

CD44-Regulated Intracellular Proliferation of *Listeria monocytogenes*

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CD44 has been implicated in immune and inflammatory processes. We have analyzed the role of CD44 in the outcome of *Listeria monocytogenes* infection in murine bone marrow-derived macrophages (BMM). Surprisingly, a dramatically decreased intracellular survival of *L. monocytogenes* was observed in CD44^{-/-} BMM. CD44^{-/-} heart or lung fibroblast cultures also showed reduced bacterial levels. Moreover, livers from CD44^{-/-} infected mice showed diminished levels of *L. monocytogenes*. In contrast, intracellular growth of *Salmonella enterica* serovar Typhimurium was the same in CD44^{-/-} and control BMM. The CD44-mediated increased bacterial proliferation was not linked to altered BMM differentiation or to secretion of soluble factors. CD44 did not mediate listerial uptake, and it played no role in bacterial escape from the primary phagosome or formation of actin tails. Furthermore, CD44-enhanced listerial proliferation occurred in the absence of intracellular bacterial spreading. Interestingly, coincubation of BMM with hyaluronidase or anti-CD44 antibodies that selectively inhibit hyaluronan binding increased intracellular listerial proliferation. Treatment of cells with hyaluronan, in contrast, diminished listerial growth and induced proinflammatory transcript levels. We suggest that *L. monocytogenes* takes advantage of the CD44-mediated signaling to proliferate intracellularly, although binding of CD44 to certain ligands will inhibit such response.

CD44 glycoprotein is found on the surface of many cell types, including lymphocytes, macrophages, and epithelial cells. Expression levels vary depending on origin and activation status of the cell. CD44-dependent processes are known to include organ development, neuronal axon guidance, hematopoiesis, and numerous immune functions. Among the latter, CD44 participates in lymphocyte adhesion to inflamed endothelium, lymphocyte homing, and tumor metastasis (31). Hyaluronan (HA), a main carbohydrate component of the extracellular matrix, is the principal but not sole ligand of CD44. CD44 is also the major receptor for HA. HA is normally a glycosaminoglycan of high molecular weight. At sites of inflammation, low-molecular-weight (LMW) HA accumulates, most likely due to the presence of hyaluronidases (HA'ses) and/or reactive oxygen species. Binding of LMW HA to CD44 can induce expression of cytokines, chemokines, and adhesion and effector molecules and can induce translocation of transcription factors in cell lines or primary cell cultures (2, 22–24, 26, 28, 29, 41). Thus, besides tethering cells to extracellular ligands, CD44 has broader functions in cellular signaling cascades. CD44 also provides a link between the plasma membrane and the actin cytoskeleton. CD44 can have coreceptor functions mediating the signaling of receptor tyrosine kinases, such as Met.

The impact of CD44 in the regulation of immune responses and inflammation has been broadly studied (27, 31, 40, 42), but few studies have addressed the potential role of CD44 in the control of pathogens (4, 12, 13, 15, 39).

The gram-positive bacterium *Listeria monocytogenes* is a hu-

man pathogen that causes severe disease in immunocompromised individuals and will induce abortions in pregnant women. *L. monocytogenes* is known to invade a variety of cells, including macrophages. After cellular uptake, the bacterium escapes from the primary phagosome into cytoplasm, where it starts to multiply and then spread to nearby cells (45). The presence of an inducible listerial hexose phosphate transporter mediating rapid intracellular replication has been recently described (17). In the cytoplasm *L. monocytogenes* expresses ActA protein, a cofactor for the nucleation of actin filaments. The bacterium polymerizes actin filaments around itself, creating a long actin tail. Such tails will propel listeria to the cell membrane, where projections involved in listerial cell-to-cell spread will be formed (11).

Immune resistance to *L. monocytogenes* depends on the ability of the host to mount a Th1-like immune response (43). Cytokines such as gamma interferon (IFN- γ) will activate macrophage bactericidal mechanisms, which play a crucial role in the control of listerial infection in vivo (20, 32).

We initially hypothesized that signals through HA and CD44 could inhibit the intracellular growth of *L. monocytogenes* by upregulating the expression of inflammatory genes and by controlling the cytoskeleton rearrangements. Instead, our studies revealed that *L. monocytogenes* makes use of CD44 signaling to grow efficiently intracellularly.

MATERIALS AND METHODS

Reagents. Anti-CD44 (KM 703, KM 81), anti-CD4, and anti-major histocompatibility complex (MHC) class I monoclonal antibodies were purified from the supernatant of hybridomas CRL-1896, TIB-241, L3T4, and HB51, respectively (American Type Culture Collection, Manassas, Va.), by using protein G-Sepharose (Amersham-Pharmacia, Uppsala, Sweden). Hyaluronidase (HA'se) from *Streptomyces* species was purchased from Calbiochem (San Diego, Calif.). HA'se

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TABLE 1. Primer sequences used in this study

Primer	Sequence
Sense iNOS.....5'	CCC TTC CGA AGT TTC TGG CAG CAG CAG C 3'
Antisense	
iNOS.....5'	GGC TGT CAG AGC CTC GTG GCT TTG G 3'
Sense IFN-γ.....5'	TGG ACC TGT GGG TTG TTG ACC TCA AAC TTG GC 3'
Antisense	
IFN-γ.....5'	TCG ATC TTG GCT TTG CAG CTC TTC CTC ATG GC 3'
Sense IL-12 p40.....5'	CGT GCT CAT GGC TGG TGC AAA G 3'
Antisense	
IL-12 p40.....5'	CTT CAT CTG CAA GTT CTT GGG C 3'
Sense IL-1α.....5'	ATG GCC AAA GTT CCT GAC TTG TTT 3'
Antisense	
IL-1α.....5'	C CTT CAG CAA CAC GGG CTG GTC 3'
Sense IL-6.....5'	ATG AAG TTC CTC TCT GCA AGA GAC T 3'
Antisense	
IL-6.....5'	CA CTA GGT TTG CCG AGT AGA TCT C 3'
Sense TNF-α.....5'	ATG AGC ACA GAA AGC ATG ATC CGC 3'
Antisense	
TNF-α.....5'	CC AAA GTA GAC CTG CCC GGA CTC 3'
Sense IFN-α.....5'	GAC TCA TCT GCT GCT TGG AAT GCA ACC CTC 3'
Antisense	
IFN-α.....5'	GAC TCA CTC CTT CTC CTC ACT CAG TCT TGC C 3'
Sense β-actin.....5'	GTG GGC CGC TCT AGG CAC CAA 3'
Antisense	
β-actin.....5'	CTC TTT GAT GTC ACG CAC GAT TTC 3'

type III from sheep testes, chondroitinase ABC from *Proteus vulgaris*, heparinase (HS'se) III from *Flavobacterium heparinum*, and heparan sulfate (HS) from bovine kidneys were all purchased from Sigma-Aldrich (St. Louis, Mo.). Hyaluronans (HA), with the sizes of 190 and 250 kDa, were kindly provided by Ove Vik (Q-Med AB, Uppsala, Sweden).

