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Absence of All Components of the Flagellar Export and Synthesis Machinery Differentially Alters Virulence of *Salmonella enterica* Serovar Typhimurium in Models of Typhoid Fever, Survival in Macrophages, Tissue Culture Invasiveness, and Calf Enterocolitis

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In this study, we constructed an *flhD* (the master flagellar regulator gene) mutant of *Salmonella enterica* serovar Typhimurium and compared the virulence of the strain to that of the wild-type strain in a series of assays that included the mouse model of typhoid fever, the mouse macrophage survival assay, an intestinal epithelial cell adherence and invasion assay, and the calf model of enterocolitis. We found that the *flhD* mutant was more virulent than its parent in the mouse and displayed slightly faster net growth between 4 and 24 h of infection in mouse macrophages. Conversely, the *flhD* mutant exhibited diminished invasiveness for human and mouse intestinal epithelial cells, as well as a reduced capacity to induce fluid secretion and evoke a polymorphonuclear leukocyte response in the calf ligated-loop assay. These findings, taken with the results from virulence assessment assays done on an *fljB fliC* mutant of serovar Typhimurium that does not produce flagellin but does synthesize the flagellar secretory apparatus, indicate that neither the presence of flagella (as previously reported) nor the synthesis of the flagellar export machinery are necessary for pathogenicity of the organism in the mouse. Conversely, the presence of flagella is required for the full invasive potential of the bacterium in tissue culture and for the influx of polymorphonuclear leukocytes in the calf intestine, while the flagellar secretory components are also necessary for the induction of maximum fluid secretion in that enterocolitis model. A corollary to this conclusion is that, as has previously been surmised but not demonstrated in a comparative investigation of the same mutant strains, the mouse systemic infection and macrophage assays measure aspects of virulence different from those of the tissue culture invasion assay, and the latter is more predictive of findings in the calf enterocolitis model.

Over 40 genes are required for the structure, assembly, and function of flagella (25). These genes are categorized into three classes that are temporally expressed in a cascade-like manner. Class 1 genes include the master regulatory genes (*flhD* and *flhC*) which are required for the activation of transcription from class 2 promoters. Class 2 genes encode hook-basal body proteins (which make up the flagellar secretory apparatus), as well as the alternate sigma factor (FliA) which transcribes class 3 genes involved in motor and chemotaxis functions and filament structures. Regulation of flagellar synthesis is accomplished through an interaction between FliA and FlgM, an antisigma factor.

Previously, we and others demonstrated that flagella are not required for *Salmonella enterica* serovar Typhimurium virulence in the murine typhoid model (3, 23). Rather, we showed that some aspects of flagellar regulation, namely the FlgM-FliA regulatory system, are involved in the in vivo pathogenicity of serovar Typhimurium. Specifically, we found that the *flgM* gene, which encodes a negative regulator of flagellar synthesis (12), is required for the virulence of serovar Typhimurium in the mouse (33). We also reported that mutation of *fliA*, a locus which encodes the flagellin-specific sigma factor (σ^{28} [31]), restores virulence to a *flgM* mutant (33). Furthermore, we found that inactivation of *fliC*, the gene that encodes the phase-1 flagellin, but not *fljB*, the gene that encodes the phase-2 flagellin, reverses the attenuated phenotype of the *flgM* mutant (32). Based on this series of observations, we hypothesized that there is a link in serovar Typhimurium between the flagellar synthesis and regulation, but not flagella per se, and the virulence of the microbe for the mouse. One theoretical explanation for this association is that export of certain virulence-associated proteins may occur through the flagellar secretory components (13, 26), a suggestion based on the striking homologies between the flagellar secretory machinery and the type III virulence-associated export systems (9, 10, 13, 16, 21, 26, 35). However, in spite of the fact that several flagellar and virulence-related proteins are secreted by serovar Typhimurium (19), a role for the flagellar apparatus in the export of virulence proteins has not been demonstrated for *Salmonella*.

