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Received 26 March 2004/Returned for modification 29 April 2004/Accepted 3 June 2004

Yersinia pestis **and** *Yersinia pseudotuberculosis* **are closely related facultative intracellular pathogens. The response regulator PhoP was previously shown to be important for** *Y. pestis* **survival in macrophages and for virulence in a murine bubonic plague infection assay. Here the importance of PhoP for** *Y. pseudotuberculosis* **pathogenesis was investigated.** *Y. pseudotuberculosis phoP* **mutants were unable to replicate in low-Mg2 medium or in macrophages.** *phoP Y. pseudotuberculosis* **strains initiated replication in macrophages after a lag period of** -**5 h, as shown by fluorescence microscopy and viable count assays.** *Y. pseudotuberculosis phoP* **mutants died at a low rate in macrophages; there was no decrease in viability over the first 5 h of infection, and there was a 10-fold decrease in viability between 5 and 24 h of infection. Trafficking of phagosomes containing** *phoP* **or** *phoP* **mutant** *Y. pseudotuberculosis* **was studied by using immunofluorescence microscopy and cathepsin D as a marker for lysosomes. Phagosomes containing** *phoP* **mutant** *Y. pseudotuberculosis* **acquired cathepsin D** at a higher rate than phagosomes containing $pho\overline{P}^+$ bacteria. However, the increased rate of marker **acquisition for phagosomes containing mutant bacteria was only evident** -**5 h after infection, suggesting that** *phoP* **mutants are able to retard phagosome maturation during the lag phase of intracellular growth. The results obtained with a** *Y. pestis phoP* **mutant were similar to those described above, except that the rates of intracellular killing and trafficking to cathepsin D-positive vacuoles were significantly higher. A** *Y. pseudotuberculosis phoP* **mutant was 100-fold less virulent than the wild-type strain in a murine intestinal infection model, suggesting that survival and replication in macrophages are important for** *Y. pseudotuberculosis* **pathogenesis.**

Yersinia pestis and *Yersinia pseudotuberculosis* are closely related gram-negative bacterial pathogens. *Y. pseudotuberculosis* is an enteropathogenic bacterium typically associated with acute infections of mesenteric lymph nodes (5). Twenty-one different O antigen serogroups of *Y. pseudotuberculosis* have been identified (41). *Y. pestis*, the agent of bubonic and pneumonic plagues, lacks O antigen and is considered a subspecies of *Y. pseudotuberculosis* (1, 41). Three biovars of *Y. pestis* have been identified, biovars Antiqua, Mediaevalis, and Orientalis, each of which has been associated with a major pandemic (32). *Y. pseudotuberculosis* and *Y. pestis* encode a number of common virulence determinants, including factors that promote coordinated gene expression and acquisition of iron (32). Another virulence factor common to *Y. pestis* and *Y. pseudotuberculosis* is a type III secretion system (TTSS) encoded on a 70-kb plasmid (pCD1 in *Y. pestis* and pYV in *Y. pseudotuberculosis*) (34). The plasmid-encoded TTSS is required for sustained bacterial replication in host tissues and exports multiple proteins known as the Yops and LcrV. Several of the Yops are delivered into host cells, where they inhibit phagocytosis and proinflammatory immune responses (22). In addition, LcrV is

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secreted into the extracellular milieu, where it downregulates inflammation (6).

Y. pseudotuberculosis is typically transmitted via the fecaloral route (13). The bacteria can cross the intestinal barrier through M cells and reach the underlying Peyer's patches. From the Peyer's patches *Y. pseudotuberculosis* disseminates to mesenteric lymph nodes, where the bacteria replicate and cause mesenteric lymphadenitis. *Y. pseudotuberculosis* has been shown to replicate as an extracellular pathogen in lymphoid tissues of mice at late stages of infection (40).

Y. pestis infections are typically transmitted by fleas. Fleas feeding on infected rodents transmit the bacteria to other animals or humans (32). The bacteria disseminate from the site of the fleabite to nearby lymph nodes, where they replicate, resulting in a swollen lymph node (or bubo) that is characteristic of the disease. From the lymph nodes the bacteria spread systemically and colonize organs such as the liver, lungs, and spleen. Infected hosts die of septic shock when large amounts of bacteria from infected organs are released into the bloodstream. *Y. pestis* can also infect the lungs directly via aerosols and cause pneumonic plague. Pneumonic plague is highly infectious, can be spread from human to human, and is typically fatal.

It is thought that *Y. pestis* survives and replicates in macrophages at early stages of the infection process in humans and animals (8). In agreement with this idea, *Y. pestis* can replicate

in cultured primary mouse macrophages (8, 36, 43). It is not known if *Y. pseudotuberculosis* survives or replicates in macrophages in vivo, although a number of different *Y. pseudotuberculosis* strains can replicate in cultured primary murine macrophages (36, 45). A notable exception to this finding is *Y. pseudotuberculosis* strain YPIII, which is unable to replicate in macrophages (28, 36). The genetic basis for the difference between YPIII and other *Y. pseudotuberculosis* strains with respect to the intracellular growth phenotype is not known. Histological analysis of Peyer's patches from rabbits infected with *Y. pseudotuberculosis* has shown that the bacteria can be internalized by intraepithelial macrophages at the early stages of the infection (15). Therefore, survival and replication in macrophages might play an important role in the *Y. pseudotuberculosis* infection process.

Macrophages are an important first line of defense of the innate immune system. One of their main functions is to take up bacteria by phagocytosis, digest them, and present their antigens to cells of the adaptive immune system. After a bacterium is internalized in a phagosome, the phagosome undergoes fusion with organelles of the endocytic pathway (e.g., late endosomes and lysosomes) to acquire degradative enzymes (e.g., cathepsin D). As the phagosome matures, the pH drops and ions (e.g., Mg^{2+}) are pumped out of the vacuole. In the fully mature phagosome, or phagolysosome, the bacterium is destroyed and digested (46). A number of pathogenic bacteria that survive and replicate in macrophages have evolved distinct and intricate mechanisms to avoid destruction in macrophage phagolysosomes. Pathogens that can prevent maturation of the phagosome into a phagolysosome include members of *Salmonella* and *Mycobacterium* species (12).

The genetic basis for the ability of *Y. pestis* and *Y. pseudotuberculosis* to survive and replicate in macrophage phagosomes is largely unknown. Strains cured of the virulence plasmid can replicate in cultured macrophages (36, 43, 45), indicating that the TTSS encoded on the virulence plasmid is not required for intracellular survival. A second distinct TTSS encoded on the chromosomes of *Y. pestis* and *Y. pseudotuberculosis* is also dispensable for bacterial replication in cultured macrophages (36). One gene that is known to be required for survival of *Y. pestis* in macrophages is *phoP*, which encodes the PhoP response regulator (29).

The importance of PhoP for bacterial survival and replication in macrophages has been extensively studied in *Salmonella enterica* serovar Typhimurium (14, 18). PhoP is part of a twocomponent system in which PhoQ is the sensor kinase and PhoP is the response regulator. At low Mg^{2+} concentrations PhoQ is phosphorylated and the phosphate is transferred to PhoP. PhoP activates a variety of genes, many of which are important for virulence. PhoP-regulated genes allow serovar Typhimurium to adapt to the low- Mg^{2+} environment of the phagosome (3) and to prevent vacuole maturation to a phagolysosome (17). Genes such as *mgtA* and *mgtB* are directly involved in the adaptation to a low- Mg^{2+} environment since they encode P-type ATPases that function as putative Mg^{2+} transporters (14, 18).

