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tenuate ghrelin production in the stomach and may consequently reduce plasma ghrelin concentrations in *H. pylori*positive subjects. To examine this hypothesis, we next examined the specific changes of gastric ghrelin production in association with *H. pylori* infection.

Ghrelin mRNA in gastric mucosa is lower in H. pyloripositive subjects

In an effort to examine the effects of *H. pylori* infection on ghrelin production in the gastric mucosa, we compared gastric ghrelin mRNA expression levels between H. pyloripositive and -negative subjects using real-time quantitative RT-PCR using corpus mucosa because gastric ghrelin is predominantly produced in the corpus rather than in the antrum (10). As shown in Fig. 2, gastric ghrelin mRNA levels of corpus mucosa were significantly lower in H. pylori-positive patients than H. pylori-negative controls. A similar difference was also significantly observed between patients with chronic gastritis alone and H. pylori-negative controls (data not shown). These results suggest that the expression of ghrelin mRNA in the gastric mucosa is markedly decreased in association with H. pylori infection. It is important to note that the average of gastric ghrelin mRNA expression levels in H. pylori-positive subjects was less than one 45th of that in H. pylori-negative controls. Moreover, as shown in Fig. 3, plasma ghrelin concentrations were in parallel with the ghrelin mRNA expression levels in H. pylori-positive subjects. Taken together, these results suggest that the attenuation of the ghrelin production in the gastric mucosa accounts for the decrease in the plasma ghrelin concentrations in *H. pylori*-positive individuals.

Ghrelin-producing cells in the gastric mucosa are lower in H. pylori-positive subjects

As an independent test to examine the effect of H. pylori infection on gastric ghrelin production and its relation with plasma ghrelin concentrations, we next investigated the numbers of ghrelin-producing cells of the corpus mucosa in H. pylori-positive and -negative subjects. For this purpose, biopsy samples taken from gastric mucosa were immunostained using an antighrelin polyclonal antibody. Immunoreactive cells were seen in the lower half of fundic epithelial glands as described previously (10). No immunoreactivity was detected in the tissue when control serum was used for staining (data not shown). Immunoreactivity was concentrated in the basal cytoplasm of the positive cells as shown in Fig. 4, A and B. As shown in Fig. 4C, the number of ghrelin-positive cells in the gastric mucosa of H. pyloripositive individuals was significantly lower than those of *H. pylori*-negative individuals. Furthermore, the numbers of ghrelin-positive cells in the gastric mucosa fell significantly in accompaniment to the decrease in plasma ghrelin concentrations in H. pylori-positive subjects (Fig. 5). These results reinforce that the attenuation of the gastric ghrelin production caused by H. pylori infection accounts for the decrease in the plasma ghrelin concentrations in H. pylori-positive individuals.



FIG. 4. Immunostaining of ghrelin in the fundic glands. A, Normal epithelium without *H. pylori* infection. B, Inflammatory gastric mucosa with *H. pylori* infection. Ghrelin-producing cells were abundant in the mucosa from *H. pylori*-negative subject, whereas they were seldom observed the mucosa from *H. pylori*-positive subject (magnification, $\times 200$). C, Numbers of the immunoreactive cells in *H. pylori*-positive and -negative subjects. Immunoreactive cells in the gastric mucosa were counted and presented as the number of positive cells per branch of oxyntic glands. Values are expressed as the mean \pm SE. The numbers of the immunoreactive cells in the gastric mucosa were in the *H. pylori*-negative controls (P < 0.0001 by unpaired two-tailed *t* test).



FIG. 5. Comparison of the frequencies of the immunoreactive cells in *H. pylori*-positive subjects among different levels of plasma ghrelin groups. Values are expressed as the mean \pm SE. There was a correlation between plasma ghrelin concentrations and the frequencies of the immunoreactive cells with a correlation coefficient value of 0.41 (P < 0.0001). The numbers of the positive cells were significantly lower in the low and middle ghrelin groups than those in the high ghrelin group (low vs. high: P = 0.018; middle vs. high: P = 0.038 by variance (ANOVA) based on Fisher's protected least significant difference test).

Plasma ghrelin concentrations are associated with serum pepsinogen concentrations in H. pylori-positive subjects

In the last sets of examinations, we further attempted to demonstrate the association between *H. pylori* infection and plasma ghrelin concentrations. Because *H. pylori* infection first induces gastric atrophy in its pathological course, we compared plasma ghrelin concentration with serum pepsinogen concentrations in *H. pylori*-positive patients. Pepsinogen I and pepsinogen II differ in their location in the stomach. Both are located in the chief and mucous neck cells of the oxyntic gland mucosa in the gastric corpus, but only pepsinogen II is present in the gastric antrum. A pepsinogen I to II ratio less than 3 is considered to be a reliable marker for severe atrophic gastritis (19, 20). Therefore, the plasma ghrelin levels in *H. pylori*-positive patients were compared with serum pepsinogen concentrations and serum pepsinogen I to II ratios. As shown in Fig. 6, serum levels of pepOsawa et al. • Ghrelin and Helicobacter pylori Infection

sinogen I and the ratio of pepsinogen I to II fell significantly as plasma ghrelin concentrations decreased, indicating the positive association between plasma ghrelin and pepsinogen I concentrations as well as pepsinogen I to II ratios in *H. pylori*-positive patients. Collectively, these results reveal that plasma ghrelin concentrations are associated with the progression of gastric atrophy.

