



Membrane-associated Activation of Cholesterol α -Glucosyltransferase, an Enzyme Responsible for Biosynthesis of Cholesteryl- α -D-Glucopyranoside in *Helicobacter pylori* Critical for Its Survival

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Summary

Helicobacter pylori (*H. pylori*) is the causative pathogen underlying gastric diseases such as chronic gastritis and gastric cancer. Previously, the authors revealed that α 1,4-linked *N*-acetylglucosamine-capped *O*-glycan (α GlcNAc) found in gland mucin suppresses *H. pylori* growth and motility by inhibiting catalytic activity of cholesterol α -glucosyltransferase (CHL α GcT), the enzyme responsible for biosynthesis of the major cell wall component cholesteryl- α -D-glucopyranoside (CGL). Here, the authors developed a polyclonal antibody specific for CHL α GcT and then undertook quantitative ultrastructural analysis of the enzyme's localization in *H. pylori*. They show that 66.3% of CHL α GcT is detected in the cytoplasm beneath the *H. pylori* inner membrane, whereas 24.7% is present on the inner membrane. In addition, 2.6%, 5.0%, and 1.4% of the protein were detected in the periplasm, on the outer membrane, and outside microbes, respectively. By using an *in vitro* CHL α GcT assay with fractionated *H. pylori* proteins, which were used as an enzyme source for CHL α GcT, the authors demonstrated that the membrane fraction formed CGL, whereas other fractions did not. These data combined together indicate that CHL α GcT is originally synthesized in the cytoplasm of *H. pylori* as an inactive form and then activated when it is associated with the cell membrane. This article contains online supplemental material at <http://www.jhc.org>. Please visit this article online to view these materials. (J Histochem Cytochem 59:98–105, 2011)

Keywords

cell wall, glycosyltransferase, *O*-glycan, *Helicobacter pylori*, immunocytochemistry

In the two decades since Marshall and Warren (1984) discovered *Helicobacter pylori* in the human stomach, it has been accepted that *H. pylori* is associated with pathogenesis of gastritis, peptic ulcer, gastric carcinoma, and gastric lymphoma of mucosa-associated lymphoid tissues (Peek and Blaser 2002). Although *H. pylori* infects more than half of the world's population (Bruce and Maarros 2008), most infections are asymptomatic, and only a fraction of individuals infected develop serious gastric diseases such as gastric ulcer and cancer, suggesting that the stomach harbors protective mechanisms.

H. pylori colonizes the surface mucin by binding blood group Lewis b and type H carbohydrates via the blood group

antigen-binding adhesin (BabA; Ilver et al. 1998; van de Bovenkamp et al. 2003). By contrast, *H. pylori* is barely detectable in gland mucin secreted from gland mucous cells

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such as mucous neck cells and pyloric gland cells, which contain unique *O*-glycans having terminal α 1,4-linked *N*-acetylglucosamine (α GlcNAc; Hidaka et al. 2001). Previously, we revealed that cholesteryl- α -D-glucopyranoside (CGL), a major cell wall lipid component of *H. pylori*, plays a critical role in bacterial survival and that α GlcNAc in gland mucin serves as a naturally occurring antibiotic against *H. pylori* by suppressing CGL biosynthesis, thus protecting gastric mucosa from *H. pylori* infection (Kawakubo et al. 2004).

Cholesterol α -glucosyltransferase (CHL α GcT), which we and others molecularly cloned from *H. pylori*, is the glycosyltransferase that transfers glucose from UDP-glucose to a carbon atom at the third position of cholesterol with an α 1,3-linkage (Lee et al. 2006; Lebrun et al. 2006). High conservation of CHL α GcT amino acid sequence among *Helicobacter* species suggests that this enzyme plays an important role in *Helicobacter* growth (Lee et al. 2008). Recently, we have also shown that the suppressive effect of α GlcNAc on CGL biosynthesis is due to its competition with CHL α GcT in vitro (Lee et al. 2006). These results suggest that *O*-glycans exhibiting α GlcNAc in the gland mucin interact with CHL α GcT expressed by *H. pylori*, thus protecting the gastric mucosa from infection. However, the exact localization of CHL α GcT in *H. pylori* remains to be determined.