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In this study, we sought to test the theory that a connection exists between the serovar Typhimurium flagellar synthesis apparatus and the virulence of the microbe. For that purpose, we constructed an *flhD* (the master flagellar regulatory gene) mutant of serovar Typhimurium that is incapable of producing any of the products of the flagellar cascade including regulatory elements, flagellar components, or flagellar export machinery. We also prepared an *fljB fliC* mutant of serovar Typhimurium that cannot produce flagellin and hence does not make flagella. We then compared the pathogenicities of the *flhD* mutant and the *fljB fliC* mutant with that of the wild-type strain (and in some cases a previously constructed *fliA* mutant) in a series of established virulence assays that included the mouse model of murine typhoid and the mouse macrophage survival assay, as well as in models thought to better reflect steps in the pathogenesis of gastroenteritis in humans, such as intestinal cell adherence and invasion assays and the calf loop model of enterocolitis.

MATERIALS AND METHODS

Bacterial strains and growth conditions. The following strains of *Salmonella* serovar Typhimurium were used in this study: the virulent wild-type strain SL3201 (15), KK2040 (*flhD*::Tn*10*, kind gift of K. Kutsukake [20]), SL3201 *fliA*::Tn*10* (virulent, nonmotile [33]), and SL3201 *fljB*::Mud*J fliC*::Tn*10* (32). Bacteria were grown at 37°C in Luria-Bertani (LB) broth (Difco Laboratories, Detroit, Mich.) with shaking or on LB agar, except where noted.

Transduction. Bacteriophage P22HT*int* was used as previously described (5) to transduce the *flhD*::Tn*10* mutation from strain KK2040 into strain SL3201.

Motility assay. Motility was assessed by stabbing motility agar with the bacterial strain and observing for migration away from the point of inoculation following incubation at 37°C for 18 h. Motility agar consisted of 10 g of tryptone per liter, 5 g of NaCl per liter, and 0.35% (wt/vol) agar, pH 7.4.

Assays for mouse virulence, kinetics of infection, and serology. Mouse experiments were conducted according to the principles outlined by the National Institutes of Health (30). Virulence was assessed by 50% lethal dose (LD_{50}) and determined with groups of 4 to 10 6- to 8-week-old female mice (C57BL/6J; Jackson Laboratories, Bar Harbor, Maine) inoculated orally with different doses of *Salmonella*. Food was removed from cages approximately 16 h before oral inoculation, and mice were given 25 μl of 2% $NaHCO_3$ just prior to being fed the inoculum. Mice were monitored for death for up to 32 days, and the LD_{50} values were calculated by the Miller Tainter method of probit analysis (29) as detailed in reference 40. Virulence was also assessed by monitoring for death of C57BL/6J mice following intraperitoneal injection with approximately 700 CFU of organisms (33). This dose and this route of infection have been used previously to discern attenuated mutants from wild-type serovar Typhimurium strain SL3201 (33). The kinetics of infection were examined by orally infecting groups of C57BL/6J mice with approximately 10^7 CFU of bacteria per mouse. Five mice from each group were sacrificed at various times postinfection for the determination of CFU per spleen.

Survival in macrophages. Survival in mouse resident peritoneal macrophages was measured as described previously (22, 33). Briefly, bacteria were grown overnight in LB broth. Bacteria were opsonized with normal mouse serum and allowed to infect resident mouse peritoneal macrophages at a multiplicity of infection (MOI) of approximately 5. Five wells per time point (T0 h, 4 h, and 24 h) were infected with each strain. After 50 min, the wells were washed three times. At 0 h, bacteria were enumerated from three wells for each strain, and the average number of macrophages was determined from the remaining two wells. Medium that contained gentamicin was added to the remaining wells. This procedure was repeated at 4 and 24 h postinfection. Controls included uninfected macrophages.

Assay for adherence and invasion of epithelial cells. The flagellar regulatory mutants were tested in an in vitro model for the capacity to adhere to and invade two small intestinal epithelial cell lines: mouse MODE-K cells (36) and human Henle-407 cells (14). The bacteria were grown in a high salt concentration (0.3 M NaCl) for optimal adherence and invasion (34, 39). Assays were conducted with mid-log-phase to late log phase bacterial cells as described previously (7, 34). Briefly, bacteria were added to cells at an MOI of ~20, and then the microtiter plates were gently centrifuged for 10 min at 2,000 × *g*. The purpose of this centrifugation step was to compensate for the lack of motility of some strains and to enhance physiological interactions of the organisms with the eucaryotic cells. The adherence assay was conducted for 90 min, and then the plates were washed. For the invasion assay, a subset of wells with washed infected cells was then incubated for an additional 90 min in the presence of media supplemented with 100 μg of gentamicin per ml. Results were calculated as mean percent adherence (number of bacteria after washes/number of bacteria in the initial inoculum × 100) or mean percent invasion (number of bacteria surviving gentamicin treatment/number of bacteria in initial inoculum × 100) from three to six samples.