Although PhoP plays an important role in *Y. pestis* virulence (29), it is not known if PhoP has a comparable function in *Y. pseudotuberculosis*. This study was undertaken to determine the importance of PhoP for *Y. pseudotuberculosis* pathogenesis.

We obtained evidence that PhoP in *Y. pseudotuberculosis* is important for growth in a low- Mg^{2+} environment and for survival and replication in macrophages. PhoP function appears to be important for *Y. pseudotuberculosis* to retard the maturation of its vacuole into a phagolysosome. A *Y. pseudotuberculosis phoP* mutant is 100-fold less virulent than a wild-type strain in a murine intestinal infection model, suggesting that replication in macrophages is important for *Y. pseudotuberculosis* pathogenesis. Based on similar results obtained with *Y. pestis* in this study and in another study (29), we concluded that the PhoP function is conserved in *Y. pseudotuberculosis* and *Y. pestis*.

MATERIALS AND METHODS

Bacterial strains and growth conditions. The *Y. pseudotuberculosis* and *Y. pestis* strains used in this study are listed in Table 1. YPIII(p^-) and YPIII(p^+) were obtained from the strain collection of Stanley Falkow. *Y. pestis* and *Y. pseudotuberculosis* strains were cultivated at 28°C on heart infusion (HI) (Difco) or Luria-Bertani (LB) agar plates. Cultures were grown in broth with aeration overnight at 28°C. *Y. pestis* was grown in HI broth. *Y. pseudotuberculosis* was grown in LB broth or HI broth depending on the experimental conditions. The *Escherichia coli* strains NovaBlue (Novagen), SM10, and S17-1, the latter two lysogenized with *pir*, were grown in LB broth or on LB agar plates at 37°C. Bacterial growth media were supplemented with ampicillin at a concentration of 100 μ g/ml, kanamycin at a concentration of 25 μ g/ml, or tetracycline at a concentration of 20 μ g/ml when appropriate.

To measure growth of *Y. pestis* and *Y. pseudotuberculosis* under Mg²⁺-limiting conditions, the defined medium TMH (42) was used with the following modifications: $MgCl₂$ was omitted, and the final pH was adjusted to 5.5 (this medium is referred to below as low- Mg^{2+} TMH). A pH of 5.5 was initially used in an attempt to mimic the low pH of the phagolysosome; however, subsequent experiments indicated that low pH was not an important variable since *phoP* mutants were able to replicate at pH 5.5 when the medium described above was supplemented with 10 mM $MgCl₂$ (high- $Mg²⁺$ TMH). Bacterial cultures grown overnight in HI medium were diluted into low- or high-Mg²⁺ TMH to obtain an optical density at 600 nm of \sim 0.1. Growth at 37°C was recorded by measuring the optical density at 600 nm at different times. The following modifications were made for strains carrying *phoP* expression vectors. Cultures grown overnight were diluted 1:10 into fresh HI medium containing 500 μ M isopropyl- β -Dthiogalactopyranoside (IPTG). The cultures were incubated for 2 h at 28°C to induce *phoP* expression. The cultures were then diluted into low-Mg²⁺ TMH supplemented with 100 μ M IPTG, and growth at 37°C was recorded as described above.

DNA methods. *Taq* polymerase and deoxynucleoside triphosphates (dNTPs) purchased from Roche were used for PCRs. For purification of plasmid DNA, QIAGEN Mini and Midi Prep DNA purification kits were used according to the manufacturer's protocols. Fast Link ligase (Epicentre) was used for ligation of DNA. Vectors were dephosphorylated with shrimp alkaline phosphatase from Roche when necessary. DNA restriction enzymes were purchased from Roche and were used according to the manufacturer's protocol. For cycle sequencing an ABI Prism BigDye Terminator v3.0 Ready Reaction cycle sequencing kit was used.

Sequence analysis of *phoP* **and construction of** *phoP* **expression vectors.** The oligonucleotides 5-GAATTCATGCGGGTTCTGGTTGTGG-3 (PhoPF) and 5'-CTAGTTGACGTCAAAACGATATCCC-3' (PhoPR) were used to amplify the *phoP* open reading frame (ORF) from *Y. pestis* KIM10⁺ (*phoP^{KIM}*) and *Y*. *pseudotuberculosis* YPIII(p⁻) (*phoP^{YPIII}*) by PCR with an Eppendorf Mastercycler. The PhoPF primer was designed to introduce an EcoRI restriction site (underlined nucleotides in the primer sequence) in front of the start codon of the *phoP* ORF. The reactions were performed with genomic DNA or whole bacteria as a template in 100 μ l of 1 × PCR buffer (10 mM Tris-HCl, 1.5 mM MgCl₂, 50) mM KCl; pH 8.3) containing 2.5 U of *Taq* DNA polymerase (Roche), each primer at a concentration of 0.5 μ M, and 160 μ M dNTPs. The PCR products were cloned into the EcoRV site of the pETBlue-2 cloning vector by using a Perfectly Blunt cloning kit (Novagen). The sequences of the inserts were verified by cycle sequencing by using pETBlueUP and pETBlueDOWN primers as specified in the manufacturer's manual. Additionally, the *phoP* ORFs from YPIII(p^-) (*phoP^{YPIII}*) and YPIII(p^+) [*phoP^{YPIII(* p^+)] were amplified by using} primers PhoP2F (5' CGTTTACATTGTGACGGCGTC 3') and PhoP2R (5' GTTATGCCACTGTGCCAGACTG 3) and whole bacteria as the template. The PCR product was purified and used for cycle sequencing with the same set

TABLE 1. *Yersinia* strains used in this study

Strain	Relevant characteristics	Reference
$KIM10+$	Biovar Mediaevalis, pCD1 ⁻ , pPCP1 ⁻	33
$KIM10^{+}/GFP$	KIM 10^+ , p67GFP3.1, Ap ^r	36
KIM10 ⁺ $phoP\Delta$	KIM 10^+ phoP Δ 127-429	This study
KIM10 ⁺ $phosP\Delta$ /PhoP ^{KIM}	KIM10 ⁺ PhoP Δ , pPhoP ^{KIM} , Ap ^r	This study
KIM10 ⁺ $phoP\Delta/MMB67EH$	KIM10 ⁺ phoP Δ , pMMB67HE, Ap ^r	This study
KIM10 ⁺ $phoP\Delta/GFP$	KIM10 ⁺ phoP Δ , p67GFP3.1, Ap ^r	This study
IP2666c	Serogroup $O3$, pYV^-	39
IP2666c/GFP	IP2666c, p67GFP3.1, Apr	36
IP2666cphoP Δ	IP2666c phoPΔ127-429	This study
IP2666cphoP Δ /PhoPKIM	IP2666c phoP Δ , pPhoP KIM , Apr	This study
IP2666cphoP Δ /PhoP Y PIII	IP2666c phoP Δ , pPhoP Y PIII, Apr	This study
IP2666cphoP Δ /MMB67EH	IP2666c phoP Δ , pMMB67EH, Ap ^r	This study
IP2666cphoP Δ /GFP	IP2666cphoP Δ , p67GFP3.1, Ap ^r	This study
$YPIII(p^+)$	Serogroup O3, pYV ⁺ , phoPM116H, R117Stop	
$YPIII(p^-)$	Serogroup O3, pYV ⁻ , phoP ^{T160P}	
$YPIII(p^-)/PhoP^{KIM}$	$YPIII(p^-)$, pPhoP ^{KIM} , Ap ^r	This study
$YPIII(p^-)/MMB67EH$	$YPIII(p^-)$, pMMB67EH, Ap ^r	This study
32777	Serogroup O1, pYV	10
32777/GFP	32777, p67GFP3.1, Ap ^r	This study
32777phoP::kan	32777 phoP:: kan , Knr	24
32777phoP::kan/GFP	32777phoP::kan, p67GFP3.1, Kn ^r Ap ^r	This study
32777phoP::kan/MMB67EH	32777phoP::kan, pMMB67EH, Kn ^r Ap ^r	This study
32777phoP::kan/PhoP	$32777phoP$::kan, pPhoP, Kn ^r , Ap ^r	This study

