# Recombinant PorA, the Major Outer Membrane Protein of *Campylobacter jejuni*, Provides Heterologous Protection in an Adult Mouse Intestinal Colonization Model<sup>⊽</sup>

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Immunity against *Campylobacter jejuni*, a major food-borne pathogen causing diarrhea, is largely serotype specific. The major outer membrane protein (MOMP) of *C. jejuni*, PorA, is a common antigen with the potential to provide broad protection. Adult BALB/c mice were orally immunized with a recombinant glutathione *S*-transferase (GST) fused to PorA prepared from *Campylobacter jejuni* C31 (O:6,7) (GST-PorA) combined with a modified heat-labile enterotoxin of *Escherichia coli* as an adjuvant and later orally challenged with C31 strain or three heterologous strains: 48 (O:19), 75 (O:3), and 111 (O:1,44). Protection from colonization with the challenge organism was studied by fecal screening daily for 9 days. Serum and intestinal lavage fluid antibodies against the vaccine and Sarkosyl-purified MOMP from C31 were measured by using an enzyme-linked immunosorbent assay. The vaccine produced robust antibody responses against both antigens in serum and secretion. Since strain C31 was a poor colonizer, homologous protection could not be studied. The protective efficacies of heterologous strains were 43% (for strain 48, P < 0.001), 29% (for strain 75, P < 0.005), and 42% (for strain 111, P < 0.001) for the 9-day period compared to control mice given phosphate-buffered saline. Thus, PorA provided appreciable protection against colonization with heterologous serotypes.

Campylobacter jejuni is a food-borne pathogen and a leading cause of diarrhea worldwide (19, 33, 34). Campylobacteriosis is associated with a number of important sequelae, including Guillain-Barre syndrome, Reiter's syndrome, reactive arthritis, and irritable bowel syndrome (10, 37). C. jejuni also contributes to significant mortality of children in developing countries (18). In view of significant morbidity, mortality, and economic burden associated with C. jejuni infection, the control and prevention of campylobacteriosis are a priority. One possible tool for the prevention of campylobacteriosis is vaccination. It has been observed that C. jejuni infection results in the acquisition of immunity. However, immunity in humans seems to be serotype (Penner serotype) specific (13, 36). There are numerous serotypes of C. jejuni according to the Penner serotyping scheme, which is based on the lipopolysaccharide capsule (25). Therefore, vaccines based on common antigens that are shared by all serotypes seem attractive for broad protection. One such antigen is the major outer membrane protein (MOMP) of C. jejuni, PorA, which is present in abundant quantity in the organism. There are three conformational forms of MOMP: folded monomer (35 kDa), denatured monomer (45 kDa), and the native trimer (120 to 140 kDa). Only the folded monomer and the native trimer have pore-forming activities (44). The MOMP is involved in ion transport across the bacterial cell wall and adhesion of the bacterium to the intestinal mucosa (30, 41). It consists of 18  $\beta$ -strands connected by short periplasmic turns and nine external loops. The external loops are antigenically variable (44). This is evident in the extreme genetic diversity of the porA gene encoding the MOMP (17, 44).

\* Corresponding author. Mailing address: Department of Microbiology, Faculty of Medicine, Kuwait University, P.O. Box 24923, Safat 13110, Kuwait. Phone and fax: (965) 2533 2719. E-mail: john@hsc.edu.kw. However, the  $\beta$ -strands representing the conserved regions have common antigenic epitopes for different strains (16, 44). The sera of patients recovering from *C. jejuni* infection (8, 12, 35), as well as those of animals immunized with *C. jejuni* (21), possess antibodies to MOMP. Therefore, PorA seems to be an appropriate antigen that could be investigated as a possible subunit vaccine.

There are no easily handled, widely available, and inexpensive natural animal models of *C. jejuni* diarrhea. Laboratory mice are widely available, inexpensive, and easy to handle. Oral feeding of the adult mouse with *C. jejuni* does not result in diarrhea in the animal. However, the organism colonizes the intestine, with shedding of the organism for several days in the stool. The organism can be isolated from the blood as well (7, 9). Thus, the adult mouse can be used as a colonization model of *C. jejuni* infection.