Bovine ligated-ileal-loop assay for enteropathogenesis. All bovine experiments were conducted according to the requirements of the Animal Scientific Procedures Act (United Kingdom, 1986). The assay has been described in detail elsewhere (38). Briefly, calves were terminally anesthetized with pentobarbitol, and intestinal loops were constructed in the ileum. Bacterial strains were grown overnight at 25°C, with shaking. The cultures were diluted approximately 1:3 in fresh LB medium and incubated at 37°C for 90 min, with shaking. The optical density at 600 nm was adjusted by addition of LB broth to give a concentration of approximately 3×10^8 CFU/ml. A total of 5 ml of this suspension was injected into each loop. The same volume of sterile LB broth was used as a negative control. All bacterial strains and controls were tested in three loops per animal. Polymorphonuclear leukocytes (PMNs) were isolated from 50 ml of blood removed from the calves, labeled with ^{111}Indium (^{111}In), and reinjected into the jugular vein. Twelve hours after inoculation, the anesthetized animals were humanely killed and all loops were exteriorized. Fluid secretion was measured as the ratio of volume of accumulated fluid to loop length. PMN influx was measured as the ratio of the ^{111}In activity in test loops to that in control loops.

Statistical analyses. For LD_{50} studies, probit analysis (29, 40) was used to calculate the standard error (SE) of the LD_{50}. The 95% confidence limits (CL) of the LD_{50} were determined according to the following formula: $LD_{50} \pm 1.96 \times$ SE of the LD_{50} (40). For studies of the number of *Salmonella* organisms per spleen over time, geometric mean (GM) values were determined for each group of five infected mice. The 95% confidence intervals about each GM were determined according to the following formula: GM $\pm$ $t_{[95\ \text{for}\ n\text{-}1]}$ (2.78 for 5 animals) × SE of the GM. Means in these two sets of experiments for which 95% CL did not overlap were considered statistically significantly different at $P = 0.05$. For the number of *Salmonella* organisms per macrophage for which there were three assays per time point, the SE of the arithmetic mean was calculated, but the 95% confidence interval was not determined because of the small sample size. For data from tissue culture adherence and invasion assays that were presented as mean percentages, calculation of SE was deemed inappropriate. Rather, the range of the samples ($n = 3$ to 6) from which the mean percentile was calculated was given. For enteropathogenic responses induced by *Salmonella* in the calf loop model, Student's unpaired *t* test was used to compare means of groups.

RESULTS

Construction and characterization of a flagellar regulatory mutant. The gene encoding the master regulator of flagella synthesis, *flhD*, was mutated in the virulent strain SL3201 by transduction of *flhD*::Tn*10* from strain KK2040. As expected, SL3201 *flhD*::Tn*10* was nonmotile in motility agar, and sodium dodecyl sulfate-polyacrylamide gel electrophoresis analysis of cell lysates and supernatants from the *flhD* mutant showed no flagellin bands (data not shown). The in vitro growth rate of the *flhD* strain in rich medium was similar to that of SL3201 (data not shown).

Mouse virulence of flagellar mutants. We compared the virulences of SL3201, the wild-type strain, SL3201 *fljB*::Mud*J fliC*::Tn*10*, a mutant capable of forming the basal body-hook complex but incapable of producing flagellin, and SL3201 *flhD*::Tn*10*, a mutant that lacks any products of the flagellar cascade including the flagellar secretory apparatus. As expected from previous studies (32), the virulence of the Fla$^-$ strain SL3201 *fljB*::Mud*J fliC*::Tn*10* was very similar to that of the wild-type parent strain (Table 1). Mutation of the gene encoding the master regulator of flagellar synthesis, *flhD*, did not decrease the virulence of serovar Typhimurium. In fact, in

TABLE 1. Virulence of *Salmonella enterica* serovar Typhimurium flagellar mutants