In the present study, we developed polyclonal antibodies specific to *H. pylori* CHL α GcT and used them to demonstrate the subcellular localization of this enzyme in *H. pylori* using immunoelectron microscopy. Then, we carried out an in vitro CHL α GcT assay using fractionated proteins isolated from *H. pylori* to test which fraction contained an active form of the enzyme.

Materials and Methods

Bacterial Strain and Culture Conditions

Three standard *H. pylori* strains—26695 (ATCC 700392), NCTC 11637 (ATCC 43504), and J99 (ATCC 700824)—obtained from American Type Culture Collection (Manassas, VA) were cultured on trypticase soy agar with 5% sheep blood (Becton Dickinson; Franklin Lakes, NJ) for 3 to 4 days at 35°C in 15% CO₂. Bacteria were then inoculated in Brucella broth (Becton Dickinson) supplemented with 10% horse serum (Invitrogen; Carlsbad, CA) and cultured at 35°C in 15% CO₂. Bacteria were subsequently cultured in Mueller Hinton broth (Eiken Chemical; Tokyo, Japan) supplemented with 5.5% horse serum.

Antibody Production

Polyclonal antibodies directed to CHL α GcT of the *H. pylori* 26695 strain were generated using synthetic peptides as immunogens based on published sequences (Lee et al. 2006). Peptide sequences with high antigenicity were

Table 1. Synthetic Peptides Used to Develop Anti-CHL α GcT Antibodies

Peptide Name	Amino Acid Sequence	Amino Acid Residues
E17S	CEAIRDFKNNPHLFKTLS	373-389
V17F	CVVDSFKDTSNGTSMTAF	6-22
P17K	CPHVDNLGSEEEGYYNLK	40-56

C indicates cysteine residue at the *N*-terminus for coupling with keyhole limpet hemocyanin.

analyzed by database searches, and three peptides unique to *H. pylori* CHL α GcT, designated E17S, V17F, and P17K, were selected and synthesized (Table 1). The combined peptides were mixed with Freund's complete adjuvant and intradermally injected into two Japanese white rabbits, followed by eight additional injections with Freund's incomplete adjuvant. Sera were separately purified using each synthetic peptide conjugated to agarose gel (Pierce; Rockford, IL) for Western blot analysis to evaluate specificity as well as suitability for immunocytochemical staining. The experiment protocol was approved by the Institutional Animal Care and Use Committee at Medical and Biological Laboratories, Co., Ltd. (Nagoya, Japan).

Western Blot Analysis

Histidine (His)-tagged CHL α GcT recombinant protein expressed by *Escherichia coli* (*E. coli*) HB101 competent cells was purified using a HisTrap HP column (GE Healthcare; Buckinghamshire, UK) as described (Lee et al. 2006). In parallel, *H. pylori* 26695, NCTC 11637, and J99 strains harvested in Brucella broth with 10% horse serum as well as *E. coli* transformed with pTKNd6xH-CHL α GcT or pTKNd6xH harvested in LB medium were centrifuged and sonicated in 50 mM Tris-HCl (pH 7.4) containing 300 mM NaCl and 0.5% Triton X-100. After centrifugation at 16000 \times g for 10 min, the supernatant served as a source for whole protein. To evaluate specificity of antibodies, Western blot analysis was carried out on His-tagged CHL α GcT recombinant protein and whole proteins from the three *H. pylori* strains. Equal amounts of protein were separated on 14% SDS-PAGE and transferred to a PVDF membrane (Millipore; Billerica, MA). Nonspecific binding was blocked with 5% skim milk in TBS, and the membrane was incubated with the three anti-CHL α GcT antibodies (anti-E17S, -V17F, and -P17K; 1:1000 dilution each) or anti-His tag (1:500 dilution; Santa Cruz Biotechnology, Santa Cruz, CA) as a primary antibody. After rinsing with TBS, membranes were incubated with horseradish peroxidase (HRP)-conjugated goat anti-rabbit antibody (Jackson ImmunoResearch; West Grove, PA), and immunoreactions were detected using the ECL Western blotting detection system (GE Healthcare). Specificity of the anti-E17S antibody

was further evaluated by an antigen absorption test using various peptides: specifically, antibody was preincubated with 10 µg/ml of synthetic E17S peptide or the irrelevant peptides V17F and P17K and then immunoblotted using absorbed anti-E17S antibodies. Prestained protein ladder BenchMark (Invitrogen) was used as a size marker.

