

## The *rovA* Mutant of *Yersinia enterocolitica* Displays Differential Degrees of Virulence Depending on the Route of Infection

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***Yersinia enterocolitica* is an invasive enteric pathogen that causes significant inflammatory disease. Recently, we identified and characterized a global regulator of virulence (*rovA*). When mice are infected orally with the *rovA* mutant they are attenuated by 50% lethal dose (LD<sub>50</sub>) analysis and have altered kinetics of infection. Most significantly, mice orally infected with the *rovA* mutant have greatly reduced inflammation in the Peyer's patches compared to those infected with wild-type *Y. enterocolitica*. However, we present data here indicating that when the *rovA* mutant bacteria are delivered intraperitoneally (i.p.), they are significantly more virulent than when delivered orally. The i.p. LD<sub>50</sub> for the *rovA* mutant is only 10-fold higher than that of the wild-type *Y. enterocolitica*, and there are significant inflammatory responses to the *rovA* mutant that are evident in the liver and spleen. Altogether, these data suggest that the RovA regulon may be required for the early events of the infection that occur in the Peyer's patches. Furthermore, these data suggest that the RovA regulon may be dispensable for *Y. enterocolitica* systemic disease and inflammatory responses if the Peyer's patches are bypassed.**

*Yersinia enterocolitica* is one of three species of *Yersinia* that are pathogenic for humans (7). The other two pathogenic species of *Yersinia* are *Yersinia pseudotuberculosis* and *Yersinia pestis*. Both *Y. enterocolitica* and *Y. pseudotuberculosis* are enteric pathogens, whereas *Y. pestis* is the causative agent of plague. *Y. enterocolitica* infection is usually the result of consuming contaminated food or water and usually results in a self-limiting infection that presents as gastroenteritis and/or lymphadenitis (26).

The initial site of *Y. enterocolitica* infection is the secondary lymphoid tissues of the small intestine called Peyer's patches (PP). To gain entry to the PP, the bacteria must first bind to and then cross the specialized intestinal epithelium (M cells) that overlay the PP. The binding and translocation event is mediated by the bacterial invasin protein (Inv) (14, 18). Inv is an outer membrane protein that binds to  $\beta_1$ -integrins expressed on the apical surface of the M cell (9, 13, 15). Inv is required for the efficient translocation of the bacteria across the intestinal epithelium (22). In fact, when an *inv* mutant is examined in the mouse model of infection it has an altered pathogenesis phenotype compared to the wild-type (WT) bacteria (22). Interestingly, even though there is a delay in the colonization of the PP of mice infected with the *inv* mutant, there is no difference in the 50% lethal dose (LD<sub>50</sub>) (22).

When the expression characteristics of the Inv protein were examined *in vitro* it was found that expression was modulated by environmental factors such as temperature, pH, and growth phase (21). These data suggested that there was a regulator of

Inv expression. Recently, we identified the Inv regulator and named it RovA (24). A homologue of RovA has been identified in *Y. pseudotuberculosis* and was shown to regulate the expression of *inv* in that organism as well (20). RovA is a member of the MarR family of transcriptional regulators and is responsible for the regulation of Inv expression both *in vitro* and *in vivo* (24). Subsequent analysis of the *rovA* mutant in the mouse provided several lines of evidence that suggested RovA plays a key role in the pathogenesis of a *Y. enterocolitica* infection. First, a *rovA* mutant is attenuated by LD<sub>50</sub> analysis in BALB/cj mice and shows altered kinetics of infection in this mouse (24). Second, a *rovA* mutant is deficient in the initiation of inflammation in the PP, a phenotype that is not a consequence of the decreased levels of Inv observed in the *rovA* mutant. Further analysis of the *rovA* mutant *in vivo* suggested that the defect in the initiation of PP inflammation was linked to a defect in the ability of the host to produce interleukin-1 $\alpha$  (IL-1 $\alpha$ ) in response to bacterial infection (12). Altogether, these data suggested that RovA was playing a central role in the infectious process that included the regulation of Inv and other as-yet-undetermined gene products. One or more of these unidentified genes are most likely responsible for the inflammatory pathologies observed during a *Y. enterocolitica* infection.

When the phenotypes of both the *inv* and the *rovA* mutant were considered together, it was reasonable to assume that RovA was important for the regulation of genes that influence the early stages of infection that occur in the PP. Therefore, it was of interest to reexamine the phenotype of the *rovA* mutant *in vivo* when the natural route of infection is bypassed. In this report we provide data suggesting that RovA is less critical for virulence in the mouse model when the bacteria are delivered by the intraperitoneal (i.p.) route of infection. Furthermore, when mice are infected i.p. with the *rovA* mutant they are

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capable of mounting a significant inflammatory response that is not observed during oral infection.

#### MATERIALS AND METHODS

**Mice.** Female C57BL/6J mice 6 to 8 weeks of age were purchased from Jackson Laboratory and maintained in the barrier facility at Washington University School of Medicine. Mice were given free access to food and water throughout all experiments. Animals were sacrificed by carbon dioxide asphyxiation. The Washington University committee on animal studies approved all animal experiments.

**Bacteria.** *Y. enterocolitica* strains used in the present study are WT strain (JB580v) (16) and the *rovA* mutant (YVM641) (22). Both strains are derivatives of the serogroup O8 strain 8081 and carry the pYV virulence plasmid. For inoculation of the mice, bacteria were grown overnight in Luria-Bertani (LB) broth at 26°C. Actual numbers of CFU were determined by serial dilutions of the overnight culture, followed by plating on LB agar containing 20 µg of nalidixic acid/ml; both JB580v and YVM641 are resistant to nalidixic acid.

**LD<sub>50</sub> and kinetics analysis.** Five groups of five mice were infected orally by gavage with successive 10-fold dilutions of the bacterial suspension (10<sup>4</sup> to 10<sup>9</sup> bacteria) or i.p. with 10 to 10<sup>6</sup> bacteria for WT or 10<sup>2</sup> to 10<sup>7</sup> bacteria for the *rovA* mutant. The mice were monitored twice daily for 14 days. This analysis was done in duplicate in two independent experiments. The LD<sub>50</sub> values were determined according to the method of Reed and Muench (23). Kinetic analysis was done by infecting mice i.p. with the indicated dose of the WT *Y. enterocolitica* strain JB580v (16) or the *rovA* mutant YVM641. At various times postinfection (1, 3, or 5 days), mice from each group were sacrificed and tissues (PP, mesenteric lymph nodes, spleen, and liver) were removed. The bacterial load recovered from the infected organs was determined by plating dilutions of the macerated tissues onto LB plates containing 20 µg of nalidixic acid/ml to select for *Yersinia* and was reported as the CFU per gram of tissue. This kinetics analysis was done in duplicate in two independent experiments.

