

## Role of *rpoS* in Acid Resistance and Fecal Shedding of *Escherichia coli* O157:H7

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**Acid resistance (AR) is important to survival of *Escherichia coli* O157:H7 in acidic foods and may play a role during passage through the bovine host. In this study, we examined the role in AR of the *rpoS*-encoded global stress response regulator  $\sigma^S$  and its effect on shedding of *E. coli* O157:H7 in mice and calves. When assayed for each of the three AR systems identified in *E. coli*, an *rpoS* mutant (*rpoS*::pRR10) of *E. coli* O157:H7 lacked the glucose-repressed system and possessed reduced levels of both the arginine- and glutamate-dependent AR systems. After administration of the *rpoS* mutant and the wild-type strain (ATCC 43895) to ICR mice at doses ranging from  $10^1$  to  $10^4$  CFU, we found the wild-type strain in feces of mice given lower doses ( $10^2$  versus  $10^3$  CFU) and at a greater frequency (80% versus 13%) than the mutant strain. The reduction in passage of the *rpoS* mutant was due to decreased AR, as administration of the mutant in 0.05 M phosphate buffer facilitated passage and increased the frequency of recovery in feces from 27 to 67% at a dose of  $10^4$  CFU. Enumeration of *E. coli* O157:H7 in feces from calves inoculated with an equal mixture of the wild-type strain and the *rpoS* mutant demonstrated shedding of the mutant to be 10- to 100-fold lower than wild-type numbers. This difference in shedding between the wild-type strain and the *rpoS* mutant was statistically significant ( $P \leq 0.05$ ). Thus,  $\sigma^S$  appears to play a role in *E. coli* O157:H7 passage in mice and shedding from calves, possibly by inducing expression of the glucose-repressed RpoS-dependent AR determinant and thus increasing resistance to gastrointestinal stress. These findings may provide clues for future efforts aimed at reducing or eliminating this pathogen from cattle herds.**

*Escherichia coli* O157:H7 is a member of the enterohemorrhagic group of pathogenic *E. coli* (EHEC). This agent has emerged as a food-borne and waterborne pathogen of humans that causes hemorrhagic colitis, hemolytic-uremic syndrome, and thrombotic thrombocytopenic purpura. Outbreaks involving undercooked ground beef (1) and a variety of other foods including salami (7) and apple cider (3) have been documented.

Most human disease outbreaks caused by this organism have been linked to bovine fecal contamination of food or water. *E. coli* O157:H7 inhabits the intestinal tract of cattle, but it is unclear whether it actually colonizes the bovine intestinal tract (31). This organism seldom causes disease in cattle (10). Prevalence in individual beef and dairy cattle in the United States is low, from 0.3 to 2.2% (14, 20, 21). *E. coli* O157:H7 exists on about 7% of farms (14). Typically, shedding of serotype O157:H7 strains from cattle is sporadic and of limited duration, lasting for approximately 1 month (2, 31).

A number of animal models have been used to assess the pathogenesis and shedding of *E. coli* O157:H7 strains, including greyhounds (15), mice (33), neonatal calves (11), gnotobiotic pigs (16), and sheep (24). A modification of the streptomycin-treated mouse model described by Myhal and coworkers

(29, 30) has been used to compare the colonization of *E. coli* O157:H7 strain 933 with that of strain 933cu, which was cured of pO157 (33). Results from these animal models have been useful in elucidating the mechanisms of virulence and pathology.

Most *E. coli* O157:H7 strains contain the *eae* locus, which encodes the adhesin protein intimin (12). The *eae* gene also is found in the enteropathogenic group of *E. coli* (23). The recent finding that intimin is required for *E. coli* O157:H7 attachment in the bovine intestinal tract in neonatal calves has led one group of investigators to suggest that an anti-intimin vaccine might be effective for reducing the level of this pathogen in herds (12). In addition to intimin, *E. coli* O157:H7 also produces Shiga-like toxins, which may play a role in the lesions seen in the gut and kidneys of infected humans. *E. coli* O157:H7 is thought to have a low infectious dose ( $\leq 200$  organisms) in humans (19). As with other enteric pathogens, the infectious dose for *E. coli* O157:H7 is thought to correlate with acid resistance (AR) (18). Because *E. coli* O157:H7 survives in acidic foods and has a low infectious dose, we hypothesize that it can resist gastric acidity in cattle and that AR is another virulence mechanism of this pathogen.