Mice. C57Bl/6 mice were bred under specific-pathogen-free conditions. A mutant mouse strain without CD44 was generated by homologous recombination in embryonic stem cells and was backcrossed with C57Bl/6 (36).

Bacteria. *L. monocytogenes* wild-type (WT) strain EGD (BUG600, serotype 1/2a) and the $\Delta plcB2$ mutant (35) with a defective lecithinase were used. The *actA*-defective mutants, deficient in actin polymerization strain LO28 $\Delta ActA$ (BUG 875) (18) and the BUG314 strain, containing a Tn917 transposon inserted in *actA* (25) and the parental control strain LO28, all obtained from the Pasteur Institute (Paris, France), were used. To study intracellular bacterial localization of *L. monocytogenes* NF-L357, which contains a transcriptional fusion between *actA-plcB* and the green fluorescent protein gene (*gfp*) was used (16). In all cases the bacteria were grown in brain heart infusion (BHI) broth and BHI agar (Difco, Becton Dickinson, Md.) at 37°C. Before infection bacteria were grown at 37°C in BHI broth to late-exponential phase (optical density at 600 nm, 0.8). The bacteria were washed once with phosphate-buffered saline (PBS), resuspended in BHI broth with 15% glycerol, plated on duplicate BHI agar plates, and quantified after an overnight incubation at 37°C. Bacterial suspensions were stored at -70°C until use.

The plasmid pAUL-A (8), which replicates at 30°C but not at 37°C and possesses an erythromycin resistance gene, was used to distinguish replication rate from death rate of *L. monocytogenes* in bone marrow-derived macrophages (BMM). *L. monocytogenes* EGD was transformed with pAUL-A by electroporation and was grown at 30°C in BHI medium containing 5 µg of erythromycin per ml overnight. To produce cultures containing less than one copy of the plasmid per bacterium, 20 ml of the culture was inoculated in 180 ml of BHI and was grown at 37°C for 6 h. Numbers of total and plasmid-containing *L. monocytogenes* and the total number of plasmids were calculated during this incubation by plating bacteria on BHI agar in the presence or absence of erythromycin or by further incubating bacteria at 42°C to stationary phase before plating on BHI agar in the presence of erythromycin. An initial 5.3-fold increase in the number of plasmid-containing bacteria during the first 4.5 h in culture was determined. The mean copy number at start was calculated to be 3.6 plasmids per

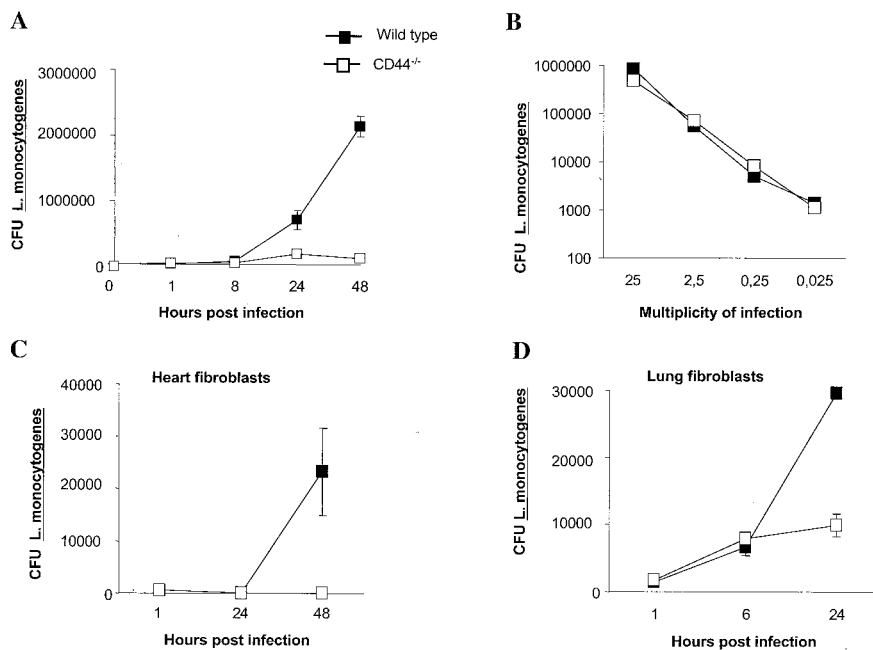


FIG. 1. Reduced intracellular growth of *L. monocytogenes* in CD44^{-/-} BMM, lung, and heart fibroblasts. (A) CD44^{-/-} and WT BMM were infected with *L. monocytogenes* at a multiplicity of infection of 0.2 bacteria per cell. One hour after infection BMM were extensively washed with PBS, and DMEM supplemented with 5% FCS containing 5 µg gentamicin per ml was added. BMM were lysed at the indicated time points, and CFU were quantified after plating. The experiment shown is representative of at least 10 independent experiments. (B) BMM from CD44^{-/-} and WT mice were infected with different numbers of *L. monocytogenes*. One hour after infection cells were extensively washed to remove extracellular bacteria and were lysed; CFU were then measured. Similarly, CFU were measured for *L. monocytogenes*-infected BMM further incubated with or without 5 µg of gentamicin per ml for 15 min, a dose that was lethal for extracellular *L. monocytogenes* (data not shown). Thus, the depicted CFU originate from intracellular *L. monocytogenes*. One representative of several independent experiments is shown. (C and D) Fibroblasts derived from heart (C) and lung (D) from CD44^{-/-} and WT mice were infected with *L. monocytogenes* as described in Materials and Methods. Cells were washed and lysed at the indicated times after infection, and the amounts of CFU were quantified. An MOI of 1:1 was used in the depicted experiments. Independent experiments that used MOIs of 10:1 and 2:1 also showed similar results. Bars represent the standard errors.

TABLE 2. Growth and death rates of BMM infected with *L. monocytogenes*^a

MOI	H after infection	Growth rate for:					
		WT BMM			CD44 ^{-/-} BMM		
		Total CFU	CFU with pAUL-A	Proportion with pAUL-A	Total CFU	CFU with pAUL-A	Proportion with pAUL-A
0.32	1	4,050	1,300	0.32	4,939	1,600	0.33
0.32	24	190,000	340	0.0018	106,500	230	0.0022
0.32	48	255,000	0	0	14,650	0	0
3.2	1	40,500	15,300	0.38	58,075	16,800	0.29
3.2	24	1,160,000	2,320	0.0020	665,000	1,330	0.0020

^a BMM were infected with *L. monocytogenes* containing pAUL-A as described in Materials and Methods. At the indicated time points after infection total CFU and CFU containing pAUL-A (bacteria grown in presence of erythromycin) were measured. The proportion of *L. monocytogenes* carrying pAUL-A was calculated. The proportion of bacteria maintaining pAUL-A is a measurement of the growth rate, while the total number of bacteria with pAUL-A is a measurement of killing by the host cell. Listerial growth is thus the outcome of bacterial death and proliferation of few surviving, in agreement with previous electron microscopy data (30). The total number of bacteria results from the balance between growth rate and killing. Note that although after 48 h of infection CD44^{-/-} BMM contain 17-fold less *L. monocytogenes* than WT BMM, neither CD44^{-/-} nor WT BMM contain any pAUL-A, indicating that the reduced number of bacteria in CD44^{-/-} BMM is the result of diminished survival and is not the consequence of a decreased growth rate. One representative of two independent experiments is shown.

