Role of *Helicobacter pylori cag* Region Genes in Colonization and Gastritis in Two Animal Models

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The *Helicobacter pylori* chromosomal region known as the cytotoxin-gene associated pathogenicity island (*cag* PAI) is associated with severe disease and encodes proteins that are believed to induce interleukin (IL-8) secretion by cultured epithelial cells. The objective of this study was to evaluate the relationship between the *cag* PAI, induction of IL-8, and induction of neutrophilic gastric inflammation. Germ-free neonatal piglets and conventional C57BL/6 mice were given wild-type or *cag* deficient mutant derivatives of *H. pylori* strain 26695 or SS1. Bacterial colonization was determined by plate count, gastritis and neutrophilic inflammation were quantified, and IL-8 induction in AGS cells was determined by enzyme-linked immunosorbent assay. Deletion of the entire *cag* region or interruption of the *virB10* or *virB11* homolog had no effect on bacterial colonization, gastritis, or neutrophilic inflammation. In contrast, these mutations had variable effects on IL-8 induction, depending on the *H. pylori* strain. In the piglet-adapated strain 26695, which induced IL-8 secretion by AGS cells, deletion of the *cag* PAI decreased induction. In the mouse-adapted strain SS1, which did not induce IL-8 induction. These results indicate that in mice and piglets (i) neither the *cag* PAI nor the ability to induce IL-8 invitro is essential for colonization or neutrophilic inflammation and (ii) there is no direct relationship between the presence of the *cag* PAI, IL-8 induction, and neutrophilic gastritis.

The gastric bacterium *Helicobacter pylori*, first cultured in 1983, is now a well-established cause of gastritis and peptic ulcer disease (22, 41). Infection is associated with production of proinflammatory cytokines which lead to chronic or chronic active gastritis. The activity of the gastritis (i.e., the intensity of neutrophil influx) is commonly considered an indicator of severity, and chronic active gastritis is associated with more severe manifestations of disease, such as peptic ulceration and neoplasia (9, 14, 47).

Among the cytokines associated with H. pylori colonization, the CXC family of chemokines may be involved in inducing neutrophilic influx. Two of these CXC chemokines and neutrophil chemoattractants, interleukin-8 (IL-8) and GRO- α , are increased in the gastric mucosa of patients with chronic active gastritis and ulceration (14-16, 18, 47, 53, 54, 56, 61). In addition, coculture of H. pylori with gastric cancer cell lines (15, 32, 52), normal human gastric mucosal cells (44), and immortalized mouse gastric mucus cells (40) results in induction of proinflammatory cytokines IL-8 and IL-6 or, in murine cells, CINC-2 β , the rodent equivalent of GRO- α . In vivo, IL-8 secretion is most prominent in the superficial gastric mucosa (18), and it is likely that mucosal epithelial cells are a source of neutrophil-inducing cytokines, although other sources likely exist (10, 26, 34, 62). Thus, it is commonly supposed that severe manifestations of H. pylori-related disease are promoted by H.

neutrophilic inflammation. *H. pylori*-induced chemokine secretion by gastric epithelial cells appears to be facilitated at least in part by genes in the *cag*

pylori-induced epithelial secretion of IL-8 and the attendant

cells appears to be facilitated at least in part by genes in the *cag* pathogenicity island (PAI). This stretch of approximately 30 open reading frames is present in many but not all strains of *H. pylori*, and many of the predicted genes are homologous to virulence-associated genes of other bacterial pathogens (2, 11, 12). Homology to the *vir* genes of *Agrobacterium tumefaciens* and the *ptl* genes of *Bordetella pertussis* suggests that *H. pylori cag* PAI genes may code for a type IV secretion complex important in delivery of proinflammatory or other pathogenic molecules to host cells (12, 43).

In addition to genetic homologies, there is functional evidence of the role of the *cag* PAI in virulence. Its presence is associated with increased severity of disease in the human host (9, 47). In addition, induction of IL-8 in cultured cells in vitro appears to depend at least in part on *cag* genes, at least in some *H. pylori* strains (2, 11). In spite of genetic, epidemiologic, and in vitro evidence supporting the link between *cag* genes, IL-8 induction in vitro, and neutrophilic gastric inflammation, however, few animal model studies have been done to investigate the role of *cag* PAI genes in vivo.