Discussion

Plasma ghrelin levels have been associated with several clinical factors including BMI, food intake, and serum insulin levels (9, 21, 22). Although ghrelin-producing endocrine cells have been found mainly in the oxyntic mucosa of the stomach, ghrelin is also released from other tissues including small and large intestines, lung, kidney, the nucleus of the hypothalamus, and A cells of the pancreatic islet (23, 24). In fact, plasma ghrelin concentrations in gastrectomized patients still remain about one third of those in normal subjects (6). Thus, it is important to clarify which organ primarily influences changes in plasma ghrelin concentrations in each disease. In this study, we have demonstrated that plasma ghrelin concentrations are influenced by H. pylori infection. In particular, we focused on the gastric mucosa to better understand the effects of H. pylori infection on the alteration of ghrelin expression. The expression levels of ghrelin mRNA and the numbers of ghrelin-producing cells in the gastric mucosa were much lower in patients with H. pylori infection. Plasma ghrelin concentrations correlated with the gastric ghrelin mRNA as well as the frequency of ghrelin-immunoreactive cells in the gastric mucosa. Finally, we compared plasma ghrelin concentrations with serum pepsinogen levels, a marker for gastric atrophy. Plasma ghrelin concentrations in H. pylori-positive patients correlated with serum pepsinogen I concentration as well as pepsinogen I to II ratio. In addition, we demonstrated that groups with histologically higher degrees of gastric atrophy in the H. pylori-positive subjects tend to have lower plasma ghrelin concentrations (data not shown). These findings strongly suggest that the reduction of ghrelin-producing cells in the gastric mucosa by H. pylori infection results in the lower plasma ghrelin concentration in *H. pylori*-positive patients.

FIG. 6. Comparison of plasma ghrelin levels and serum pepsinogen levels in *H. pylori*-infected subjects. Values are expressed as the mean \pm SE. Both serum pepsinogen I levels and pepsinogen I to II ratio significantly fell with the decrease of plasma ghrelin levels (A and B). An ANOVA based on Fisher's protected least significant difference test was used.



Our current data are consistent with the report of Nwokolo's group (17) that plasma ghrelin concentrations increased after *H. pylori* eradication. In their study, however, gastric ghrelin before and after *H. pylori* eradication was not measured. Moreover, effects of a change in BMI before and after *H. pylori* eradication on plasma ghrelin concentrations could not be excluded. In addition, the number of the enrolled subjects in their study was relatively small (10 subjects). Therefore, our present study expanded their observations by enrolling many more subjects with comparable BMI and showing a direct association between *H. pylori* infection and lower gastric ghrelin production.

On the other hand, Gokcel's group compared plasma ghrelin concentrations between *H. pylori*-positive and -negative subjects, and, opposed to our results, they found no differences. Although their study design was similar to ours, they did not provide any data on the gastric atrophy or gastric ghrelin production in their subjects. It is, therefore, only a speculation, but the discrepancy of the results may be due to the different features of gastric atrophy between Western and Japanese populations including disease frequency and severity. The earlier age of acquiring *H. pylori* infection in Japan, compared with Western countries, may also explain the high incidence of atrophic gastritis in Japanese adults and lower concentrations of plasma ghrelin concentrations as well.

It would be intriguing to clarify how a persistent decrease in plasma ghrelin concentration influences human growth and body weight. Recently several reports demonstrated that H. pylori-positive children have a high incidence of growth retardation (25, 26). H. pylori is acquired early in life in most of the developing world. Together with our results, the decrease of plasma ghrelin levels accompanied by H. pylori gastritis may have considerable influences on growth retardation in childhood. In our study, the plasma ghrelin levels in H. pylori-positive subjects were lower than H. pylori-negative subjects, even in patients with mild atrophic changes, implying that even mild gastric inflammation by H. pylori infection in children may reduce the production of gastric ghrelin. Further study in children including plasma ghrelin levels, degree of atrophy in the stomach, and presence of H. pylori infection may clarify these relationships.

In conclusion, our study indicates that plasma ghrelin concentrations are influenced by the presence of chronic gastritis in association with *H. pylori* infection. Decreases in gastric ghrelin production may account for lower concentrations of plasma ghrelin in *H. pylori*-positive individuals. These observations provide novel insights for understanding the physiological function of ghrelin and its relation to various diseases.

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宿主サイドからのアプローチ(宿主側因子)

H. *pylori* 感染における血漿グレリン濃度と 胃粘膜グレリン発現量

Plasma and gastric ghrelin levels in subjects with Helicobacter pylori infection

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 Key works
 : 血漿グレリン, 胃粘膜グレリン, H. pylori 感染, 萎縮性胃炎

はじめに

成長ホルモン(GH)の分泌は、これまで視床下 部ホルモンである成長ホルモン放出ホルモン (GHRH)およびソマトスタチンによりそれぞれ 促進性、抑制性に制御されていると考えられて いた. 1996年にヒト, ラット, ブタの視床下部 と下垂体でのGHS-Rの存在と1次構造が明ら かにされ", それまで存在が疑問視されていた内 在性リガンドの探索が、国内外で競って行われ た. 1999年に児島, 寒川ら²は新しい成長ホルモ ン分泌促進ペプチド、グレリンを発見・構造決 定に成功した.彼らはGHS-Rを安定発現する 培養細胞を調製し、細胞内カルシウムイオン濃 度の上昇を指標とするアッセイ系を確立した. これを用いて胃抽出物中に GHS-Rに対する非 常に強い活性分画を見いだし、グレリン(ghrelin)と名づけた.現在では、哺乳類以外の動物で もその構造が決定され、グレリンが進化上、保 存されてきた機能的に重要なホルモンであるこ とがうかがえる. グレリンは成長ホルモン分泌 促進作用, 強力な摂食促進作用³, 消化管運動促 進作用4, 胃酸分泌促進作用9を有し, また視床 下部に作用して体重増加を促す". 血漿グレリン 濃度は食前に上昇し、食後に低下する.このペ プチドは成長や脂肪組織量の制御に関与し,短期の食欲促進や長期的な体重の制御物質でもある^{4,6,7}.