of primers. Cycle sequencing was performed with an ABI 3100 automated genetic analyzer by using an ABI PRISM BigDye Terminator cycle sequencing kit (Applied Biosystems). The sequencing data were compared to *Yersinia phoP* sequences available in GenBank by using BLASTN. The sequences of *phoPYPIII* and $phoP^{YPIII(p+)}$ were submitted to GenBank.

pETBlue-2 plasmid DNAs containing *phoPKIM* or *phoPYPIII* were purified and digested with BamHI and EcoRI to excise the *phoP* fragments. The resulting *phoP* fragments were ligated into BamHI- and EcoRI-digested pMMB67EH (16) to place *phoP* under control of the IPTG-inducible *tac* promoter. The resulting
plasmids were designated pPhoP^{KIM} and pPhoP^{YPIII}. pPhoP^{KIM} and pPhoP^{YPIII} were electroporated into S17 *Npir*, and transformants were selected with ampicillin. S17 *pir* was used to transfer pPhoPKIM and pPhoPYPIII or their parent, pMMB67EH, into different *Y. pestis* and *Y. pseudotuberculosis* strains by conjugation. Transconjugants were selected on *Yersinia* selective medium (Oxoid) supplemented with ampicillin.

Creation of *phoP* **deletion mutants.** To create an in-frame deletion mutation in *phoP*, pETBlue-2 plasmid DNA containing *phoPKIM* was digested with HpaI and SphI to remove nucleotides 127 to 429 of the *phoP* sequence. After restriction digestion, 1 U of T4 DNA polymerase and 100μ M dNTPs were added to the reaction mixture and incubated for 15 min at 12°C to remove 3' single-stranded regions. The larger restriction fragment containing pETBlue-2 and the 5' and 3' segments of *phoP^{KIM}* was purified and ligated. The resulting plasmid, pETBlue-2phoP Δ , was digested with BamHI and XbaI. The restriction fragment containing $phoP\Delta$ was inserted between the BamHI and XbaI sites of the suicide vector pSB890 (30). The resulting plasmid, pSB890phoP Δ , was electroporated into S17 ρ *pir*, and transformants were selected with tetracycline. S17 ρ *pir*/pSB890*phoP* Δ was used to transfer the plasmid to different *Yersinia* strains by conjugation. Transconjugants were selected with tetracycline on *Yersinia* selective medium and streaked on LB plates supplemented with tetracycline. Tet^r transconjugants were grown for several generations in the absence of tetracycline and then plated on LB agar plates lacking NaCl and supplemented with 5% sucrose to select against the *sacB* gene carried on pSB890. Sucrose-resistant colonies were tested for the Tet^s phenotype and for the presence of $phoP\Delta$ by using colony PCR and primers for the *phoP* ORF as specified above. *Yersinia* strains containing the $phoP\Delta$ allele were defective for growth in low-Mg²⁺ TMH at 37°C.

Analysis of intracellular survival and replication by CFU assay. The mouse macrophage-like cell line J774A.1 was cultured in Dulbecco's modified Eagle's medium (Gibco) with a high glucose concentration (4,500 μ g/ml) supplemented with 10% heat-inactivated fetal bovine serum (HyClone), 2 mM L-glutamine, and 1 mM sodium pyruvate at 37° C with 5% CO₂. Cells were grown to confluence and detached from the dishes, and 2×10^5 cells were seeded into wells of 24-well tissue culture plates and incubated overnight. Bacterial cultures were grown overnight in HI broth at 28°C with shaking. The cultures were centrifuged for 5

min at $2,000 \times g$ at room temperature, and the pellets were resuspended in phosphate-buffered saline (PBS) (Gibco). Dilutions were made in PBS as needed, and the final dilution was made in complete tissue culture medium (see above). The bacterial numbers were adjusted so that the multiplicity of infection (MOI) was two to five bacteria per cell. After addition of the bacteria to the cells, the 24-well plate was centrifuged for 5 min at 200 \times g at room temperature to facilitate bacterial contact with the cells. After 15 min of infection the cells were washed twice with PBS. Some of the wells were used to measure the initial number of cell-associated bacteria by a CFU assay as described below. To the remaining wells, tissue culture medium containing 8μ g of gentamicin per ml was added to kill extracellular bacteria. After 1 h, the medium was removed and replaced with fresh complete tissue culture medium containing 2μ g of gentamicin per ml (for KIM10⁺-infected cells) or 4.5 µg of gentamicin per ml (for *Y*. *pseudotuberculosis*-infected cells), and incubation was continued. At different times thereafter the cells were washed, and the number of intracellular bacteria was determined by a CFU assay. To each well 0.5 ml of 0.1% Triton X-100 in double-distilled H_2O was added. After incubation for 10 min at 37°C, the lysates were removed, and the wells were rinsed with 0.5 ml of HI broth. The lysates and HI rinses from each well were pooled in a microcentrifuge tube. The samples were briefly sonicated with a Microson XL200 ultrasonic cell disruptor equipped with a P-1 microprobe (Misonix) to reduce bacterial aggregation. Dilutions were plated on LB agar plates. The plates were incubated at 28°C for 2 days, and the output CFU were enumerated.

Analysis of intracellular survival and replication by immunofluorescence microscopy. Infection of J774A.1 cells was carried out as described above for the CFU assay, except that macrophages were seeded on glass coverslips. The bacterial strains used for infection contained the p67GFP3.1 plasmid (36), which encodes *gfp*mut3.1 under the control of an IPTG-inducible *tac* promoter. One hour prior to fixation green fluorescent protein (GFP) expression was induced with 500 μ M IPTG where indicated below. At the end of the incubation period the cells were fixed with 4% paraformaldehyde in PBS for 10 min. This and all subsequent steps were done at room temperature. The fixed cells were incubated in PBS containing 0.2% Triton X-100 for 10 min to permeate membranes and then in PBS containing 3% bovine serum albumin (BSA) to reduce nonspecific binding of antibodies. The fixed cells were incubated in rabbit anti-*Yersinia* antiserum SB349 (2) diluted 1:1,000 in PBS containing 3% BSA for 40 min to label the bacteria. After two washes with PBS, goat anti-rabbit secondary antibody conjugated to Alexa 594 (Molecular Probes) diluted 1:1,500 in 3% BSA– PBS was added to the fixed cells and incubated for 40 min in the dark. Cells were washed three times with PBS and mounted on glass slides in Airvol (Air Products and Chemicals, Inc.) containing 5% 1,4-diazobicyclo-[2.2.2]-octane (DABCO; Sigma). The slides were examined by phase-contrast and epifluorescence microscopy by using a Zeiss Axioplan2 microscope equipped with a $\times 100$ oil immersion

objective (NA 1.3), and images were captured by using a Spot camera (Diagnostic Instruments, Inc.). Alternatively, slides were analyzed by confocal microscopy by using a Leica DM IRE2. Images of the red and green emission signals, as well as the transmission light at a magnification of \times 40, were captured separately with Leica's LCS software package. Images were processed by using Adobe Photoshop.