The goal of this study was to use two animal models to evaluate both the *cag* PAI as a whole and selected *cag* PAI genes in induction of gastritis in vivo. The germ-free piglet model of infection uses strain 26695, a pig-adapted human isolate (23). Strain 26695 colonizes piglets well and induces gastritis which is largely lymphocytic and lymphofollicular, with a variable neutrophilic component. Strain 26695 induces IL-8 in vitro (2, 38), its genomic DNA sequence is known (57), and its *cag* PAI is uninterrupted (the *cag*I and *cag*II regions are

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TABLE 1. Number of animals examined at each sacrifice interval

	No. at each sacrifice interval (time postinfection)					
Bacterial strain	2 days	3 wk	7 days	14 days	13 wk	
Germ-free piglets						
None	0		4	0		
26695	4		6	2		
$26695\Delta cag$	4		6	2		
Mice						
None		0			6	
SS1		4			12	
SS1virB11::cat		5			5	
SS1virB10::cat		3			0	

adjacent), allowing deletion of the entire region by simple reverse genetics (2). Mice are poorly colonized by strain 26695 but are well colonized by strain SS1, the other strain used in this study (24, 37).

MATERIALS AND METHODS

Bacterial strains. Strain 26695 is a human isolate which was adapted to pig colonization by serial passage (3). Strain SS1 is a mouse-adapted human isolate (37). Both strains contain the *cag* PAI, which is continuous in 26695 (2) but divided into *cag*I and *cag*II in SS1 as described for other *H. pylori* strains (2). Construction of mutant strains by transformation was performed as previously described (2). The mutated genes were HP0524, HP0525, and HP0527 (57). These genes are located in the *cag*II region of the *cag* PAI and have previously been referred to as *cag10*, *cag11*, and *cag13* (2). They are homologous to *A. tumefaciens virD4*, *virB11*, and *virB10*, respectively. In *A. tumefaciens*, these genes code for proteins which form part of a type IV secretion apparatus, and they are presumed to have a similar function in *H. pylori* (12, 43). Their presence has been associated with tyrosine phosphorylation of host and bacterial proteins (4, 38, 43, 50, 55) as well as secretion of IL-8 (15) upon contact with cultured cells.

The genetic constructs used have been previously described (2). To inactivate individual genes, *H. pylori* strain SS1 or 26695 was transformed with a DNA construct containing a chloramphenicol resistence cassette (*cat*) inserted into the coding region of the gene to be interrupted. To delete either *cag*II or the entire *cag* region, strain SS1 or 26695 was transformed with a DNA construct containing *cag* flanking sequences surrounding a chloramphenicol resistance cassette. Gastritis and colonization potential of bacterial mutants were evaluated in germ-free piglets inoculated with 26695 or 26695 Δ *cag* and in mice given SS1, SS1*virB10::cat*.

IL-8 assay. IL-8 induction by *H. pylori* cocultured with AGS cells was determined by the method of Crabtree et al. (15). Briefly, AGS cells (a gastric carcinoma cell line) were grown in 24-well plates to 60 to 75% confluence in RPMI 1640 supplemented with fetal calf serum. Broth-cultured *H. pylori* was washed in RPMI 1640 and resuspended to $10^7/ml$, and 1 ml of the suspension was added to AGS cell monolayers. After incubation for 24 h at 37°C, media were collected, bacteria were removed by centrifugation, and media were stored at -70° C until assay. IL-8 was detected with a capture enzyme-linked immunosorbent assay kit (Pharmingen) according to the manufacturer's instructions. Values reported were normalized by subtraction of endogenous IL-8 production (production by unstimulated AGS cells). Specificity of the assay was verified by inhibition with 1 µg of recombinant human IL-8 per ml.

Animal studies. Germ-free piglets were derived and maintained as previously described (23). They were orally inoculated with 10^9 CFU of broth-cultured *H. pylori* at 3 days of age and killed 2, 7, or 14 days after inoculation. Female 6 to 8-week-old helicobacter-free C57BL/6 mice (from Jackson Laboratory) were maintained in microisolator cages and fed sterile lab chow and water ad libitum. They were orally inoculated with 10^7 CFU of broth-cultured *H. pylori* and killed 3 or 13 weeks after inoculation. The number of animals in each sacrifice group is given in Table 1.

At sacrifice, stomachs were removed and bisected along the greater and lesser curvatures, and bacterial colonization in one half of the stomach was quantified by plate dilution on blood agar plates with and without 20 μ g of chloramphenicol per ml. Sections of piglet stomach from cardia, fundus, and antrum were formalin fixed and paraffin embedded for histologic examination. Mouse stomach was



FIG. 1. Induction of IL-8 by *H. pylori* strains 26695, SS1, and *cag*deficient mutants. *, significantly different from isogenic mutants, P < 0.05; †, significantly different from strain 26695, P < 0.05.

sectioned in approximately 1-mm-wide slices and then fixed and embedded as described above.