Two recent studies have investigated the effect of diet on AR of *E. coli* shed from cattle. Diez-Gonzalez and coworkers (13) examined the effect of high-grain diets on shedding of acid-resistant *E. coli* from cattle. They found that the proportion of acid-resistant *E. coli* increased in cattle fed a high-grain diet, but that a switch to a diet of hay decreased the number of acid-resistant bacteria. It is unclear if the results observed for *E. coli* in these experiments are relevant to serotype O157

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TABLE 1. *E. coli* strains used in this study

Strain	Serotype or genotype <sup>a</sup>	Reference
ATCC 43895	O157:H7	34
EK274	ATCC 43895 Na <sup>r</sup> Rf <sup>r</sup>	This work
FRIK 816-3	ATCC 43895 <i>rpoS</i> ::pRR10 (Ap)	8
EK275	FRIK 816-3 Na <sup>r</sup> Rf <sup>r</sup>	This work
EF501	43895 <i>gadA</i> ::pRR10 (Ap) Na <sup>r</sup> Rf <sup>r</sup>	This work
EK272 (B2F1)	O91:H21	27

<sup>a</sup> Abbreviations for antibiotics: Na, nalidixic acid; Rf, rifampin; Ap, ampicillin.

strains of *E. coli*. Hovde et al. (22) examined the AR and the duration of shedding of *E. coli* O157:H7 from experimentally inoculated cattle fed hay or grain. They found no difference in the proportion of acid-resistant *E. coli* O157:H7 between these two groups but observed that animals fed hay shed *E. coli* O157:H7 longer than animals fed grain. Further studies are needed to clarify the role played by AR on *E. coli* O157:H7 shedding from cattle.

EHEC strains contain three distinct AR systems (26). The oxidative or glucose-repressed system (AR system 1) is active when the cells are growing aerobically or anaerobically in the absence of glucose. In contrast, the glutamate-dependent (AR system 2) and arginine-dependent (AR system 3) systems are active during fermentation. All three systems are active during stationary-phase growth, which suggests the involvement of the alternate, stationary-phase sigma factor  $\sigma^S$ , which is encoded by *rpoS*.  $\sigma^S$  is now known to play a protective role when *E. coli* O157:H7 is exposed to a variety of environmental stresses, including acid (8).

In the present study, the role of resistance to acid stress in *E. coli* O157:H7 gastrointestinal passage in mice and shedding in calves was examined. We tested the hypothesis that *rpoS* regulates one or more of the AR systems in *E. coli* O157:H7 and that a mutation in *rpoS* would affect the ability of the organism to survive in mice and calves. The results indicate that *rpoS* regulates especially the glucose-repressed AR system and is important to passage and shedding of *E. coli* O157:H7 in mice and calves, respectively, possibly by inducing resistance to gastrointestinal acid stress.

(Portions of this work have been presented elsewhere [J. Minter, S. C. Richardson, F. J. DeGraves, J. C. Wright, T. A. Penfound, J. W. Foster, and S. B. Price, Abstr. 98th Gen. Meet. Am. Soc. for Microbiol. 1998, abstr. B-169, p. 81, 1998].)

#### MATERIALS AND METHODS

**Bacterial strains.** The *E. coli* strains used in this study are listed in Table 1. *E. coli* O157:H7 strain ATCC 43895, originally isolated from raw hamburger meat implicated in a hemorrhagic colitis outbreak (strain 933 [34]), was used throughout the study. An *rpoS* mutant of this strain (FRIK 816-3 [8]) was constructed by insertional inactivation with pRR10, which carries the gene encoding  $\beta$ -lactamase. A *gadA* mutant of ATCC 43895 was constructed in a similar manner as described by Castanie-Cornet et al. (6). The location of the *gadA* insertion was confirmed using PCR and primers that amplified the correct fragment size only if the insertion had occurred in *gadA*. Spontaneous mutants of the *rpoS* and *gadA* strains resistant to nalidixic acid and rifampin were generated to aid in recovery from calf fecal specimens. Ampicillin (50  $\mu$ g/ml) was added to cultures of these mutants to maintain pRR10.

**AR assays.** Cells were grown overnight in one of several media: LBG (Luria-Bertani [LB] plus 0.4% glucose), BHIG (brain heart infusion [BHI] plus 0.4% glucose), LB or BHI buffered with either 100 mM morpholinepropanesulfonic acid [pH 8] or 100 mM morpholineethanesulfonic acid [pH 5.5] [28]), and minimal E glucose (EG [32]). Cultures were grown in 3 ml of the appropriate medium in 13-mm test tubes with shaking (240 rpm) at 37°C to stationary phase (22 h). The glucose-repressed system was tested using cells grown overnight in pH 5.5 buffered LB or BHI followed by 1:1,000 dilution into prewarmed (37°C) pH 2.5 EG. The glutamate and arginine systems were tested using stationary-phase cells grown in LBG followed by 1:1,000 dilution into prewarmed pH 2.5 EG supplemented with 1.5 mM glutamate or 0.6 mM arginine, respectively.