Bacterial strain	No. of dead mice/no. infected[a]	Oral LD_{50}[b] (CL)[c]
SL3201	10/10	1.5×10^5 (9×10^5–2.5×10^4)
SL3201 *fljB*::Mud*J* *fliC*::Tn*10*	10/10	1.0×10^5 (3.0×10^5–3.3×10^4)
SL3201 *flhD*::Tn*10*	10/10	1.4×10^4 (2.4×10^4–8.8×10^3)

[a] Intraperitoneal injection of ~700 CFU. The data are from two experiments.
[b] The LD_{50} values of each strain administered orally were determined at least twice by probit analysis (29, 40), and the mean values are presented in CFU.
[c] CL, 95% confidence limits.

each experiment the oral LD_{50} value of SL3201 *flhD*::Tn*10* was at least 10-fold lower than that of SL3201, and the 95% CL of the mean oral LD_{50} value for the mice given SL3201 did not overlap with the CL of the mean oral LD_{50} for the mice given SL3201 *flhD*::Tn*10*. Therefore, the oral LD_{50} for the mice given SL3201 *flhD*::Tn*10* was significantly lower than that of wild-type SL3201 ($P = 0.05$). These in vivo results suggest that products secreted solely by the serovar Typhimurium flagellar secretory apparatus are not required for invasion of the intestinal mucosa or persistence in tissues of mice.

Kinetics of infection after oral inoculation. The capacities of the Fla^- *flhD* and *fljBfliC* mutants to reach and multiply within the spleen were compared to those of the Fla^+ wild-type strain. Mice were orally inoculated with approximately 10^7 CFU of SL3201, SL3201 *flhD*::Tn*10*, or SL3201 *fljB*::Mud*J* *fliC*::Tn*10*. By day 9 postinfection, slightly higher numbers of the *flhD* strain than of the wild-type strain were recovered from the spleens of infected mice, and two of five mice infected with the *fljB fliC* mutant had died (data not shown). We repeated and extended the kinetic study with the wild-type parent and the *flhD* mutant (Fig. 1). Groups of mice were infected and examined up to 14 days postinfection. Although the 95% CL of the GM of the two groups overlapped at all time points (see the legend to Fig. 2), certain trends were noted. First, *Salmonella* organisms were recovered from the spleens of only two mice during the first 2 days of infection. Second, the wild-type strain SL3201 was isolated from the spleens at earlier time points than was the *flhD* mutant (days 1 and 3). Third, four of the five *flhD*-infected mice died by day 14 of infection, compared to only one of the mice from the SL3201 group. All together, this pattern may indicate that the *flhD* mutant was less capable of reaching the spleen than was the wild-type strain, but once the organism reached the spleen it either replicated more rapidly or survived better than the wild-type strain.

Survival in mouse macrophages. We have shown a correlation between the degree of survival of serovar Typhimurium in resident peritoneal macrophages and the rate of net growth of wild-type serovar Typhimurium in spleens (22). Therefore, we compared the survival rates of the *flhD* mutant, the *fljB fliC* mutant, and wild-type *Salmonella* in resident peritoneal mac-

