circulating phagocytes. In terms of their intracellular niches, *O. tsutsugamushi* and the rickettsiae lyse the phagosome and replicate predominantly in the cytoplasm of the host cell, whereas *E. chaffeensis*, *A. phagocytophila*, and *C. burnetii* remain in specialized vacuoles.

***Rickettsia* spp.** Rickettsiae are small (0.3 by 1 μm) gram-positive coccobacilli that are divided into the *R. rickettsii* and typhus groups, with *R. rickettsii* and *R. prowazekii*, respectively, being the prototypes of these groups. Rocky Mountain spotted fever (RMSF) is the clinical entity caused by *R. rickettsii*. The first identified case of RMSF was thought to have occurred in 1873 in the Bitter Root Valley in Montana, and by the end of the 19th century, Wood (1896) and Maxey (1899) described this disease as an entity occurring predominantly in Idaho and Montana (cited in references 161 and 409). More recently, however, RMSF has been identified in 42 states, with North Carolina and Oklahoma reporting the greatest number of

TABLE 2. Key parameters of intracellular parasitism and CNS invasion by members of the *Rickettsiaceae*

Microbe	Intracellular niche	Cellular target in vivo		Predominant CNS manifestation
		Endothelium	Leukocyte	
<i>R. rickettsii</i>	Cytoplasm	×		Encephalitis
<i>R. prowazekii</i>	Cytoplasm	×		Encephalitis
<i>O. tsutsugamushi</i>	Cytoplasm	×	×	Meningoencephalitis
<i>C. burnetii</i>	Phagosome		×	Meningitis and meningoencephalitis
<i>E. chaffeensis</i>	Phagosome		×	Meningitis

cases during the period from 1993 to 1996 (384). The etiologic agent of the disease was identified in 1909 by Howard Ricketts, who found *R. rickettsii* in the eggs of ticks and in blood from infected humans and demonstrated its transmissibility from ticks to guinea pigs (320).

RMSF is a systemic disease that typically manifests as fever, headache, myalgias, gastrointestinal disturbances, and severe dysfunction of multiple organ systems (379). Headache is the most common CNS symptom and is one of the most consistent clinical findings in RMSF. This was noted in an early report by Wilson and Chowning (409), and headache is present in 38 to 93% of cases reviewed by others (260, 379). The presence of headache is not pathognomonic of actual CNS invasion by the organism. However, less frequent symptoms such as lethargy, confusion, stupor, and coma and physical signs including meningismus, focal neurological deficits, and behavioral changes consistent with encephalitis are also present and indicate actual brain involvement. CSF abnormalities are often present but usually are not as impressive as those found in bacterial meningitis (160, 261). In one series, 27 of 33 patients with RMSF had a modest (<50 white blood cells) mononuclear pleocytosis, 28 of 33 showed small but abnormal numbers of erythrocytes, and 7 of 18 had an elevated CSF protein concentration (160). By comparison, boutonneuse fever, also known as Mediterranean spotted fever or tick typhus, which is caused by *R. conorii*, tends to result in fewer CNS findings and also has a lower case fatality rate than RMSF. For example, headache was reported as present in only 15% of 645 pediatric patients from Sicily, and only 2 children had lymphocytic meningitis (55).

R. prowazekii, the etiological agent of louse-borne epidemic typhus, is the prototype of the typhus group of the genus *Rickettsia* and was identified in the early 20th century by Rocha-Lima (cited in reference 127). This disease has a long history and is typically associated with war, mass movements of people, especially refugees, and squalid living conditions (422). Humans are the main reservoir for the disease, and the role of the body louse in its transmission was discovered by Nicolle in 1909 (153, 309). Like RMSF, typhus is a systemic disease that can be quite severe and is frequently fatal. A severe headache is nearly invariably present in patients with typhus and has been used as a key clinical criterion for identifying suspected cases during epidemics (127, 288, 308). More severe CNS features such as neck stiffness, neurological weakness, seizures, delirium, and coma are reported in 3 to 78% of cases (127, 288, 308, 310). Delirium and coma were reported in 35 and 39% of fatal cases, respectively (187). Another member of the typhus group, *R. typhi*, is the etiologic agent of murine typhus. This is a less severe disease than epidemic typhus, and symptoms of

severe CNS disease such as seizures, stupor, and ataxia are infrequent, being found in <5% of patients in one study (98); meningitis was reported in only 3 of 137 patients in a different study (361). Despite the low prevalence of CNS symptoms, *R. typhi* has been demonstrated by immunofluorescence in the brain of an elderly woman who died of murine typhus, indicating that this organism has the potential for CNS invasion (398).