Viable-cell counts were determined at 0, 2, and 4 h post-acid challenge by diluting cells in LB, plating cells onto LB agar, and incubating plates for 20 h at 37°C. Values given are representative of the results of triplicate experiments reproducible to within 50%.

**Mouse inoculation studies.** A modified version of a previously described procedure (33) was used for oral administration of *E. coli* O157:H7 strains 43895 (RpoS<sup>+</sup>) and FRIK 816-3 (RpoS<sup>-</sup>). Bacteria were grown in Trypticase soy broth (BBL Microbiology Systems, Cockeysville, Md.) at 37°C with shaking (150 rpm) to stationary phase. The cells were harvested by centrifugation (10,000  $\times$  g, 10 min) and diluted to the appropriate concentration in a 10% sucrose solution or 0.05 M phosphate buffer (7.2 g of Na<sub>2</sub>HPO<sub>4</sub>, 1.2 g of KH<sub>2</sub>PO<sub>4</sub> [pH 7.4]) containing 10% (wt/vol) sucrose. ICR mice (ca. 20 g; Harlan Sprague Dawley Inc., Madison, Wis.) were housed individually in cages with grated floors so that feces could be collected from sterile liners on trays below the cages. The mice were deprived of feed for 24 h preinoculation and then provided a sterile plate containing the sucrose cell suspension (0.5 ml) with numbers of *E. coli* O157:H7 cells ranging from 0 cells (control) to 10<sup>4</sup> CFU. Some mice received strain FRIK 816-3 in 10% sucrose containing 0.05 M phosphate buffer to ensure that susceptibility to acid and not another factor was the cause for reduction in gastrointestinal passage in mice. The mice were observed to make sure that they consumed the entire inoculum. To allow for clearance of the inoculum through the stomach and to avoid any protection to gastric acidity, the mice were not provided feed for 4 h after administration of the inoculum.

For each trial, five mice were administered the specified strain and dosage of *E. coli* O157:H7. Three trials were conducted. Fecal samples were collected on the 3 consecutive days after inoculation and tested for the presence or absence of *E. coli* O157:H7 cells. A 1:10 suspension was made (0.5 g of feces/4.5 ml of peptone [0.1%]) and mixed thoroughly using a vortex mixer. A 0.5-ml portion of this suspension was then spread plated onto five plates (0.1 ml/plate) of MacConkey sorbitol agar (MSA; Difco, Detroit, Mich.) supplemented with potassium tellurite (2.5  $\mu$ g/ml; Sigma Chemical Co., St. Louis, Mo.) and cefixime (0.05  $\mu$ g/ml; Lederle Laboratories, Pearl River, N.Y.) (MSA+ [35]). The plates were incubated at 42°C for 24 h and examined for sorbitol-negative *E. coli* O157:H7 colonies. Suspect colonies were confirmed as *E. coli* O157 by agglutination (RIM *E. coli* O157:H7; Remel, Inc., Lenexa, Kans.). The minimum detection level of this procedure was approximately 20 CFU/g. Mice with a positive fecal sample on any of the 3 days postinoculation were scored positive, meaning that the O157:H7 strain had survived passage through the gastrointestinal tract at the dose used.

**Calf inoculation studies.** Weaned 6- to 8-week-old calves were acclimated outdoors for 1 week and then moved to an indoor, climate-controlled BL-2 containment facility, where they were acclimated for an additional week prior to inoculation. Pairs of calves were housed together on pine bedding and fed grain and hay in the morning and evening, with water provided ad libitum. Outdoor and indoor pens were cleaned twice daily. Each calf was cultured for *E. coli* O157:H7 three times before inoculation to ensure that any calves shedding wild strains of the organism were excluded from the study. At the completion of each experiment, the calves were euthanized with sodium pentobarbital and incinerated. Housing and care of the calves followed the guidelines of the American Association for Laboratory Animal Care.

Bacterial strains for inoculation into calves were grown in 12.5 ml of BHI broth (pH 5.5) to stationary phase. Cell pellets were harvested by centrifugation, washed, and suspended in 0.85% NaCl. Calves were inoculated by gastric lavage with a 50-ml inoculum of 0.85% NaCl containing 10<sup>10</sup> CFU of an equal mixture of EK274 (RpoS<sup>+</sup>) and EK275 (RpoS<sup>-</sup>), followed by 500 ml of 0.85% NaCl. In a control experiment designed to confirm that pRR10 had no effect on shedding, a mixture of equal amounts of EK274 and EF501 (*gadA*) was inoculated into four calves.