bacterium, and growth of plasmid-containing bacteria represents the segregation of preexisting plasmids into daughter bacteria. After 4.5 h the total number of plasmid-containing bacteria stopped increasing and was identical to the total number of plasmids. Moreover, the total number of plasmids was not altered during the 5-h culture period. Thus, after 4.5 h each plasmid-containing bacteria had one copy of the plasmid. At this time, 20 to 30% of total bacteria contained

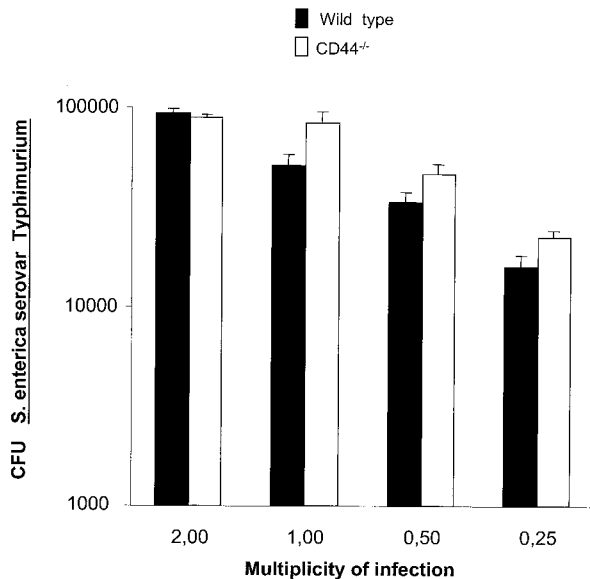


FIG. 2. CD44 does not mediate intracellular growth of *S. enterica* serovar Typhimurium. CD44^{-/-} and WT BMM were infected with *S. enterica* serovar Typhimurium at an MOI of 0.25, 0.5, 1.0, and 2.0 bacteria per cell. One hour after infection cells were extensively washed and incubated in DMEM-5% FCS containing 5 µg of gentamicin per ml. CFU were measured 24 h after infection. Differences in the uptake of *S. enterica* serovar Typhimurium by BMM after 1 h were measured and were the same in both WT and CD44^{-/-} samples (data not shown).

pAUL-A. Aliquots of the culture grown for 4.5 h were frozen at -70°C and were used later for infection of BMM. BMM were infected as described below, and CFU in lysates were measured in plates containing 5 µg of erythromycin/ml (measuring plasmid-containing bacteria) and in plates not containing erythromycin (measuring total number of bacteria). The relative proportion of pAUL-A-containing bacteria is a measurement of the growth rate, while the total number of bacteria with pAUL-A is a measurement of killing by the host cell. A similar strategy has been previously used to evaluate toxicity during in vitro and in vivo infection with *Salmonella enterica* serovar Typhimurium (3, 21).

Differentiation of BMM. BMM were established as described previously (33). In brief, mice were killed by cervical dislocation, and femurs and tibias of the hind legs were dissected and flushed with 5 ml of ice-cold, sterile PBS. Bone marrow cells (typically 2×10^7 to 3×10^7 per mouse) were washed and resuspended in Dulbecco's minimal essential medium (DMEM) containing glucose and supplemented with 10% fetal calf serum (FCS), 20% L929 cell-conditioned medium (as a source of macrophage-colony stimulating factor), 100 µg of streptomycin/ml, 100 U of penicillin/ml, and 10 mM HEPES buffer. The cells were cultured at 5×10^4 cells per well in 24-well plates for 7 days at 37°C, 5% CO₂. BMM cultures were then created by washing vigorously with PBS to remove nonadherent cells. Adherent cells were also harvested and counted by trypan blue exclusion. Typically, the yield of bone marrow cells was 2.0×10^5 to 4.0×10^5 BMM per well (24-well plates) after 7 days in culture, which is when they were infected.

Fibroblast primary cultures. Fibroblasts were obtained by enzymatic digestion of hearts and lungs from 6- to 10-week-old CD44^{-/-} or WT mice as described previously (7). In brief, organs were minced in Iscove's modified medium (IMDM) supplemented with 10% FCS, 100 µg of streptomycin/ml, 100 U of penicillin/ml, and 10 mM HEPES buffer and were incubated in IMDM containing 1 mg of collagenase (type IV; Sigma, St. Louis, Mo.) per ml at 37°C for 10 min. Released cells in the supernatant were collected, and undigested tissue was further enzymatically treated for two more cycles. Recovered cells were pooled and plated, and highly enriched cultures of adherent cardiac and pulmonary cells were recovered during the plating procedure. After the third passage, nearly all cells appeared to be fibroblasts by morphology, as also previously determined by immunostaining (7). Such cultures were trypsinized and resuspended in IMDM, and 5×10^4 cells per well were seeded on a 24-well plate. Infection was performed 24 h after plating.

Infection and infectivity assay. BMM and fibroblasts cultured in 24-well plates were cocultured with *L. monocytogenes* for 1 h at 37°C, 5% CO₂. Cells were then extensively washed with PBS to remove the extracellular bacteria. Triplicate wells were used for all conditions. To prevent extracellular bacterial growth, cells were then cultivated in DMEM containing 5% FCS and 5 µg of gentamicin per ml. At different time points after infection cells were washed with PBS and lysed with 0.1% Triton X-100 in PBS, and aliquots of lysates were plated on BHI agar plates. Bacteria in two or three tenfold dilutions of the lysates were quantified. Plates were incubated overnight at 37°C and CFU were determined.

BMM were also infected in parallel with *S. enterica* serovar Typhimurium strain 14028 (American Type Culture Collection). *S. enterica* serovar Typhimurium was grown in Luria broth (LB) agar plates overnight. For this purpose, bacterial colonies were diluted in PBS at a concentration of 5×10^8 bacteria per ml. Ninety microliters of bacteria was coincubated with 10 µl of C57Bl/6 normal mouse serum for 30 min at 37°C. Bacteria were further diluted at 10^6 per ml in DMEM-5% FCS. BMM were cocultured with *S. enterica* serovar Typhimurium and were centrifuged at $500 \times g$ for 5 min at room temperature. Cells were incubated for 1 h at 37°C and were extensively washed with PBS, and DMEM containing 5 µg of gentamicin per ml was then added. Cells were lysed at the indicated time points after infection as described above, and CFU in cell lysates were quantified on LB agar plates after overnight incubation.