Pathological involvement of the brain is common in fatal cases of rickettsial diseases, with lesions being found in 14 of 15 brains from patients with RMSF (160, 261) and in 63 of 66 brains from patients dying from epidemic typhus (223, 415). Moreover, histological studies have confirmed rickettsial invasion of vascular endothelial cells in the brains of humans (396, 398) and experimentally infected mice (399, 400). Characteristic CNS lesions in rickettsial infections are termed typhus nodules; they are proliferative vascular lesions that are similar to those found in the periphery (223). In the brain, they are associated with blood vessels and usually are 120 to 180 μ m in diameter. They are composed of vascular endothelium and neuroglia and undergo a characteristic evolution throughout the course of the disease (223). In addition to nodules, other frequent findings include perivascular accumulations of mononuclear cells and microinfarcts, the latter being the most common lesions in RMSF, but they are also found with epidemic typhus (5, 223). Postmortem examination of patients dying of RMSF shows widely scattered microinfarcts throughout the brain and spinal cord that are usually associated with small to medium-sized blood vessels and necrotic arterioles (5, 261). In some series, these inflammatory lesions were found only in the white matter (5), but others have also reported that they were present in gray matter as well (160, 261). In epidemic typhus, however, the lesions spare the white matter and show a striking predilection for the inferior olivary nuclei; they are also common in the gray matter of the cerebral cortex, pons, medulla, basal ganglia, and cord, presumably reflecting the vascularity and cellularity of these tissues (5, 223, 415). Meningeal lesions in the pia arachnoid are mild in patients with RMSF and consist of lymphocytic infiltrate without frank purulent meningitis or vascular occlusion, whereas acute inflammatory cell infiltrates are frequently present in patients with epidemic typhus (5, 223, 261). *R. prowazekii* also has been identified by PCR in the CSF of a human patient (244).

The central pathophysiological event of *Rickettsia* infections, including CNS infection, has long been identified as parasitism of vascular endothelial cells by blood-borne bacteria (415). Hematogenous dissemination of bacteria was demonstrated by inoculating animals and tissue culture cells with blood or plasma from patients infected with *R. rickettsii* (185, 320) and

R. prowazekii (40). In addition, *R. rickettsii* has been isolated from bone marrow monocytes and from peripheral blood monocytes in experimentally infected animals (48, 84, 195). Nevertheless, whether infected phagocytes also contribute to systemic dissemination is not known. Rickettsial invasion of endothelial cells (362, 397) and other nonprofessional phagocytes, such as chicken embryo cells, Vero cells, and fibroblasts (414), occurs by parasite-directed endocytosis, which requires bacterial metabolic activity and involves bacterial phospholipase A₂ (363). Candidate invasion proteins for this are rOmpA (221) and the protein encoded by the recently described *R. prowazekii* invasion gene homologue *invA* (125). This protein is a member of the Nidux hydrolase subfamily and is conserved among pathogenic rickettsiae. The *invA* gene was identified by sequence analysis of the *R. prowazekii* genome as a gene that encodes a putative protein with significant homology to invasion proteins of other bacterial pathogens (125). Transcriptional analysis indicates that *invA* is upregulated at several different times during the pathogenic life cycle, including the early entry phase, and may improve bacterial survival in the cytoplasm of the host cell (124). Internalized rickettsiae lyse the phagosomal membrane through an incompletely understood process in which bacterial phospholipase A₂ could play a role and then gain access to the cytoplasm. Subsequently, the pathogens proliferate in the cytoplasm and, in the case of *R. rickettsii*, also in the nucleus (363, 401, 412). *R. rickettsii* group bacteria exhibit actin-based motility in the cytoplasm similar to that used by *L. monocytogenes* but with notable differences in the ultrastructure and protein components of the actin tail (139, 163, 388). By comparison, *R. prowazekii* does not associate with F-actin, whereas *R. typhi* does form a very short F-actin tail (388). Rickettsial parasitism of endothelial cells as described above induces numerous intracellular events including inhibition of apoptosis, expression of proinflammatory cytokines and chemokines, upregulation of surface adhesion molecule expression, and oxidative stress. Together, these changes lead to altered hemostatic properties that then contribute to the formation of pathological lesions (109).