**Histopathology.** Mice of the specified group were infected i.p. with indicated amounts of WT *Y. enterocolitica* strain JB580v or *rovA* mutant YVM641. On days 1, 3, 5, 14, and 24 postinfection the mice were sacrificed and the small intestines, mesenteric lymph nodes, spleen, liver, kidney, and lungs were removed. The lumen of the intestine was flushed with phosphate-buffered saline (PBS), and then tissues were fixed in either 10% neutral buffered formaldehyde or Bouin's fixative prior to being embedded in paraffin and stained with hematoxylin and eosin. Mice infected orally by gavage for 14 or 24 days were treated in a similar manner. Slides were investigated in a blind fashion by two independent investigators. Tissue sections were given an arbitrary score based on the frequency and severity of the lesions. Tissues were scored on a scale of 0 to 4, with 0 representing no histopathologic abnormality (consistent with the mock-infected control tissues) and 4 being the most severe pathology observed. Sections were scored for increased numbers of macrophages and neutrophils, granuloma formation, inflammatory infiltrates, necrosis, fibrin thrombi, bacterial colonization, and adhesions. A minimum of five mice per time point were examined. Tissues from dead mice were excluded from analysis.

**Immunohistochemistry of paraffin-embedded sections.** Detection of *Y. enterocolitica* within the tissues of infected mice was performed by using the avidin-biotin complex procedure as described previously (19). *Y. enterocolitica* was detected by using purified rabbit polyclonal antibodies directed against membranes from the WT strain (JB580v) at a dilution of (1:500). Antibodies were prepared by the immunization of a New Zealand White rabbit with *Y. enterocolitica* membrane preparations. After exsanguination, serum was separated from whole blood. Serum was absorbed with acetone powders derived from *E. coli* strains DH5α and HB101 and *Salmonella enterica* serovar Typhimurium strain 14028s. Immunoglobulin G was then purified from the absorbed serum by affinity chromatography by using protein A-Sepharose. Positive staining was detected by using diaminobenzidine, and sections were counterstained with Mayer's hematoxylin.

**Immunohistochemistry of frozen sections.** Mice were infected i.p. with 10<sup>3</sup> CFU of WT or *rovA* mutant bacteria and then sacrificed after 14 or 24 days, and the indicated tissues were removed, embedded in OCT compound, and flash frozen. Sections were cut on a cryostat and fixed in acetone or methanol prior to staining. Tissues were rehydrated in PBS. Peroxidase activity was blocked by incubation with 0.3% hydrogen peroxide and then blocked with bovine serum albumin and milk, as well as avidin-biotin block (Vector Laboratories). Sections were incubated overnight with biotinylated antibody at the indicated concentrations. The antibodies used were anti-murine macrophage antigen F4/80 (1:250; Serotec), anti-TCRβ (1:1,000; Pharmingen), and anti-*Y. enterocolitica* (1:1,000). Granulocytes were detected with anti-GR-1 antigen (1:250; Pharmingen). Sec-

tions were washed in PBS and then incubated with 1 µg of streptavidin-horse-radish peroxidase and goat anti-rabbit-fluorescein isothiocyanate (1:1,000)/ml for 1 h. Sections were washed in PBS, and then positive staining was detected by using tyramide signal amplification 3' cyanine (Dupont-NEN) counterstained with Hoechst stain and visualized with fluorescence microscopy.

**Statistical analysis.** All statistical analysis was done by using GraphPad Prism v.3.03 (Graphpad Software). Kinetic data was log<sub>10</sub> transformed and then subjected to analysis of variance by using a nonparametric Mann-Whitney test to generate a two-tailed *P* value. Comparisons were made between a given tissue within a given time point. Statistically significant comparisons are denoted with an asterisk.

#### RESULTS

**The *rovA* mutant is more virulent by the i.p. route of infection than by the oral route.** The normal route of infection for *Y. enterocolitica* is via the oral route. Previously, we reported that when BALB/cj mice were orally infected with the *rovA* mutant they were found to be 70-fold attenuated by LD<sub>50</sub> analysis compared to the WT bacteria (24). When the oral LD<sub>50</sub> was determined for the *rovA* mutant in C57BL/6j mice, the bacteria were even less virulent in this mouse background. In the C57BL/6j background the *rovA* mutant was 500-fold attenuated compared to the WT bacteria in an oral LD<sub>50</sub> analysis (Fig. 1A). Interestingly, when this analysis was extended to include the i.p. route of infection the *rovA* mutant was almost as virulent as the WT bacteria, with only an 11-fold change in LD<sub>50</sub>, (Fig. 1A). These data suggest that *rovA* may play a more important role in the establishment of colonization of the PP and/or dissemination from the PP than in the elaboration of a systemic infection.

When the kinetics of oral infection were examined for the *rovA* mutant in the BALB/cj mouse model, it was apparent that there was a defect in the ability of the *rovA* mutant to disseminate (24). The BALB/cj mice infected orally with the *rovA* mutant showed a defect in the dissemination to the mesenteric lymph nodes and, to a lesser extent, the spleen compared to mice infected with WT bacteria. C57BL/6j mice infected orally with the WT bacteria or the *rovA* mutant showed a similar but more pronounced dissemination defect than that observed with oral infection of BALB/cj mice (data not shown). Analysis of an i.p. *rovA* infection of C57BL/6j mice displayed a different kinetic profile than that for the oral infection (Fig. 1B). There was significant colonization of all of the tissues by day 1 when the mice were infected with the WT bacteria. However, when the mice were infected i.p. with the *rovA* mutant there were fewer bacteria colonizing the deeper tissues at earlier time points (especially days 1 and 3) compared to the WT. Unlike the data from the oral infection studies, there appears to be no bias in the colonization of the deeper tissues by the *rovA* mutant (i.e., all tissues are equally affected) nor does there appear to be a significant virulence defect once infection is established. There only appears to be a delay in efficient colonization. Statistical analysis suggests that there were no significant differences between mice infected i.p. with WT bacteria and mice infected with the *rovA* mutant. When subjected to statistical analysis, only the data from spleens infected for 5 days were statistically significant (*P* = 0.0055). Altogether, these data, considered along with the LD<sub>50</sub> data, suggest that *RovA* may be required early in i.p. infection but then, as the infectious process proceeds, the *rovA* mutant behaves more like the WT bacteria.