Since the PorA of *C. jejuni* has never been investigated as a vaccine candidate, we studied a recombinant PorA in the form of a fusion protein in the adult mouse colonization model for broad protection against colonization. We studied fecal excretion of the challenge organisms and serum and intestinal antibody responses to the vaccine.

## MATERIALS AND METHODS

Bacterial strains and growth conditions. The *C. jejuni* C31 strain, isolated from a patient with bloody diarrhea in the United States (20), was donated by Richard L. Guerrant, University of Virginia School of Medicine, Charlottesville. The other three *C. jejuni* strains—48, 75, and 111—were isolated from diarrheal stools of patients treated at the Mubarak Al-Kabir Hospital, Kuwait, during 2000 to 2005. The organisms were grown on campylobacter-selective agar consisting of agar base (Oxoid, Basingstoke, Hampshire, England) supplemented with 7% defibrinated horse blood SR0050 (Oxoid), growth supplement SR0232 (Oxoid), and selective supplement SR0098E (Oxoid). The culture was incubated at 42°C for 48 h in an atmosphere of 85% N<sub>2</sub>, 10% CO<sub>2</sub>, and 5% O<sub>2</sub> in a water-jacketed incubator (Nuaire, US Autoflow, Plymouth, MN). The organisms were con-

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firmed by Gram staining, motility, and oxidase, catalase, and hippurate hydrolysis tests (31). Molecular confirmation of species was carried out by PCR as described previously (1). The molecular fingerprint of each *C. jejuni* strain was identified by *flaA*-restriction fragment length polymorphism (RFLP) described elsewhere (2). Each of the four strains had unique fingerprints. All strains were motile by spot inoculation by the Mueller-Hinton soft agar method (24).

Animals. BALB/c female mice (6 to 8 weeks old) were obtained from the Animal Resources Centre, Health Sciences Centre, Kuwait University, and housed separately for 1 week prior to experimentation. The animals were fed a standard laboratory chow (Special Diet Food Services, Ltd., Essex, United Kingdom) and water *ad libitum*. Mice used in the present study were free from campylobacter infection. The studies were conducted according to institutional guidelines.

**Serotyping.** The *C. jejuni* strains were serotyped by the Penner serotyping scheme by a passive hemagglutination test using a commercial kit (Denka Seiken Company, Ltd., Tokyo, Japan) according to the procedures recommended by the manufacturer.

**Preparation of GST-PorA fusion protein of** *C. jejuni.* A recombinant clone, pGSTP7RIL, containing the *porA* gene, which encodes the MOMP, from the C31 strain was used (26). This clone expressed PorA as a fusion with glutathione *S*-transferase (GST). Urea-solubilized GST-PorA fusion protein was affinity purified by use of a glutathione-Sepharose B column. The identity of the protein was confirmed by immunoblotting with GST-specific antibody (26). The fusion protein was in denatured form (26). The protein concentration was estimated by using the Bio-Rad (Hercules, CA) protein assay.

Preparation of enriched MOMP (PorA). The MOMP of C. jejuni strain C31 was enriched by the Sarkosyl method (11). Briefly, C. jejuni C31 was grown on blood agar at 42°C for 48 h in a microaerobic atmosphere as described previously. Bacterial cells from plates were harvested in 50 ml of sterile distilled water, centrifuged twice at 5,000  $\times$  g for 10 min at 25°C, and then suspended in 0.01 M Tris buffer (pH 7.4) (Sigma, St. Louis, MO). The cells were disrupted by sonication and centrifuged twice at 5,000  $\times$  g for 20 min to remove whole cells. The supernatant was centrifuged at 100,000  $\times$  g at 4°C for 1 h in an L8-70 ultracentrifuge (Beckman Instruments, Fullerton, CA). The pellet was suspended in 20 ml of 1% N-lauryl sarcocinate (Sigma) containing 7 mM EDTA (Sigma), incubated at 37°C for 20 min, and centrifuged at 100,000  $\times g$  for 2 h. The pellet was then suspended in 0.01 M Tris buffer (pH 7.4) (Sigma) and centrifuged at 100,000  $\times$  g for 2 h. The pellet produced was suspended in 1 ml of sterile distilled water and used as the Sarkosyl-enriched MOMP. The concentration of the protein was measured by the Bio-Rad method. The preparation was checked by immunoblotting as described previously (26) using rabbit antibody to GST-PorA fusion prepared previously (26) and then stored at  $-70^{\circ}$ C.