Orientia tsutsugamushi. *O. tsutsugamushi*, the etiologic agent of scrub typhus, was formerly a member of the spotted fever group of *Rickettsia* species but was placed in its own genus based on molecular phylogenetic studies (376). In vitro experiments show that *O. tsutsugamushi* infects neutrophils (322) and macrophages (272) as well as nonprofessional phagocytes including fibroblasts and endothelial cells (275, 387). After internalization, it escapes from phagosomes by an unknown mechanism and then proliferates in the cytoplasm (112, 323, 387). In contrast to other rickettsiae, intracellular movement of *O. tsutsugamushi* depends on microtubular and dynein function rather than actin polymerization (199). The mechanism for cell-to-cell spread is through budding of membrane-coated bacteria from infected cells (112).

These organisms are spread to humans through the bite of the larval stage of trombiculid mites and cause a systemic disease with features like those of other *Rickettsia* infections (351). The neurological manifestations are similar in many respects to other rickettsial diseases in that headache is nearly always present (35, 284). Signs of more severe CNS disease such as meningismus or meningitis have been found in 5.7 to

13.5% of patients (35, 340, 361). However, in a recent series of 25 patients who underwent lumbar puncture in the absence of overt CNS signs, 48% had a reactive spinal fluid showing a mild mononuclear pleocytosis, and *O. tsutsugamushi* was identified by PCR in 24% (284). These data suggest that CNS invasion is much more common in this disease than is suggested by symptoms alone. Necropsy studies show brain parenchymal lesions with a similar appearance and distribution to those in epidemic typhus, but the number of lesions is generally smaller (5). In contrast, the meninges are more commonly involved by *O. tsutsugamushi* than by other rickettsial infections, and the overall histological picture in the CNS is best described as a meningoencephalitis (5).

Dissemination of bacteria from the periphery to the CNS is hematogenous. Compared with other rickettsiae, *O. tsutsugamushi* is more frequently found in circulating mononuclear cells during naturally acquired infection of humans (402) and during experimental infection of dogs and nonhuman primates (355). Moreover, infection can be transmitted by blood transfusion due to prolonged microbial survival in leukocytes (56, 257). These findings suggest that phagocyte-facilitated infection could play a role in CNS invasion.

O. tsutsugamushi parasitizes endothelial cells in the periphery as well as in the brain, but it also can be found in macrophages of the liver and spleen (267). The mechanism of cellular invasion is not known, but binding to nonprofessional phagocytes may involve interaction between cell surface heparan sulfate proteoglycans and bacterial lectins (176). Infection of host cells triggers a response including activation of the transcription factors NF- κ B (in macrophages) and NF- κ B and AP-1 (in endothelial cells), leading to subsequent expression of chemokine genes for MIP-1 α / β , MIP-2, and MCP-1 in macrophages and MCP-1, IL-8, and RANTES in endothelial cells (61–63). In an interesting contrast to *R. rickettsii*, which inhibits endothelial apoptosis, *O. tsutsugamushi* was shown to induce apoptosis in an endothelial cell line. However, it inhibited the apoptotic process in monocyte-like THP-1 cells (194, 198) and might also actively suppress cytokine production by infected macrophages (197).

Ehrlichia chaffeensis. *E. chaffeensis* is an obligate intracellular parasite of mononuclear phagocytes and is the etiologic agent of human monocytotropic ehrlichiosis (282). Individual organisms are termed elementary bodies and are small (approximately 0.5 μ m in diameter) gram-negative bacteria that are found within infected cells (250). Monocytes are the primary cellular targets of *E. chaffeensis* in vivo; however, bacteria are also found in liver endothelial cells in experimentally infected mice (367, 413) and can infect nonprofessional phagocytes in vitro (60). Ehrlichiae have a characteristic intracellular life cycle that progresses through three developmental stages: elementary bodies, initial bodies, and morulae (reviewed in reference 250). Elementary bodies enter monocytes by phagocytosis. The receptors and ligands for *E. chaffeensis* invasion of mononuclear phagocytes are not well established. A bacterial 120-kDa outer membrane protein is probably important in this respect, since expression of it by *Escherichia coli* allows the *E. coli* cells to bind and invade HeLa cells (297). Once internalized, *E. chaffeensis* localizes to early endosomal compartments and causes upregulated expression of mRNA for the transferrin receptor with subsequent accumulation of transferrin

receptors in phagosomes (30, 268). Bacteria replicate in the phagosomal compartment by binary fission, so that over a period of 3 to 5 days the individual organisms become tightly packed and form structures that are 1.0 to 2.5 μm in diameter, referred to as initial bodies. Over the ensuing 7 to 10 days, initial bodies coalesce, forming a more mature inclusion body known as a morula. When the infected cell ruptures, morulae break up into elementary bodies and the infective cycle repeats itself.