Following inoculation, calf fecal samples were cultured daily for 16 days for the presence of *E. coli* O157:H7. Fifty-gram specimens were collected each morning and immediately transported back to the laboratory for culture. Quantitative culture of the specimens was performed by adding 1 g of feces to 9 ml of phosphate buffer followed by serial 10-fold dilution. A 0.1-ml volume of each dilution was plated in duplicate onto MSA plates containing nalidixic acid (35  $\mu$ g/ml), which selected for EK274 and EK275 (or EF501), and MSA plates containing ampicillin (50  $\mu$ g/ml), which selected for EK275 or EF501. Following 16 to 20 h of incubation at 37°C, sorbitol-negative colonies were counted manually. To determine the quantity of EK274 present in specimens, the number of ampicillin-resistant colonies was subtracted from the total number of nalidixic acid-resistant colonies.

To detect the organism in feces containing <10<sup>3</sup> CFU/g, fecal swabs were enriched in BHI containing novobiocin (20  $\mu$ g/ml), potassium tellurite (2.5  $\mu$ g/ml), and rifampin (25  $\mu$ g/ml). Following overnight incubation, the BHI cultures were streaked onto Rainbow Agar O157 (Biolog, Inc., Hercules, Calif.) containing ampicillin or nalidixic acid. This medium is an *E. coli* O157-selective and differential medium on which colonies of this pathogen turn grey-black. Using this enrichment technique, the presence in fecal samples of as few as three organisms per fecal swab could be detected. Serologic confirmation of *E. coli* O157:H7 suspect colonies in both the direct plating and enrichment culture protocols was made using a commercial latex agglutination kit (Remel).

**Statistical analysis.** Data from the mouse experiment were analyzed by analysis of variance using the SAS statistical analysis system (SAS Institute, Inc.,

TABLE 2. Effect of *rpoS* on each of the three AR systems in *E. coli* O157:H7

Adaptation medium <sup>a</sup>	Challenge medium (pH 2.5)	% Survival <sup>b</sup>		
		EK274 ( <i>rpoS</i> <sup>+</sup> )	EK275 ( <i>rpoS</i> )	EF501 ( <i>gadA</i> )
LB				
pH 8	EG	<0.005	<0.006	<0.004
pH 5.5	EG	13.6	<0.005	13
LBG	EG	<0.004	<0.007	<0.005
	EG + Glu	35	0.4	75
	EG + Arg	<0.004	<0.007	ND
BHI				
pH 8	EG	<0.004	<0.004	<0.003
pH 5.5	EG	4.2	0.09	5.8
BHIG	EG	<0.004	<0.006	<0.004
	EG + Glu	72	23	60
	EG + Arg	2	1.3	10

<sup>a</sup> Adaptation involved overnight growth in the medium indicated.

<sup>b</sup> Measured by determining the number of surviving cells after 4 h of acid treatment. Experiments were conducted in triplicate, and results varied within 50% of the stated value. ND, not determined.

Cary, N.C.). Fecal shedding data from the calf study were entered into a spreadsheet program (Microsoft Excel 97 for Windows, version 4.0) and analyzed using SAS. Daily values (CFU per gram of feces) for each strain were converted to log<sub>10</sub> for analysis. Total CFU per gram shed over days 1 to 7 was calculated using area under the curve following the trapezoidal rule (17). EK275 (*rpoS*) shedding as a percentage of EK274 shedding was calculated by dividing area under the curve results from EK275 shedding by area under the curve EK274 shedding and multiplying the result by 100. EF501 (*gadA*) shedding as a percentage of EK274 shedding was calculated in a similar manner. Differences in percent shedding between EK275 and EF501 were analyzed using Student's *t* test (36).