Cross-reactivity of GST with C. jejuni. The cross-reactivity of GST with C. jejuni proteins was checked by immunoblotting. The agar-grown C. jejuni was suspended in phosphate-buffered saline (PBS; pH 7.2) to a concentration of 109 CFU/ml and disrupted by sonication. The bacterial proteins were separated by discontinuous sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) with a 5% stacking gel and a 12.0% separating gel according to the method of Laemmli (27) and stained with Coomassie blue. For immunoblotting, the separated proteins were transferred electrophoretically onto nitrocellulose (Bio-Rad) and then blocked with 5% skim milk in PBS (pH 7.2). The membrane was reacted with goat GST antibody (Pharmacia Biotech, Piscataway, NJ) or normal goat serum control, as appropriate, both diluted to 1:5,000. The secondary antibody (peroxidase-conjugated anti-goat IgG raised in rabbit; Sigma) diluted to 1:5,000 was added, after which the gel was developed with enhanced chemiluminescence Western blotting detection reagents according to the instructions of the manufacturer (Amersham Pharmacia Biotech). GST-PorA fusion protein was used as a positive control.

**Vaccination.** Vaccination of mice was carried out as described by Baqar et al. (7). Briefly, BALB/c mice were orally vaccinated twice at a weekly interval with the purified GST-PorA fusion protein combined with a modified heat-labile enterotoxin of *Escherichia coli* toxin, LT R192G, as an adjuvant (15) (provided by John D. Clements, Department of Microbiology and Immunology, Tulane University Medical Center, New Orleans, LA). Mice immunized with LT R192G or PBS (pH 7.2) alone were included as controls. Vaccination was done by using a stainless steel, curved-ball-tip feeding needle (20 gauge, 1.5-in. long; Popper and Sons, Inc., New Hyde Park, NY). Just prior to vaccination, gastric acidity was neutralized with two doses (0.5 ml each) of a 5% sodium bicarbonate (pH 8.5) solution given as an oral gavage at an interval of 15 min. An amount of 300 µg of the GST-PorA fusion protein mixed with 5 µg of LT R192G in a total volume of 300 µl in PBS (pH 7.2) was given to each of nine animals. Another group of mice was each fed with 5 µg of LT R192G and yet another group each fed with

 $1 \times$  PBS (pH 7.2). The mouse groups were immunized separately on different days for challenge with different *C. jejuni* strains.

Collection of intestinal lavage fluid. Intestinal lavage fluid from individual mice was collected 7 days after the second vaccine dose according to the procedure described previously by Baqar et al. (7). Four doses of 0.5 ml of the lavage solution (25 mM NaCl, 40 mM Na2SO4, 10 mM KCl, 20 mM NaHCO3, 48.5 mM polyethylene glycol; Sigma) were given orally to the mice at 15-min intervals. At 30 min after administration of the last dose, the mice were given 0.1 ml of PBS (pH 7.2) containing 0.1 mg of pilocarpine (Sigma) intraperitoneally. The intestinal discharge was collected in a petri dish over the following 20 min. It was transferred to a 15-ml conical centrifuge tube, and the volume brought up to 5 ml with PBS (pH 7.2). The sample was vigorously vortexed and then centrifuged at  $650 \times g$  for 15 min. A total of 40 µl of phenylmethylsulfonyl fluoride (35 mg/ml in 95% ethanol; Sigma) and 200 µl of fetal bovine serum (Gibco/Invitrogen Life Technologies, Carlsbad, CA) were added to 4 ml of the supernatant. The protein concentration was determined by using the Bio-Rad method and adjusted to 1 mg/ml with PBS (pH 7.2) for samples from each mouse. The samples were stored at -70°C until used.

**Collection of blood.** Blood was collected from individual mice 3 weeks after the second vaccine dose by snipping the tail as described previously (7). A 50-µl portion of blood was mixed with 950 µl of PBS (pH 7.2) containing 0.1% Tween 20 (PBS-T). After one cycle of freezing and thawing, the sample was centrifuged at  $400 \times g$  for 15 min, and the supernatant was stored at  $-70^{\circ}$ C until used.