Immunostaining and flow cytometry. BMM were phenotypically characterized by fluorescence-activated cell sorter analysis. For this purpose, BMM were detached from plates by using a cell scraper and were incubated with PBS containing 10% normal goat serum (Sigma) to block unspecific binding of the secondary antibody. Cells were then incubated with purified monoclonal rat antibody to F4/80, Mac-3, CD80, or GR-1 for 30 min on ice. Cells were then washed and incubated with biotinylated goat anti-rat serum for 30 min on ice. Anti-CD44, -CD19, -CD3, and -CD8 antibodies were directly labeled with fluorescein isothiocyanate. Anti-CD4, -CD14, and -NK1.1 antibodies were directly labeled with phycoerythrin. Anti-MHC class II (Ia) and -ICAM-1 (MALA-2) antibodies were biotinylated. Isotype-matched control rat immunoglobulins (Igs) directly labeled with fluorescein isothiocyanate, phycoerythrin, or biotin were used as controls. When staining with biotinylated or directly labeled antibodies, BMM were preincubated with anti-CD16/CD32 antibodies to block Fcγ receptors. After incubation with biotinylated antibodies, cells were washed, stained with neutravidin

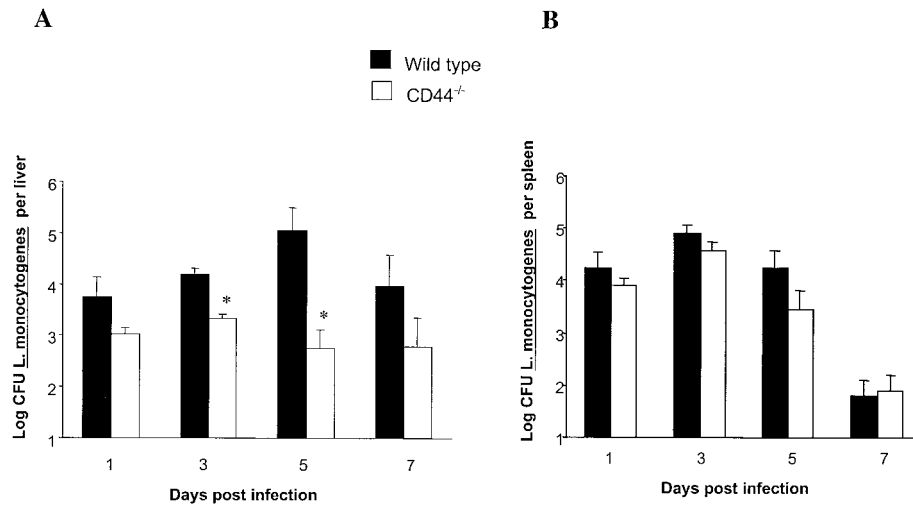


FIG. 3. Diminished susceptibility of CD44^{-/-} mice to infection with *L. monocytogenes*. CD44^{-/-} and WT mice were infected intraperitoneally with 10⁵ CFU of *L. monocytogenes* (0.1 50% lethal infective dose). Mice (eight animals per group) were sacrificed at the indicated time points after infection, and bacterial loads (CFU) were measured in spleens and livers. Mortality was not observed in WT or mutant mouse strains. Asterisks indicate that differences in log₁₀ CFU versus that of WT controls are significant (*P* < 0.05, Student's *t* test).

Alexa (Molecular Probes, Eugene, Oreg.), and analyzed with a FACScan (Becton-Dickinson). All primary antibodies were purchased from Pharmingen (San Diego, Calif.), except for F4/80 (Serotec, Raleigh, N.C.).

Immunostaining of intracellular bacteria. To analyze intracellular bacterial localization, BMM were infected with the *L. monocytogenes* NF-L357 strain, which contains a chromosomal *actA-gfp-plcB* transcriptional fusion, and were immunostained with anti-*L. monocytogenes* serum. *actA/plcB* expression (and, therefore, *gfp*) occurs shortly after *L. monocytogenes* gains access to the host cell cytosol (16). BMM grown on 13-mm²-diameter coverslips were infected with strain NF-L357 as described above. At different time points after infection cells were washed with PBS and fixed with 4% paraformaldehyde in PBS for 15 min, and the cell membrane was lysed with 0.5% Triton X-100 in PBS for 15 min at room temperature. Thereafter cells were washed with PBS, and unspecific bind-

ing was blocked by incubation with PBS containing 5% bovine serum albumin (BSA) and 0.5% Triton X-100. The cultures were then incubated with rabbit anti-*L. monocytogenes* serum (Difco Laboratories, Detroit, Mich.) diluted 1:800 in PBS supplemented with 1% BSA, 0.5% Triton X-100, and 0.01% NaN₃ for 1 h at room temperature. The cells were then washed with PBS and were stained with a Texas-Red-conjugated donkey anti-rabbit IgG (H+L) (Jackson ImmunoResearch Laboratories, West Baltimore Pike, Pa.) diluted 1:100 in PBS containing 1% BSA for 1 h at room temperature and were washed again. Cover slides were analyzed with a fluorescent microscope.

Fluorescent staining of actin filaments. When residing in cytoplasm, *L. monocytogenes* generates an actin comet tail which confers movement to the bacteria, enabling it to enter adjacent cells. Listerial actin tails were stained with fluorochrome-labeled phalloidin. In brief, BMM grown on 13-mm²-diameter cover-

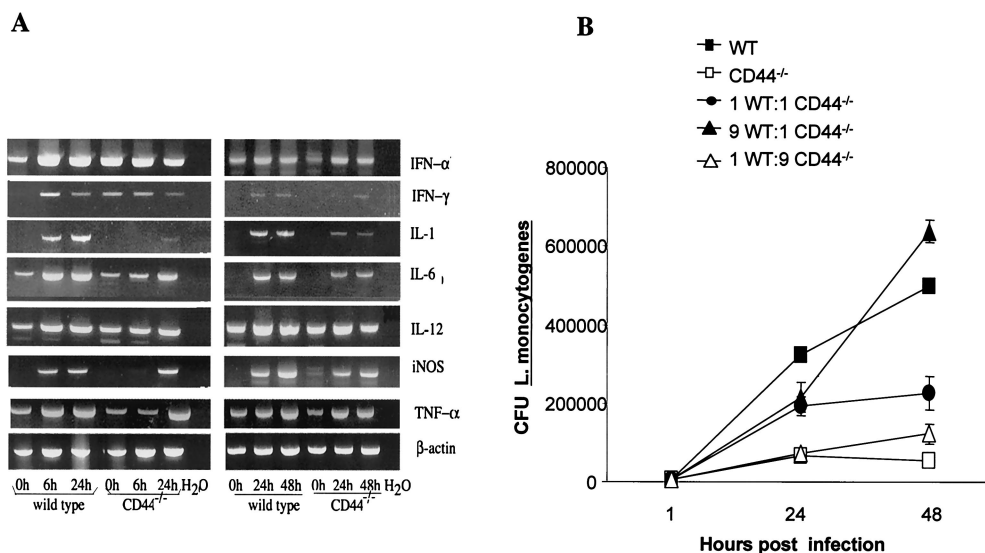


FIG. 4. Role for soluble factors in the reduced intracellular survival of *L. monocytogenes* in CD44^{-/-} BMM. (A) Total RNA was extracted from *L. monocytogenes*-infected CD44^{-/-} and WT BMM at the indicated time points after infection. The accumulation of IFN- α , IFN- γ , IL-1, IL-6, IL-12, iNOS, TNF- α , and β -actin in these samples was visualized by RT-PCR. Two independent experiments are shown. (B) WT and CD44^{-/-} BMM were cocultured at the indicated ratios, and the coculture was infected with *L. monocytogenes*. Controls containing only CD44^{-/-} or WT BMM were also included. All wells contained 2 \times 10⁵ total BMM. CFU were measured at the indicated time points as described in Materials and Methods. A representative from two independent experiments is shown.