1. グレリンの構造

単離されたグレリンのペプチドは28アミノ酸 残基よりなり,興味深いことに3番目のセリン 残基の側鎖は炭素数8個の脂肪酸,オクタン酸 によってエステル化されていた.このオクタン 酸によるアシル化修飾はグレリンの生物活性発 現に重要であり,アシル化修飾のないペプチド 結合鎖のみでは全くGH分泌作用は示さないと 報告されてきた.しかしながら,最近では非ア シル化グレリンは,アシル化グレリンと反対の 作用を示すという内容が我が国の学会で報告さ れ,今後の詳細な報告を待ちたい.

2. グレリンの摂食促進作用

グレリンは脳内の視床下部弓状核のニューロ ンでも産生され、グレリン受容体は脳の様々な 部位で発現しているので、種々の中枢生理機能 に関与していると考えられていた、グレリンを ラットの脳室内に投与すると、摂食が促進され て体重増加をもたらし、この効果は、遺伝的に 成長ホルモンを欠くラットにもみられる、逆に

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グレリン抗体を投与すると, 摂食が強く抑制さ れる. グレリンの脳室内投与後に, ニューロン 活性化の指標となる Fos 蛋白質の発現が, 神経 ペプチドY(NPY)産生ニューロンやアグーチ関 連蛋白質(AGRP)産生ニューロンを含む摂食 制御に重要な複数の脳領域で起こる. NPYや AGRPに対する抗体を投与すると, グレリン誘 発性の摂食行動は消失する³⁾. グレリンはNPY 遺伝子の発現を増強し, レプチンで誘発される 摂食低下を抑えることから, グレリンとレプチ ンは摂食行動に関して拮抗的に作用するといえ る. このように, グレリンは摂食行動の生理的 信号物質であり, 成長ホルモンの分泌と摂食を 促進して成長を制御する機能をもつ.

3. グレリンの分布

ヒトのグレリンは、ペプチドとmRNAともに 胃に最も多く,腸,膵臓,視床下部,胎盤,腎 臓などでも産生される. 胃グレリンは胃体部の 内分泌細胞で主に産生される。胃体部には、ヒ スタミンとウログアニリンを産生する enterochromaffin-like(ECL)細胞, ソマトスタチンを 産生するD細胞,セロトニンを産生する enterochromaffin (EC)細胞に加え、膵臓でグルカゴン を産生するA細胞と形態が類似しているがグル カゴンを含まず、また産生物質が不明なことか らA-like細胞(またはX細胞)と呼ばれていた内 分泌細胞の4種類がある.これらの既知の細胞 が産生する物質とグレリンとの二重免疫染色の 結果, グレリン産生細胞は前三者の内分泌細胞 とは異なっていた. グレリン抗体を用いた免疫 電顕により、グレリンは直径 120nm でほぼ均一 なサイズの電子密度の高い分子顆粒に貯蔵され ていることが明らかになった⁸. グレリン細胞は 過去の消化管内分泌細胞の命名法に準じ、Gr細 胞と呼ばれている.

4. 血漿グレリン濃度

新生児では血漿グレリン濃度は性差や体重に よって影響されないが⁹,小児や成人の肥満者で は正常体重者や痩せ型の人より低下している¹⁰. 血漿グレリンの大半は胃で産生されるので,慢 性胃粘膜障害では、その産生が障害される可能 性がある. H. pylori 感染は胃十二指腸潰瘍、慢 性萎縮性胃炎に関連しているので、この菌が胃 グレリン産生だけでなく血漿グレリン濃度に影 響を及ぼすかどうか検討することは重要である.

欧州では Nwokolo ら¹¹⁾は,血漿グレリン濃度 は H. pylori 除菌後に上昇すると報告し,一方, Gokcel ら¹²⁾は H. pylori 感染の有無は血漿グレリ ン濃度に影響しないと報告した.しかし,これ までの報告では, H. pylori 感染胃粘膜のグレリ ン発現について検討されてはこなかった.

5. *H. pylori* 感染による胃グレリンの 産生障害

日本人ではH. bylori感染に伴う重度の萎縮性 胃炎がみられることが多い. 著者らは、内視鏡 生検材料を用いて H. pylori 感染胃粘膜のグレリ ンmRNA量とペプチド発現,更に血漿グレリン 濃度を比較検討した13).体重の影響を最小限に するため,正常肥満指数(18.5-25)を有するH. pylori 陽性者 110 人と陰性者 50 人において空腹 時血漿グレリン濃度を測定し、更に胃体部大彎 粘膜から RNA を抽出して real-time RT-PCR法 にて胃グレリンmRNA量を測定した. また, H. pylori陽性者の血漿グレリン濃度を高グレリン群 (≥150 fmol/ml, n=40), 中グレリン群(70-150 fmol/ml, n=36), 低グレリン群 (<70 fmol/ ml, n=34)に分類し、それらを慢性胃炎の萎縮 の程度、胃グレリンmRNA量、胃グレリン陽性 細胞数と比較した.

H. pylori陽性者の血漿グレリン濃度および胃 グレリンmRNA量はH. pylori陰性者より低下し ていた(図1,2). H. pylori陽性者では血漿グレ リン濃度が低い症例ほど胃グレリンmRNA量の 低下(図3)と組織学的に強い萎縮を認め、血清 ペプシノゲンIおよびI-II比(図4)も有意に低下 していた.また、グレリン陽性細胞数はH. pylori陽性粘膜で減少し(図5)、血漿グレリン濃度 が低い症例ほどその低下は顕著となった.