Immunohistochemistry. Gastric sections from piglets killed 7 days after inoculation were examined for the presence of cells bearing CD4 and CD8 surface markers. OCT-embedded frozen tissue was cut in 3- μ m sections and stained as previously described (25). Primary antibody (mouse monoclonal anti-pig CD4 or CD8; VMRL, Inc., Pullman, Wash.) was diluted 1:100. In control slides, normal mouse serum was substituted for the primary antibody. Sections were scored by enumerating the number of cells per 20× field. For each tissue block, replicate sections stained for CD4 and CD8 were examined to identify fields which were present in both slides. Of these, 6 to 12 fields which contained CD4⁺ or CD8⁺ cells were selected and scored. All sections were scored blind without knowledge of their source or antibody stain.

Histopathology. Hematoxylin-and-eosin stained sections were scored for the extent of gastritis as previously described (24). Briefly, microscopic fields containing gastritis severe enough to displace glands, and fields containing neutrophilic infiltrate were enumerated and expressed as a percentage of the gastric mucosa. All available fields were scored, and slides were examined blind, without knowledge of their source.

Statistics. Means were compared by *t*-test or by Mann-Whitney *U* test. Statistical significance was set at P < 0.05.

RESULTS

Induction of IL-8 in AGS cells. Strain 26695 induced IL-8 secretion when cocultured for 24 h with AGS cells, and deletion of the *cag* PAI in this strain significantly decreased IL-8 induction to about 20% of the wild-type level (P = 0.02 [Fig. 1]). In contrast to 26695, and despite the presence of the *cag* PAI, wild-type strain SS1 did not induce IL-8 secretion by AGS cells. IL-8 induction by SS1 was 3% of induction by 26695 (P = 0.02) and not significantly different from background (P = 0.25). Interestingly, however, deletion of the entire *cag*II region as well as any of the *cag* genes tested, the *virB10*, *virB11*, and *virD4* homologues, resulted in slightly increased IL-8 induction by this strain. These differences were consistent among all mutants, and IL-8 induction by all mutant groups combined was significantly greater than induction by wild-type SS1 (P = 0.009 [Fig. 1]).

Bacterial colonization. Colonization rates were similar in piglets (Fig. 2) and mice (Fig. 3). In both species, all inoculated animals became colonized with all challenge strains, and the extent of colonization ranged from approximately 10^5 to 10^8 CFU/g of gastric mucosa. In piglets, colonization by 26695 decreased over time from greater than 10^8 CFU/g of gastric



FIG. 2. Colonization of piglets by *H. pylori* strain 26695 and *cag*deficient mutants 2, 7, and 14 days postinoculation (PI). Two days after inoculation, colonization by $26695\Delta cag$ was significantly less than colonization by 26695 (*, P < 0.05), but the two strains did not differ at the other sacrifice intervals.

mucosa 2 days after inoculation to 4×10^6 CFU/g by 14 days after inoculation. In mice, colonization increased somewhat between 3 and 13 weeks after inoculation. Overall, there was no consistent difference in colonization between wild-type and mutant bacteria. Two days after inoculation, extent of colonization by $26695\Delta cag$ was lower than that by 26695, but the difference did not persist to the other sacrifice intervals. Colonization by SS1, SS1 *virB11::cat*, and SS1*virB10::cat* differed somewhat from each other, but there were no statistically significant differences.

Gastritis. In piglets, gastritis was present 7 and 14 days after inoculation (P = 0.01 and 0.021 compared to uninfected piglets [Fig. 4]). It was characterized primarily by lymphocytic and plasmacytic infiltrate (Fig. 5); neutrophilic infiltrate was variable and mild (Fig. 4). In mice, gastritis was present 13 weeks after inoculation (P = 0.0015, SS1-infected compared to uninfected mice [Fig. 6]) and was characterized by mixed inflam-



FIG. 3. Colonization of mice by *H. pylori* strain SS1, and *cag*-deficient mutants 3 and 13 weeks postinoculation (PI). Colonization ranged from approximately 10^6 to 10^7 CFU/g of gastric mucosa, but there were no significant differences between strains or sacrifice intervals.



FIG. 4. Gastritis and neutrophilic infiltration (PMN) in piglets colonized by *H. pylori* strain 26695 and *cag*-deficient mutants 2, 7, and 14 days postinoculation (PI). Seven and 14 days after inoculation gastritis was significantly greater than in uninfected piglets (*, P < 0.05), but there were no differences between strains.

matory infiltrate including lymphocytes, macrophages, plasma cells, and neutrophils as well as multifocal loss of fundic gland morphology with replacement by mucus-type glands (metaplasia) (Fig. 7). Neutrophilic infiltrate was significantly elevated in these mice (P = 0.005, SS1 infected compared to uninfected), and was more prominent than in piglets. In both mice and piglets there was no difference in gastritis or neutrophilic infiltrate between groups infected with wild-type *H. pylori* and groups infected with *cag* mutant strains.