Phagosome trafficking assay. Infection was carried out as described above for the immunofluorescence microscopy assays. Bacteria were transformed with p67GFP3.1, and GFP expression was maintained in the overnight cultures and during the infection by addition of $100 \mu M$ IPTG. The MOIs used were 2 for *Y*. *pestis*, 5 for 32777, and 10 for IP2666c, which accounted for the less efficient uptake observed with the latter strain. Cells were fixed at different times with 1 ml of fixing solution (200 μ l of 5× PLP buffer, which contained 100 mM morpholineethanesulfonic acid [MES], 350 mM NaCl, 25 mM KCl, 350 mM lysine-HCl, 25 mM $MgCl₂$, and 10 mM EGTA, 45 mg of sucrose, 505 µl of doubledistilled H₂O, 250 µl of 10% paraformaldehyde, 2.14 mg of NaIO₄) for 30 min. Cells were washed with PBS containing 5% sucrose. The fixed cells were incubated in PBS containing 0.2% Triton X-100 for 10 min and in PBS containing 3% goat serum to reduce nonspecific binding of antibodies. Cathepsin D was labeled by incubating the cells in anti-cathepsin D antibody (Scripps Laboratories) diluted 1:200 in PBS containing 3% goat serum for 40 min. After two washes with PBS, goat anti-rabbit secondary antibody conjugated to Alexa 594 (Molecular Probes) diluted 1:1,500 in PBS containing 3% goat serum was added to the fixed cells and incubated for 40 min in the dark. Cells were washed three times with PBS and mounted on glass slides in Airvol containing 5% DABCO. Coverslips were analyzed by confocal microscopy by using a Leica DM IRE2. Images of the red emission and the green emission at a magnification of \times 40 were captured separately and combined in one image file by using Leica's LCS software package. The images were analyzed with Adobe Photoshop. Approximately 100 phagosomes were scored for each infection. Phagosomes were scored as cathepsin D positive when the shape of the phagosome was clearly visible in the red channel and colocalized with the GFP emission from the bacteria in the green channel. Data from three independent experiments were collected, and averages and standard deviations were calculated with Microsoft Excel.

Mouse infection experiments. Eight-week-old female C57BL/6 mice (Iffa Credo, L'Arbresle, France) were challenged by the intragastric route by using a feeding needle and 32777 or 32777*phoP*::*kan* (referred to as 32777 PhoPA in reference 24) suspended in a 0.2-ml volume. Bacterial inocula were prepared from overnight cultures grown in LB medium at 28°C. The cultures were centrifuged, and the bacterial pellets were washed once and suspended in distilled water. C57BL/6 mice were fasted for 18 h prior to infection. Groups of 10 animals were infected with 10^7 , 10^8 , or 10^9 CFU of 32777 or 10^8 , 10^9 , or 10^{10} CFU of 32777*phoP*::*kan*. Animals were kept in positive-pressure cabinets, and mortality was monitored daily over a 3-week period. These experiments were carried out in compliance with animal care and use regulations at the Université Lille II. The 50% lethal dose (LD_{50}) was calculated by the method of Reed and Muench (37).

Nucleotide sequence accession numbers. The GenBank accession numbers assigned to the *phoP* gene sequences in YPIII(p^-) and YPIII (p^+) are AY649416 and AY649417, repectively.

RESULTS

Y. pseudotuberculosis **YPIII strains carry nonfunctional alleles of** *phoP*. *Y. pseudotuberculosis* $YPIII(p^-)$ (Table 1), unlike other *Y. pseudotuberculosis* strains, is defective for replication in murine macrophages (36). Serovar Typhimurium mutants that are defective for growth in macrophages (e.g., *phoP* mutants) also do not replicate in a low- Mg^{2+} environment (14, 18). To determine if $YPIII(p^-)$ is defective for growth in a low-Mg²⁺ environment, a culture of YPIII(p^-) was diluted into a defined medium lacking magnesium (low- Mg^{2+} TMH) (see Materials and Methods), and bacterial growth was monitored over time by determining the optical density. The same medium was inoculated with IP2666c (Table 1), another *Y. pseudotuberculosis* strain belonging to the same serogroup (serogroup O3), which is able to replicate in macrophages (36). $YPIII(p^-)$ did not grow in low-Mg²⁺ TMH, while IP2666c did grow in this medium (Fig. 1A and data not shown). $YPIII(p^-)$

grew in TMH containing 10 mM MgCl₂ (high-Mg²⁺ TMH), indicating that $YPIII(p^-)$ is specifically defective for growth in a low-Mg²⁺ environment (Fig. 1B).

To investigate if $YPIII(p^-)$ encodes a defective allele of *phoP*, a DNA fragment containing the *phoP* ORF from *Y. pestis* KIM10^{$+$} (Table 1) was inserted into a plasmid under the control of an IPTG-inducible promoter. The resulting plasmid $(pPhoP^{KIM})$ or the empty expression plasmid $(pMMB67EH)$ was introduced into $YPIII(p^-)$. Strains containing these plasmids (Table 1) were assayed for growth in low- Mg^{2+} TMH. $YPIII(p^-)/PhoP^{KIM}$ grew in low- Mg^{2+} TMH containing IPTG but not in low-Mg²⁺ TMH lacking IPTG (Fig. 1C). YPIII(p^-)/ MMB67EH did not grow in low- Mg^{2+} TMH containing IPTG. Therefore, the phenotype of $YPIII(p^-)$ (inability to grow in a low-Mg²⁺ environment) was complemented by expression of *phoPKIM* in *trans*.

To directly show that PhoP is required for growth of *Y. pseudotuberculosis* in a low- Mg^{2+} environment, a deletion was introduced into *phoP* in IP2666c, creating IP2666c*phoP* (Table 1). IP2666cphoP Δ grew in high-Mg²⁺ TMH (Fig. 2B) but not in low- Mg^{2+} TMH (Fig. 2D). The inability of IP2666cphoP Δ to grow in low-Mg²⁺ TMH was rescued by expression of *phoPKIM* in *trans* (Fig. 1D), indicating that the mutation introduced into IP2666cphoP Δ was nonpolar. To determine if the *phoP* ORF in $YPIII(p^-)$ encodes a defective product, a DNA fragment containing this ORF (*phoPYPIII*) was inserted into pMMB67EH under control of the IPTG-inducible promoter. The resulting plasmid (pPhoP^{YPIII}) was transferred into IP2666cphoP Δ , and the resulting strain (Table 1) was assayed for growth in low- Mg^{2+} TMH containing IPTG. Expression of PhoP^{YPIII} did not rescue growth of IP2666cphoP Δ in low-Mg²⁺ TMH (Fig. 1D). We therefore concluded that *phoPYPIII* is a nonfunctional allele of *phoP*.