Immunocytochemistry

H. pylori 26695 strain was obtained from a 3-day culture in trypticase soy agar supplemented with 5% sheep blood (Beckton Dickinson). Bacteria were placed on MAS-coated glass (Matsunami glass; Osaka, Japan), air-dried, and fixed with 20% formalin neutral buffer solution, pH 7.4 (Wako; Osaka, Japan). Slides were incubated with 0.3% H₂O₂ in absolute methanol for 30 min and then permeabilized with 1% saponin dissolved in TBS containing 0.02% BSA. Because Western blot analysis described above revealed that the anti-E17S antibody was the most specific in recognizing recombinant and native CHLαGcT proteins, we used it for subsequent immunocytochemical analysis. After incubation with anti-E17S antibody (1:200 dilution) overnight at 4C, anti-rabbit immunoglobulins conjugated with HRP-labeled polymer Envision (DakoCytomation; Carpinteria, CA) were used as secondary antibody. HRP activity was visualized using 3,3'-diaminobenzidine (DAB; Dojindo, Kumamoto, Japan) with H₂O₂. As a control, anti-E17S antibody was omitted from the procedure, and no specific staining was found. Gram stain was carried out to observe the morphology of *H. pylori*.

Immunoelectron Microscopy

The *H. pylori* 26695 strain was washed with phosphate buffer and plated onto poly-L-lysine-coated specimen carrier (Leica; Vienna, Austria) (Sawaguchi et al. 2003). Samples were quick frozen using a high-pressure freezing system EM PACT (Leica) and fixed with 1% glutaraldehyde in acetone, followed by gradual warming from -85C to -3C by auto-freeze substitution using EM AFS (Leica). After washing with cold acetone, fixed cells were embedded in LR White Resin (London Resin Company; Berkshire, UK) and polymerized with UV at -20C for 24 hr. Immunogold staining was carried out as described (Kasai et al. 2003). Briefly, ultrathin sections with 0.1-µm thickness cut by ultramicrotome were mounted on nickel grids coated with formvar (Nacalai Tesque; Kyoto, Japan). To prevent non-specific reaction, sections were treated with 1% BSA in PBS and then incubated with anti-E17S antibody (1:50 dilution). After washing with PBS, sections were incubated with goat anti-rabbit IgG conjugated with 15 nm colloidal gold (1:50 dilution) purchased from BBIInternational (Cardiff, UK). After washing with PBS and distilled water (DW), sections were poststained with 2% uranyl acetate in

DW and coated with carbon and observed using conventional transmission electron microscopy (TEM), JEM-1200EX (JEOL; Tokyo, Japan), at an accelerating voltage of 80 kV. Control samples omitting primary antibody showed no specific signal.

Fractionation of Bacterial Proteins

The *H. pylori* 26695 strain harvested in Brucella broth with 10% horse serum was washed with 10 mM Tris-HCl (pH 7.5) containing 30 mM NaCl and suspended in 30 mM Tris-HCl (pH 7.5) containing 20% sucrose. EDTA was added to 1 mM and the sample incubated at room temperature with agitation. The pellet obtained by centrifugation was immediately suspended in 0.5 mM MgCl₂ and incubated at 4C with gentle agitation. After centrifugation at 8000 × g, the supernatant was carefully removed and used as the periplasmic fraction. The pellet was resuspended in 10 mM HEPES buffer (pH 7.0) and sonicated. To remove unbroken cells, the cell suspension was centrifuged at 3000 × g for 20 min and supernatant ultracentrifuged at 100 000 × g for 1 hr. The supernatant was collected as the cytoplasmic fraction, and the pellet was dissolved in 10 mM HEPES buffer containing 0.5% Triton X-100 and served as the membrane fraction. All samples contained a protease inhibitor cocktail (Roche; Mannheim, Germany). To validate the purity of fractionated proteins, the enzymatic activity of a cytoplasmic protein, malate dehydrogenase, was measured as described (Pitson et al. 1999; Appels and Haaker 1988). Briefly, mixtures of each fractionated protein with 0.1 M potassium phosphate buffer (pH 7.5) and 0.33 mM oxaloacetate were incubated at 25C for 5 min. After adding 143 µM NADH, the tube was mixed immediately and then incubated at 25C for 60 min. The decrease in absorbance at 340 nm was monitored. The molecular sizes of each fractionated protein were determined by Western blot analysis with anti-E17S antibody as described before.