Immunohistochemistry. Both CD4⁺ and CD8⁺ cells were significantly increased in the gastric mucosa of infected piglets compared to uninfected piglets (P < 0.05 [Table 2]). However, both cell types were present in all infected piglets, and there were no differences between piglets given 26695 and those given 26695 Δ cag. In all piglets, more gastric lamina propria Imphocytes expressed CD8 than CD4 (Table 2).

DISCUSSION

The results reported here seem to contradict current dogma in two ways. First, we have shown that in at least one strain of H. pylori, SS1, the ability to induce IL-8 secretion by cultured gastric epithelial cells is not directly correlated with the presence of a cag-PAI. In contrast to strain 26695, strain SS1 did not induce IL-8 secretion despite the presence of the PAI, and, surprisingly, inactivation of several genes in that region actually promoted, rather than inhibited, IL-8 induction. While these finding contrast with many published studies (1, 11, 17, 19, 27, 38, 44, 58), they may actually represent an accurate assessment of the phenotypic diversity of H. pylori. Most published studies used only one or a few well-characterized strains to demonstrate dependence of IL-8 secretion on the cag PAI. Recently published studies, however, suggest that over a broad range of strains, IL-8 induction may not always correlate with presence or absence of the cag PAI. In one study in which 80 clinical isolates were examined, IL-8 induction by *cag*-positive strains varied widely, and some *cag*-negative strains were able to induce IL-8 in cultured cells (61). In another study, 153 clinical isolates were examined, and in many cases strains with the cag PAI failed to induce IL-8, while strains without the PAI



FIG. 5. Tissue from the gastric cardia of an uninfected gnotobiotic piglet (A) and a piglet infected with *H. pylori* strain 26695 for 14 days (B). Gastritis is characterized by lymphocytes and plasma cells (arrowheads) with scattered neutrophils (arrow). Hematoxylin and eosin stain. Bar = $75 \ \mu m$.

did induce secretion (7). A third study failed to find either an association between IL-8 induction and *cag* or an association between in vivo IL-8 levels and in vitro IL-8 induction by the same strain (45). In addition, non-*cag*-containing bacterial species other than *H. pylori* have been shown to induce IL-8 in cultured cells in some studies (31). Finally, several *H. pylori* genes unrelated to *cag* have been associated with IL-8 induction. In one study cited above, IL-8 induction was partly dependent on *oipA*, a gene which is unrelated to *cag* (61). In other studies IL-8 induction has been associated with flagellar morphology (46), heat shock proteins (60), and other *H. pylori* products (21).

Our demonstration that inactivation of *cag* genes leads to promotion of IL-8 induction in SS1 has not been previously reported. However, this finding may be explained by diversity in secretory function as reported for other bacterial species. The *cag* genes examined are presumed to encode part of a type IV secretion system involved in injection of bacterial proteins into the host cell (12, 43, 55), and it is widely assumed that this



FIG. 6. Gastritis, neutrophilic infiltration (PMN), and epithelial metaplasia in mice colonized by *H. pylori* strain SS1 and *cag*-deficient mutants 3 and 13 weeks postinoculation (PI). Three weeks after inoculation, neutrophilic inflammation was more prominent in mice colonized by the *cag*-deficient mutants than in mice colonized by SS1 (†, P < 0.05), but this difference did not persist. Thirteen weeks after inoculation, PMN, gastritis, and metaplasia were all significantly elevated in all mice (*, P < 0.05), but there were no differences between bacterial strains.

secretion function is exclusively proinflammatory (1, 27, 51). However, other bacterial species possess secretion systems which can induce either proinflammatory or anti-inflammatory signals to host cells (42, 59). It has been suggested, in fact, that secretion of anti-inflammatory signals may be one way in which nonpathogenic enteric bacteria down-regulate a host immune response. Conceivably, an equivalent mechanism in *H. pylori* contributes to the virulence of specific strains. That is, the *cag*-related secretion system may function to deliver anti- as well as proinflammatory signals to target tissues, thereby determining in part the level of the host response.