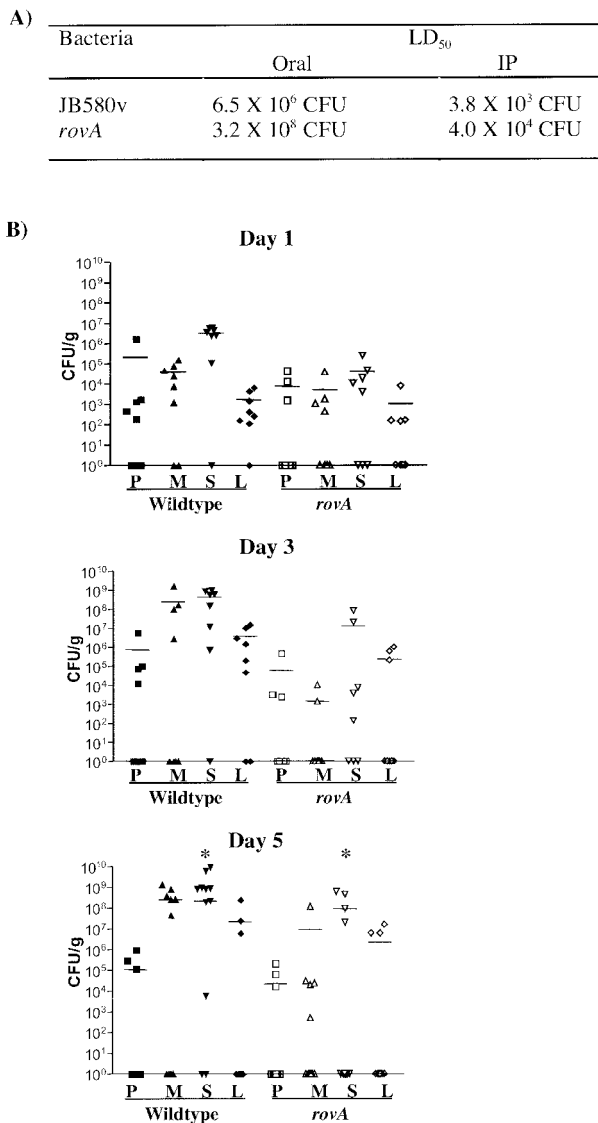


FIG. 1. Analysis of the *rovA* mutant by the i.p. route of infection. (A) Five C57BL/6j mice per group were infected orally or i.p. with 10-fold serial dilutions of WT virulent *Y. enterocolitica* (JB580v) or *rovA* mutant (YVM641) and then monitored for death over a 14-day period. The LD<sub>50</sub> was calculated by the method of Reed and Muench (21). The results are the average of two independent experiments. (B) Kinetics of infection. C57BL/6j mice were infected i.p. with 10<sup>5</sup> CFU of WT *Y. enterocolitica* or *rovA* mutant. The infection was allowed to proceed for the indicated amount of time. Mice were then sacrificed, and the indicated organs were harvested. Bacterial load was determined by enumerating the number of viable bacteria in each organ (P, PP; M, mesenteric lymph nodes; S, spleen; L, liver). Each individual symbol on the histogram represents an individual mouse. Symbols on the x axis represent tissues that did not have recoverable CFU. The results are the composite of two independent experiments. Asterisks indicate statistically significant comparisons ( $P = 0.0055$ ). Eight mice were analyzed for days 1 and 3. Thirteen mice were analyzed on day 5. Solid symbols represent the WT *Y. enterocolitica*-infected animals, and the open symbols represent mice infected with the *rovA* mutant. At day 5 (the bottom panel), three of the animals for the WT day 5 time point died prior to analysis.

**Histopathology of i.p. infection with either WT *Y. enterocolitica* or *rovA* mutant reveals inflammatory changes.** The most striking phenotype of an oral infection with the *rovA* mutant is a lack of PP inflammation due to decreased levels of IL-1 $\alpha$  (12); this is in contrast to infection with WT *Y. enterocolitica*, which shows extensive inflammation as early as 2 days postinfection. The initial studies only examined a 7-day time course of infection. To further study the inflammatory response to the *rovA* mutant, an analysis of the tissues after an extended oral infection was conducted. C57BL/6j mice were infected orally with either *rovA* mutant or WT *Y. enterocolitica* for 14 or 24 days. Tissues (small intestine, spleen, liver, kidney, and mesenteric lymph nodes) were removed and prepared for hematoxylin and eosin staining, followed by histological examination. Histopathology data are summarized in Tables 1 and 2. The histopathology of earlier time points of *Y. enterocolitica* infection has been described (2, 6, 12). Interestingly, as previously reported for earlier time points, mice infected orally with the *rovA* mutant did not have a strong inflammatory response even at the 24-day time point in any of the tissues. However, an acute inflammatory response to infection with WT *Y. enterocolitica* was apparent by day 3 in the PP and mesenteric lymph nodes and in all tissues examined by day 7 (12). The mice infected orally with the WT *Y. enterocolitica* for 14 days had more severe inflammatory pathologies than the mice infected orally for 24 days. This is probably due to the mice with the most severe pathologies dying prior to the 24-day time point.

To further examine the inflammatory response to the *rovA* mutant, an analysis of the inflammatory response to an i.p. infection was conducted. C57BL/6j mice were infected i.p. with either the *rovA* mutant or WT *Y. enterocolitica* for 1, 3, 5, 14, or 24 days, and then tissues (small intestine, spleen, liver, lung, kidney, and mesenteric lymph nodes) were removed and prepared for hematoxylin and eosin staining, followed by histological examination. Consistent with the kinetic data, there appeared to be a slight delay in histopathological evidence of infection by the *rovA* mutant compared to WT *Y. enterocolitica* (Tables 1 and 2). This was readily apparent during gross examination of the animals during necropsy. The animals infected with WT *Y. enterocolitica* developed colonization of the serosal surfaces of the peritoneum as early as day 1, whereas the *rovA* mutant did not show this phenotype until day 3 or later (data not shown). This same phenotype was evident during histological examination of intestinal sections (Fig. 2). Distinct microcolonies were evident on the serosal surface of the small intestine (Fig. 2B); this was not observed during an oral infection. The presence of *Y. enterocolitica* in the tissues was confirmed by using a *Y. enterocolitica* specific antibody (Fig. 2C).