## RESULTS

**Effect of *rpoS* on acid resistance of O157.** The *E. coli* O157:H7 parent (EK274) and *rpoS* mutant (EK275) were tested for the AR systems previously identified (25). The data presented in Table 2 show that insertional inactivation of *rpoS* eliminated the glucose-repressed system (AR system 1) and reduced the arginine- and glutamate-dependent systems to various levels when cells were adapted in LB (pH 5.5) and LBG, respectively. Previous results obtained with an *E. coli* K12 *rpoS* mutant were similar to the results reported here with the serotype O157:H7 *rpoS* mutant, EK275 (26). Although an *rpoS* mutation caused decreased glutamate- and arginine-dependent AR, these systems do not have an absolute requirement for RpoS. This was evident when adaptation was made in BHIG rather than LBG. After growth in BHIG, EK275 exhibited near-normal levels of glutamate- and arginine-dependent AR (Table 2). Yet, when the *rpoS* mutant was adapted in BHI (pH 5.5), it remained defective in glucose-repressed AR. Clearly, *rpoS* inactivation selectively prevents the induction of AR system 1, showing that this glucose-repressed system is also RpoS dependent. The effect of adaptation in BHI on AR was also observed in another EHEC strain (O91:H21) originally thought to lack acid resistance (27). Strain O91:H21 exhibited no AR when adaptations were performed in LB (Table 3). However, when adaptations were performed using BHI medium, this strain exhibited significant levels of all three AR systems (Table 3).

**Passage of *E. coli* O157:H7 RpoS<sup>+</sup> and RpoS<sup>-</sup> strains in mice.** Oral administration of *E. coli* O157:H7 strains ATCC 43895 (RpoS<sup>+</sup> wild-type strain) and FRIK 816-3 (RpoS<sup>-</sup>) did not result in visible signs of disease in ICR mice. Feed intake remained constant (average, 8.3 ± 0.5 g/day), and mice produced a daily average of 2.2 g of feces, of which 0.5 g was tested

TABLE 3. AR of *E. coli* O91:H21 strain

Adaptation medium <sup>a</sup>	Challenge medium (pH 2.5)	% Survival <sup>b</sup>
LB, pH 5.5	EG	<0.005
BHI, pH 5.5	EG	9
LBG	EG	<0.006
	EG + Glu	<0.06
	EG + Arg	<0.01
BHIG	EG	<0.004
	EG + Glu	29
	EG + Arg	6

<sup>a</sup> Adaptation involved overnight growth in the medium indicated.

<sup>b</sup> Measured by determining the number of surviving cells after 4 h of acid treatment. Experiments were conducted in triplicate, and results varied within 50% of the stated value.

for the presence of the *E. coli* O157:H7 strains. The plating of fecal samples from mice (control and preinoculation) on MSA+ resulted in low numbers of background bacteria that were similar to colonies of serotype O157:H7 strains (i.e., sorbitol negative).

Of the 405 fecal samples cultured from inoculated mice, 38 samples (9%) tested positive for *E. coli* O157:H7 at day 1 after dosage, and two of these mice were still fecal positive at day 2. No fecal samples tested positive at day 3 after dosage. The results from the mouse inoculation studies are shown in Table 4. Control mice which received the sucrose or sucrose-phosphate buffer solution tested negative for *E. coli* O157:H7. Strain 43895 was recovered from 5 of 15 mice (33%; total from three trials) which received 100 CFU, while 12 of 15 mice (80%) administered 1,000 CFU tested positive. None of the

TABLE 4. Recovery of *E. coli* O157:H7 strains ATCC 43895 and FRIK 816-3 from feces following oral administration to ICR mice<sup>a</sup>

<i>E. coli</i> dose (CFU)	No. of mice with ≥1 fecal sample positive for <i>E. coli</i> O157:H7/5 mice examined in trial (no. of mice with ≥1 fecal sample positive for <i>E. coli</i> O157:H7/15 mice examined in 3 trials)						
	Trial with 10% sucrose			Trial with 0.05 M PB-10% sucrose			
	I	II	III	I	II	III	
Control	0	0	0 (0)	0	0	0 (0)	
Strain 43895	10 <sup>1</sup>	0	0	0 (0)	ND	ND	ND ND
	10 <sup>2</sup>	0	3	2 (5)	ND	ND	ND ND
	10 <sup>3</sup>	5	4	3 (12)	ND	ND	ND ND
Strain 816-3	10 <sup>2</sup>	0	0	0 (0) <sup>b</sup>	0	0	0 0
	10 <sup>3</sup>	1	0	1 (2) <sup>c</sup>	2	2	1 (5)
	10 <sup>4</sup>	1	2	1 (4) <sup>d</sup>	3	4	3 (10)

<sup>a</sup> Cell pellets (strain ATCC 43895 or FRIK 816-3) were suspended and diluted to the appropriate concentration in 10% sucrose or 0.05 M phosphate buffer (PB) in 10% sucrose. Each mouse was administered 0.5 ml of the sucrose-cell suspension or sucrose-phosphate-cell suspension. Control mice were fed 0.5 ml of 10% sucrose or 0.05 M phosphate buffer containing 10% sucrose with no *E. coli* O157:H7. Results are from fecal samples collected 24 h after administration. ND, not determined.