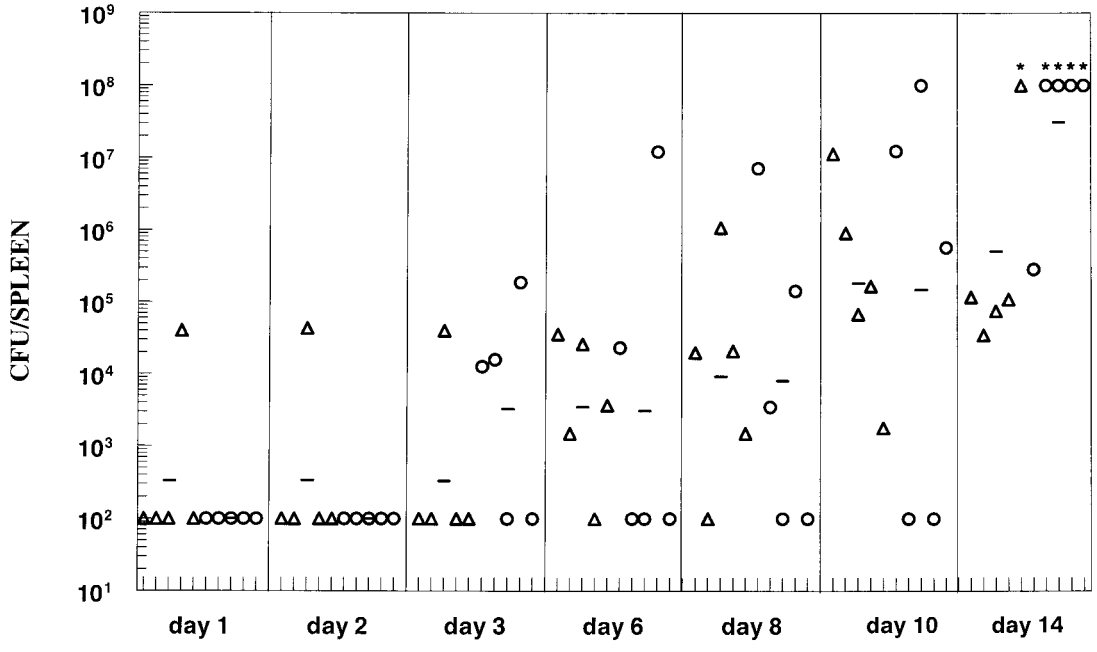

FIG. 1. Kinetic study of flagellar mutants recovered from the spleen following oral inoculation of mice. C57BL/6J mice were inoculated with SL3201 (triangles) or SL3201 *flhD*::Tn*10* (circles) at a dose of approximately 10^7 CFU per mouse. At various times postinoculation up to 14 days, mice were sacrificed and the number of CFU per spleen was determined. Each symbol represents data from an individual animal. The horizontal line indicates the GM for the group of five mice. Asterisks indicate mice that had died; values of 10^8 CFU/spleen were assigned to these mice. The 95% CL of the GMs for the two groups overlapped at all time points including day 14 (data not shown). Limit of detection = 10^2 CFU/spleen.

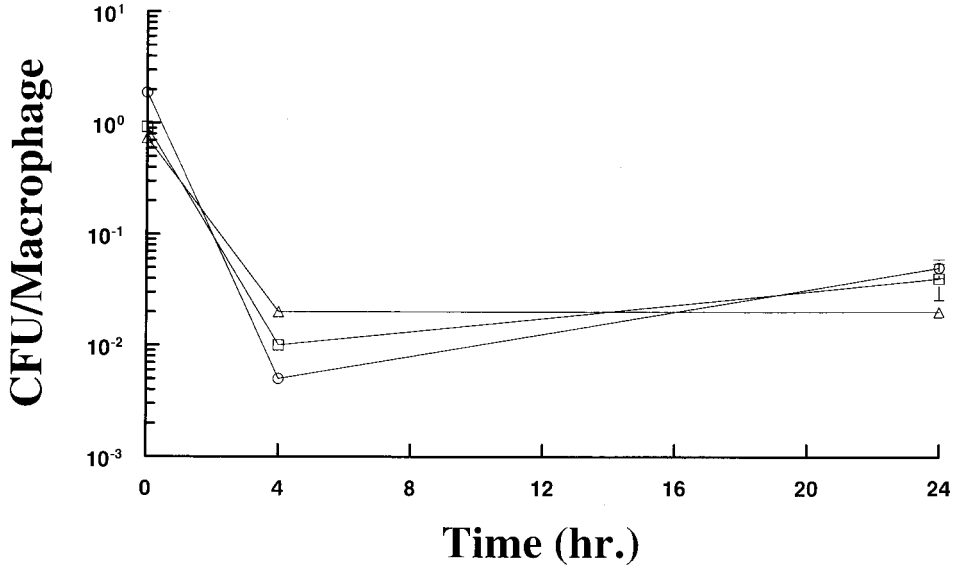

FIG. 2. Survival in mouse resident peritoneal macrophages. Macrophages were infected in vitro with SL3201 (triangles), SL3201 *flhD*::Tn*10* (circles), or SL3201 *fljB*::Mud*J fliC*::Tn*10* (squares) at a MOI of approximately five opsonized bacteria per cell. The numbers of viable bacteria and macrophages per well were determined at 0, 4, and 24 h postinfection. The ratio of CFU/macrophage and one SE of the mean are shown. Note that the SE bars are covered by the symbols at 0 and 4 h postinfection.