 ドグレリンとの関連で Cummings らⁿは、胃切

 除後の血漿グレリン濃度が約 1/3 に低下すると

 報告した、著者らの検討では H. pylori 慢性胃炎





H. pylori 陽性者の血漿グレリン濃度は *H. pylori* 陰性者と比較して有意に低下していた(128±8 vs 194±15 fmol/ml).



図2 胃グレリン mRNA 量¹³⁾

real-time RT-PCR 法にて胃体部粘膜内グレリン mRNAを定量した. *H. pylori* 陽性者の胃粘膜内グ レリン mRNA 量は *H. pylori* 陰性者と比較して著明 に低下していた.



図3 *H. pylori* 感染胃粘膜グレリン mRNA 量¹³ *H. pylori* 陽性者では血漿グレリン濃度が低い症例ほど 胃粘膜グレリン mRNA 量は低下していた.

では血漿グレリン濃度は H. pylori 陰性者に比べ て約 2/3 に低下していた.これは胃グレリン産 生が H. pylori 感染による胃粘膜の炎症や萎縮と 関連して血漿グレリン濃度に大きく影響するこ とを示している.逆に,除菌治療によって胃グ レリン産生が回復するのかもしれない.萎縮性 変化との検討では血清ペプシノゲンの低下と血 漿グレリン濃度および胃グレリン mRNA 産生の

低下が相関しており,胃体部粘膜の組織学的な 萎縮の程度とも相関していた.更に萎縮のない *H. pylori*陽性症例と*H. pylori*陰性症例の比較で は,陽性症例で血漿グレリン濃度の有意な低下 がみられ,炎症だけでも胃グレリン産生は低下 すると考えられた.

欧米では*H. pylori* 感染は血漿グレリン濃度に 影響しないとの報告¹²⁾もみられる.しかし,研









究対象症例が少なすぎる点,肥満指数によるグ レリン濃度の変化を考慮していない点,*H. pylori*感染の胃粘膜への影響を検討していない点 において,その結果には疑問が残る.

6. 今後の課題

以前からH. pylori 感染児の発育遅延が報告さ

れている¹⁴. *H. pylori* は発展途上国では幼児期に 感染している. グレリンは強力な成長ホルモン 分泌促進作用を有しており,小児における胃グ レリン産生が *H. pylori* 感染によって影響を受け ている可能性が高く,今後の検討課題である. また,最近グレリンの粘膜防御作用が報告され ている¹⁵⁾. 逆に *H. pylori* 感染によるグレリン産 生低下が胃粘膜防御機能低下に関連している可 能性も否定できない. 更にグレリンは消化管運 動とも関連しており,運動機能異常を来す消化 管疾患についても検討が必要である.

おわりに

グレリンの発見により胃が消化機能だけでな く,成長ホルモンの分泌調節や摂食調節にも機 能していることが明らかになった.更に H. pylori感染は胃グレリン産生を低下させ、最終的に 血漿グレリン濃度を低下させる.血漿グレリン 濃度は、胃粘膜内グレリンmRNA量と正の相関 性を示しており、また組織学的および血清学的 検討から慢性胃炎、特に萎縮の程度を反映して いた. 酮文

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Histamine-2 receptor expression in gastric mucosa before and after Helicobacter pylori *cure*

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SUMMARY

Background: Helicobacter pylori infection prevents the occurrence of the tolerance phenomenon of Histamine-2 (H2) receptor antagonists. Gastro-esophageal reflux disease develops in some cases with the restoration of acid secretion after *H. pylori* eradication therapy.

Aim: To clarify the mechanisms of H2 receptor restoration after the eradication of *H. pylori* on parietal cells. *Methods:* We enrolled 80 consecutive asymptomatic male patients with *H. pylori* infection, having chronic gastritis with or without the presence of peptic ulcers. Biopsy specimens from the greater curvatures at the mid-corpus of the stomach were obtained endoscopically from all subjects before and 12 weeks after the eradication of *H. pylori*. Degrees of gastric atrophy were evaluated by serum pepsinogen levels. The amounts of mRNA expression of H2 receptor were evaluated in each subject's gastric mucosa by real time reverse transcriptase-polymerase chain reaction (RT-PCR).

Results: H2 receptor mRNA expression levels significantly correlated with serum pepsinogens I and II ratios. The expression level of H2 receptor mRNA was lower in subjects with hypergastrinemia. The median expression level of H2 receptor after *H. pylori* eradication was threefold greater than prior to treatment. In addition, its restoration became more pronounced in subjects with severe gastric atrophy. However, a comparatively low restoration of H2 receptor mRNA was found in subjects with hypergastrinemia.

Conclusions: H2 receptor mRNA levels decrease with the progression of gastric atrophy induced by *H. pylori* infection, and are restored after *H. pylori* eradication. Such expression levels of H2 receptor may explain a part of the tolerance phenomenon to H2 receptor antagonists.

INTRODUCTION

Histamine-2 receptor antagonists (H2RAs) show a potent and quick acid-suppressing effect, especially during the nocturnal period.¹⁻³ However, the antisecretory activity of H2RAs was reported to decrease

during continuous administration, leading to rebound hypersecretion of gastric acid after withdrawal of H2RAs.³⁻¹¹ This attenuation of the antisecretory activity of H2RAs has been described as tolerance. In patients with gastro-oesophageal reflux disease (GERD), such rebound hypersecretion may increase acid reflux, making it difficult to stop treatment.^{12,13}

Helicobacter pylori infection directly and indirectly influences the physiological functions of gastric parietal cells.¹⁴ Treatment with proton pump inhibitors induces tolerance to H2RAs in *H. pylori*-negative patients.¹⁵ On the other hand, *H. pylori* infection prevents the tolerance phenomenon of H2RAs.¹⁶ Therefore, *H. pylori* infection has a strong effect on tolerance to H2RAs. However, the

Abbreviations: RT-PCR, reverse transcriptase-polymerase chain reaction; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; GERD, gastro-oesophageal reflux disease; H2, Histamine-2; H2RAs, Histamine-2 receptor antagonists..