Challenge of vaccinated mice. Mice were challenged with bacterial culture 3 weeks after the second vaccine dose, immediately after tail vein blood collection. A 48-h bacterial culture grown on campylobacter agar was suspended in PBS (pH 7.2) to a concentration of  $2 \times 10^9$  CFU per ml, and 0.5 ml of the suspension was orally fed to the mice immediately after neutralization of the gastric acidity, as described above. Fecal shedding of the organism by each mouse was monitored for nine consecutive days according to the method described by Bagar et al. (7) with some modifications. Instead of a few fresh pellets of feces, all fecal pellets shed every day were collected by using a sterile stainless steel spoon and placed in a sterile 50-ml centrifuge tube. Feces was diluted to 10% with PBS (pH 7.2), followed by 5-fold serial dilutions. An aliquot of 50 µl of each dilution was spread onto campylobacter agar by using a sterile glass spreader. Plates were incubated at 42°C under microaerobic condition for 48 h and examined for the presence of C. jejuni colonies. C. jejuni colonies appeared as gray, moist colonies about 2 to 3 mm in diameter, which were easily distinguishable from normal fecal flora. The colonies were further confirmed by Gram staining, motility, and catalase and oxidase tests. The strain identity was confirmed by flaA-RFLP as described above.

**Passage of** *C. jejuni* **strain C31 through mouse intestine.** To increase the colonization potential of *C. jejuni* C31 for mice, six BALB/c mice were orally inoculated with *C. jejuni* C31 as described in the preceding section. Feces was cultured on campylobacter agar every day for the next 4 days. Several colonies from each mouse were pooled and used as the seed culture for preparation of challenge inoculum for another 6 mice. This way, the strain was passaged four times. After each passage, the identity of the strain was monitored by *flaA*-RFLP (2).

Measurement of antibody responses. C. jejuni-specific antibodies (IgA, IgG, and IgM isotypes combined) were measured in mouse intestinal lavage fluid and blood samples by enzyme-linked immunosorbent assay using the modified method of Baqar et al. (7). The wells of microtiter Maxisorp plates (Nunc, Rochester, NY) were coated with carbonate buffer (pH 9.6) containing either 1 µg of purified GST-PorA fusion protein or enriched PorA of C. jejuni C31 at 4°C for 24 h. The wells were washed four times with PBS (pH 7.2) containing 0.1% Tween 20 (PBS-T) and blocked with 5% milk powder (Regilait, Saint-Martin-Belle-Roche, France) in PBS (pH 7.2) at 37°C for 1 h. Serial doubling dilutions of lavage fluid or lysed blood samples in PBS–0.05% Tween 20 were added to the wells, followed by incubation at 37°C for 1 h. The wells were washed four times with PBS-T and then incubated with horseradish peroxidase-conjugated goat anti-mouse secondary antibodies (that recognize IgA, IgG, and IgM isotypes) (Kirkegaard & Perry Laboratories, Gaithersburg, MD) diluted 1:5,000 with 1× PBS (pH 7.2) at 37°C for 1 h. The wells were washed four times with PBS-T. The substrate 2,2'-azino-di-(3-ethyl-benzthiazoline sulfonate) (ABTS; Sigma) was added, and after 30 min of incubation at 37°C, the optical density at 405 nm was measured in a Powerwave Microplate spectrophotometer (Biotek Instruments, Inc., Bedfordshire, United Kingdom). Endpoint titers were expressed as the reciprocal of the dilution giving an absorbance value ≥2 standard deviations above the background absorbance in wells containing intestinal lavage fluid or lysed blood samples from normal mice (7).

**Vaccine efficacy.** Vaccine efficacy was calculated as follows: ([rate of control mice – rate of vaccinated mice]/rate of control mice)  $\times$  100, where the rate is the



FIG. 1. Coomassie blue-stained SDS–12% PAGE gel after electrophoresis (lanes 1 to 3) and immunoblot gel of the purified MOMP from *C. jejuni* C31 (lane 4). Lane 1 shows molecular mass markers, lane 2 shows crude MOMP (~48 kDa), and lane 3 shows Sarkosylpurified MOMP. Lane 4 shows the Sarkosyl-purified MOMP detected using anti-GST-PorA antibody by immunoblotting. The picture of the Coomassie blue-stained gel containing lanes 1 to 3 and the picture of the immunoblot containing lane 4 were combined for the composite picture. The values on the left of the figure indicate the molecular masses (in kilodaltons) of standard markers, and the arrow on the right indicates the MOMP.

percentage of mice free of *C. jejuni* colonization (5). Protection for the total duration of 9 days was calculated by taking into account the sum of vaccinated mice free of colonization for each of 9 days in comparison to similar data in the control group of mice.