<sup>b</sup> Significantly different ( $P < 0.05$ ) from results obtained with the parent strain (ATCC 43895) inoculated at a dose of 10<sup>2</sup> CFU.

<sup>c</sup> Significantly different ( $P < 0.001$ ) from results obtained with the parent strain (ATCC 43895) inoculated at a dose of 10<sup>3</sup> CFU.

<sup>d</sup> Significantly different ( $P < 0.05$ ) from results obtained with strain 816-3 inoculated at a dose of 10<sup>4</sup> CFU in phosphate buffer.

mice given 10 CFU tested positive. In comparison, the *rpoS* mutant strain FRIK 816-3 was not detected in mice receiving 100 CFU and was found in only 2 of 15 (13%) and 4 of 15 (27%) mice administered 1,000 CFU and 10,000 CFU, respectively. Results from inoculation of 100 CFU and 1,000 CFU of strain 816-3 were significantly different ( $P < 0.05$  and  $P < 0.001$ , respectively) from the results obtained with the parent strain 43895 administered at the same dose (Table 4). These results indicate that there was a reduction in the ability of strain 816-3 to survive passage through the gastrointestinal tract of mice in comparison to strain 43895.

To determine if acid susceptibility rather than another factor was responsible for the reduced passage noted with strain 816-3, cells were suspended in 10% sucrose containing phosphate buffer. Suspension of strain 816-3 in phosphate buffer significantly increased ( $P < 0.05$ ) the number of mice shedding at a dose of 10,000 CFU (Table 4). At a dose of 1,000 CFU, suspension of 816-3 in phosphate buffer increased the number of mice shedding the pathogen from 2 of 15 to 5 of 15, but this increase was not statistically significant. Although there was a significant increase in passage when strain 816-3 was administered in phosphate buffer with sucrose in comparison to sucrose alone, the wild-type strain 43895 was detected at similar frequencies following administration of 10-fold-fewer cells (e.g., 100 and 1,000 CFU). Suspension and administration of strain 816-3 in bicarbonate buffer as described previously (5) was detrimental to its survival (data not shown).

**Shedding of *E. coli* O157:H7 *RpoS*<sup>+</sup> and *RpoS*<sup>-</sup> strains in calves.** A quantitative approach was taken to examine the effect of *rpoS* on *E. coli* O157:H7 shedding patterns from experimentally inoculated calves. In these experiments, equal amounts of the wild-type (EK274; *RpoS*<sup>+</sup>) and *RpoS*<sup>-</sup> (EK275) strains were inoculated into three calves. The animals remained healthy throughout the experiment. As has been observed previously with inoculation of *E. coli* O157:H7 into calves, there was considerable variation of shedding among the calves (10). The reason for this variation is unknown.

In each calf, the wild-type (*RpoS*<sup>+</sup>) strain was shed in large amounts within 24 h from all three calves, with a minimum of  $2.7 \times 10^5$  to a maximum of  $2.6 \times 10^7$  CFU/g of feces (Fig. 1). Shedding of  $>10^3$  CFU of this strain per g of feces continued for a variable amount of time (from 6 to 16 days) in each calf. Generally, the number of organisms shed decreased over time (Fig. 1).

The *RpoS*<sup>-</sup> mutant strain was shed in significantly decreased amounts from all three calves compared to wild-type shedding ( $P \leq 0.05$ ). Although shedding of the *RpoS*<sup>-</sup> strain dropped below  $10^3$  CFU/g 8 days earlier than did the *RpoS*<sup>+</sup> strain (Fig. 1), the mutant could be detected in feces from experimentally infected calves for 15 to 39 days, comparable to the length of parent strain shedding, which ranged from 17 to 43 days (data not shown).

To confirm that shedding differences observed were due to the inactivation of *rpoS* and not the plasmid (pRR10) insert, a mutant containing pRR10 inserted into the gene for glutamate decarboxylase (*gadA*) was constructed. The *gadA* mutant (EF501) was fully acid resistant (Table 2). The difference in shedding in four calves between the parent and the *gadA* mutant was not significantly different ( $P > 0.05$ ; Fig. 1), indicating that pRR10 itself had no effect on shedding. This finding also illustrates the consistency of results when one is comparing two strains with equal ability to survive passage through the bovine gastrointestinal tract.