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effect of H2 receptor expression on the restoration of acid secretion after *H. pylori* eradication has not been clearly identified. We therefore conducted this study to investigate the relationship between expression levels of H2 receptor in the oxyntic gland before and after eradication therapy. To this end, we evaluated H2 receptor mRNA expression levels in detail using quantitative real time reverse transcriptase-polymerase chain reaction (RT-PCR) methods. We report here that the expression levels of H2 receptor mRNA decreases with the progression of gastric atrophy in *H. pylori*-infected subjects, and restoration after *H. pylori* cure is threefold greater than the levels prior to treatment.

MATERIALS AND METHODS

Patients

Of those who participated in the gastric cancer surveillance program in Tochigi, Japan, 85 consecutive men with H. pylori infection had received eradication therapy; treatment was successful in 80 subjects (mean age \pm S.E.; 49.2 \pm 0.5 years) enrolled in the present study. Indications of the eradication therapy in these patients included chronic gastritis alone (four patients), chronic gastritis accompanied by either adenoma, a family history of gastric cancer, hyperplastic polyp, severe atrophic gastritis, chronic urticaria, or gastric ulcer (66 patients), duodenal ulcer (10 patients). All subjects were clinically stable at the time of evaluation and had no history of eradication therapy before the study. Written informed consent was obtained from the participants in accordance with the Declaration of Helsinki and its later revision. The Ethics Committee of the Jichi Medical School, Japan, approved this study.

Eradication therapy for H. pylori-infected subjects

A triple regimen, composed of lansoprazole 30 mg twice daily, clarithromycin 200 mg twice daily, and amoxicillin 750 mg twice daily, was given for 7 days following endoscopic examination. No other medications were given during the course of the study.

Specimens

Five adjacent biopsy specimens from the greater curvatures at the mid-corpus of the stomach as well

as five from the antrum were obtained endoscopically from all subjects. One biopsy specimen from the corpus of the stomach and one from the antrum were cultured individually to assess the presence of H. *pylori* infection. Three biopsy specimens from the corpus and three from the antrum were immediately snap-frozen and stored in liquid nitrogen for later use. The remaining corpus and antral specimens were fixed and stained with hematoxylin and eosin, and Giemsa. Histological assessments were performed by a single observer (H.Os.). The presence of H. *pylori* infection was diagnosed by either positive bacterial culture or positive Giemsa staining.

Blood sample

Before and 12 weeks after the eradication of *H. pylori*, venous blood samples after fasting overnight were obtained for the determination of serum gastrin and serum pepsinogens I and II levels.

RNA extraction and RT-PCR

Total RNA was isolated from biopsy specimens with ISOGEN (Nippon Gene, Tokyo, Japan), and 2 μ g of total RNA was reverse-transcribed with random nanomers and reverse transcriptase (TOYOBO, Osaka, Japan), which includes a DNase incubation step. The expression level of H2 receptor mRNA was evaluated using a realtime quantitative RT-PCR method with an ABI 7700 sequence detector system (PE Applied Biosystems, Foster City, CA, USA). The sense primer for H2 receptor was 5'-CCACCATCAGGGAGCACAA-3' and the antisense primer was 5'-AGGGAAACCAGCAGATGATGAA-3'. The reaction mixture was prepared according to the manufacturer's protocol using TaqMan PCR kits (PE Applied Biosystems). The reactions also contained target hybridization H2 receptor probe labeled with a reporter fluorescent dye, 6-carboxyfluorescein, at the 5' end (5'-CACAGTGACACTGGCCGCCGTC-3'). Thermal cycling conditions for all reactions included 50 °C for 2 minutes and 95 °C for 10 minutes, followed by 40 cycles of 15 seconds. of denaturation at 95 °C and 1 minute of annealing and extension at 60 °C.

As a control, the mRNA was also subjected to realtime quantitative RT-PCR for the measurement of glyceraldehyde-3-phosphate dehydrogenase (GAPDH) using TaqMan GAPDH control reagents (PE Applied Biosystems). For relative quantification of H2 receptor

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marker expression, calibration curves were constructed using the mRNA obtained from *H. pylori*-negative gastric mucosa. The levels of H2 receptor mRNA were normalized to that of GAPDH.

Statistical analyses

The level of H2 receptor mRNA was expressed as the median (first quartile to third quartile). The Wilcoxon rank sum test was used to compare H2 receptor mRNA levels before and after *H. pylori* eradication. Differences at P < 0.05 were considered significant.

RESULTS

H2 receptor mRNA levels decrease with the progression of gastric atrophy in H. pylori-infected subjects

Since the relationship between H2 receptor mRNA and H. pylori infection has not yet been adequately elucidated, we first attempted to assess the effect of H. pylori infection on H2 receptor mRNA expression. To this end, we compared H2 receptor mRNA expression in oxyntic glands with serum pepsinogen levels. Pepsinogens I and II differ in their location in the stomach. Both are located in the chief and mucous neck cells of the oxyntic gland mucosa in the gastric corpus, but only pepsinogen II is present in the gastric antrum. Serum pepsinogen I significantly correlate with peak acid output.¹⁷ A pepsinogen I/II ratio of <3 is considered to be a reliable marker for severe atrophic gastritis.^{18,19} As shown in Figure 1a and b. expression levels of H2 receptor mRNA significantly correlated with serum pepsinogen I/II ratios, but did not correlate with serum pepsinogen I levels. These results reveal that expression levels of H2 receptor mRNA were associated with the progression of gastric atrophy.