A CFU assay (see Materials and Methods) was carried out to determine if the growth defect of $YPIII(p^-)$ in macrophages could be rescued by expression of *phoPKIM*. YPIII(p^-)/PhoPKIM and $YPIII(p^-)/MMB67EH$ were used to infect J774A.1 macrophages. The infected macrophages were incubated in the presence of a low concentration of gentamicin to inhibit growth of extracellular bacteria and in the presence of IPTG to induce expression of *phoP^{KIM}*. As shown in Fig. 2, YPIII(p^{-})/ PhoPKIM grew in macrophages, while the viability of $YPIII(p^-)/MMB67EH$ decreased in macrophages. These results suggest that $YPIII(p^-)$ is defective for survival and replication in macrophages because it carries a nonfunctional *phoP* allele.

The *phoP^{YPIII}* ORF was sequenced, and the deduced amino acid sequence of PhoPYPIII was compared with the sequences of PhoPKIM and PhoPIP2666 available in the GenBank database. As shown in Fig. 3, there is one amino acid difference unique to PhoPYPIII, corresponding to a threonine-to-proline substitution at amino acid 160 (T160P). PhoP belongs to the family of winged-helix transcription factors (25). Analysis of the amino acid sequence of PhoP^{KIM} by using PSIPRED structure predictions (26) showed that residue 160 is predicted to lie within an alpha helix (Fig. 3). Additional analysis with SWISS-MODEL (19, 38) to create a three-dimensional model of PhoPKIM revealed that this helix corresponds to the α 1 helix found in other winged-helix transcription factors (25) . The α 1 helix forms the hydrophobic core of the DNA binding domain

FIG. 1. Analysis of *Y. pseudotuberculosis* strains for growth in low- or high-Mg²⁺ medium. Bacterial cultures of the strains indicated were diluted into low-Mg²⁺ TMH (A, C, and D) or high-Mg²⁺ TMH (B), and growth at 37°C was measured by determining the optical density (OD) at 600 nm at different times. The presence of IPTG in the culture medium is indicated. The results are representative of at least two independent experiments.

in winged-helix transcription factors. It forms a structural core, aiding in the positioning of the β -sheets in the winged-helix motif. A T160P substitution in the α 1 helix is expected to cause a change in the positioning of residues in the helix and in the contacts that these residues make within the domain. We hypothesize that the T160P substitution results in a conformational change in the positioning of the α 2 and α 3 helices that make contact with the DNA, thereby disrupting the function of PhoP^{YPIII}. Taken together, our results suggest that $YPIII(p^-)$ is defective for survival and replication in macrophages because it encodes a nonfunctional PhoP protein.

PhoP is important for survival and replication of *Y. pseudotuberculosis* **in macrophages.** Macrophages infected with IP2666c or IP2666cphoP Δ were analyzed by fluorescence microscopy (see Materials and Methods) to obtain further evidence that PhoP is important for growth of *Y. pseudotuberculosis* in macrophages. Both strains were transformed with p67GFP3.1, a plasmid that encodes GFP under control of an

IPTG-inducible promoter (36). The resulting strains (Table 1) were used to infect J774A.1 macrophages for different periods of time in the presence of a low concentration of gentamicin to inhibit the growth of extracellular bacteria. One hour prior to fixation, IPTG was added to the medium to induce GFP expression in intracellular bacteria. After fixation, the total intracellular bacteria were labeled with anti-*Yersinia* antibodies and secondary antibodies conjugated to Alexa 594. In a previous study it was found that the numbers of viable *Y. pseudotuberculosis* cells in macrophages remain constant over the first 4 to 5 h of infection, that bacterial replication begins within 8 h of infection, and that between 8 and 24 h the numbers increase \geq 10-fold (36). As expected, IP2666c/GFP underwent significant replication between 5 and 24 h postinfection (Fig. 4, panels A1, A2, C1 to C3, and E1 to E3). By comparison, IP2666c*phoP* /GFP was clearly defective for replication in macrophages, although GFP-positive intracellular bacteria could be detected at 5 and 24 h (Fig. 4, panels B1, B2, D1 to

FIG. 2. Expression of *phoPKIM* in *trans* promotes the survival and growth of YPIII (p^-) in macrophages. J774A.1 macrophages were infected with YPIII(p⁻)/MMB67EH (gray bar) or YPIII(p⁻)/PhoP^{KIM} (solid bar) at an MOI of 5 in the presence of IPTG. The numbers of CFU recovered from the infected macrophages after 0.25 and 24 h of infection were determined as described in Materials and Methods. The percent increase in CFU was calculated by using the following formula: (CFU at T_{24} – CFU at $T_{0.25}$)/(CFU at T_{24}) × 100, where T_{24} and T_{24} are 0.25 and 24 h after infection, respectively. The data are the averages \pm standard deviations of values obtained from two independent experiments with duplicate wells.

D3, and F1 to F3). No obvious cytotoxic effect of bacterial replication on the macrophages was observed in these experiments (Pujol, unpublished data). A CFU assay was carried out to measure bacterial replication. J774A.1 macrophages were infected with IP2666c, IP2666cphoP Δ /PhoPKIM, or IP2666cphoP Δ

MMB67EH in the presence of IPTG, and viable intracellular bacteria were quantified after 0.25 or 24 h of incubation. The resulting data were used to calculate a survival index (Table 2). As shown in Table 2, the *phoP* mutant was defective for survival in macrophages, and the defect could be rescued by expression of *phoPKIM* in *trans*.

In order to compare directly the functional importance of PhoP in *Y. pseudotuberculosis* and *Y. pestis*, the *phoP* deletion mutation was introduced into *Y. pestis* KIM10⁺. The resulting strain, KIM10⁺ $phoP\Delta$ (Table 1), and its parent were assayed for growth in low- or high- Mg^{2+} TMH. KIM10⁺ grew in both media, while KIM10⁺phoP Δ grew only in high-Mg²⁺ TMH (data not shown). Therefore, PhoP is required for growth of *Y. pestis* in a low- Mg^{2+} environment. As determined by fluorescence microscopy, KIM10⁺/GFP grew in macrophages (Fig. 4, panels G1, G2, I1 to I3, and K1 to K3), while most KIM10⁺phoP Δ /GFP cells were not viable or destroyed by 5 h (Fig. 4, panels H1, H2, J1 to J3, and L1 to L3). The results of a CFU assay confirmed that the *phoP* mutant of KIM10⁺ was defective for intracellular survival and more sensitive to intracellular killing (-10-fold more sensitive) than the *phoP* mutant of IP2666c (Table 2). Therefore, *phoP* mutants of *Y. pseudotuberculosis* and *Y. pestis* are both defective for survival and replication in macrophages, although these mutants differ in that the latter is more sensitive to killing in macrophage phagosomes.