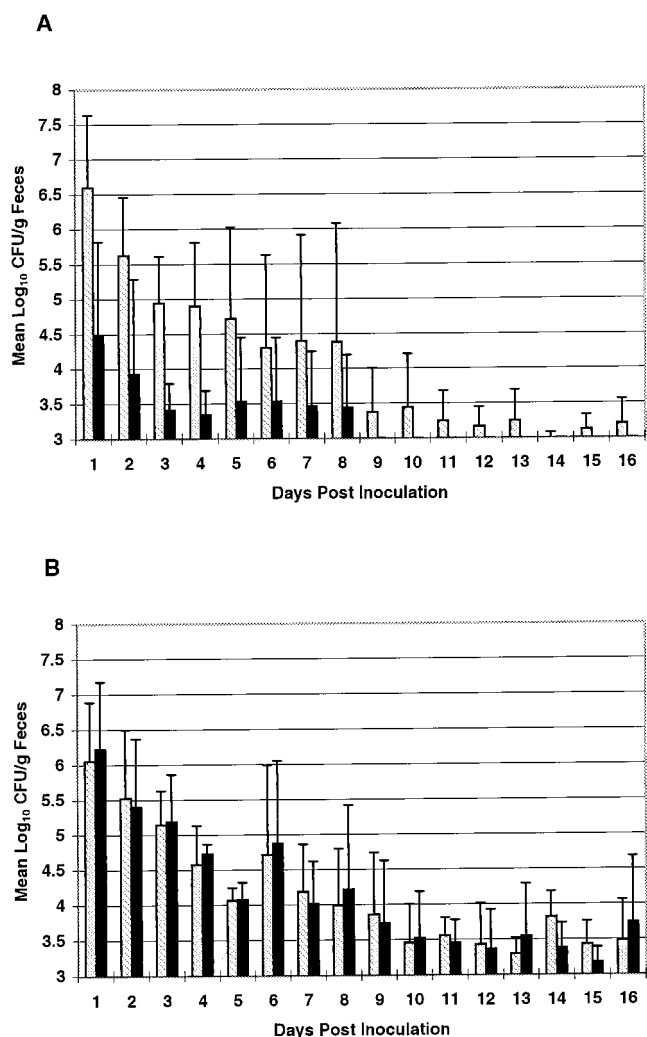


FIG. 1. Shedding levels over time of *E. coli* O157:H7 wild-type (□) and *RpoS*<sup>-</sup> (■) (A) or wild-type (□) and *GadA*<sup>-</sup> (■) (B) strains. Calves were inoculated with  $10^{10}$  to  $10^{11}$  total CFU containing equal numbers of wild-type and mutant strains on day 0. Fecal samples were cultured daily beginning 1 day postinoculation. Colony counts are displayed as the means for three (A) or four (B) calves. Bars indicate standard deviation of shedding among calves. Specimens containing  $<10^3$  CFU/g of feces were below the level needed for accurate enumeration ( $<10$  CFU/ $10^{-1}$  dilution plate).

## DISCUSSION

In the present study, the role of *rpoS* in AR and the subsequent impact on shedding of *E. coli* O157:H7 from mice and calves were examined. Our working hypothesis is that AR in *E. coli* O157:H7 is induced during passage and growth of the organism in the intestinal tract of cattle (26). This hypothesis has been supported by recent data from Hovde et al. (22), who showed that acid-resistant *E. coli* O157:H7 was shed from cattle. We have shown previously that once induced, all three AR systems (*rpoS* dependent, glutamate dependent, and arginine dependent) will persist for at least a month at 4°C (26). Combined, the published data indicate that AR can be induced in the intestinal tract, and growth of acid-resistant *E. coli* O157:H7 on a contaminated beef carcass is unnecessary for persistence of the phenotype.

The *in vitro* AR studies reported here extend the findings of a previous study by Chevillon et al. (8), which described the

construction and characterization of acid tolerance in an *rpoS* mutant of *E. coli* O157:H7 (FRIK 816-3). A marked difference in the glucose-repressed AR system between the wild-type and RpoS mutant was observed, confirming a role for *rpoS* in induction of this system (Table 2). Interestingly, an *rpoS* effect on the arginine- and glutamate-dependent AR systems in LBG was not found when the wild-type and RpoS mutant strains were acid challenged in BHIG. This finding was not limited to *E. coli* O157:H7 EHEC, as another EHEC strain, O91:H2, also showed differences in AR in LBG and BHIG.