The lower expression levels of H2 receptor mRNA in subjects with hypergastrinemia

Reduced gastric acidity induces hypergastrinemia. Therefore, in an additional effort to examine the relationship between gastric atrophy and the expression levels of H2 receptor mRNA, we further attempted to investigate its levels in subjects with hypergastrinemia (>200 pg/mL) who were considered to have severe gastric atrophy.^{20,21} As shown in Figure 2, there was a slightly negative correlation of H2 receptor mRNA and serum gastrin levels before *H. pylori* eradication, and extremely lower expression levels of H2 receptor mRNA were found in subjects with hypergastrinemia.

H2 receptor mRNA restore significantly after H. pylori eradication

In the next series of examinations, we investigated the restoration of H2 receptor mRNA after *H. pylori* eradication. As shown in Figure 3, the median expression levels after *H. pylori* treatment increased to 2.9 times levels in comparison with the prior levels.

Restoration of H2 receptor mRNA after H. pylori eradication is enhanced in most subjects, especially with a lower serum pepsinogens I and II ratios

In an additional effort to examine the relationship between gastric atrophy and restoration of H2 receptor mRNA expression, we investigated the restoration of H2 receptor mRNA in association with the degree of gastric atrophy that *H. pylori*-infection induces in its pathological course. Therefore, H2 receptor mRNA expression levels before and after *H. pylori* cure were compared relative to serum pepsinogen I/II ratios. As shown in Figure 4, H2 receptor mRNA expression levels rose significantly despite the serum pepsinogen I/II ratio



Figure 1. Relationship between H2 receptor mRNA expression in the oxyntic mucosa and serum pepsinogen levels. H2 receptor mRNA expression levels significantly correlate with serum pepsinogen I/II ratios (n = 80) (a), but not with serum pepsinogen-I levels (n = 80) (b).

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Figure 2. Relationship between H2 receptor mRNA expression in the oxyntic mucosa and serum gastrin levels. There is a slightly negative correlation of H2 receptor mRNA and serum gastrin levels (n = 80), and extremely lower expression levels of H2 receptor mRNA are found in subjects with hypergastrinemia (>200 pg/mL).



Figure 3. Comparison of H2 receptor mRNA expression levels in the oxyntic mucosa before and after *H. pylori* eradication. The median expression levels after *H. pylori* treatment increased to 2.9 times levels in comparison with prior levels [median (first quartile to third quartile); 0.42 (0.20–0.97) and 1.19 (0.88–1.91), n = 80; P < 0.0001 by Wilcoxon rank sum test].

levels, in particular showing an increase of 4.2 times in subjects with a lower pepsinogen I/II ratios.

Restoration of H2 receptor mRNA after H. pylori eradication is less enhanced in subjects with hypergastrinemia

In the last sets of examinations, we further attempted to investigate H2 receptor mRNA restoration in subjects with hypergastrinemia (>200 pg/mL). As shown in Figure 5, a relatively low restoration of H2 receptor

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mRNA was found in such subjects, compared with those having normal serum gastrin levels.

DISCUSSION

The present study shows that H2 receptor mRNA levels correlated positively with pepsinogens I and II ratios, suggesting that its levels decreased with the progression of gastric atrophy induced by *H. pylori* infection. Moreover, we found that the restoration of H2 receptor expression after *H. pylori* eradication was threefold



Figure 4. Comparison of H2 receptor mRNA expression levels in the oxyntic mucosa and serum pepsinogen levels before and after *H. pylori* eradication. H2 receptor mRNA expression levels rose despite the levels of serum pepsinogen I/II ratios, especially showing an increase of 4.2 times in subjects with a lower pepsinogen I/II ratios [median (first quartile to third quartile); 0.24 (0.11–0.57) and 1.00 (0.77–1.76) in 43 subjects with pepsinogen I/II ratios <3; P < 0.0001, 0.75 (0.31–1.68) and 1.51 (1.13–2.20) in 37 subjects with I/II ratio >3; P < 0.0001 by Wilcoxon rank sum test].

Figure 5. Comparison of H2 receptor mRNA expression levels in the oxyntic mucosa and serum gastrin levels before and after *H. pylori* eradication. A weak restoration of H2 receptor mRNA was found in subjects with hypergastrinemia, compared with subjects having normal serum gastrin levels [median (first quartile to third quartile); 0.50 (0.23–1.09) and 1.48 (0.94– 2.09) in 71 subjects with serum gastrin <200 pg/mL; *P* < 0.0001, 0.17 (0.89– 0.31) and 0.76 (0.29–0.95) in nine subjects with serum gastrin >200 pg/mL; *P* = 0.021 by Wilcoxon rank sum test].

greater than prior levels in most subjects with successful eradication therapy. This restoration was enhanced in subjects with more severe gastric atrophy showing lower serum pepsinogens I and II ratios.

To date, the evaluation of H2 receptor mRNA expression levels on parietal cells was difficult and contentious because RT-PCR studies using non-intron spanning primers were unable to differentiate between genomic DNA and mRNA. The present study found that gastric atrophy induced by *H. pylori*-infection affected strongly the expression levels of H2 receptor mRNA on parietal cells, because there was a positive correlation between pepsinogens I and II ratios and H2 receptor mRNA. These results suggested that decreased mRNA expression in atrophic mucosa was partially associated with the hyposecretion of gastric acid, accompanied by an attenuation of parietal cell mass.