A second surprising finding of this study was the lack of congruence between IL-8 induction in vitro and neutrophilic gastritis in vivo in mice. It is widely assumed that IL-8 induction in gastric epithelial cells is the mechanism whereby neutrophilic gastritis is induced by *H. pylori*. However, this hypothesis has not been tested in animal models. In humans, IL-8 and neutrophils are both present in gastric mucosa colonized by *H. pylori* (18, 47). The presumption of causation is based on two types of studies, the in vitro induction of IL-8 by *H. pylori* and the epidemiologic association between *cag*-containing strains and disease.

The relationship between in vivo studies in cultured cells and in vivo pathogenesis must be interpreted with caution. Gastric cancer cell lines used for in vitro assays differ from gastric surface mucus cells in many ways, including surface markers and receptors, lack of polarity, and synthetic capability. Thus, their interactions with bacteria are likely to differ. Even freshly isolated gastric epithelial cells may not reflect in vivo conditions, since these cells are mixtures of parietal cells, chief cells, neck cells, and surface mucus cells. Only a minority of cells may be of the same phenotype and response to *H. pylori* as the surface mucus cells that encounter *H. pylori* in vivo. Another critical difference between in vivo and in vitro interactions between *H. pylori* and gastric epithelial cells is the multiplicity



FIG. 7. Tissue from the gastric fundus of an uninfected mouse (A) and a mouse infected with *H. pylori* strain SS1 for 13 weeks (B). Gastritis is characterized by a mixed infiltrate of lymphocytes and neutrophils. Note the paucity of parietal and chief cells characteristic of epithelial metaplasia in *H. pylori*-infected mice (see reference 24). The arrows indicate three remaining parietal cells. Hematoxylin and eosin stain. Bar = $50 \ \mu m$.

of infection. Cell culture studies typically use ratios of 50 to 100 bacteria per gastric epithelial cell, several orders of magnitude greater than the ratio found in vivo (10^5 to 10^6 CFU/biopsy or more than 10 epithelial cells to one bacterium) (6). Further, bacteria in vitro surround and closely adhere to the gastric

TABLE 2.	T-cell subsets in germ-free piglets infected with wild-
	type and cag-deficient H. pylori

Group	No. of cells/20× microscopic field (mean \pm SE)		
	CD4 ⁺ cells	CD8 ⁺ cells	
Uninfected All infected Infected with 26695 Infected with 26695∆ <i>cag</i>	$0.15 \pm 0.15 \ 4.5 \pm 1.6^a \ 5.6 \pm 2.9 \ 3.2 \pm 1.2^b$	$\begin{array}{c} 2.1 \pm 0.5 \\ 14.6 \pm 2.1^a \\ 13.2 \pm 4.0 \\ 15.9 \pm 1.8^b \end{array}$	

^{*a*} Significantly greater than uninfected piglets, P < 0.05.

^b Not significantly different from 26695 group, P > 0.1.

epithelial cells in the absence of intervening mucus. In vivo, bacteria are mostly within the mucus, a minority of bacteria actually adhere to the epithelial cells, and those only adhere to the luminal surface. Thus, interactions between bacteria and epithelial cells in vivo and in vitro are likely to differ markedly, and induction of IL-8 synthesis by bacterial contact in vitro does not necessarily indicate that such induction occurs in vivo or is of sufficient magnitude to have a primary effect.

Epidemiologic studies provide stronger evidence of an association between *cagA* or the *cag* PAI and increased severity of disease. Many such studies are published (for recent reviews, see references 5 and 13), and most indicate that the cag PAI has a role in the pathogenicity of H. pylori. However, the epidemiologic association between cag and disease is not absolute, suggesting that the relationship is likely not direct. Not all published studies have shown a relationship between the cag PAI and increased severity of disease (8, 28, 29, 35, 36, 39, 48, 63), and no study has shown an absolute relationship. Even when there is an overall statistical association between infection by strains which express *cag* and increased severity of disease, many patients examined have severe disease but were colonized by cag-negative strains (28, 30, 33). Thus, although the cag PAI clearly contributes to disease due to H. pylori, cag-negative strains can also be pathogenic.

Taken together, the occurrence of *cag*-negative disease in humans and the absence of an effect of *cag* in animal models are most easily explained by the role of the host response in disease outcome. Data are accumulating that in both humans and experimental animals, the outcome of *H. pylori*-related disease is strongly dependent on host immune response (20, 49). Different hosts may differ in their response to *H. pylori* antigens, thus leading to different outcomes in individuals infected with strains of similar antigenicity. Greater antigenicity of *cag*-dependent antigens in some individual hosts, host strains, or host species could explain the tendency toward greater pathogenicity of these strains. Clearly, however, *cag*-independent antigens can induce disease, at least in animals, and *cag*-negative strains are not necessarily innocuous.

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