In Vitro CHLαGcT Assay

The enzyme activity of CHLαGcT was measured using an in vitro CHLαGcT assay as described (Lee et al. 2006). Briefly, recombinant CHLαGcT was prepared by transforming *E. coli* JM109 cells with pTKNd6xH-CHLαGcT, followed by purification with a ProtinoR Ni-TED 2000 packed column (Macherey-Nagel; Duren, Germany) according to the manufacturer's protocol. The eluted protein was dialyzed against 50 mM HEPES buffer (pH 7.5) containing 15% glycerol and 5 mM dithiothreitol (DTT) by using an ultrafiltration unit (Millipore) at 4C and then used as an enzyme source for positive control. In parallel, the fractionated proteins isolated from *H. pylori* as described before were concentrated to 20 times, and 10 µl of the solution was incubated with 25 mM Tris-HCl buffer (pH 8.0) containing

3.7 μ M UDP-[14 C]glucose (111 000 dpm/ μ l), 500 μ M cholesterol, and 0.1% Triton X-100. In addition, 0.5 μ l of the recombinant CHL α GcT was used as well. After an 18-hr incubation at 30C, the reaction was stopped by adding 3 μ l of 1 N HCl, and then 10 volumes of ethyl acetate were added and vortexed. After brief centrifugation, radioactivity in the ethyl acetate (upper) layer, which contained α -glucosyl cholesterol, was measured using a liquid scintillation counter (Beckman Coulter; Brea, CA). In addition, these samples were also subjected to TLC with ethyl acetate layer. Briefly, the ethyl acetate layer was laid on preparative TLC (HPTLC LiChrospher Silica gel 60 F₂₅₄; Merck, Darmstadt, Germany) and separated in chloroform/methanol (85:15), and radioactive materials were visualized by fluorography (Typhoon 8600; GE Healthcare).

Statistical Analysis

All statistical data are presented as means \pm SD. Statistical analysis of enzyme activity in multiple comparisons was determined by one-way ANOVA with a Bonferroni multiple comparisons test using InStat 3 software (GraphPad Software; San Diego, CA). $p < 0.05$ was considered significant.

Results

CHL α GcT Antibody Specificity

The reactivity of anti-CHL α GcT antibodies raised against three CHL α GcT-specific peptides, E17S, V17F, and P17K, was verified by Western blot analysis using whole-protein lysates of *E. coli* transformed with CHL α GcT vector or vehicle alone. Both anti-E17S and anti-P17K antibodies reacted with recombinant CHL α GcT protein, whereas the immunoreaction of the anti-V17F antibody was less potent (Fig. 1A).

To determine whether anti-E17S and anti-P17K antibodies react with native CHL α GcT protein extracted from *H. pylori*, Western blot analysis using whole-protein lysates from 3 strains of *H. pylori* (26695, NCTC 11637, and J99) was carried out. The anti-E17S antibody clearly reacted with native CHL α GcT isolated from all *H. pylori* strains, but its immunoreactivity was less potent against NCTC 11637 and J99 strains than against the 26695 strain (Fig. 1A). By contrast, anti-P17K antibody did not react with native CHL α GcT proteins extracted from any *H. pylori* strain tested, although it reacted strongly with recombinant CHL α GcT protein (Fig. 1A).

Next, the specificity of anti-E17S antibody for CHL α GcT was confirmed using an antigen absorption test. Anti-E17S immunoreactivity was apparently decreased when antibody was preincubated with the synthetic E17S peptide used as its immunogen. On the other hand, a distinct CHL α GcT band was detected when anti-E17S antibody was preincubated with the irrelevant peptides V17F and P17K (Fig. 1B).

Last, the specificity of the anti-E17S antibody was confirmed by immunocytochemistry. *H. pylori* strain 26695 was positive for the anti-E17S antibody (Fig. 1C). These results collectively indicate that the anti-E17S antibody specifically recognized recombinant and native CHL α GcT proteins and was suitable for immunocytochemical analysis of *H. pylori*.