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H. pylori infection is associated with various disturbances of gastric secretory function, ranging from a marked increase to marked decrease in acid output. El-Omar *et al.* reported that the degree of recovery acid secretion was variable, with some regaining normal or near-normal values while others remained significantly hypochlorhydric over 18 months after eradication.²² In this study, the degree of recovery H2 receptor mRNA was stronger in subjects with severe gastric atrophy showing lower serum pepsinogens I and II ratios, consistent with a previous report, in which the increase in acid secretion after *H. pylori* eradication was greatest in subjects with severe gastritis.²³ However, subjects with hypergastrinemia were observed to have only a mild recovery of H2 receptor mRNA. These subjects may have had persisting hypochlorhydria, which may not resolve the atrophy and intestinal metaplasia after H. pylori eradication, resulting in an irreversible stage of hyposecretion of gastric acid.^{22, 24}

The exact mechanism by which the eradication of H. *pylori* increases gastric acid secretion remains unclear. El-Omar *et al.* reported that both the inflammation and subsequent inhibition of acid secretion are readily reversed by the eradication of H. *pylori*.²² Another investigator reported that atrophic gastritis induced by chronic infection of H. *pylori* decreases gastric acid secretion, simply by reducing the number and mass of acid-secreting parietal cells and oxyntic glands of the stomach.²⁵ The restoration of H2 receptor mRNA expression on parietal cells in the present study may be partially associated with the increase of gastric acid secretion.

Several previous studies have demonstrated that long-term therapeutic effects of H2RAs are inferior to those of proton pump inhibitors, particularly when these drugs are administered to patients with GERD whose infection rate of *H. pylori* is low.^{2,26–29} Whether this effect is simply due to an increase in parietal cell mass, an increase in H2 receptor sensitivity, or a true biological H2 receptor upregulation is unclear. This tolerance phenomenon is not likely to occur in *H. pylori*-positive subjects.¹⁶ The precise mechanism as to why H. pylori-infection prevents the occurrence of the tolerance phenomenon was not clarified. The attenuated expression of H2 receptor mRNA on parietal cells in H. pylori-infected mucosa in the present study may explain the phenomenon of lack of tolerance to H2RAs. Furthermore, higher levels of H2 receptor on parietal cells

after *H. pylori* cure may explain the tolerance phenomenon in *H. pylori*-negative subjects.

In summary, H2 receptor mRNA levels decrease with the progression of gastric atrophy induced by *H. pylori* infection. The restoration of H2 receptor expression after *H. pylori* eradication was threefold greater than the previous levels, which occurred in most subjects with successful eradication therapy. These expression levels of H2 receptor may explain a part of the tolerance phenomenon of H2RAs.

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STOMACH

Pericryptal fibroblast sheath in intestinal metaplasia and gastric carcinoma

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Revised version received 27 May 2004 Accepted for publication 3 June 2004 **Background and aims:** In the progression of chronic gastritis, gastric mucosal cells deviate from the normal pathway of gastric differentiation to an intestinal phenotype which is closely related to gastric carcinoma. However, to date, it has not been elucidated whether the intestinal metaplasia is merely a change in the epithelium or whether the underlying mesenchyme also changes from gastric type to intestinal type. We have investigated the relationship between intestinal metaplasia and the pericryptal fibroblast sheath (PCFS) in the mesenchyme. In addition, we also examined PCFS in gastric carcinoma. **Methods:** We determined the existence of PCFS in the intestinal metaplastic mucosa and carcinoma of both human and Cdx2 transgenic mouse stomach. PCFS was determined using the antibody against α-smooth muscle actin and electron microscopic observations.

Results: PCFS formed an almost complete layer around the small and large intestinal crypts while it did not exist around the normal gastric glands in both mice and humans. PCFS was seen around the glands of intestinal metaplastic mucosa in both Cdx2 transgenic mouse and human stomachs. However, PCFS was virtually absent in the intestinal-type gastric adenocarcinoma area.

Conclusion: We successfully demonstrated that the epithelium as well as the mesenchyme changed from the gastric type to the intestinal type in intestinal metaplasia and that PCFS disappeared in intestinal-type gastric carcinoma.

uman intestinal-type gastric carcinoma is associated with gastric atrophy and intestinal metaplasia that are caused mainly by *Helicobacter pylori* infection. Correa presented a hypothesis with respect to the mechanism of gastric carcinogenesis due to *H pylori* infection.¹ *H pylori* infection is involved in the process of progression from normal gastric mucosa to superficial gastritis, chronic active gastritis, atrophic gastritis, and finally to intestinal metaplasia.¹ The terminal stage of this process is gastric carcinoma.

We and others have reported that the intestinal specific transcription factor Cdx2 is expressed in human gastric intestinal metaplastic mucosa.²⁻⁶ Furthermore, we established Cdx2 transgenic mice expressing the transcription factor Cdx2 exclusively in the gastric epithelium.⁷ The gastric fundic mucosa of the Cdx2 transgenic mouse was completely changed into intestinal metaplastic mucosa. However, whether intestinal metaplasia is limited to the epithelium or influences the underlying mesenchyme has not been elucidated as it is difficult to discriminate intestinal mesenchyme from gastric mesenchyme.