rophages. The CFU of all strains tested consistently decreased at 4 h postinfection of the macrophages (Fig. 2). However, between 4 and 24 h postinfection, the rate of net growth (difference between multiplication and death) of the *flhD* and *fljB fliC* mutants appeared to be higher than that of the wild-type strain, as revealed by the slopes of the lines between these time points which were steeper for the mutants than for the wild-type strain. By 24 h postinfection the numbers of all three bacterial strains were similar. No significant differences were seen in the numbers of macrophages present, regardless of the bacterial strain tested.

Adherence and invasion of intestinal epithelial cells. The flagellar regulatory mutants were also tested in an in vitro model for the capacity to adhere to and invade mouse and human small intestinal epithelial cell lines (MODE-K and Henle-407). In these assays, bacteria were gently centrifuged onto the cell monolayers; this centrifugation step was included to compensate for the fact, which others have reported, that nonmotile strains of serovar Typhimurium are less invasive in vitro than their motile counterparts (8, 18). The mean percents (calculated from sample sizes of three to six) of *Salmonella* organisms that adhered to MODE-K cells were similar for all strains tested: for SL3201, the mean was 33% and the range was 23 to 48%; for SL3201 *fliA*::Tn*10*, the mean was 29% and the range was 20 to 36%; for SL3201 *fljB*::Mud*J fliC*::Tn*10*, the mean was 31% and the range was 16 to 47%; and for SL3201 *flhD*::Tn*10*, the mean was 32%, and the range was 29 to 35%. However, the wild-type serovar Typhimurium strain SL3201 invaded mouse MODE-K cells in greater numbers (mean, 20%; range, 19 to 29%) than did *fliA* (mean, 7%; range, 5 to 8%), *fljB fliC* (mean, 10%; range, 7 to 13%), and *flhD* flagellar mutants (mean, 6%; range, 5 to 8%), even following centrifugation of the bacteria onto the cells. The wild-type serovar Typhimurium strain SL3201 also invaded human Henle-407 cells in greater numbers (mean, 9.4%; range, 7.6 to 11%) than did the *fliA* (mean, 1.1%; range, 0.9 to 1.3%) and *flhD* (mean, 1.3%; range, 1.2 to 1.3%) flagellar mutants. These results are similar to those reported by Eichelberg and Galán (6), who found that *fliA* and *flhD* mutants of serovar Typhimurium invaded Henle-407 cells at reduced levels (~35% of wild-type). These in vitro results suggest that the flagellar mutants are capable of binding to but are defective in penetrating the intestinal epithelial cells.

Induction of enteropathogenic responses. *Salmonella* infection can also cause enteritis in several animal hosts. Therefore, the contribution of serovar Typhimurium flagella or the flagellar secretory apparatus to enteropathogenesis was assessed in the well-established bovine ileal-loop model (Fig. 3). In that model, the secretory response induced by the *flhD* mutant was significantly less than that of the wild-type strain ($P < 0.04$ by Student's unpaired *t* test), as was the PMN influx evoked by the *flhD* mutant compared to that of the parent ($P < 0.02$). Moreover, the average fluid volume induced by the *fljB fliC* mutant and the average PMN influx in response to that mutant were reduced compared to what was observed with the parent strain, but these differences were not statistically significant.

DISCUSSION

In this investigation, flagellar mutants were tested in in vivo and in vitro assays to evaluate the importance of flagellar components in the virulence of serovar Typhimurium. An *flhD* mutant does not synthesize any flagellar components and therefore has no flagellar secretory apparatus. Both *fliA* and *fljB fliC* mutants produce functional flagellar secretory apparatuses, but the former is incapable of expressing certain class 3 gene products while the latter does not make flagellin. We and others have previously demonstrated that flagella are not