The first monocytotropic *Ehrlichia* species discovered was *E. canis*, which was found in tick-infected dogs in 1935 in Algeria (87). Human monocytotropic ehrlichiosis was first identified in 1987, and the infecting agent was subsequently named *E. chaffeensis* (235). Ehrlichial infections are zoonoses transmitted to humans and other mammals through the bite of infected ticks, principally the lone star tick, *Amblyomma americanum*. Animal reservoirs of *E. chaffeensis* include white-tailed deer and coyotes (75, 113, 203). Human infection with *E. chaffeensis* is clinically similar to RMSF, with the major exception that most patients with ehrlichiosis do not develop a rash. The manifestations of infection are nonspecific and include fever, myalgia, and malaise, with headache being the main symptom referable to the CNS and which is found in approximately 80% of patients (97).

A review of CNS manifestations of ehrlichiosis by Ratnasamy et al. identified 22 cases of which 21 were attributed to *E. chaffeensis* and 1 was attributed to the closely related *Anaplasma phagocytophila*, formerly known as the agent of human granulocytic ehrlichiosis (312). Patients with severe CNS involvement had headache, nuchal rigidity, confusion, ataxia, hyperreflexia, seizures, photophobia, and cranial nerve palsies. Findings in the CSF were nonspecific and consisted of a lymphocytic pleocytosis with a mildly elevated protein level in almost 75% of cases. The presence of organisms in the CSF has been established by PCR and by microscopic identification of parasitized cells (312). In addition, *E. chaffeensis* has been isolated from CSF of two patients by primary culture (368). Pathological findings in the CNS from patients with fatal cases either were unremarkable (312) or consisted only of a lymphocytic infiltrate in the leptomeninges with involvement of blood vessel walls and extension into the Virchow-Robin space (144). Thus, the brain parenchyma appears to be spared from direct infection.

As noted above, *E. chaffeensis* is more frequently found in the CNS in humans than is *A. phagocytophila*. This relative difference is also noted in animals infected with related organisms. For example, dogs that are experimentally infected with *E. canis*, a monocytotropic organism closely related to *E. chaffeensis*, commonly develop meningitis with a mild vascular cuffing and gliosis in the parenchyma (285). In contrast, brain lesions are not found in horses and sheep experimentally infected with the organism formerly known as *E. equi*, now termed *A. phagocytophila* comb. nov. (96). The mechanism used by *E. chaffeensis* to invade the CNS is unknown. Given that its *in vivo* niche is within monocytes, phagocyte-facilitated invasion is the most likely route by which these bacteria enter the CNS. Nevertheless, recent data by Li and Winslow show that approximately 10% of blood-borne bacteria in *E. chaffeensis*-infected mice are actually extracellular and that these cell-free organisms are virulent if they are immediately transferred

into other mice or are cocultured with the DH82 macrophage-like cell line (222). These authors suggest that cell-free *E. chaffeensis* organisms are liberated from infected cells by necrotic cell death and are an important target for antibody-mediated clearance mechanisms. Hence, it is possible that these extracellular bacteria could invade endothelial cells of the CNS directly, but the lack of widespread endothelial cell involvement *in vivo* suggests that this is unlikely. In addition, the greater incidence of CNS infection observed with monocytotropic ehrlichiae compared with granulocytotropic *Anaplasma* is consistent with the greater ability of monocytes to migrate across the blood-brain barrier under steady-state conditions (reviewed in reference 165).