Role of PhoP in retarding phagosome maturation. Mills and Finlay have reported that phagosomes containing $YPIII(p^+)$ acquire the lysosomal marker cathepsin D and are trafficked to phagolysosomes in J774A.1 macrophages (28). As our findings

FIG. 3. Predicted secondary structure of PhoP^{KIM} and ClustalW alignment of PhoP sequences (single-letter code) from *Yersinia* species and serovar Typhimurium. The secondary structure was predicted by using PSIPRED. H, α-helix; E, β-sheet; C, coil; *Y. pseudotb, Y. pseudotuberculosis*; *S. typhimurium*, *S. enterica* serovar Typhimurium. The confidence levels for secondary structure assignments are indicated by bar graphs. The amino acid sequences of PhoP proteins for KIM10⁺ (accession no. NP_669111), IP2666 (accession no. AAK54056), and serovar Typhimurium (accession no. RGEBFT) were obtained from GenBank. The sequence of PhoPYPIII was deduced from the nucleotide sequence of the *phoP* gene from $YPIII(p^-)$ (*phoP^{YPIII}*) obtained in this study. Sequences were aligned by using the ClustalW alignment function in MacVector. The T160P substitution in PhoP^{YPIII} is enclosed in a box. The arrow at position 116 indicates the first codon in $phoP^{YPII}(P^+)$ that is changed (from M to H) due to a frameshift mutation corresponding to a repeat of nucleotides 341 to 345. The asterisk indicates the position of the stop codon in $phoP^{YPIII}(P^+)$ which results from the frameshift mutation.

FIG. 4. Analysis of *phoP* and *phoP* mutant *Y. pseudotuberculosis* and *Y. pestis* strains for survival and replication in macrophages by microscopy. J774A.1 macrophages were infected with IP2666c/GFP (panels A1, A2, C1 to C3, and E1 to E3), IP2666c*phoP* \triangle /GFP (panels B1, B2, D1 to D3, and F1 and F3), KIM10⁺/GFP (panels G1, G2, I1 to I3, and K1 to K3), or KIM10⁺*phoP* Δ /GFP (panels H1, H2, J1 to J3, and L1 to L3). Infections were carried out at MOIs of 2 for *Y. pestis* and 5 for *Y. pseudotuberculosis*. The infected cells were fixed after 0.25, 5, or 24 h of infection. One hour before the 5- and 24-h samples were processed, the cells were incubated in the presence of IPTG to induce GFP expression in intracellular bacteria. After fixation, the samples were stained with anti-*Yersinia* antibody and an Alexa 594-conjugated secondary antibody. The samples were viewed by phase-contrast and epifluorescence microscopy. Representative images were captured by using a digital camera. The GFP and anti-*Yersinia* signals were merged with phase-contrast images (merged + phase).

suggested that $YPIII(p^-)$ is a *phoP* mutant, we used immunofluorescence microscopy and cathepsin D as a marker for lysosomes to determine if phagosomes containing *phoP*⁺ or *phoP* mutant *Y. pseudotuberculosis* are trafficked differently in macrophages. IP2666c/GFP and IP2666cphoP Δ /GFP were grown in the presence of IPTG and used to infect J774A.1 macrophages in the presence of IPTG to ensure continuous GFP expression during the experiment. At 15 min postinfection the macrophages were incubated in medium containing gentamicin. At different times thereafter, the cells were fixed

and stained with an anti-cathepsin D antibody and an Alexa 594 secondary antibody. The samples were examined by confocal microscopy, and cathepsin D-positive phagosomes were identified by an overlap in the green GFP and red Alexa 594 emission signals (Fig. 5A). Very few phagosomes containing either IP2666c/GFP or IP2666cphoP Δ /GFP had acquired cathepsin D at 1.5 h postinfection (data not shown). By 8 h postinfection there was a twofold difference between the percentage of cathepsin D-positive phagosomes in macrophages infected with IP2666c/GFP and the percentage of cathepsin

TABLE 2. PhoP is important for survival of *Y. pseudotuberculosis* and *Y. pestis* in macrophages

 a J774A.1 cells were infected with parental strains (IP2666c, 32777, KIM10⁺) or the *phoP* mutant strains at an MOI of 5 (IP2666c, 32777) or 2 (KIM10⁺). Bacterial cultures were grown at 28°C prior to infection. IPTG (500 μ M) was present in the medium throughout the infection to induce *phoP* expression. CFU counts were determined after 0.25 or 25 h of infection as described in Materials and Methods. The values are values from a representative experiment with duplicate wells. The survival index was calculated from the CFU counts by using the following formula: (CFU of mutant at T_{24}/CFU of parent at T_{24}/CFU of mutant at $\overline{T}_{0.25}/\text{CFU}$ of parent at $T_{0.25}$), where T_{24} and $\overline{T}_{0.25}$ are 24 and 0.25 h after infection, respectively.

D-positive phagosomes in macrophages infected with IP2666c*phoP* /GFP (4 versus 8%) (Fig. 5B). A CFU assay was carried out in parallel with the trafficking assay to enumerate viable intracellular bacteria. The results are summarized in Fig. 5C. Between 0.25 and 5 h after infection, the number of viable bacteria remained constant for both IP2666c and IP2666cphoP Δ . Between 5 and 24 h, the number of viable IP2666cphoP Δ cells decreased 10-fold, while the number of viable IP2666c cells increased 10-fold. These results suggest that a *Y. pseudotuberculosis phoP* mutant can retard phagosome maturation and maintain viability during the several-hour lag phase that precedes intracellular growth. Following the lag phase, mutant bacteria begin to loose the ability to retard phagosome maturation and are killed within phagolysosomes.

To determine if PhoP influences maturation of phagosomes containing *Y. pestis*, the trafficking experiments described above were repeated by using $KIM10^{+}/GFP$ and $KIM10^{+}$ *phoP* /GFP. Interestingly, 60% of the phagosomes containing KIM10⁺ $phoP\Delta/GFP$ were positive for cathepsin D at 1.5 h postinfection (Fig. 5D and E). In contrast, only 5% of the phagosomes containing $KIM10⁺/GFP$ were positive for cathepsin D at 1.5 h postinfection (Fig. 5E). The rapid acquisition of cathepsin D by phagosomes containing KIM10⁺ $phoP\Delta$ was paralleled by a rapid decrease in viability, and there was a 100-fold decrease in the number of viable bacteria recovered between 0.25 and 5 h after infection (Fig. 5F). These results suggest that *phoP* mutant *Y. pestis* cells are rapidly killed in phagolysosomes because they are unable to retard phagosome maturation during the intracellular lag phase.

PhoP is important for *Y. pseudotuberculosis* **virulence.** Oyston et al. (29) have shown that a *Y. pestis phoP* mutant is less virulent (75-fold less virulent as determined by an LD_{50} assay) in a murine model of bubonic plague. To determine if PhoP is important for *Y. pseudotuberculosis* pathogenesis, we utilized 32777, a pYV-containing serogroup O1 strain of *Y. pseudotuberculosis*. This strain has previously been shown to be highly virulent in a mouse oral infection assay (10). 32777*phoP*::*kan* is an isogenic mutant containing a kanamycin resistance gene inserted into *phoP* (Table 1). Since all previous assays were carried out with pYV-cured *Y. pseudotuberculosis phoP* mutants, the characteristics of 32777 and 32777*phoP*::*kan* were studied by using the different assays described above to confirm that the presence of pYV did not alter the phenotype of a *phoP* mutant in low- Mg^{2+} medium or macrophages. 32777 and 32777*phoP*::*kan* grew at the same rate in high-Mg²⁺ TMH, while 32777*phoP*::*kan* was specifically defective for growth in low-Mg2 TMH (J. P. Grabenstein, unpublished data). 32777*phoP*::*kan* showed a defect in intramacrophage survival comparable to that observed with other *Y. pseudotuberculosis phoP* mutants (Table 2; Grabenstein, unpublished). Additionally, expression of PhoPKIM rescued the intracellular survival defect of 32777*phoP*::*kan* (Table 2), demonstrating that the *phoP*::*kan* mutation was nonpolar. In two independent trafficking assays 27.7% of the phagosomes containing 32777*phoP*::*kan* were cathepsin D positive at 8 h postinfection, while 3% of phagosomes containing 32777 were positive at this time (Grabenstein, unpublished). These data demonstrate that a pYV-containing *Y. pseudotuberculosis phoP* mutant is phenotypically similar to a pYV-cured *Y. pseudotuberculosis phoP* mutant with respect to growth in low- Mg^{2+} medium and macrophages.