Although there appeared to be an initial delay in the colonization of the peritoneal organs and the onset of inflammatory responses when mice were infected i.p. with the *rovA* mutant, severe inflammatory pathologies were evident at the 14- and 24-day time points. This finding is in contrast to an oral infection with the *rovA* mutant that showed no significant inflammatory response even at 24 days (data not shown). The most severe pathologies involved the liver. There were multiple liver abscesses in the 24-day *rovA* mutant i.p.-infected mouse. Within the liver, lesions included well-defined granulomas (Fig. 2I and Fig. 3), areas of necrosis (Fig. 2F), and

TABLE 1. Histopathology of *rovA* mutant-infected mice

Source	Avg pathological score (frequency [%]) of mice infected with <i>rovA</i> mutant:											
	Orally at (day, dose [CFU], no. of mice):					i.p. at (day, dose [CFU], no. of mice):						
	14, 10 <sup>9</sup> , 5	24, 10 <sup>9</sup> , 10	1, 10 <sup>5</sup> , 5	3, 10 <sup>5</sup> , 5	5, 10 <sup>5</sup> , 5	14, 10 <sup>4</sup> , 8	24					
						10 <sup>3</sup> , 3	10 <sup>4</sup> , 3	10 <sup>5</sup> , 1	10 <sup>6</sup> , 4	10 <sup>7</sup> , 2	10 <sup>8</sup> , 2	
<b>Liver</b>												
Granuloma	0	0	0	0	0	4 (0.5)	4 (1)	3 (1)	* <sup>e</sup>	4 (0.5)	0	4 (0.5)
Microcolony <sup>a</sup>	0	0	3 (0.2)	3 (0.4)	4 (0.4)	3.6 (0.75)	4 (1)	0	*	4 (0.5)	0	3.5 (1)
Microabscess <sup>b</sup>	0	0	3 (0.2)	3 (0.4)	4 (0.4)	3.6 (0.75)	3.6 (1)	0	*	4 (0.5)	0	3.5 (1)
Necrosis	3 (0.2)	0	0	0	0	3.6 (0.75)	3.3 (1)	3 (1)	*	4 (0.5)	0	3.5 (1)
Infiltrate <sup>c</sup>	3 (0.2)	0	2 (0.2)	2.5 (0.4)	4 (0.4)	3.5 (0.1)	4 (0.7)	4 (1)	*	2.8 (1)	1 (1)	4 (0.5)
Fibrin thrombi	3 (0.2)	0	0	3 (0.4)	0	0	3 (0.7)	0	*	4 (0.5)	0	3 (0.5)
Adhesion	0	0	0	0	0	4 (0.5)	3.5 (0.7)	0	*	4 (0.5)	0	0
<b>Spleen</b>												
Splenomegaly	0	0	0	0	0	2.8 (0.1)	2.5 (1)	0	*	3 (0.5)	0	3.5 (1)
Granuloma	0	0	0	0	0	0	0	0	*	0	0	0
Microcolony	0	0	3 (0.2)	3.5 (0.4)	0	0	0	0	*	0	0	0
Microabscess	0	0	0	0	0	0	0	0	*	0	0	0
Necrosis	0	0	0	2 (0.4)	0	3 (0.3)	2 (0.3)	0	*	0	0	0
Infiltrate	0	0	2 (0.2)	2 (0.2)	0	0	0	0	*	0	0	0
Fibrin thrombi	0	0	0	0	0	0	0	0	*	0	0	0
Adhesion	0	0	0	0	0	0	0	0	*	0	0	0
Inflammation <sup>d</sup>	0	0	0	2.5 (0.6)	0	2.8 (0.75)	0	0	*	3 (0.5)	0	4 (1)
<b>Mesenteric lymph nodes</b>												
Granuloma	0	0	0	0	0	3.8 (1)	4 (0.3)	0	*	4 (1)	0	0
Microcolony	0	0	0	2.5 (0.6)	0	3.8 (1)	4 (0.3)	0	*	3.5 (1)	0	0
Microabscess	0	0	0	3 (0.2)	0	3.8 (1)	4 (0.3)	0	*	4 (1)	0	0
Necrosis	0	0	0	0	0	4 (0.75)	4 (0.3)	0	*	0	0	0
Infiltrate	0	2 (0.1)	2 (0.1)	0	0	0	0	0	*	0	0	0
Adhesion	0	0	0	0	0	4 (0.2)	0	0	*	0	0	0
Inflammation	0	0	0	2.5 (0.6)	0	4 (0.75)	3.5 (0.7)	0	*	3 (1)	0	0
<b>PP</b>												
Microcolony	0	0	0	3 (0.2)	0	0	0	0	*	0	0	0
Microabscess	0	0	1.5 (0.5)	0	0	0	0	0	*	0	0	0
Inflammation	2.5 (0.5)	2 (0.2)	0	0	0	2.8 (1)	2.5 (1)	0	*	2 (0.5)	0	0
Edema	0	0	2 (0.2)	0	2 (0.2)	3.8 (1)	2.5 (1)	0	*	2 (0.3)	0	0
Infiltrate	0	0	0	0	0	3 (0.75)	2 (1)	0	*	0	0	0

<sup>a</sup> Microcolony refers to a defined infiltrate with microscopic evidence of bacterial colonization.

<sup>b</sup> Microabscess refers to a defined infiltrate without evidence of bacterial colonization.

<sup>c</sup> Infiltrate refers to a small defined infiltration of PMNs that is distinct from a microabscess or microcolony based upon size and organization of the lesion.

<sup>d</sup> Inflammation is scored as an undefined congestion of PMNs and macrophages in the tissues; this is distinct from the defined lesions that constitute a microabscess, microcolony, or infiltrate.

<sup>e</sup> An asterisk denotes that an insufficient number of animals survived to evaluate. Examples are shown in Fig. 2. Average pathological scores are reported from tissues that had pathology and the frequency that the finding was observed relative to the total number of animals in the experimental group is reported as a fraction in the parentheses (i.e., 1 = 100%).

infiltrates of polymorphonuclear leukocytes (PMNs) (Fig. 2E). The liver lesions often had microabscesses (defined infiltrates without evidence of bacterial colonization) (Fig. 2F), as well as microcolonies (defined infiltrates with evidence of bacterial colonization) (Fig. 2G and H). Adhesions between the liver and the kidney or the liver and the lung were frequently observed (Fig. 2I). In both cases, normal tissue architecture was severely disrupted. Inflammatory pathologies were also evident in other tissues as well, including granulomas of the inguinal lymph nodes, fibrin thrombi, ischemic necrosis in the kidneys and liver, and consolidation of the lungs, as well as edema and increased macrophages observed in the PP (data not shown).