***Coxiella burnetii*.** *C. burnetii* is an obligate intracellular bacterium that is the etiologic agent of Q fever (249). Q fever is a zoonosis that was first described as an outbreak of a febrile disease in an abattoir in Queensland, Australia (83). The pathogenesis of *C. burnetii* infection in humans is dependent on this microbe parasitizing macrophages and monocytes, although it also can enter and replicate within nonprofessional phagocytes such as HeLa cells and fibroblasts (162, 248). Once internalized by nonprofessional phagocytes, *C. burnetii* resides in an acidified endosomal compartment characterized as a phagolysosome (162). However, it has been proposed recently that intracellular bacteria delay (170) or completely inhibit (133) phagolysosome fusion in macrophage-like and monocytic cell lines, respectively. Intracellular *C. burnetii* organisms undergo a complex intracellular life cycle involving a sporulation-like process that leads to the formation of intracellular forms termed "large-cell variants" and "small-cell variants" (reviewed in reference 158).

A recent review of 1,383 cases of Q fever by Raoult et al. found that isolated fever, hepatitis, and pneumonia with or without associated hepatitis accounted for 91% of cases (311). The most common neurological sign is headache, which has been found in 65 to 90% of cases (339). A review of neurological involvement in Q fever identified headache severe enough to warrant lumbar puncture to rule out meningitis in 3.5% of hospitalized patients. After exclusion of patients with a normal CSF examination, the authors concluded that the prevalence of serious neurological symptoms was 2.2% (38). By comparison, a different series reported an incidence of neurological symptoms of 22% (314). A variety of manifestations of CNS disease in Q fever have been reported, including meningitis and meningoencephalitis in 0.5 and 1% of patients, respectively (311), neck stiffness in 5%, and confusion and apathy in 7% (64). Behavioral and psychiatric disturbances, seizures, and peripheral neuritis have also been reported (38). Approximately two-thirds of CSF analyses show increased numbers of leukocytes, typically mononuclear, and one-third of specimens had increased protein levels (38). *C. burnetii* is difficult to culture from the CSF but has been identified by PCR in one patient (338). One necropsy of a patient who died of Q fever pneumonia showed small perivascular hemorrhages in the brain, with capillary endothelial swelling without perivascular accumulation of leukocytes (406). Rickettsial forms were identified extracellularly and inside neuroglia and endothelial cells by Giemsa staining.

The method by which this organism enters the CNS is not known, but its tropism for monocytes and the ability to survive

for extended periods within them suggests that trafficking of infected cells into the CNS is the most likely mechanism. Chronic persistence of *C. burnetii* within human monocytes is well described and is associated with increased cellular secretion of TNF- α (80), IL-10, and transforming growth factor β 1 (53). Overproduction of IL-10 may play a role in inhibiting the bactericidal activity of the host cell by altering cellular responses to TNF- α (130, 132). Interestingly, *C. burnetii* is one of the few organisms studied from the perspective of the way in which intracellular infection of monocytes alters their interactions with endothelial cells. Dellacasagrande et al. reported that infection of THP-1 cells with a virulent strain of *C. burnetii* increased cellular adhesion to a TNF- α -stimulated human microvascular cell line but transmigration across the cell monolayers was significantly inhibited (81). The authors suggested that decreased transmigration might interfere with localization of the bacterium to target tissues and/or may contribute to the formation of poorly formed granulomas that are found in chronic Q fever. From the standpoint of CNS invasion, the inability of infected monocytes to migrate across endothelial barriers could contribute to the relatively low incidence of CNS infection found in Q fever in spite of their long-term presence in the circulation.

Tropheryma whipplei

T. whipplei, the causative agent of Whipple's disease, is a gram-positive, periodic acid-Schiff (PAS)-positive, diastase-resistant bacterium that is a member of the high-G+C phylum of gram-positive bacteria (315). In vitro study of this obligate intracellular organism and understanding of how it lives intracellularly have been complicated by the fact that sustainable ex vivo culture of the organism was achieved only in 2000 (306). Studies using infection of HeLa cells for a model show that *T. whipplei* survives by altering the phagosomal environment. Phagosomes containing bacteria are acidic and colocalize with lysosome-associated protein 1 but not with cathepsin D, indicating that phagolysosome fusion and maturation are incomplete (131). Interestingly, acidification of the vacuole is critical to the survival of the organism, as shown by the finding that agents that increase the intravacuolar pH decrease bacterial viability.