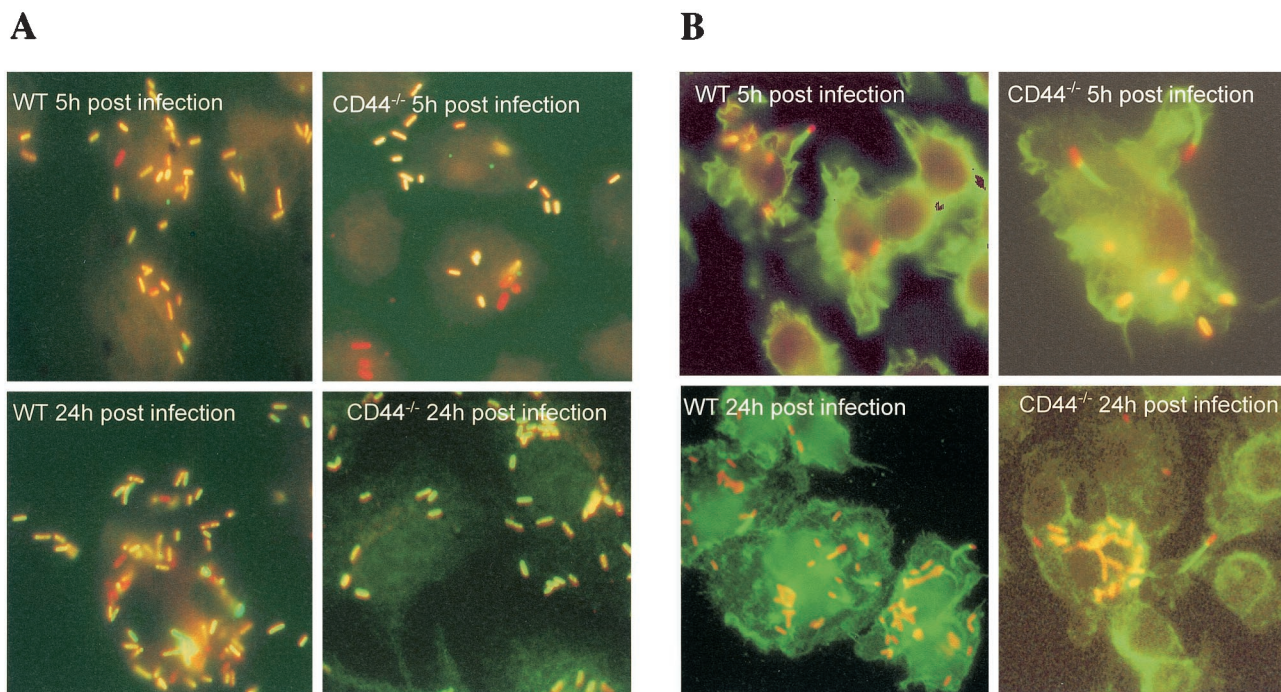


FIG. 5. Role of CD44 in the listerial escape into the cytoplasm and actin tail formation. (A) CD44^{-/-} and WT BMM were infected as indicated in the text with *L. monocytogenes* NF-L357, which contains *gfp* that is expressed in the cytoplasm of the infected cell. At the indicated time points after infection BMM were washed, fixed, incubated with antilisteria sera, and stained with Texas-Red-conjugated donkey anti-rabbit IgG as described in Materials and Methods. Bacteria expressing GFP (in green or yellow) or not expressing GFP (in red) were detected in both WT and CD44^{-/-} BMM. No qualitative difference in the ratio of GFP-positive and -negative bacteria in CD44^{-/-} and WT BMM was observed. On the other hand, lower amounts of *L. monocytogenes* were visualized in CD44^{-/-} BMM. Magnification, 400 \times . (B) *L. monocytogenes*-infected CD44^{-/-} and WT BMM were washed, fixed, and stained with antilisteria-specific antibodies (as indicated in the legend to Fig. 4A) and with Alexa Fluor 488-labeled phalloidin to visualize actin tails. Actin tails were observed on bacteria infecting both CD44^{-/-} and WT BMM. Magnification, 400 \times .

slides and infected with *L. monocytogenes* were used. At indicated time points after infection cells were washed with PBS and were fixed with PBS containing 4% formaldehyde, and the cell membrane was lysed with PBS containing 0.5% Triton X-100. Cells were washed and incubated with anti-*L. monocytogenes* serum as described above, washed, and stained with Alexa Fluor 594-labeled goat anti-rabbit IgG (H+L) (Molecular Probes). The cells were then washed and stained with Alexa Fluor 488-labeled phalloidin (Molecular Probes) for 20 min at room temperature, and coverslips were read with a fluorescent microscope.

RT-PCR assay. Tumor necrosis factor alpha (TNF- α), IFN- γ , IFN- α , interleukin-1 (IL-1), IL-6, IL-12p40, inducible nitric oxide synthase (iNOS), and β -actin mRNA in freshly extracted RNA from *L. monocytogenes*-infected or HA-treated WT or CD44^{-/-} BMM were visualized by reverse transcription (RT)-PCR assays as previously described (34).

Reactions were carried out for 37 cycles (38 cycles for IL-12 and 30 cycles for β -actin) in a thermal cycler with an annealing step at 60 $^{\circ}$ C (67 $^{\circ}$ C for IL-12).

The primer sequences for the amplification of the cDNA are listed in Table 1. IFN- α primer sequences were obtained from Clontech. These primers recognize a consensus sequence present in IFN- α 1, IFN- α 2, and IFN- α 7 genes.

RESULTS

We first compared the outcome of *L. monocytogenes* infection in CD44^{-/-} and WT BMM. Surprisingly, CD44^{-/-} BMM showed a strikingly reduced growth of intracellular *L. monocytogenes* compared to that of WT cells (Fig. 1A). The reduced numbers of *L. monocytogenes* in CD44^{-/-} BMM was not due to a diminished bacterial uptake by the mutant cells, since similar numbers of intracellular *L. monocytogenes* were measured in mutant and WT BMM at 1 h after infection (Fig. 1B). By using *L. monocytogenes* containing one copy of the temper-

ature-sensitive, erythromycin-resistant gene containing plasmid pAUL-A per bacteria, it was shown that the reduced intracellular number of *L. monocytogenes* in CD44^{-/-} BMM was due to increased rate of death rather than to reduced growth rate of the bacteria (Table 2).

To test for the presence of CD44-facilitated listerial growth in cell types other than BMM, we studied *L. monocytogenes* infection in primary lung and heart fibroblast cultures from CD44^{-/-} and WT mice. Listerial number was again lower in CD44^{-/-} cells from both sources (Fig. 1C and D).