**Statistical analysis.** Statistical analysis was done by using SPSS version 17 (SPSS, Chicago, IL). Comparison of rates of mice free from colonization with *C. jejuni* was done by the Fisher exact test. The comparisons of endpoint antibody titers were done by Mann-Whitney test. Values differing with  $P \le 0.05$  were considered statistically significant.

# RESULTS

**Serotypes of** *C. jejuni* **strains.** The Penner serotypes of *C. jejuni* strains were determined by passive hemagglutination test. Strain C31 belonged to serotype 6,7, strain 48 belonged to serotype 19, strain 75 belonged to serotype 3, and strain 111 belonged to serotype 1,44. Since the *porA* gene was cloned from strain C31 to make the GST-PorA fusion protein for vaccination, strain C31 was considered the homologous strain, and the other strains were considered heterologous strains.

**GST-PorA fusion protein.** The recombinant *porA* gene clone pGSTP7RIL expressed a fusion protein of 76 kDa, as expected (26). The identity of the protein was confirmed by immunoblotting with GST-specific antibody (data not shown).

**Sarkosyl-enriched MOMP (PorA).** The Sarkosyl-enriched MOMP of strain C31 had a molecular mass of ~48 kDa. Its identity was confirmed by immunoblotting with anti-GST-PorA antibody (Fig. 1).

**Cross-reactivity of GST with** *C. jejuni*. No cross-reactivity of GST with antigens from the four *C. jejuni* strains C31, 48, 75, and 111 was seen. However, the positive control produced a 76-kDa band, as expected (data not shown).

**Colonization of** *C. jejuni* **in vaccinated and control mice and vaccine efficacy.** Reduced colonization in vaccinated mice on certain days, which achieved significance in comparison to the control mice, and colonization only for the 9-day period are shown below for each strain.

Strain C31. With the homologous strain C31, colonization was seen on day 1 in both the vaccinated and control groups of mice (Fig. 2A). Attempts at improving the colonization potential of the strain by repeated passage through mice did not improve colonization. The organism was not shed in the feces beyond the first day postinfection after each of the four passages. The percentage of mice colonized on day 1 in the vaccinated group was significantly reduced (P = 0.029 versus both PBS and LT R192G control groups), with a high vaccine efficacy (56%).

**Strain 48.** This strain colonized vaccinated and control mice on all 9 days (Fig. 2B), but colonization was significantly reduced in the vaccinated group compared to the PBS group on days 8 (P = 0.029) and 9 (P = 0.009) and for the 9-day period (P = 0.001). Reduction was also significant against the LT control group on day 9 (P = 0.015) and for the 9-day period (P = 0.001). The vaccine efficacy was appreciable for days 8 (56%) and 9 (78%) and for the 9-day period (43%).

**Strain 75.** This strain colonized the vaccinated and control mice throughout the 9 days, but a lower proportion of vaccinated mice experienced colonization (Fig. 2C). Reduction in colonization was significant for day 7 (P = 0.029 versus the PBS control group and P = 0.009 versus the LT control group) and for the 9-day period (P = 0.002 versus the PBS control group and P = 0.001 versus the LT control group). The vaccine efficacy for day 7 (72%) was greater than the efficacy for the 9-day period (29%).

**Strain 111.** Vaccinated mice were colonized for 8 days and the control mice for all 9 days by this strain. A lower proportion of vaccinated mice than control mice was colonized (Fig. 2D). Reduction in colonization was significant on days 2 (P = 0.029 versus the PBS control group), 7 (P = 0.015 versus the PBS control group and P = 0.003 versus the LT control group), and 9 (P = 0.029 versus the PBS control group only) and for the 9-day period (P = 0.001 versus both the PBS and the LT control groups). The vaccine efficacies for these different days (56 to 100%) were higher than the efficacy for the 9-day period (42%).