C57BL/6 mice were infected via the intragastric route with different doses of 32777 or 32777*phoP*:: kan , and the LD_{50} for each strain was determined after 21 days. The LD_{50} of 32777 in C57BL/6 mice was calculated to be 5×10^8 CFU, while the LD_{50} of the *phoP* mutant was at least 100-fold higher. When mice were challenged with 10^9 CFU of either strain, 70% of the mice infected with 32777 died during the observation period, while 20% of the mice infected with the *phoP* mutant died (Fig. 6). Therefore, PhoP is important for *Y. pseudotuberculosis* virulence.

DISCUSSION

Y. pseudotuberculosis and *Y. pestis* share a number of characteristics that are important for virulence as a result of their close genetic relatedness (7, 32). A characteristic that is common to *Y. pseudotuberculosis* and *Y. pestis* is the ability to replicate in macrophages (36). It was suggested previously that *Y. pseudotuberculosis* and *Y. pestis* are likely to have a common genetic basis for the ability to replicate in macrophages (36). As the response regulator PhoP has been shown to be important for survival of *Y. pestis* in macrophages and for *Y. pestis* virulence (29), this study was undertaken to evaluate the importance of PhoP for the pathogenesis of *Y. pseudotuberculosis*. The data presented here show that PhoP is important for survival and replication of *Y. pseudotuberculosis* in macrophages. Furthermore, we found that PhoP is an important virulence factor of *Y. pseudotuberculosis*. Our results indicate that survival and replication in macrophages are important for pathogenesis in *Y. pseudotuberculosis* and *Y. pestis* infections.

Y. pseudotuberculosis strain $YPIII(p^-)$ is unable to grow in macrophages or low-Mg²⁺ medium. In this regard, YPIII(p^-) behaves phenotypically like a *phoP* mutant of serovar Typhimurium. The results of several complementation tests suggest that $YPIII(p^-)$ carries a nonfunctional allele of *phoP*. Sequence analysis of *phoPYPIII* revealed that a nucleotide change in this ORF generates a T-to-P substitution at codon 160. A three-dimensional model of PhoPKIM, which belongs to the class of winged-helix transcription factors (25), predicts that the T160P substitution lies within a key α -helix $(\alpha 1)$, which forms the hydrophobic core of the DNA binding domain. We

FIG. 5. Analysis of phagosome trafficking in macrophages infected with *Y. pseudotuberculosis* or *Y. pestis* by anti-cathepsin D staining and confocal microscopy. IP2666c/GFP, IP2666cphoP Δ /GFP, KIM10⁺/GFP, and KIM10⁺phoP Δ /GFP were grown in the presence of IPTG and used to infect J774A.1 macrophages at an MOI of 10 for *Y. pseudotuberculosis* or 2 for *Y. pestis*. At different times postinfection the cells were fixed and stained with anti-cathepsin D antibody and a secondary antibody conjugated to Alexa 594. The samples were analyzed by confocal microscopy. (A and D) Representative images of the merged anti-cathepsin D and GFP signals obtained with cells infected for 8 h with IP2666c*phoP* Δ /GFP (A) or for 1.5 h with KIM10⁺phoP Δ /GFP (D). The arrows indicate cathepsin D-positive phagosomes. (Insets) Enlargement of the region indicated by the white rectangle. The left inset shows merged GFP and cathepsin D signals, and the right inset shows cathepsin D signals alone. (B) Percentages of cathepsin D-positive phagosomes in macrophages after 8 h of infection with IP2666c/GFP or IP2666c*phoP* /GFP calculated by using merged images captured at a magnification of \times 40. (E) Percentages of cathepsin D-positive phagosomes in macrophages containing KIM10⁺/GFP or KIM10⁺phoP Δ /GFP at 1.5 h postinfection determined in the same manner. The data in panels B and E are the averages of three independent experiments. The error bars indicate the standard deviations. (C and F) Numbers of CFU recovered from J774A.1 macrophages infected with IP2666c/GFP or KIM10⁺/GFP or the corresponding *phoP* mutants at the times indicated. Each data point represents the average \pm standard deviation of the values recovered from triplicate wells. The results are representative of two independent experiments.

speculate that the T160P substitution results in a global conformational change in PhoPYPIII and loss of function. It is currently not clear how and when this codon substitution originated in YPIII. This codon substitution results from an A-to-C transversion, which is a relatively uncommon mutation. $YPIII(p^+)$, the presumed parent of $YPIII(p^-)$, is also unable to replicate in macrophages (28; Grabenstein, unpublished data). The *phoP* ORF from YPIII(p^+) (Table 1) was sequenced to determine if it carries a nonfunctional *phoP* allele. Interestingly, we found that $phoP^{YPHI(p+)}$ contains a different mutation, corresponding to a repeat of nucleotides 341 to 345 of *phoP*, which creates a frameshift mutation and a stop codon at predicted residue 117 (Fig. 3). Therefore, the YPIII(p^-) and $YPIII(p^+)$ strains used in our study both have nonfunctional alleles of *phoP* and are unable to replicate in macrophages. Additionally, $YPIII(p^-)$ is most likely not derived from $YPIII(p^+)$ because of the different nature of the *phoP* mutations that are present in these strains. A question raised by these observations is whether there is some selective pressure on YPIII strains cultivated in the laboratory to inactivate *phoP*.

As we have shown that PhoP is important for virulence of *Y. pseudotuberuculosis*, it is possible that $YPIII(p^+)$ is partially attenuated due to the mutation in *phoP*, and we speculate that the virulence of this strain would be increased by restoration of the wild-type *phoP* sequence.

Y. pseudotuberculosis and *Y. pestis* mutants carrying sitedirected mutations in *phoP* were also unable to grow in a low-Mg²⁺ defined medium. In serovar Typhimurium, PhoP activates the expression of high-affinity Mg^{2+} transporters encoded by the *mgtA* and *mgtB* genes. In addition, PhoP-regulated gene expression in serovar Typhimurium is required for downregulation of CorA activity, which prevents the influx of toxic metals, such as Fe(II) (9). The inability of yersinia *phoP* mutants to grow in low- Mg^{2+} TMH may therefore result from a combination of Mg^{2+} starvation and metal toxicity. Bacteria growing in TMH are exposed to high concentrations of Fe(II) because this medium contains 100 μ M FeSO₄ (42).