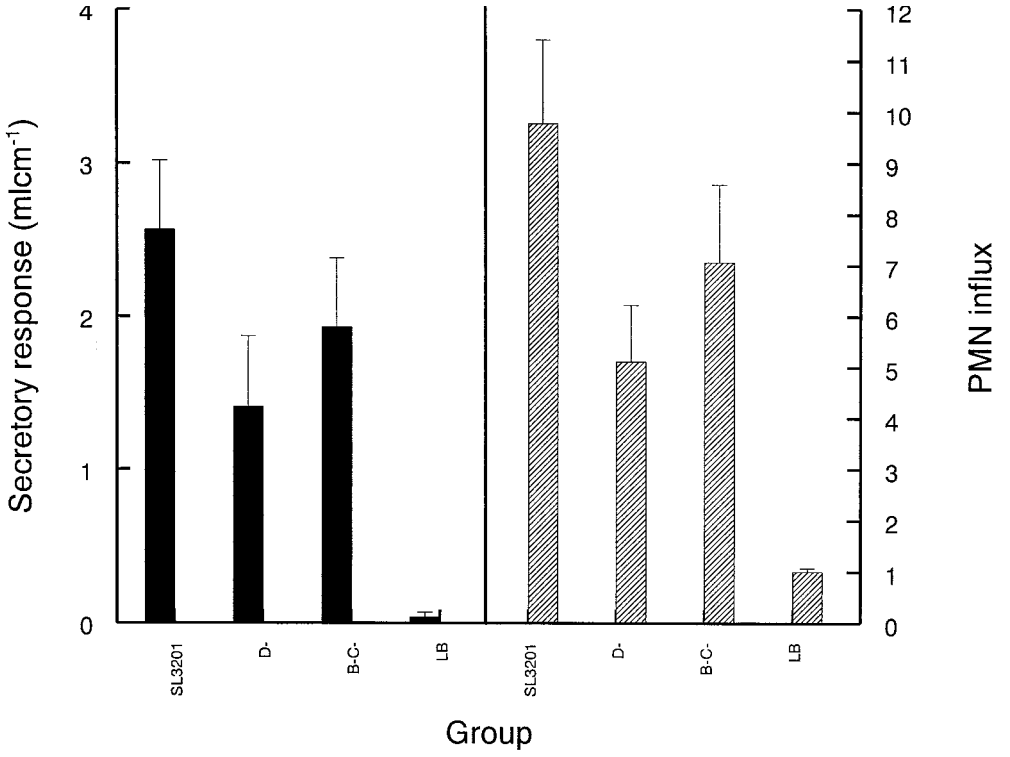

FIG. 3. Secretory response and PMN influx elicited by serovar Typhimurium strains in bovine intestinal loops. Approximately 1.5×10^9 CFU of the wild-type strain SL3201 (WT), SL3201 *flhD*::Tn*10*(D-) or SL3201 *fljB*::Mud*J fliC*::Tn*10* (B-C-) was injected into each loop and left for 12 h before analysis. LB broth (LB) was also used as a negative control. The secretory response is shown as volume per length of loop. The PMN influx is a ratio of the PMN radioactivity within the infected loops to the PMN radioactivity within the control loops. Each mean is calculated from either nine loops in three calves (secretory response) or six loops in two of the three calves (PMN influx) and is presented with the SE.

necessary for the virulence of serovar Typhimurium in the mouse model of typhoid (23, 32). Consistent with the previous work, we found in this study that a mutation in the master regulator of flagellar synthesis (*flhD*) did not decrease virulence of the strain. The *flhD* mutant was virulent in mice whether administered intraperitoneally or orally (Table 1), which suggests that there are no factors secreted solely by the flagellar secretory apparatus necessary for virulence in the mouse model. In fact, the *flhD* mutant was slightly more virulent (the oral LD_{50} value was 10-fold lower than for the wild-type strain), and in kinetic studies (Fig. 1) more mice infected with the *flhD* mutant died than did those infected with the wild-type strain. Although the growth rate of the mutant was similar to that of the wild-type strain in rich media, it is possible that the slight increase in virulence by the *flhD* mutant may be due to the decreased metabolic burden resulting from the incapacity to synthesize any flagellar components. Consistent with this hypothesis is that both flagella-less mutants (*flhD* and *fljB fliC*) grew at a slightly faster net rate (*flhD* slightly faster than *fljB fliC*) in peritoneal macrophages between 4 and 24 h postinfection (Fig. 2).