Interestingly, when the localization of the *Yersinia* was examined at later time points (14 or 24 days) by immunohistochemistry, the bacteria were almost exclusively con-

tained within the granulomas (Fig. 3). This suggests that granuloma formation may provide the host partial protection during a *Yersinia* infection. Furthermore, these pathologies were often present in mice that received relatively low inocula (10<sup>3</sup> CFU i.p.) of either WT bacteria or *rovA* mutant. The architecture of granulomas formed in response to *Y. enterocolitica* infection is consistent with other types of granulomas. The center of the granuloma contains bacteria, PMNs (confirmed by positive GR-1 staining), and cellular debris (Fig. 3C). A concentric layer of epithelioid macrophages detected by F4/80 antigen staining and T cells detected by staining of the  $\beta$  chain of the T-cell receptor surrounds the center of the granuloma (Fig. 3A and B). Formation of granulomas in the mice infected i.p. with the *rovA* mutant was especially significant because it is indica-



TABLE 2. Histopathology of WT *Y. enterocolitica*-infected mice

Source	Avg pathological score (frequency [%]) of mice infected with WT <i>Y. enterocolitica</i> :											
	Orally at (day, dose [CFU], no. of mice):		i.p. at (day, dose [CFU], no. of mice):									
	14, 10 <sup>7</sup> , 5	24, 10 <sup>7</sup> , 5	1, 10 <sup>5</sup> , 5	3, 10 <sup>5</sup> , 5	5, 10 <sup>5</sup> , 5	14, 10 <sup>3</sup> , 5	24					
							10 <sup>2</sup> , 4	10 <sup>3</sup> , 3	10 <sup>4</sup> , 4	10 <sup>5</sup> , 4	10 <sup>6</sup> , 1	10 <sup>7</sup> , 1
<b>Liver</b>												
Granuloma	4 (0.2)	0	0	0	4 (0.2)	3.5 (0.8)	0	0	0	0	**e	*
Microcolony <sup>a</sup>	3 (0.2)	0	0	4 (0.2)	3.5 (0.6)	4 (0.8)	0	0	0	0	*	*
Microabscess <sup>b</sup>	3 (0.2)	4 (0.8)	0	4 (0.2)	3.8 (0.8)	4 (0.8)	3 (0.3)	0	0	0	*	*
Necrosis	2.8 (0.8)	3.5 (0.8)	0	0	3.2 (0.8)	3 (0.8)	3 (0.3)	0	0	2 (0.5)	*	*
Infiltrate <sup>c</sup>	4 (0.4)	3 (0.8)	2 (0.2)	3 (0.6)	0	3.5 (1)	4 (0.3)	0	3 (0.5)	2.5 (1)	*	*
Fibrin thrombi	3 (0.4)	3.2 (0.6)	0	3 (0.2)	2 (0.2)	3 (0.4)	3 (0.3)	0	0	0	*	*
Adhesion	0	0	0	0	0	4 (0.8)	0	0	0	0	*	*
<b>Spleen</b>												
Splenomegaly	3.2 (1)	4 (1)	0	0	0	3.8 (1)	4 (1)	4 (1)	4 (1)	4 (1)	*	*
Granuloma	4 (0.2)	0	0	3 (0.6)	4 (0.6)	3 (0.2)	0	0	0	0	*	*
Microcolony	4 (0.2)	0	0	4 (0.6)	4 (0.6)	3.2 (1)	0	0	0	0	*	*
Microabscess	4 (0.2)	0	2 (0.2)	4 (0.6)	4 (0.6)	3.2 (1)	3 (0.3)	0	3 (0.3)	0	*	*
Necrosis	4 (0.2)	0	0	0	0	0	3 (0.3)	0	3 (0.3)	0	*	*
Infiltrate	4 (0.2)	0	2 (0.2)	3 (0.6)	4 (0.6)	4 (1)	0	0	3 (0.3)	0	*	*
Fibrin thrombi	4 (0.2)	0	0	0	0	0	0	0	0	0	*	*
Adhesion	0	0	0	0	0	0	0	0	0	0	*	*
Inflammation <sup>d</sup>	2.5 (0.5)	3.8 (0.8)	2 (0.2)	3 (0.6)	4 (0.6)	3.5 (1)	3 (0.3)	0	3 (0.3)	4 (0.3)	*	*
<b>Mesenteric lymph nodes</b>												
Granuloma	3 (0.8)	4 (0.4)	0	0	3 (0.2)	3.8 (1)	4 (0.5)	0	0	4 (1)	*	*
Microcolony	3.5 (0.8)	4 (0.4)	0	3 (0.2)	3 (0.6)	4 (1)	4 (0.5)	0	0	4 (1)	*	*
Microabscess	3 (0.8)	4 (0.4)	0	3 (0.2)	3 (0.2)	4 (0.6)	4 (0.5)	0	0	4 (1)	*	*
Necrosis	3 (0.8)	4 (0.4)	0	0	0	4 (0.2)	3.2 (0.8)	0	0	4 (0.5)	*	*
Infiltrate	2.5 (0.2)	0	0	3 (0.6)	3 (0.6)	4 (0.2)	3 (0.8)	0	0	4 (1)	*	*
Adhesion	0	0	0	0	0	3 (0.6)	0	0	0	4 (0.5)	*	*
Inflammation	3.5 (1)	3.8 (1)	2 (0.2)	3 (0.6)	3 (0.6)	3.8 (1)	4 (1)	0	0	3.5 (1)	*	*
<b>PP</b>												
Microcolony	4 (0.2)	0	0	3.8 (0.8)	3.8 (0.8)	4 (0.2)	0	0	0	0	*	*
Microabscess	4 (0.2)	0	0	3 (0.6)	0	0	0	0	0	0	*	*
Inflammation	3 (0.6)	3 (1)	0	3 (0.6)	3 (0.8)	2 (1)	0	0	0	0	*	*
Edema	3.8 (0.8)	3 (0.8)	0	0	0	2 (1)	0	0	0	0	*	*
Infiltrate	3 (0.6)	3 (1)	0	3 (0.8)	3.8 (0.8)	0	0	0	0	0	*	*

<sup>a</sup> See Table 1, footnote a.<sup>b</sup> See Table 1, footnote b.<sup>c</sup> See Table 1, footnote c.<sup>d</sup> See Table 1, footnote d.<sup>e</sup> See Table 1, footnote e.

tive of a strong inflammatory response. This finding was in contrast to the pathologies observed in mice infected orally with the *rovA* mutant for 24 days. Mice infected orally for 24 days even with 10<sup>9</sup> CFU of the *rovA* mutant showed very little evidence of inflammatory pathologies (data not shown).