Y. pseudotuberculosis phoP mutants were shown to be defective for survival and replication in J774A.1 macrophages. A *Y. pestis phoP* mutant was also defective for survival and replication in macrophages, as expected from a previous study (29). Therefore, a requirement for PhoP to survive in macrophages is conserved in *Y. pseudotuberculosis* and *Y. pestis*. To gain insight into the role of PhoP for survival of yersiniae in macrophages, we studied the trafficking of phagosomes containing $phoP⁺$ or *phoP* mutant bacteria using cathepsin D as a subcellular marker for lysosomes. Evidence that *phoP* mutant *Y. pseudotuberculosis* and *Y. pestis* strains are defective for retarding phagosome maturation was obtained. The concept that $phoP⁺$ yersiniae can retard phagosome maturation is supported by the data of Tsukano et al. (45), who reported that *Y. pseudotuberculosis* inhibits acidification of phagosomes. The results of a previous study indicated that phagosomes containing *Y. pestis* fuse with lysosomes (44). In this previous study the workers used a different lysosomal marker (thorium dioxide) and a different detection method (transmission electron microscopy) than we used in this study. The previous results (44) are not necessarily in conflict with those reported here, since it is possible that nascent phagosomes containing yersiniae can fuse with lysosomes but as the phagosomes are remodeled by the pathogen, they become secluded from the phagolysosomal pathway. This appears to be the fate of phagosomes containing serovar Typhimurium (21). Interestingly, it has been shown that PhoP is important for serovar Typhimurium to retard phagosome maturation (17). The technique used by Straley and Harmon (44) may be sufficiently sensitive to detect phagosomes that have undergone transient fusion events with lysosomes, while the approach which we used may only detect phagosomes that have undergone sustained fusion events with lysosomes. Cathepsin D may need to accumulate to high levels before it can be easily detected by immunofluorescence microscopy. Studies to better characterize the trafficking of phagosomes containing *phoP* or *phoP* mutant yersiniae are under way.

Although a general conclusion based on our results is that the function of PhoP is conserved in *Y. pseudotuberculosis* and *Y. pestis*, an interesting observation which we made was that a *phoP Y. pestis* mutant was \sim 10-fold more sensitive to killing in macrophages than a *phoP Y. pseudotuberculosis* mutant (Table 2). The increased rate of intracellular killing of the *Y. pestis phoP* mutant was paralleled by an increased rate of phagosomal cathepsin D staining. The data suggest that PhoP is required for *Y. pestis* to retard phagosome maturation during the intracellular lag phase, while PhoP is dispensable for *Y. pseudotuberculosis* to delay phagosome maturation during the intracellular lag phase. The mechanistic basis behind this difference is not known. It is possible that *Y. pseudotuberculosis* encodes a mechanism that can function in the absence of PhoP to retard phagosome maturation during the intracellular lag phase. After *Y. pseudotuberculosis* exits the lag phase, it appears that PhoP function becomes important for retarding phagosome maturation. This hypothetical backup mechanism could be unique to *Y. pseudotuberculosis* or may have been lost in *Y. pestis* due to genetic decay. One known difference between *Y. pseudotuberculosis* and *Y. pestis* is that the lipopolysaccharide (LPS) of *Y. pestis* lacks O antigen due to the presence of several mutations in genes required for O antigen biosynthesis (41). O antigen could play a role in retarding maturation of phagosomes containing *Y. pseudotuberculosis* during the intracellular lag phase. It has been shown that the O antigen of *Brucella suis* is involved in the ability of this intracellular pathogen to avoid

FIG. 6. Virulence of wild-type and *phoP* mutant *Y. pseudotuberculosis* in mice. Wild-type *Y. pseudotuberculosis* 32777 and the isogenic *phoP* mutant 32777*phoP*::*kan* were used to challenge C57BL/6 mice by the intragastric route. Groups of 10 animals were infected with $10⁹$ CFU of a strain, and mortality was monitored for 21 days after challenge.

phagosome maturation in macrophages (35). The presence of O antigen appears to promote entry of *B. suis* into macrophages via lipid rafts, resulting in the formation of phagosomes that avoid fusion with lysosomes (35). If O antigen allows *Y. pseudotuberculosis* to delay phagosome maturation, then mutants defective for O antigen and PhoP should be rapidly killed in macrophages. Mutants of $YPIII(p^+)$ defective for O antigen biosynthesis have been isolated and shown to be attenuated for virulence in mouse infection assays (23, 27). It is conceivable that these mutants, which are missing O antigen and a functional PhoP protein, are attenuated due to increased sensitivity to intracellular killing.

It is currently not known how PhoP retards phagosome maturation. In serovar Typhimurium, over 40 genes that have diverse functions and are directly or indirectly under the control of PhoP have been identified. Products of the *pmr* and *pag* genes modify the lipid A portion of LPS, making the outer membrane less negatively charged and less permeable for cationic antimicrobial peptides (14). Other gene products, such as the Mg^{2+} transporters encoded by the *mgt* genes, are required for the adaptation of serovar Typhimurium to the low- Mg^{2+} environment of the phagosome (3). Homologues of the *mgt*, *pmr*, and *pag* genes are present in the *Y. pestis* genome (11, 31), so it is possible that the products of these genes have similar functions in serovar Typhimurium and yersiniae. The LPS core of *Y. pestis* appears to be modified in a *phoP*-dependent fashion (20). Therefore, it is conceivable that yersinia *phoP* mutants are defective for retarding phagosome maturation due to an inability to acquire a nutrient or to modify the membranes or surfaces within the phagosome. This in turn could prevent the bacterium from synthesizing or properly secreting a factor(s) that is required to retard phagosome maturation. In this context, PhoP-regulated gene expression could be indirectly involved in the process of retarding phagosome maturation. Alternatively, it is possible that PhoP activates an unidentified gene(s) whose product functions directly to retard phagosome maturation.

A murine intestinal infection model was used to demonstrate that PhoP is an important virulence factor of *Y. pseudotuberculosis*. Inactivation of *phoP* resulted in a \sim 100-fold increase in the LD_{50} of *Y. pseudotuberculosis*. This increase is the same magnitude as that reported for a *Y. pestis phoP* mutant in a mouse bubonic plague infection model (29). The *phoP* mutant was not defective for expression or secretion of Yop proteins under standard growth conditions (Grabenstein, unpublished), suggesting that the *phoP* mutant is selectively impaired for intracellular pathogenesis. It is possible that survival and replication in macrophages allow *Y. pseudotuberculosis* and *Y. pestis* numbers to increase while the organisms are protected from the actions of neutrophils (8). Thus, survival and replication in macrophages may allow yersiniae to thwart innate immune responses. Yersiniae may also circumvent the development of adaptive immunity by surviving in macrophage phagosomes. It has been shown that a serovar Typhimurium *phoP* mutant is more efficiently processed for class II antigen presentation by macrophages than the parental *phoP*⁺ strain (47). The more efficient antigen processing of a serovar Typhimurium *phoP* mutant could be directly related to the inability of the mutant to avoid phagosome maturation (17). As we have shown that PhoP is important for yersiniae to avoid maturation of phagosomes to cathepsin D-positive vacuoles, it is possible that *Y. pseudotuberculosis* or *Y. pestis phoP* mutant strains induce an earlier adaptive immune response than parental $phoP⁺$ bacteria. The induction of an earlier adaptive immune response could contribute to the significant decrease in virulence of *phoP* mutant yersiniae. Further studies are needed to determine if there is a connection between the ability of yersiniae to replicate in macrophages and their ability to avoid innate and adaptive immune responses.

ACKNOWLEDGMENTS

We thank members of the Bliska laboratory for comments on the manuscript and Kwadwo Bonsu for helping establish the cathepsin D staining assay.

This work was supported by Public Health Service grant AI48507 awarded to J.B.B.

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