Next we analyzed whether CD44 is a general supporter of intracellular growth of bacterial pathogens. However, no differences in uptake (data not shown) or intracellular growth of *S. enterica* serovar Typhimurium in CD44^{-/-} and WT BMM were found (Fig. 2).

We next explored whether reduced in vitro growth of *L. monocytogenes* in CD44^{-/-} BMM related to a diminished susceptibility of CD44^{-/-} mice to bacterial infection in vivo. WT and CD44^{-/-} mice were infected intraperitoneally with 10⁵ *L. monocytogenes* cells, and bacterial numbers were quantified in spleen and liver at different time points after infection. The livers from CD44^{-/-} mice showed a significantly diminished number of *L. monocytogenes* compared to that of WT controls, whereas bacterial numbers recovered from the spleens of both mouse strains were similar (Fig. 3).

We found no variation in phenotype of CD44^{-/-} BMM

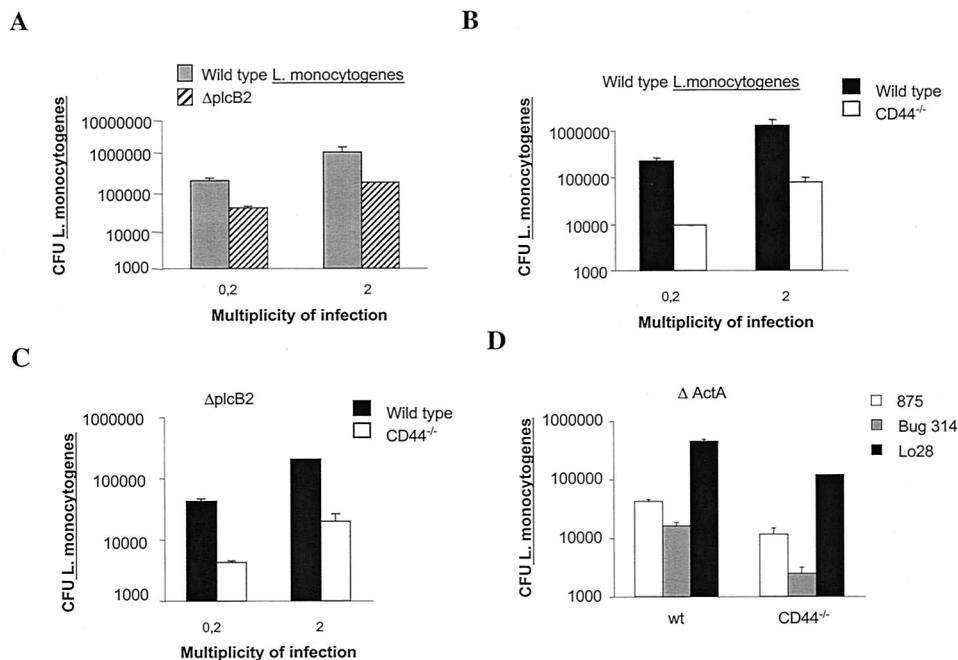


FIG. 6. Role of *plcB2* and *actA* in the reduced bacterial survival in CD44^{-/-} BMM. WT (A, B, and C) and CD44^{-/-} (B and C) BMM were infected with the Δ *plcB2* mutant (A and C) or the EGD parental strain (A and B) of *L. monocytogenes* at an MOI of 2 and 0.2 bacteria per cell, respectively. WT and CD44^{-/-} BMM were infected with the Δ *actA* mutants BUG314 or BUG875 or the LO28 parental strain (D). Twenty-four hours after infection cells were lysed and the CFU in cell lysates were quantified. A representative of two independent experiments is shown.

compared to that of WT BMM that could be related to the diminished growth of *L. monocytogenes*. CD44^{-/-} and WT BMM both expressed high levels of Mac-3, GR-1, F4/80, and MHC class II, expressed moderate levels of CD80 and CD14, and were negative for T, B, or NK cell markers, such as CD3, CD4, CD8, CD19, and NK1.1 as measured by fluorescence-activated cell sorting (data not shown).

We then studied if different levels of cellular activation could account for the diminished intracellular growth of *L. monocytogenes* in CD44^{-/-} BMM. For this purpose RNA was extracted from BMM before and at different time points after infection, and the presence of transcripts for inflammatory cytokines was evaluated by RT-PCR. The levels of IFN- γ , IFN- α , IL-12p40, IL-1, IL-6, iNOS, and TNF- α mRNA were all increased in *L. monocytogenes*-infected WT and CD44^{-/-} BMM. CD44^{-/-} BMM contained equal or lower levels of all transcripts than WT BMM (Fig. 4A). Thus, the enhanced death rate of *L. monocytogenes* in CD44^{-/-} BMM is not linked to the presence of higher levels of inflammatory cytokine transcripts.

We next analyzed intracellular bacterial growth by using various cocultures of CD44^{-/-} and WT BMM. We hypothesized that an increased secretion of inflammatory cytokines from CD44^{-/-} BMM or the secretion of soluble factors promoting bacterial growth by WT BMM could modify the listerial numbers in either direction in the coculture. No shift in the expected bacterial numbers in these cocultures was detected (Fig. 4B), suggesting that soluble factors account neither for the enhanced listerial growth in the presence of CD44 nor for the increased bacterial death rate in its absence.

CD44 might be necessary for escape of *L. monocytogenes* from phagosome to cytoplasm. To study this, BMM were in-

fectured with the NF-L357 strain of *L. monocytogenes* containing a *gfp* gene only expressed when bacteria localize in the cytoplasm (16). Staining *gfp*-expressing bacteria with anti-*L. monocytogenes* sera, we found similar ratios of *gfp*-expressing and nonexpressing bacteria in CD44^{-/-} and WT BMM. WT BMM showed higher total numbers of bacteria than CD44^{-/-} BMM. Thus, CD44 plays no detectable role in the escape of *L. monocytogenes* from the phagosome into the cytoplasm (Fig. 5A).

CD44 may participate in the formation of actin tails by *L. monocytogenes*, thereby reducing bacterial growth in CD44^{-/-} BMM. However, intracellular *L. monocytogenes* with actin tails was observed both in CD44^{-/-} and in WT BMM (Fig. 5B).

The possible importance of cell-to-cell spread in CD44-enhanced growth of *L. monocytogenes* was next examined. For this purpose cells were infected with Δ *plcB2* or Δ *actA* *L. monocytogenes* mutants (showing reduced or abolished cell-to-cell spread). As expected, WT BMM showed lower numbers of bacteria when infected with the Δ *plcB2* or the Δ *actA* mutants than with the respective WT strains of *L. monocytogenes* (Fig. 6A and D). CD44^{-/-} BMM showed lower levels of Δ *plcB2*, LO28 Δ *actA*, or BUG375 Δ *actA* *L. monocytogenes* compared to those shown by WT BMM (Fig. 6C and D). Thus, the diminished bacterial growth in CD44^{-/-} BMM is unlikely to be due to a reduction in bacterial cell-to-cell spreading.