Subcellular Distribution of CHL α GcT in *H. pylori* as Determined by Immunoelectron Microscopy

To determine the subcellular localization of CHL α GcT in *H. pylori*, the 26695 strain labeled with immunogold was observed by TEM using the anti-E17S antibody. Immunoelectron microscopy revealed that CHL α GcT was largely cytoplasmic (Fig. 2A,B) and adjacent to the inner membrane (Fig. 2C,D) and that some CHL α GcT signals were also detected in cell wall components (Fig. 2E,F). To quantitate the number of gold particles representing CHL α GcT in *H. pylori*, cellular components were divided into five regions—cytoplasm, inner membrane, periplasm, outer membrane, and extracellular space—and gold particles were counted approximately up to 1000 in total. In this analysis, particles touching the inner or outer membranes were counted as residing in the inner membrane or outer membrane, respectively (Table 2). We found that 710 (66.3%) of 1070 gold particles indicative of CHL α GcT were located in the cytoplasm, and 264 (24.7%) particles were present on the inner membrane. Notably, 28 (2.6%) and 53 (5.0%) particles were also detected in the respective periplasm and on the outer membrane. Fifteen (1.4%) particles were present outside of bacteria.

Detection of CHL α GcT Activity in the Fractionated Protein of *H. pylori*

To test which subcellular fractions contained the active form of CHL α GcT, we carried out an in vitro CHL α GcT assay using fractionated *H. pylori* compartments. To validate the method of fractionation, we monitored the activity of malate dehydrogenase, which is a marker for cytoplasmic protein, and confirmed that the malate dehydrogenase activity was detected only in the cytoplasmic fraction (data not shown).

Thus, we carried out an in vitro CHL α GcT assay using 20-fold concentrated proteins. Western blot analysis revealed that molecular weights of each fractionated protein were the same (Fig. 3A). However, the enzyme activity was detected only in the membrane fraction (Fig. 3B). By contrast, such enzyme activity for CHL α GcT was not detected from other fractions, including the cytoplasm and periplasm. Similar results were also obtained when the samples were tested by