Not only were there significant inflammatory pathologies apparent in the tissues of the mice infected i.p. for 24 days with the *rovA* mutant, but there was evidence of disseminated bacterial infection. These findings range from minor findings such as splenomegaly and small inflammatory infiltrates to very severe findings such as fibrin thrombi, necrosis, and eschars (data not shown and Fig. 2 and 3). Taken together, the data presented here suggest that the *rovA* mutant is capable of causing severe disease in the mouse model of infection if the natural route of infection is bypassed.

## DISCUSSION

*Y. enterocolitica* has been an invaluable tool for understanding the intricate interactions that occur between a bacterial pathogen and the mammalian host. The reason that the system is so powerful is multifactorial: (i) the bacteria and the mouse are genetically tractable; (ii) the disease that is caused in the mouse is similar to that caused in humans; and (iii) the pathogen can be delivered to the host by the natural route of infection (orally). By using this system, significant progress has been made in the understanding of the host-pathogen interaction (for reviews, see references 7, 10, 11, and 25). We now have a greater understanding of bacterial factors important for the invasion of host tissues, as well as of the molecules responsible for the regulation of these factors. Examination of the mouse as a model host also has provided insight into the T-cell,

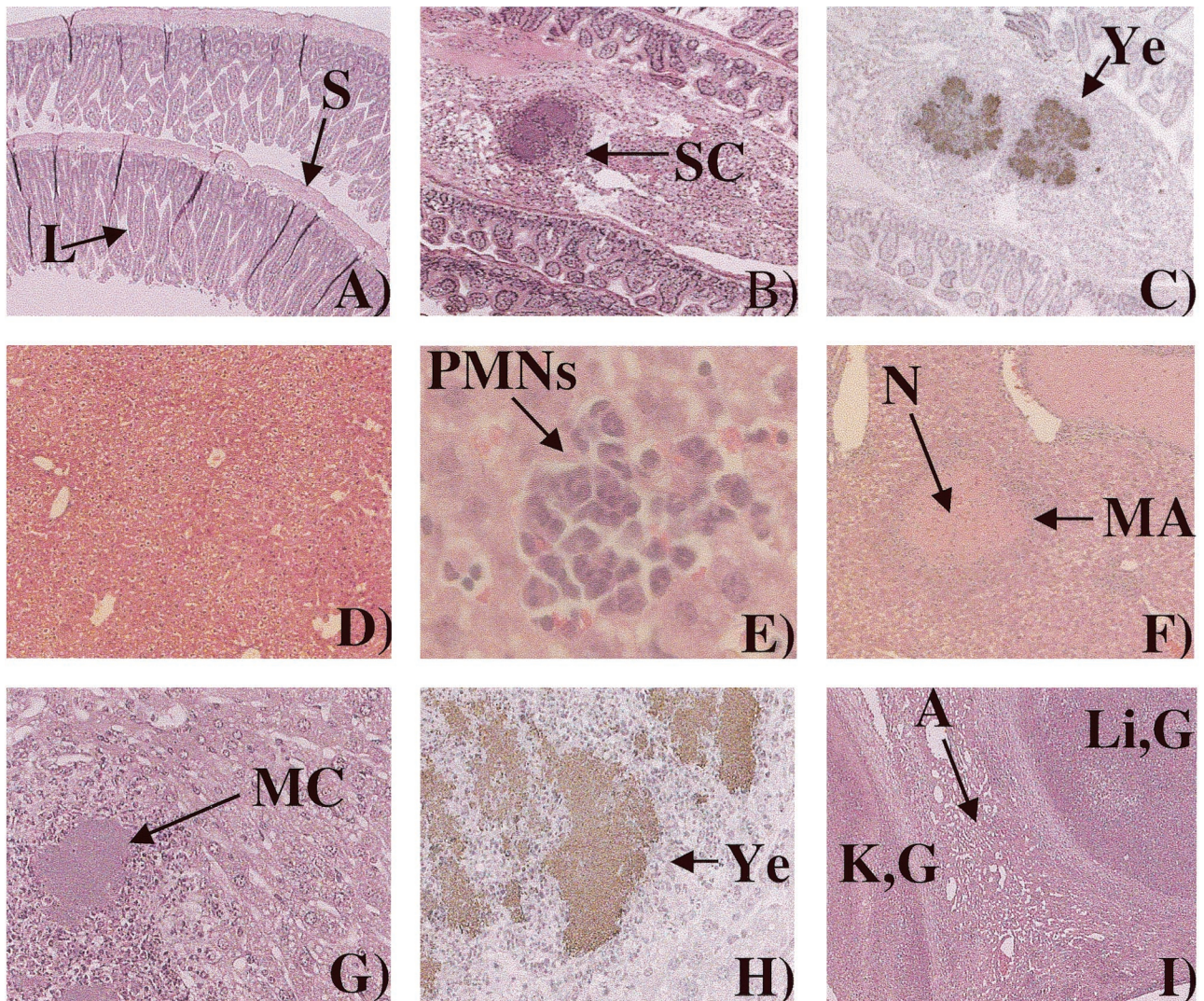


FIG. 2. Inflammatory histopathology of mice infected i.p. with the *rovA* mutant. (A) Normal mouse small intestine at  $\times 38$  magnification. "L" indicates the luminal surface, and "S" indicates the serosal surface. (B) Small intestine of a mouse infected for 5 days with  $10^3$  CFU of the *rovA* mutant at  $\times 38$  magnification. SC, serosal colony (a colony on the serosal surface of the small intestine). (C) Immunohistochemistry of the same tissue shown in panel B). Ye, positive *Y. enterocolitica* immunoreactivity. (D) Normal mouse liver at  $\times 38$  magnification. (E) Small infiltrate of the liver of a mouse infected i.p. for 14 days with  $10^2$  CFU of the *rovA* mutant at  $\times 380$  magnification. (F) Liver of a mouse infected i.p. for 14 days with  $10^2$  CFU of the *rovA* mutant at  $\times 190$  magnification. "N" indicates necrosis, and "MA" indicates microabscess (a large infiltrate without microscopic evidence of bacterial colonization). (G) Liver of a mouse infected i.p. for 24 days with  $10^2$  CFU of the *rovA* mutant at  $\times 190$  magnification. MC, microcolony (a large infiltrate with microscopic evidence of bacterial colonization). (H) Immunohistochemistry of the same tissue shown in panel G at  $\times 380$  magnification. Ye, positive *Y. enterocolitica* immunoreactivity. (I) Liver and kidney of a mouse infected for 24 days with  $10^3$  CFU of the *rovA* mutant at  $\times 38$  magnification. K,G, kidney granuloma; Li,G, liver granuloma; A, adhesion.

cytokine, and inflammatory responses to *Y. enterocolitica* infection (1, 3–5, 12).