Antibody response to vaccine. Serum and intestinal lavage fluid samples from the PBS control group, the adjuvant control group, and the vaccinated group were used to estimate antibody responses. There were nine mice in each group that were randomly selected (Table 1). There was no antibody response in the PBS control group. Some antibody response in the LT control group was noticed. There was a higher antibody response in the vaccinated group. In general, the response in lavage fluid was higher than the response in serum. There was antibody response to both MOMP and GST-PorA, although the response to the latter was somewhat higher. The antibody response in the vaccinated group was in general significantly higher than the responses in the PBS and LT control groups.

#### DISCUSSION

An effective vaccine could be based on antigens common to all strains, critical for bacterial metabolism or virulence and against which immunity can be elicited. Since PorA protein fulfills these criteria, we evaluated this antigen for its ability to provide homologous and heterologous protection. We calculated the protective efficacy of the vaccine by taking into con-



FIG. 2. Percent colonization of mice which orally received PBS ( $\bullet$ ), LT R192G ( $\blacksquare$ ), or GST-PorA + LT ( $\bigcirc$ ). Each group of mice (n = 9) was challenged with the homologous strain *C. jejuni* C31 (O:6,7) (A) or the heterologous strains *C. jejuni* 48 (O:19) (B), *C. jejuni* 75 (O:3) (C), and *C. jejuni* 111 (O:1,44) (D).

sideration the ability of the vaccine to reduce colonization, as done by others (4, 6, 28, 29).

The PorA protein used as a vaccine in the present study was expressed from a previously cloned *porA* gene from strain C31 in fusion with GST protein (26). When we tested C31 in the current study, we found the strain to be a poor colonizer of the mouse intestine. Passage of the strain by oral inoculation of

mice did not improve colonization. Motility is linked to colonization of chickens, the natural host of the organism (32). However, this may not correlate with the colonization of mouse intestine since C31 was motile and yet a mouse non-colonizer. It has been shown that the colonization potential of some strains increases by chicken passage (14, 24) but remains unchanged for others (39). We had to use C31 as the homol-

TABLE 1. Mean endpoint titers of antibody responses in sera and intestinal lavage fluid of vaccinated and control (PBS or LT R192G) mice (n = 9)

Immunizing agent(s)	Endpoint titer $(range)^a$			
	Serum		Intestinal lavage fluid	
	Plates coated with MOMP	Plates coated with GST-PorA	Plates coated with MOMP	Plates coated with GST-PorA
PBS	0 (0–0)	0 (0–0)	0 (0-0)	0 (0–0)
LT R192G	0 (0–0)	11 (0-100)	20 (0-100)	40 (0-100)
GST-PorA + LT R192G	1,444 (200–3,200)*	1,956 (800–3,200)*	2,667 (800–3,200)†‡	3,200 (3,200–3,200)*

 $a^{a}$  \*, P = 0.001 compared to the PBS and adjuvant control groups;  $\dagger$ , P = 0.004 compared to the PBS control group;  $\ddagger$ , P = 0.024 compared to the adjuvant control group.

ogous strain since the vaccine was derived from this strain. Being a poor colonizer, the strain was excreted on the first day only, and the vaccine afforded significant protection against colonization on the first day. Protection on the first day was seen against the homologous challenge strain only and not against heterologous challenge strains. This shows that homologous protection is stronger than heterologous protection. Since there was no colonization beyond the first day for strain C31, we did not compute protection for the 9-day period for this homologous challenge strain. Had we used a better colonizing strain as the homologous strain, it is likely that we would have seen colonization of the strain for more days and higher protection for the 9-day period. However, to confirm homologous protection, studies have to be done with PorA derived from a better-colonizing strain.

With the heterologous strains, the vaccine afforded significant protection against colonization on certain days and for the total duration. The variability of protection on different days among the three strains can be attributed to their different colonizing abilities and/or differences in the amount of shared antigenic epitopes present in their PorAs.