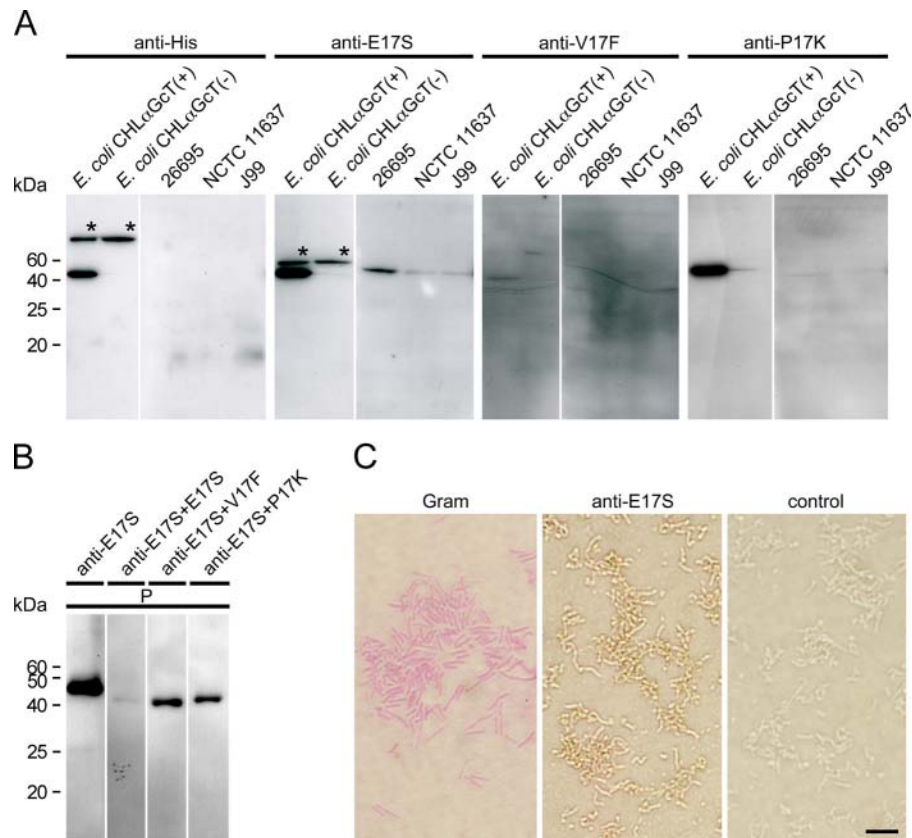


Figure 1. Characterization of three anti-CHL α GcT antibodies. (A) Immunoreactivity against various *Helicobacter pylori* strains (26695, NCTC 11637, J99) and *Escherichia coli* transformed by pTKNd6xH-CHL α GcT (CHL α GcT (+)) or pTKNd6xH (CHL α GcT (-)) reacted with anti-His (anti-His) and anti-CHL α GcT antibodies (anti-E17S, anti-V17F, and anti-P17K). Asterisks indicate extra bands for proteins other than CHL α GcT. (B) Inhibition of anti-E17S antibody using E17S peptide. Recombinant His-tagged CHL α GcT protein, which was preincubated with 10 μ g/ml of the synthetic peptides E17S, V17F, or P17K, was reacted with anti-E17S antibody. P, His-purified cell lysate prepared from *E. coli* transformed by pTKNd6xH-CHL α GcT used for a positive control. (C) Gram staining and immunocytochemistry using anti-E17S antibody against *H. pylori* strain 26695. Control indicates secondary antibody alone. Bar = 5 μ m.

TLC, and the positive signal indicative of CGL was detected only in the membrane fraction (Fig. 3C).

Discussion

CHL α GcT is the glycosyltransferase that forms CGL, a major cell wall glycolipid of *Helicobacter* species, including *H. pylori*. CGL was originally discovered as one of three major cholesteryl glucosides (CGs) that account for 25% of total cell wall lipids of *Helicobacter* species, including *H. pylori*, whereas other CGs include cholesteryl-6-*O*-tetradecanoyl- α -D-glucopyranoside (CAG) and cholesteryl-6-*O*-phosphatidyl- α -D-glucopyranoside (CPG; Hirai et al. 1995). Recently, Wunder et al. (2006) showed that CGL is required for *H. pylori* to evade the host's immune response. In addition, we have shown that CGL is critical for *H. pylori* survival, and CGL biosynthesis by *H. pylori* is significantly suppressed when microbes are cultured with α GlcNAc (Kawakubo et al. 2004). We have also

demonstrated that CHL α GcT activity is inhibited by α GlcNAc in vitro (Lee et al. 2006; Lee et al. 2008). Despite its important roles of CGL on *H. pylori* survival, the expression and activation of CHL α GcT in *H. pylori* have not been demonstrated so far.

In this study, we first developed a polyclonal antibody against CHL α GcT by immunizing the same rabbit with a cocktail of three different peptides specific for CHL α GcT. One technical benefit of our antibody production strategy is to reduce the number of animals used for immunization. Sera were then purified with each of the three peptides using affinity columns, and immunoreactivity to recombinant CHL α GcT protein and CHL α GcT isolated from *H. pylori* strains was evaluated. Anti-E17S antibody reacted with both recombinant and native CHL α GcT proteins (see Fig. 1A). An extra band with anti-E17S antibody found in both protein lysates of *E. coli* CHL α GcT (+) and *E. coli* CHL α GcT (-) was most possibly caused by the cross-reactivity of the anti-E17S antibody with unidentified protein

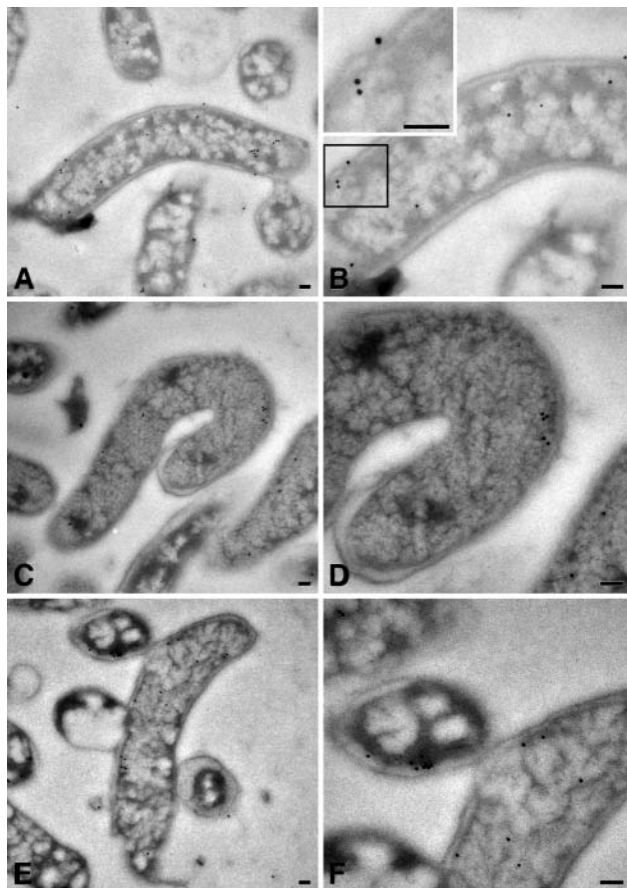


Figure 2. Ultrastructural localization of CHL α GcT in *Helicobacter pylori* strain 26695. Gold particles indicate CHL α GcT immunoreactivity. B, D, and F are enlarged images of A, C, and E, respectively. Bar = 0.1 μ m.