All flagellar mutants tested (*fliA, fljB fliC*, and *flhD*) invaded MODE-K or Henle-407 cells less well than did the wild-type strain, although strains adhered to their target cells (MODE-K cells) similarly following centrifugation onto the cells. A possible explanation reported by others for the decreased invasion by *fliA* and *flhD* mutants is that the flagellar sigma factor (FliA) also affects the expression of invasion genes (6). Another possibility suggested by Lucas et al. (24) is that *fliZ*, rather than *fliA*, may be involved in invasion gene expression. These authors reasoned that *fliZ* is located downstream of *fliA* in the same operon. Moreover, the *fliZ* product activates *hilA*, and *hilA* is involved in the expression of invasion genes (1, 2). In any case, we found that the *fliA* (33) and *flhD* mutants were still virulent in susceptible mice. Therefore, we conclude that either the epithelial invasion assay does not reflect the interactions of the bacterium with the mouse intestine in the murine typhoid oral challenge model or the assay does mimic that step in the pathogenesis but the low number of flagellar mutant cells that invade epithelial cells is sufficient to cause systemic disease like the wild-type strain.

The difference between the *flhD* mutant and the wild-type strain in fluid secretion responses in bovine ligated ileal loops (Fig. 3) suggests that flagellar components of serovar Typhimurium are involved in enteropathogenesis. One possible explanation is that *Salmonella* secretes through the flagellar apparatus a factor(s) that affects fluid secretion. Type III secretion has been shown to be important in the enteropathogenesis of *Salmonella* infection (37). *Salmonella* pathogenicity island 1 (SPI-1) effectors that are secreted may be involved in cell invasion, PMN recruitment, and fluid secretion, although the correlation of cell invasion with enteropathogenesis is not clear. We speculate that the factor(s) responsible for fluid secretion that are exported by the type III secretory apparatus

may also be secreted by the flagellar apparatus, or other factor(s) responsible for fluid secretion may be exported exclusively through the flagellar apparatus. Proteins secreted by the flagellar apparatus have been shown to affect virulence. For example, *Yersinia enterocolitica* secretes a phospholipase involved in virulence through the flagellar apparatus (42). Also, the anti-sigma factor (FlgM) regulates flagellar synthesis by secretion through the flagellar apparatus (17) and was shown to be involved in the virulence of serovar Typhimurium in the mouse model (33). However, the attenuated phenotype resulting from an *flgM* mutation can be reversed by a second mutation in *fliA* or *fliC*, a finding which suggests that unregulated expression of flagella attenuates the bacterial cell and not the lack of interaction of FlgM with the host cell (32, 33). Another explanation for the decreased fluid secretion response to an *flhD* mutant is that other flagellar proteins such as FliA or FliZ, which are not made in an *flhD* mutant, control expression of virulence determinants, as has been proposed by others (6, 24).

The PMN influx responses to both nonflagellated mutants (*flhD* and *fljB fliC* mutants) were lower than those of the wild-type strain (Fig. 3), suggesting that flagella or motility influences the recruitment of PMNs. Flagella of *Salmonella* are known to be potent inducers of cytokines (4, 41). In fact, Gewirtz et al. (11) recently reported that flagellin of serovar Typhimurium is responsible for induction of interleukin-8 (IL-8) production by intestinal epithelial cells exposed to wild-type organisms. As would be predicted from that observation, Gewirtz and colleagues also showed that serovar Typhimurium *flhD* and *fljB fliC* mutants (the same ones used in this study) failed to induce IL-8 secretion (11). IL-8 is a potent PMN attractant and is involved in PMN migration into the subepithelial and luminal compartments of the intestine by *Salmonella* (27, 28).

In conclusion, neither flagella nor components of the flagellar synthetic apparatus of serovar Typhimurium are required for the expression of virulence in the mouse model of typhoid fever or survival in mouse macrophages in vitro, but flagella are required for full virulence potential in tissue culture invasion assays and for the induction of a complete inflammatory response in the calf model of enteropathogenesis. In the latter model, the synthetic apparatus is also required for maximum fluid secretion in the calf intestine. The differential results obtained with these assays using the same mutant strains suggest that the mouse systemic infection and macrophage assays may measure aspects of virulence other than those measured by the tissue culture invasion assay and the calf enterocolitis model.

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