Whipple's disease is rare, with less than 1,000 cases reported to date (116), although this is certainly an underestimate of the number of cases that have occurred (86). It is named after George Whipple, who described an autopsy in 1907 of a 36-year-old physician with gradual loss of weight and strength, diarrhea, ill-defined abdominal signs, and arthritis in multiple joints. The disease was called "intestinal lipodystrophy" based on findings of polyserositis, aortic valve lesions, and prominent deposition of fat within intestinal mucosa and mesenteric lymph nodes. In addition, there was marked infiltration of the intestinal lamina propria by foamy macrophages that contained rod-shaped bacilli (405). In 1991, Wilson et al. reported on the 16S rRNA gene and partial sequence of bacterial DNA extracted from duodenal and lymph node biopsy specimens from patients with Whipple's disease (408). These findings were confirmed and extended the following year, and the proposed name of the new organism as *Tropheryma whipplei* (Greek *trophe* for nourishment and *eryma* for barrier) was

advanced (315). In 2001, the causative organism of Whipple's disease was officially named *Tropheryma whipplei* (216).

Whipple's disease occurs worldwide, but 55% of cases were reported from Europe and 38% were reported from North America (86). The habitat of *T. whipplei* in the environment is unknown, as is the mode of acquisition of the bacteria. However, data suggest that *T. whipplei* is an environmental human commensal, which is acquired by drinking water but rarely causes human infection. This notion is based on identification of the organism by PCR in 25 of 38 samples of sewage water from five different sewage plants in Germany (237) and also in a variety of specimens from patients with no evidence of Whipple's disease (371). The reason why this assumed commensal organism rarely causes disease might be found in underlying host conditions. Immune abnormalities have been demonstrated in patients with Whipple's disease, as reviewed in references 243 and 263. If there is an underlying immune defect, it appears to be quite specific, because patients with Whipple's disease do not acquire other opportunistic infections. These abnormalities might be primary, as is suggested by the finding of decreased intracellular killing of *Candida tropicalis* by monocytes from patients with Whipple's disease in remission compared to normal controls (24). Furthermore, peripheral blood mononuclear cells from patients with Whipple's disease had increased production of IL-4 in response to in vitro stimulation with phytohemagglutinin and reduced secretion of gamma interferon and IL-2 compared to controls (242). Thus, macrophages might be more susceptible to *T. whipplei* infection in part because of a decreased influence of classically activating cytokines and the increased alternatively activating IL-4 signal (138). The report of a patient with refractory Whipple's disease who had an apparent response to gamma interferon therapy lends credence to this hypothesis (342).

Whipple's disease is a chronic relapsing multisystem disease, and its clinical manifestations have been recently reviewed (243). Almost all reported cases (97%) were in whites (86), with a male-to-female ratio of approximately 8:1 (243). The average age at the time of diagnosis was 50 to 55 years (100, 243), although patients with neurological abnormalities as the initial manifestation may be younger (average age, 42 years) and more commonly female (male-to-female ratio, 1:1) (129). Intermittent arthropathy, which is the initial symptom in 63% of cases, precedes the diagnosis by an average of 8 years and is present at the time of diagnosis in 85% of patients (304). The cardinal symptoms include significant weight loss, intermittent diarrhea or steatorrhea, abdominal pain, fever, and lymphadenopathy. Involvement of the CNS is found in 20 to 40% of patients with Whipple's disease (11, 54, 65, 129, 210, 228, 255, 343, 373, 394). CNS involvement typically occurs as part of a systemic infection and usually develops in the later stage of disease, although cases confined to the CNS have been reported (373). Patients with CNS disease nearly always exhibit diffuse neurological signs. In a review of 122 cases, the neurological signs were classified as follows, in order of decreasing frequency: supranuclear ophthalmoplegia, confusion and dementia, psychiatric signs, myoclonic signs, seizures, hypothalamic involvement, cerebellar forms, myorhythmic forms, cranial nerve involvement, peripheral neuropathies, and spinal cord involvement (129). Meningeal involvement is often associated with all these neurological signs, so that the disease is

actually a meningoencephalitis or, in cases of ophthalmologic involvement, meningouveitis (129). The CSF was abnormal in half of all patients with Whipple's disease studied ($n = 73$) and in two-thirds of those experiencing neurological relapse ($n = 29$).