Recently, we took advantage of this system to identify the regulator (*rovA*) of the principal invasion factor (*inv*) (14, 18, 24). Analysis of the *rovA* mutant in the mouse revealed an attenuated phenotype, as well as a defect in the ability to cause inflammation of the PP (12, 24). The defect in the ability to cause PP inflammation was linked to a defect in the ability of the host to induce IL-1 $\alpha$  upon infection with the *rovA* mutant (12, 24). The phenotypes observed during the initial characterization of the *rovA* mutant in vivo utilized the natural route of infection (oral). Because RovA was identified as a regulator

of *inv* and infection with the *rovA* mutant also affected the inflammatory responses in the PP (which are not linked to *inv*), it was reasonable to assume that RovA regulates gene products that are important for host-pathogen interactions that occur in the gut.

To further evaluate the role of *rovA* in the host-pathogen interaction, we bypassed the PP and delivered the bacteria directly into the peritoneum. Surprisingly, when mice are infected i.p. with the *rovA* mutant, the bacteria are significantly more virulent than when they are delivered by the oral route of infection. During an oral infection the *rovA* mutant is  $\sim 500$ -fold attenuated by LD<sub>50</sub> analysis, but during an i.p. infection



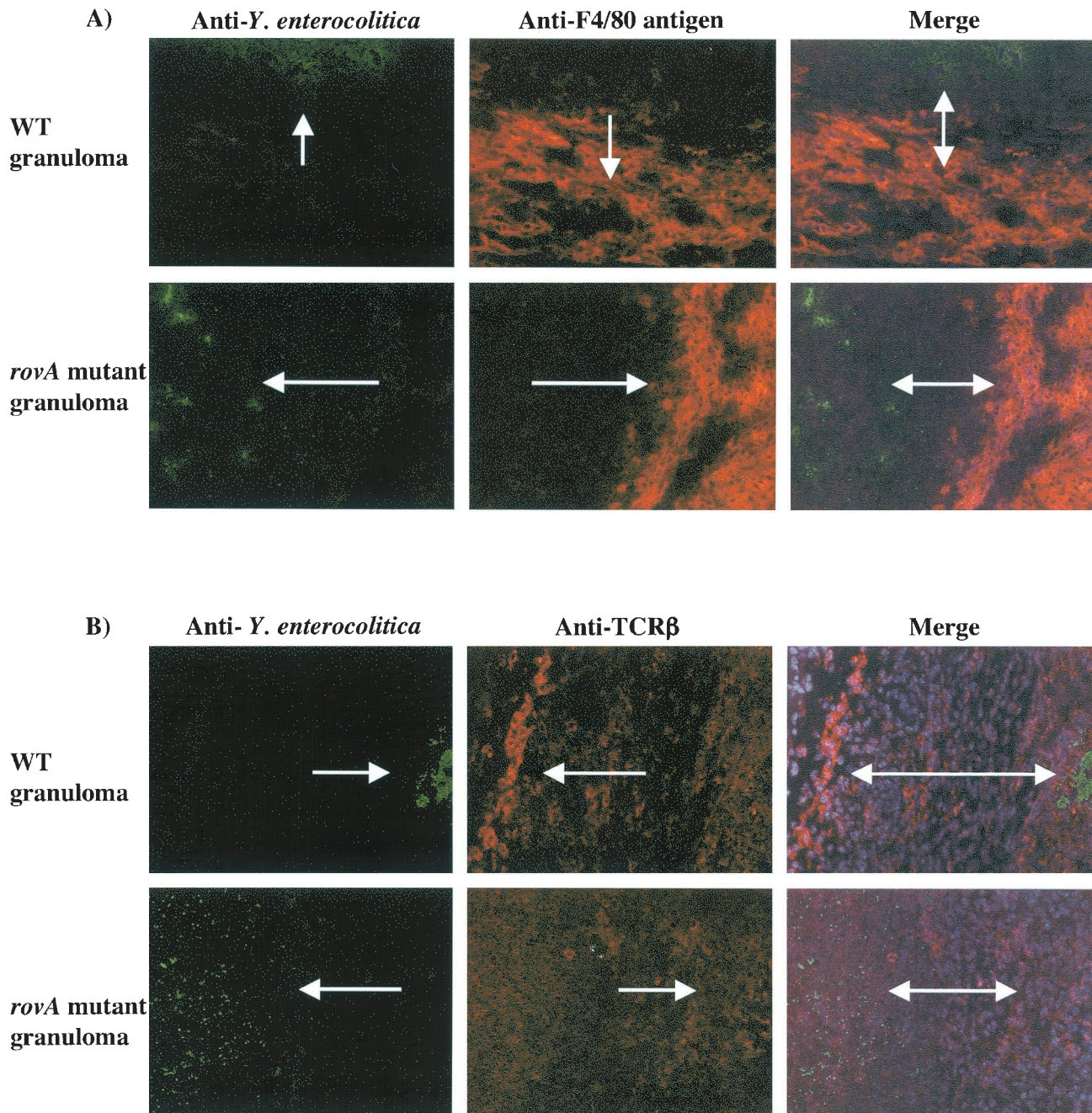


FIG. 3. i.p. infection with the *rovA* mutant induces an inflammatory response. C57BL/6j mice were infected i.p. for 14 days with  $10^3$  CFU of either the *rovA* mutant or WT *Y. enterocolitica*. Livers were removed and prepared for immunohistochemistry of fresh frozen sections. All images are either from the edge of the same large granuloma (A and B) or from the center of the granuloma (C) from the respective experimental group. (A) *Y. enterocolitica* immunoreactivity is in green, and the macrophage-specific F4/80 antigen is in red; merged images are counterstained with Hoechst dye. (B) *Y. enterocolitica* immunoreactivity is in green, and the  $\beta$  chain of the T-cell receptor is in red; merged images are counterstained with Hoechst dye. (C) *Y. enterocolitica* immunoreactivity is in green, and the granulocyte-specific GR-1 antigen is in red; merged images are not counterstained with Hoechst dye in this image to facilitate visualization of both markers. In all panels arrows indicate positive immunoreactivity. All images are a magnification of  $\times 190$ .