Secretion of HA by infected BMM might mediate the CD44-dependent enhanced intracellular listerial growth. BMM were thus treated with HA'se before and after infection with *L. monocytogenes* (Fig. 7A). Coincubation of WT BMM with HA'se, however, increased the intracellular proliferation of *L. monocytogenes*. In agreement with this result, coincubation of BMM with HA reduced intracellular bacterial growth (Fig. 7C) while enhancing accumulation of iNOS, IL-1, IL-6, and IFN- γ

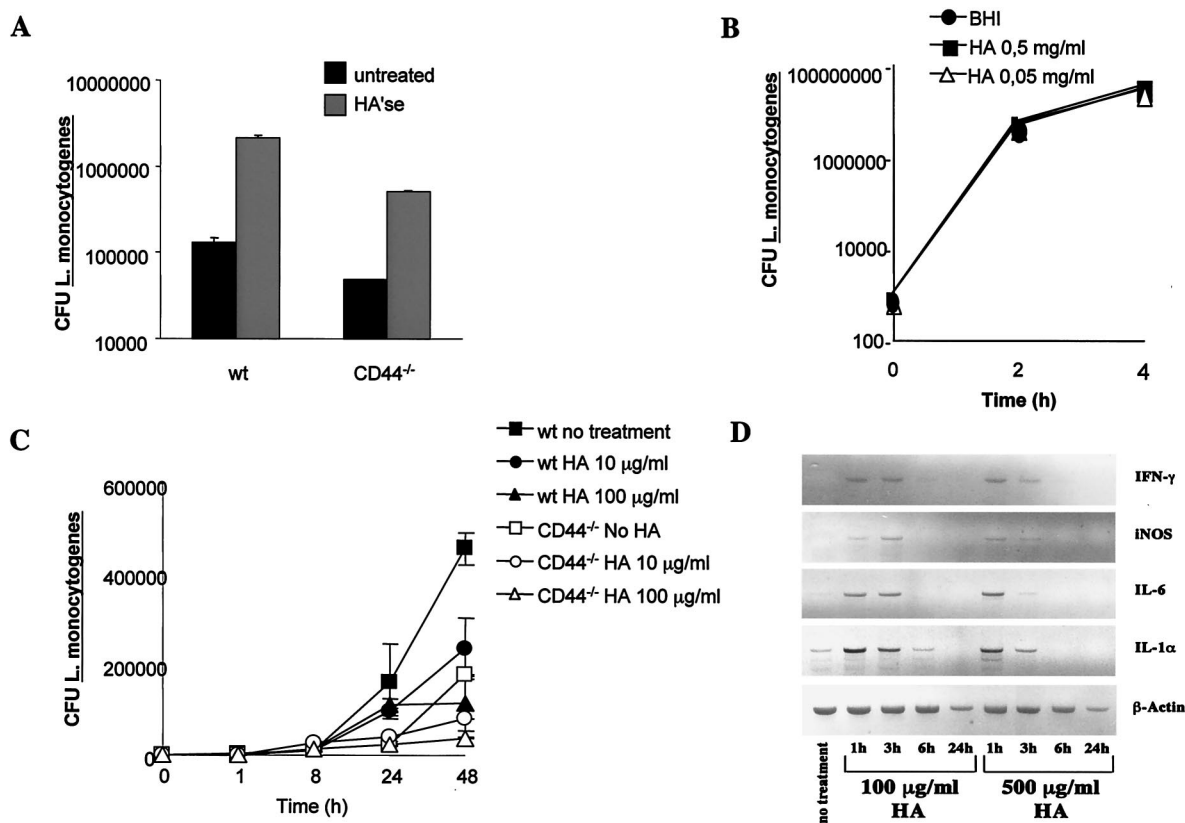


FIG. 7. Effect of addition or depletion of HA in the intracellular growth of *L. monocytogenes* in BMM. (A) CD44^{-/-} and WT BMM were coincubated with 2 U of HA'se per ml of DMEM-5% FCS at 37°C, 5% CO₂ 1 h before and during infection with *L. monocytogenes*. The bacterial load (CFU) of treated and control BMM was quantified 24 h after infection. A representative of four independent experiments is depicted. Addition of HA (250 kDa) to cocultures of HA'se and BMM completely inhibited the enzyme-mediated enhancement of bacterial proliferation, confirming the specificity of the enzymatic reaction (data not shown). (B) *L. monocytogenes* was grown at 37°C in BHI medium in the presence of 50 and 500 µg of HA (190 kDa) or in the absence of HA. CFU were recorded in culture aliquots at the indicated time points of incubation. (C) WT and CD44^{-/-} BMM were treated with 10 and 100 µg of HA (190 kDa) per ml 1 day before and during infection with *L. monocytogenes*. The bacterial load (CFU) of treated and control BMM was quantified at the indicated time points after infection. A representative from two independent experiments is shown. (D) WT BMM were treated with 100 and 500 µg of HA/ml (190 kDa) diluted in DMEM-5% FCS. The presence of IFN-γ, iNOS, IL-6, IL-1α, and β-actin mRNA in treated and control BMM was determined by RT-PCR at the indicated hours after treatment. A representative from two independent experiments is shown.

mRNA (Fig. 7D). Control experiments showed that HA as such did not affect extracellular growth of *L. monocytogenes* (Fig. 7B). However, HA and HA'se also affected listerial proliferation in CD44^{-/-} BMM in a manner similar to that of WT BMM (Fig. 7A and C). Thus, hyaloadherins other than CD44 participate in HA-mediated control of listerial infection. To analyze whether CD44 plays a role in modulation of intracellular bacterial growth by HA, BMM were treated with monoclonal antibodies neutralizing (KM 81) or not neutralizing (KM 703) the HA binding ability of CD44 (47). Anti-MHC class I and anti-CD4 monoclonal antibodies were used as controls. Incubation of WT but not CD44^{-/-} BMM with KM 81 resulted in an increased intracellular proliferation of *L. monocytogenes*, whereas no such consequence was observed after incubation of BMM with KM 703 or the control monoclonal antibodies (Fig. 8). Thus, whereas signaling via CD44 facilitates listerial growth, addition of HA can restrict growth occurring in part via binding to CD44.

Proteoglycan forms of CD44 also exhibit affinity for molecules, such as chondroitin sulfate, HS, fibronectin, and osteo-

pontin (31). Treatment of BMM with chondroitin sulfatase also enhanced intracellular listerial growth (Fig. 9A), whereas no effect on the outcome of infection was observed upon adding HS or HS'se (Fig. 9B and C).