endogenously expressed by *E. coli*, because an antigen absorption test using 10 μ g/ml and 100 μ g/m of the synthetic E17S peptide revealed that this extra band and the band indicative for CHL α GcT were diminished in a dose-dependent manner (unpublished data). Interestingly, the reactivity to CHL α GcT protein from *H. pylori* NCTC 11637 and J99 strains was much weaker than that from the 26695 strain. We thus compared the amino acid sequence of CHL α GcT of the *H. pylori* 26695 strain with those of the NCTC 11637 and J99 strains. No amino acid substitution was found in the E17S sequence of NCTC 11637, whereas two amino acid substitutions were found in the E17S sequence of J99 (Suppl. Fig. S1). Thus, weak reactivity for the anti-E17S antibody to J99 and NCTC 11637 strains might be caused by the difference of the primary structure of CHL α GcT for J99 and the weak expression level of the CHL α GcT protein for NCTC 11637, respectively. On the other hand, anti-P17K antibody reacted with recombinant CHL α GcT produced in *E. coli* but not with native protein

Table 2. Number (%) of Gold Particles Indicative of CHL α GcT in *Helicobacter pylori*

	Inner Cytoplasm	Membrane	Periplasm	Outer Membrane	Outside of Cell	Total
	710 (66.3)	264 (24.7)	28 (2.6)	53 (5.0)	15 (1.4)	1070 (100)

from *H. pylori*. It is not obvious why anti-P17K antibody could recognize only recombinant protein. It might be possible that the P17K sequence of the native CHL α GcT was cryptic due to its secondary structure. By contrast, anti-V17F antibody reacted with neither recombinant nor native CHL α GcT (see Fig. 1A). It is not clear why anti-V17F antibody was less potent compared with other antibodies, despite that the V17F sequence of recombinant and native proteins was the same (Suppl. Fig. S1). The binding site for anti-V17F antibody on both proteins might be masked due to the secondary structure of CHL α GcT. Further study will be necessary to address these problems.

After confirming the specificity for anti-E17S antibody by a peptide inhibition test (see Fig. 1B), we carried out immunocytochemistry and showed that the CHL α GcT protein was most abundant in the cytoplasm, particularly beneath the inner cell membrane (see Fig. 2). Moderate and small amounts of enzyme were also expressed in the inner cell membrane and the outer membrane/periplasm, respectively. Then, we measured the enzyme activity using fractionated proteins and revealed that the active form of CHL α GcT was present only in membrane fraction, even though the molecular sizes of each fractionated protein were the same (see Fig. 3). Although this result is consistent with the data reported before (Lebrun et al. 2006), it seems to be paradoxical because the majority of CHL α GcT found in the cytoplasm lacked its activity, whereas the membrane fraction contained the active form of CHL α GcT despite a lack of transmembrane domain. At this moment, it is unknown why CHL α GcT is inactive in the cytoplasm and how this enzyme is converted to an active form in the membrane. These questions will be important future issues to be addressed.

It is noteworthy that such an expression pattern of CHL α GcT is similar to that of *H. pylori* α 1,3/4-fucosyltransferase (α 1,3/4-FucT), the enzyme responsible for the biosynthesis of Lewis antigens attached to lipopolysaccharide on the cell wall (Ge et al. 1997). Interestingly, both α 1,3/4-FucT and CHL α GcT lack a transmembrane domain (Ge et al. 1997; Lee et al. 2006), suggesting that both are soluble proteins. Recently, Ma et al. (2003) suggested that α 1,3/4-FucT could associate with the inner membrane through C-terminal regions rich in positively charged and hydrophobic residues functioning as a membrane