the *rovA* mutant is 11-fold attenuated. This suggests that *rovA* plays a critical role in the part of the bacterial life cycle that occurs in the PP. Consistent with the data obtained from the analysis of the kinetics of an oral *rovA* infection, there appears to be a slight delay in the ability of the *rovA* mutant to colonize

the tissues of the peritoneum efficiently. This may be due to a survival defect. The closely related *rovA* homolog (78% identical) from *Salmonella* (*slyA*) is sensitive to oxidative stress (8, 17). Although this has never been formally demonstrated for the *rovA* mutant of *Y. enterocolitica*, the delay in the coloniza-



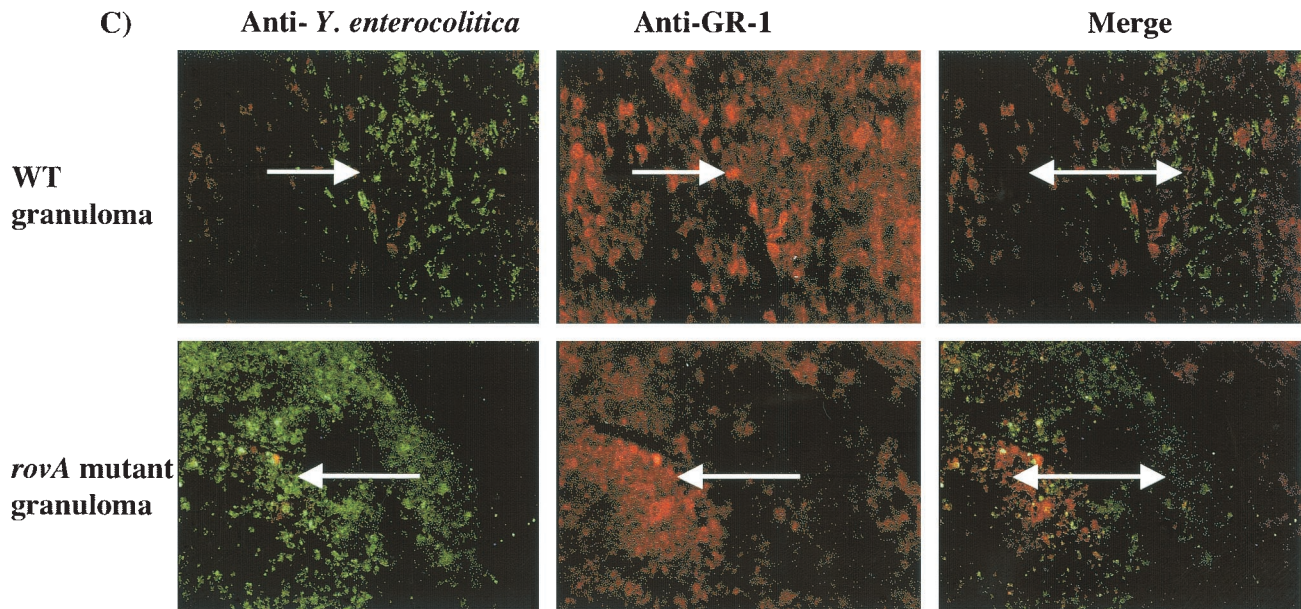


FIG. 3—Continued

tion of the tissues of the peritoneum during an i.p. infection, along with the attenuation phenotype observed during LD<sub>50</sub> analysis, is consistent with a partial survival defect.

The most striking defect of the *rovA* mutant during an oral infection is a lack of inflammation in the PP. The initial examination of this phenotype was limited to a 7-day time course. In the present study we extended the histological analysis of the oral *rovA* infection to 24 days. Consistent with our previous results there appears to be very little inflammation of the PP even after 14 or 24 days when mice are infected orally with the *rovA* mutant. Interestingly, mice infected with the WT bacteria are resolving the inflammatory pathologies of the gut at these time points but still have significant pathologies in other tissues. However, when mice are infected i.p. with the *rovA* mutant, inflammatory pathologies can be observed as early as day 1 and consistently by day 5. Consistent with the kinetic data there does appear to be a delay in the severity of the inflammatory pathologies observed in the *rovA* mutant-infected mice. At the 24-day time point very severe inflammatory pathologies can still be observed in the tissues from mice infected i.p. with the *rovA* mutant. The presence of granulomas and adhesions in the tissues of the mice infected i.p. with the *rovA* mutant for 24 days is indicative of a strong gamma interferon response. This finding suggests that when the intestine is bypassed, the *rovA* mutant is capable of initiating a strong inflammatory response. Furthermore, these data suggest that the host response in the gut and peritoneum may be different in significant ways that influence the outcome of the infection. Interestingly, when tissues from the day 24 *rovA* mutant and WT i.p. infections are stained with a *Y. enterocolitica* specific antibody, the bacteria are only apparent inside of the granulomas. This suggests that granuloma integrity may help protect the infected mice from the greater mortality and disease.

Altogether, the data presented in this report provides further evidence for the possible role of the virulence factor RovA during a *Y. enterocolitica* infection. One possible interpretation

of the data, although not the only one, is that RovA regulates several sets of genes. One set of genes is involved in the immediate-early stages of infection and is responsible for the virulence phenotypes that are observed in the PP. These genes would include *inv* and other genes that are responsible for immediate events postinvasion of the PP, as well as genes involved in the initiation of PP inflammation. A second set of genes would be a set of genes that are expressed constitutively. These genes may be responsible for helping with survival in the host and thus play a role in both oral and i.p. infections. An example would be genes responsible for protecting the bacteria from oxidative stress. Any survival defect *rovA* has would only be a partial defect because viable bacteria can be recovered from the host after both an oral infection and an i.p. infection. The differential virulence that the *rovA* mutant displays during an oral infection versus an i.p. infection has further illustrated the importance of RovA and the RovA-regulated genes during the natural course of a *Y. enterocolitica* infection.

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P.H.D. and S.A.H. contributed equally to this study.

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