We did not vaccinate mice with PorA alone, but vaccination was done with the mucosal adjuvant LT R192G, as was done with several candidate campylobacter vaccines to boost immune response (4, 6, 29). This mucosal adjuvant induces dendritic cell migration into the follicle-associated epithelium of Peyer's patches, thus promoting capture of the antigens by dendritic cells of the Peyer's patches (3). The adjuvant enhances Th1 responses and stimulates acquired immunity by activating the components of the innate immune system (40). However, control mice vaccinated with LT alone did not have protection.

Vaccinated mice responded with antibody production in the serum and intestinal secretion. We did not look for individual isotypes of antibodies. Since all three isotypes are present in the serum and the IgA isotype, along with small amounts of IgG and IgM isotypes, is present in the secretion, we assayed for the total antibodies representing all three isotypes. Since the antigen was delivered orally, the antibody response in the secretion was higher than in the serum. Since the vaccine is a combination of the PorA and the carrier protein, the antibody response would be expected to be directed against both components. To find out whether appreciable antibody production occurred against the PorA component, we tested the samples in parallel against the Sarkosyl-enriched MOMP as the antigen. This showed that most of the antibody was directed against MOMP.

In *C. jejuni* infection of chickens, the serum antibodies produced are predominantly against conformational epitopes and weakly against denatured (linear) epitopes of the MOMP. Also, rabbits immunized with conformational epitopes of MOMPs from different strains produced strain-specific antibodies, while antibodies induced by denatured MOMP were cross-reactive with MOMPs from different strains (21). At the same time, sera from naturally infected humans and from rabbits immunized with whole bacterial cells were found to have antibodies against a 45-kDa MOMP from *C. jejuni* strains, which is the denatured epitope of MOMP (12). Taken together, these data suggest that in natural *C. jejuni* infection, most of the antibodies produced are directed against conformational epitopes and some against linear epitopes of MOMP. Since the PorA vaccine used in our study is a denatured antigen, we believe that antibodies to this denatured antigen are providing heterologous protection. Boosting antibody response to linear epitopes (by use of denatured antigen) should improve heterologous protection. Comparative studies in which animals are immunized separately with the two different forms of antigen and then challenged with homologous and heterologous serotypes will clarify whether linear epitopes are involved in heterologous protection. With other organisms, both linear and conformational epitopes (45) or linear epitopes of OMPs (22) were found to impart protection.

We did not include control mice immunized with GST and measure antibodies to it in serum and intestinal lavage fluid. No such control group was needed since the four *C. jejuni* challenge strains did not cross-react with GST. Therefore, it can be concluded that antibodies to PorA, but not to GST, were responsible for the protection observed.

Different antigens have been evaluated as possible vaccines using different animal models for homologous and/or heterologous protection (reviewed in reference 23). However, vaccines evaluated in mice are pertinent to our own data. The vaccine candidates evaluated included killed whole cells (6), maltose-binding protein of Escherichia coli fused to flagellin protein (28), periplasmic binding protein (38), flagellumsecreted proteins (FlaC, FspA1, and FspA2) (4), conjugated capsular polysaccharides (29), and an adherence protein expressed on and delivered by an attenuated Salmonella enterica serovar Typhimurium strain (42). The mouse models included intranasal or intestinal colonization models. Vaccines were delivered intranasally, orally, or parenterally. The protection evaluated was homologous or both homologous and heterologous. The vaccines provided no or various degrees of protection. Further studies with some of these vaccines are required to assess their usefulness as serious candidates.

Ours is the first study in which the MOMP of *C. jejuni* in the form of a fusion protein has been evaluated as a vaccine in a mouse intestinal colonization model. Our data provide evidence that this vaccine can impart substantial heterologous protection. There is preliminary evidence to show that it could provide homologous protection as well. Our study was undertaken with a view to developing a vaccine for human use. There are several other issues that need to be addressed in the preparation of a PorA vaccine. For example, since bacterial porins can be posttranslationally modified, a recombinant protein needs to be compared to PorA enriched from bacteria to assess which constitutes a better vaccine.

It is pertinent to note that PorA antigen-based vaccines are in use against group B meningococcus. Since there is antigenic polymorphism in meningococcal PorA, many antigenic types need to be incorporated in a vaccine for optimal protection (43). Likewise, it is possible that protection can be enhanced by incorporating PorA from different strains of *C. jejuni*.

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