The histopathology of Whipple's disease in the small intestine is a readily recognizable characteristic group of findings including villi with many histiocytes containing diastase-resistant PAS-positive granular material in the lamina propria, club-shaped villi, prominent dilated lymphatic spaces, and large-droplet lipid deposits (25). Histopathology of the involved cerebral cortex shows scattered aggregates of foamy macrophages with prominent perivascular cuffing and large reactive astrocytes; the large foamy macrophages are filled with diastase-resistant PAS-positive granules. In patients with florid disease, *T. whipplei* can be seen apparently in the extracellular compartment, and it has recently been cultivated from the CSF of two patients (238). The mechanism(s) by which *T. whipplei* invades the CNS has not been studied. However, *T. whipplei* has been demonstrated by immunocytochemical staining of peripheral blood monocytes of a patient with Whipple's disease (307). This suggests that circulating mononuclear phagocytes could transport the organism into the CNS.

CONCLUDING REMARKS

The intracellular microbes causing CNS disease reviewed in this paper are a diverse group of human pathogens. One way in which this is demonstrated is the wide variety of mechanisms used to enter their human hosts: through contaminated food (e.g., *L. monocytogenes*, *Salmonella*, and *Brucella* spp.), through the bite of infected arthropods (*R. rickettsii*, *R. prowazekii*, and *E. chaffeensis*), and through inhalation (*M. tuberculosis* and *C. burnetii*). This is an interesting contrast to extracellular neuroinvasive bacteria such as *S. pneumoniae*, *N. meningitidis*, and *H. influenzae*, which infect humans via the respiratory tract. The diversity of the intracellular pathogens inflicting CNS disease also is reflected in their place in human history. For example, there is evidence of human infection with *M. tuberculosis* and *Brucella* spp. dating back several thousand years, whereas human infection with *E. chaffeensis* has been recognized only in the past two or three decades and *T. whipplei* was identified only about one decade ago. Typhus and tuberculosis are clearly associated with poverty, calamity, and social upheaval. The former emerged historically in times of war, whereas the latter now is united with a truly modern epidemic, HIV-1, to reestablish itself as the "Captain of all these men of death" (J. Bunyan in *The Life and Death of Mr. Badman*, 1680). On the other hand, none of these bacteria are particularly associated with prosperity, except that only in a prosperous society do people live long enough that a bacterium like *L. monocytogenes* can prey on the elderly. Given the limited insight into the etiopathogenesis of many of the CNS infections caused by intracellular bacteria, it is unlikely that new therapies aimed specifically at preventing CNS infection by these microorganisms will be available soon. Thus, continued research and clinical effort in the development of CNS-specific therapies are needed. In spite of profound biological differences, intracellular pathogens are united in their adaptation to the intracellular environment and in the exploitation of

host cells for their maintenance and dissemination. It should be underscored that for most of these organisms, the basic cellular and molecular mechanisms used to invade the CNS are largely unknown or are just now being explored. Of the typical routes used by blood-borne bacteria to enter the CNS, the two routes most likely to be used by intracellular organisms are direct invasion of endothelial cells and trafficking of organisms within infected mononuclear phagocytes. The main point in favor of the former mechanism is that most, if not all, of these pathogens can invade and replicate within nonprofessional phagocytes. The clearest examples of this mode of invasion are provided by *R. rickettsii* and *R. prowazekii*. In diseases caused by these organisms, endothelial cell invasion is the key pathophysiological event.

By comparison, CNS invasion by infected phagocytes is less well studied and does not leave easily identifiable footprints in vivo. Nevertheless, parasitism of mononuclear phagocytes is an essential feature of several of the organisms considered here. For example, the recent characterization of *Brucella* spp. as facultative extracellular pathogens emphasizes that these bacteria have an extreme preference for the intracellular environment despite their ability to live outside host cells. For such organisms, CNS invasion via trafficking within infected phagocytes is a logical route. By comparison, some microbes, such as *L. monocytogenes* and *O. tsutsugamushi*, may be capable of CNS invasion by more than one mechanism. In this instance, data from experimental infection with *L. monocytogenes* are helpful in defining the molecular and cellular mechanisms that are operative in vivo. Additionally, data that show an important role for transportation of *M. tuberculosis* and *S. enterica* serovar Typhimurium within mononuclear phagocytes across epithelial barriers suggest that trafficking of infected phagocytes could be a central motif for several intracellular bacterial pathogens in the same way that subversion of key intracellular processes, e.g., maturation of phagosomes, is a common feature of intracellular pathogenesis.

ADDENDUM IN PROOF

Readers are directed to new data concerning the role of infected monocytes and transportation of intracellular *L. monocytogenes* into the brain: D. A. Drevets et al., *J. Immunol.*, in press.

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