DISCUSSION

Herein we demonstrate that the presence of CD44 facilitates intracellular growth of *L. monocytogenes* in murine systems. Changes in bacterial proliferation did not relate to altered bacterial phagocytosis or secretion of proinflammatory cytokines in CD44^{-/-} BMM. The presence of CD44 increased bacterial survival without altering growth rate and was not required for bacterial escape from primary phagosome into cytoplasm or for actin tail formation. CD44-facilitated listerial growth does not depend on functional cell-to-cell spreading by *L. monocytogenes*. These results altogether suggest that CD44 signaling affects survival of *L. monocytogenes* in the cytoplasm of the primary infected cell. In agreement with this suggestion, inhibitors of tyrosine protein kinases, phosphatidylinositol 3-

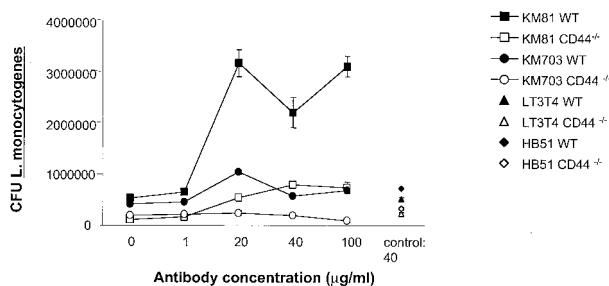


FIG. 8. Increased intracellular listerial growth in BMM coincubated with a monoclonal antibody specifically neutralizing the HA binding ability of CD44. CD44^{-/-} and WT BMM were treated with different concentrations of the anti-CD44 monoclonal antibodies KM 81 and KM 703 1 h before and during infection with *L. monocytogenes*. Anti-CD4 (LT3T4) and anti-MHC class I (HB51) antibodies were used as controls. The bacterial load (CFU) of treated and control BMM was quantified after 24 h of infection.

kinase, and actin polymerization enzymes that mediate CD44 signaling all reduced listerial growth in WT macrophages (data not shown). These inhibitors, however, do not uniquely affect CD44 activation. To our knowledge, such a role for CD44 in determining the intracellular fate of bacteria has not been described. This CD44-enhanced intracellular listerial survival was observed both in cultures of primary macrophages and fibroblasts and is likely to be directly related to diminished *L. monocytogenes* numbers in livers of CD44^{-/-}-infected mice. The presence of different immune effector cells (10) participating in listerial control in spleen and liver could account for the dissimilar involvement of CD44 in listerial control in these organs. Alternatively, CD44-mediated intracellular listerial proliferation could be more prevalent in liver compared to that in spleen cell populations.

On the other hand, CD44 did not affect the intracellular growth of another intracellular pathogen, *S. enterica* serovar Typhimurium.

We plan to use cells transfected with CD44 lacking intracellular protein domains necessary for different signaling pathways to further dissect the role of this cell receptor in listerial growth. CD44 undergoes alternative splicing (encompassing 10 variant exons) giving rise to isoforms that probably show differences in their biological functions (31), and it could relate to the regulatory function of the receptor during infection with

L. monocytogenes. Alternative splicing will not take place in the transfected molecule. Affinity of CD44 for ligands is dependent on posttranslational modifications, such as glycosylation, that are cell type and growth condition specific (38), and it could be also altered by listerial infection. CD44 has been suggested to act as a functional coreceptor for hepatocyte growth factor/scatter factor by binding and presenting growth factor to the receptor tyrosine kinase c-Met (44). The bacterial surface protein InlB promotes not only internalization (5, 14) but also the escape into the cytoplasm of *L. monocytogenes* in mammalian cells (19). In turn, the Met receptor mediates InlB-dependent internalization of *L. monocytogenes* (37). However, mutants lacking InlB (as well as InlA and InlA/B) showed diminished intracellular growth in CD44^{-/-} BMM compared to that of WT BMM (data not shown).

There is indirect evidence that the ability to replicate within host cells requires specific adaptations in the microbe. A hexose phosphate transporter (hpt) has been identified as a virulence factor involved in the listerial replication phase (9, 17). Whether listerial hpt and other virulence factor(s) specifically interact with CD44 signaling and promote listerial growth will be further studied.

CD44 signaling has been indicated to affect the outcome of other extracellular and intracellular infections. Binding of the HA capsule of *Streptococcus pyogenes* to CD44 mediates cytoskeletal rearrangements (12, 13). Such rearrangements disrupt the intercellular junctions and permit bacterial passage between epithelial cells (13). Interaction of the *Shigella flexneri* IpaB-IpaC-secreted complex with CD44 has also been suggested to be involved in the cellular uptake of shigella and in shigella-induced cytoskeletal rearrangements (39).

Inflammatory stimuli promote local accumulation of LMW HA fragments (6, 42). We confirmed previous reports (22, 23, 26, 29, 41) indicating that coincubation of cells with LMW HA increased proinflammatory transcripts and demonstrate further that such coincubation diminished listerial growth in macrophages. Eliminating HA via HA'se treatment accelerated listerial proliferation, which is in line with the possibility that macrophage-derived HA acts by binding to hyaloadherins and diminishes listerial growth. A role for CD44 in HA-mediated control of listerial growth was further suggested by our finding of an accelerated listerial growth in WT BMM treated with a monoclonal antibody that selectively interferes with HA binding to CD44 (47). However, and in line with the fact that CD44

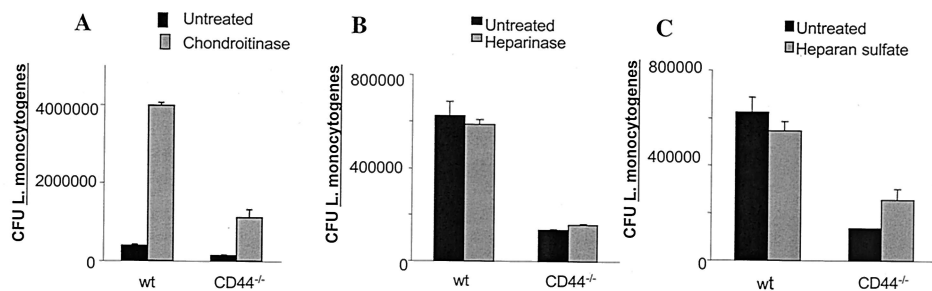


FIG. 9. Effect of treatment with chondroitinase, HS'se, and HS on the intracellular growth of *L. monocytogenes*. CD44^{-/-} and WT BMM were coincubated with 0.01 U chondroitinase (A), 0.5 U of HS'se (B), or 20 µg of HS (C) per ml of DMEM-5% FCS at 37°C 1 h before and during infection with *L. monocytogenes*. The bacterial load (CFU) of treated and control BMM was quantified at 24 h (A) and at the indicated time points (B and C) after infection. A representative from two independent experiments for each condition is shown.

is not the sole receptor reacting with HA (1, 42, 46), other cellular receptors also participate in the control of listerial infection, since HA's enhanced listerial growth in CD44^{-/-} BMM and coinubation with HA diminished listerial growth in these cells. Adding complexity to this analysis is the fact that CD44 not only binds HA but also binds HS, chondroitin sulfate, collagen, fibronectin, and osteopontin (42). In this study we found that treatment of BMM with chondroitin sulfatase but not HS'se increased listerial intracellular growth, a finding we will further explore.

Altogether our results suggest a dual role of CD44 in the outcome of infection with *L. monocytogenes*; whereas a quiescent receptor (or low levels of ligands) will deliver signals that will facilitate intracellular listerial growth, ligands binding to CD44 will diminish such response.

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