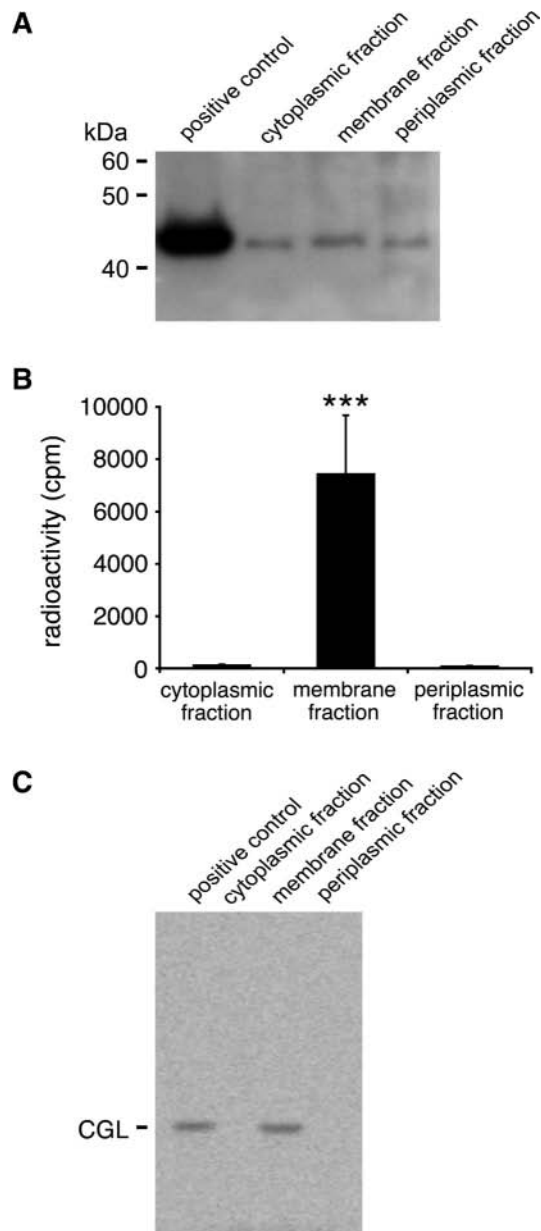


Figure 3. In vitro CHL α GcT assay using fractionated proteins of *Helicobacter pylori*. (A) Molecular weights of each fractionated protein were examined using Western blot analysis with anti-E17S antibody. (B) Radioactivity in each fractionated protein of *H. pylori* incubated with a solution containing UDP-[14 C]glucose and cholesterol was measured by a liquid scintillation counter. Radioactivity of the membrane fraction is significantly higher than cytoplasmic and membrane fractions (***) ($p < 0.001$). Data are presented as means \pm SD. (C) In parallel, these samples were analyzed by TLC. The product indicative of cholesteryl- α -D-glucopyranoside (CGL) was formed only when the membrane fraction was used as an enzyme source.

anchor. These data are consistent with our present observation that the active form of CHL α GcT is expressed in the cell membrane. Recently, we found that the *N*-terminal

region of CHL α GcT, which contains hydrophobic residues, is critical for its function because the truncated form of CHL α GcT lacking the *N*-terminal 18 amino acid residues was functionally inactive (YI and MFukuda, personal communication, 2008). It might be possible that CHL α GcT was associated with the cell membrane via the *N*-terminal hydrophobic residues.

Thus, these combined results for the subcellular localization of CHL α GcT, as demonstrated by immunoelectron microscopy, and its enzyme activity, as revealed by the in vitro CHL α GcT assay, indicated that CHL α GcT was originally synthesized in the cytoplasm of *H. pylori* as an inactive form and then activated when it was associated with the cell membrane. It is also noted that the membrane fraction thus obtained in the present study could contain protein fractions derived from both the inner and outer membranes. Although the gold particles indicative for CHL α GcT were more frequently found on the inner membrane compared with the outer membrane, it remains to be addressed which membrane is the major one associated with the active form of CHL α GcT. Future study should address this problem.

In summary, the present study developed a CHL α GcT-specific polyclonal antibody and then demonstrated that active form of CHL α GcT was associated with the cell membrane. Because CGL is critical for *H. pylori*'s survival, the present study provides useful information for developing novel and potentially safe therapeutic agents targeting the active form of CHL α GcT to treat *H. pylori* infection in humans.

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Declaration of Conflicting Interests

The author(s) declared no potential conflicts of interest with respect to the authorship and/or publication of this article.

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