ICR mice were used in studies to evaluate the influence of the *rpoS* regulon and acid tolerance on gastrointestinal passage in mice. Mice were used as one of the animals for testing our hypothesis due to the ease of handling and the increased size of groups which this species affords. However, the streptomycin-treated mouse model used by others (29, 33) for colonization and pathogenicity studies was not used because this study concerned passage through the gastric barrier and not colonization. In fact, colonization and subsequent replication would have made results more difficult to interpret. Positive fecal samples from mice were detected at day 1 after inoculation, and only two mice remained fecal positive at day 2. No mice tested positive 3 days after inoculation, which indicates that there was no colonization of mice by either *E. coli* O157:H7 strain (ATCC 43895 or FRIK 816-3).

Results from inoculation of mice with levels of wild-type strain ATCC 43895 or FRIK 816-3 ranging from  $10^1$  to  $10^4$  CFU demonstrated that a functional *rpoS* system promoted survival during gastrointestinal passage in mice. Inoculation of lower numbers of strain 43895 than of FRIK 816-3 resulted in positive fecal samples. Also, the frequency of positive fecal samples was greater in mice inoculated with 43895 when the same number of wild-type and mutant strain was administered (Table 4). For example, 12 of 15 mice (80%) had a positive fecal sample when administered  $10^3$  CFU of the wild-type strain, whereas, 2 of 15 (13%) mice given  $10^3$  CFU of the *rpoS* mutant strain FRIK 816-3 tested positive. To ensure that the reduced recovery of FRIK 816-3 was due to acid sensitivity and not another factor detrimental to passage, the mutant strain was suspended in 0.05 M phosphate buffer with 10% sucrose and used to dose mice. Suspension of FRIK 816-3 in phosphate buffer at doses of  $10^3$  and  $10^4$  CFU increased the number of positive fecal samples from 2 of 15 (13%) to 5 of 15 (33%) and from 4 of 15 (27%) to 10 of 15 (67%), respectively (Table 4). Early studies defining the infectious dose of *Vibrio cholerae* also demonstrated the importance of the gastric barrier, as suspension of *V. cholerae* in bicarbonate buffer reduced the infectious dose in humans from  $10^8$  to  $10^4$  organisms (5). Our study of *E. coli* O157:H7 used phosphate buffer instead of bicarbonate buffer because the latter reduced viable numbers of FRIK 816-3 (data not shown). The data from this study indicate that protection from acid afforded by *rpoS*-regulated systems promotes gastric passage.

A calf model of *E. coli* O157:H7 shedding was used to examine the effect of the *rpoS* system on the survival and shedding from the gastrointestinal tract of cattle. As in other studies using a calf model (4, 9, 10), an inoculum of  $10^{10}$  CFU of the organism resulted in appreciable shedding during the first week postinoculation. When the RpoS<sup>+</sup> and RpoS<sup>-</sup> strains were simultaneously administered to calves, we were able to compare numbers of each strain shed concurrently.

The RpoS<sup>-</sup> mutant (EK275) was reproducibly shed in lower numbers from the calves than was its RpoS<sup>+</sup> parent (EK274) (Fig. 1), and this difference was significant ( $P < 0.05$ ). This finding indicates that *rpoS* plays a role in *E. coli* O157:H7 shedding in calves, possibly by inducing resistance to gastroin-

testinal stress, including acid stress offset by the glucose-repressed RpoS-dependent AR system.

The observation that strains of *E. coli* O157:H7 were shed in detectable numbers for several weeks reflects the fact that these calves were inoculated with high numbers of *E. coli* O157:H7 (ca.  $10^{10}$  CFU) in order to achieve extended periods of shedding at appropriate numbers for detection. In the natural setting, cattle most likely ingest fewer organisms. Indeed, field observations indicate that exposure of cattle does not result in shedding of high numbers of *E. coli* O157:H7 strains for extended periods of time (31). What was unexpected about these results was the finding that both the parent and *rpoS* strains were shed for essentially the same length of time (data not shown). This observation may indicate that *E. coli* O157:H7 survivors that reach and colonize the lower intestinal tract, where the pH is neutral to alkaline, no longer require AR for survival.

The long-term goal of this study is to further define the role of AR systems in gastrointestinal passage and to develop intervention strategies that may inactivate one or more of these systems and possibly reduce or eliminate this pathogen from the bovine intestinal tract. Such an approach might be complementary to other strategies such as vaccination of herds with intimin, as suggested by Dean-Nystrom et al. (12), or manipulation of feed content (13, 22) as part of a complete program to exclude *E. coli* O157:H7 from cattle by reducing its ability to compete or survive in this complex digestive system. In the present work, a global regulator of acid and other stress resistance was shown to influence *E. coli* O157:H7 shedding from mice and calves. We are presently continuing experiments aimed at dissecting the role of each AR determinant in *E. coli* O157:H7.

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