The normal intestinal crypt of Lieberkühn is invested by a mesenchymal sheath, a specialised part of the lamina propria consisting of fibroblasts tightly surrounding the epithelium and of collagen fibres oriented circumferentially to the crypt.* The mesenchymal sheath is a distinct and highly organised system of fibroblasts immediately subjacent to the epithelial basement membrane located at the epithelial-mesenchymal interface. The highly specialised fibroblasts, that were reported as pericryptal fibroblasts by Kay and colleagues,⁹ form the pericryptal fibroblast sheath (PCFS). PCFS consists of a network of fibroblast cells and extracellular matrix immediately subjacent to the crypt epithelial cells.8 10 Such pericryptal fibroblasts under the basement membrane envelop the glands of the intestine, are seen as elongated cells tightly appliqued to the base of the epithelial cells of the crypt, have a fusiform appearance, and follow the contours of

the wall of the crypt. Fibroblasts of the PCFS have an important role in colonic fluid absorption.¹¹ Colonic absorptive function depends not only on crypt luminal cells but also on the fibroblast cells of the surrounding pericryptal sheath.¹¹

There is a close interactive epithelial-mesenchymal relationship between the epithelial cell system and the underlying pericryptal fibroblast system to maintain the normal structure and function of the crypts of Lieberkühn.8 1 Pericryptal fibroblasts have been shown to play a fundamental role in epithelial differentiation via epithelialmesenchymal cell interactions during both fetal and adult life.^{12 13} The PCFS is a self renewing population of mesenchymal cells in close contact with the intestinal epithelium; its cells maintain a parallel relationship in replication, migration, and differentiation with the overlying epithelium suggesting that it is involved in the maintenance of the normal structure and function of the intestinal mucosa,89 Autoradiographic studies in rabbit colon after ³H-thymidine injection demonstrate steady state renewal of pericryptal fibroblasts and migration to upper portions of the synchrony with epithelial migration. The fibroblast progenitor population, like that of the epithelium, is in the deep one third of the crypt.* The kinetics of this portion of the normal intestinal mucosa suggest that the pericryptal fibroblasts and the epithelium act as a unit to maintain the normal structure, maturation, and function of the crypt of Lieberkühn.

These findings prompted us to investigate (1) whether intestinal metaplasia influences the epithelium as well as the formation of the PCFS and (2) the relationship between PCFS and intestinal-type gastric carcinoma.

Abbreviations: PCFS, pericryptal fibroblast sheath; α -SMA, α -smooth muscle actin; PBS, phosphate buffered saline

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Figure 1 Mouse gastric glands (A), small intestinal crypts (B), and large intestinal crypts (C) shown in cross section. Cross sections of crypts of mouse small (B) and large (C) intestines show pericryptal fibroblast nuclei (arrows) immediately subjacent to the epithelial cells. The long axis of each pericryptal fibroblast is perpendicular to that of the crypt. Pericryptal fibroblast nuclei are not seen forming a sheath round the glands of mouse gastric mucosa (A). Magnification ×400.

MATERIALS AND METHODS

Cdx2 transgenic mice

We used Cdx2 transgenic mice with stomach specific expression of Cdx2 using the β -subunit gene promoter of rat H⁺/K⁺-ATPase.⁷ The gastric mucosa of Cdx2 transgenic mice was completely changed to intestinal metaplastic mucosa.⁷

Histology

Stomach tissue specimens were fixed in neutral buffered 10% formalin for 12–24 hours, washed in 70% ethanol, processed by standard methods, embedded in paraffin, sectioned at 3 μ m, and stained with haematoxylin and eosin for histological evaluation.

Immunohistochemistry

Thick sections (3 μ m) were cut, deparaffinised, rehydrated in phosphate buffered saline (PBS), placed in 10 mM citrate buffer (pH 6.0), and heated in an 850 W microwave for 15 minutes to recover antigenicity. Endogenous peroxidase activity was blocked by incubation for 30 minutes in methanol containing 0.3% H₂0₂. After washing twice with PBS, including 0.1% Triton X-100, sections were preincubated with blocking buffer (Dako, Carpinteria, California, USA) for 15 minutes at room temperature. Primary antisera, anti- α smooth muscle actin (α -SMA) (1:100; Dako), or anti-Cdx2 (1:100; BioGenex, San Ramon, California, USA) were diluted in PBS and incubated overnight at 4°C. Slides were then washed in PBS and incubated with Envision (Dako). After



Figure 2 Normal mouse small (A, B) and large (C, D) intestinal mucosa. Immunohistochemical stain for α -smooth muscle actin. The pericryptal fibroblast sheath formed by pericryptal fibroblasts is closely embracing epithelial cells of normal intestinal crypts. Magnification ×100 (A, C); ×400 (B, D).



Figure 3 Normal human small (A, B) and large (C, D) intestinal mucosa. Immunohistochemical stain for α -smooth muscle actin. Pericryptal fibroblast sheath formed by pericryptal fibroblasts is closely embracing the epithelial cells of normal intestinal crypts. Magnification ×100 (A, C); ×400 (B, D).

development with 3,3'-diaminobenzidine tetrahydrochloride (Wako Pure Chemical Industries, Osaka, Japan), slides were counterstained with haematoxylin and viewed under a light microscope.

Fixation and preparation of tissue for electron microscopy

Intestinal metaplastic mucosa from Cdx2 transgenic mouse stomach was fixed at 4° C in 2% glutaraldehyde in PBS, followed by six washes in PBS with post fixation in 1% osmium tetroxide. Sections were examined under a Hitachi H-7500 scanning electron microscope.

RESULTS

To examine whether intestinal metaplasia is merely epithelial cell metaplasia or influences the underlying mesenchyme, we focused our attention on the PCFS in the mesenchyme. Light microscopic examination of normal mouse small and large intestine revealed fibroblasts subtending the epithelial basement membrane in the crypt (fig 1B, 1C). In contrast, fibroblasts were not seen forming a sheath round the glands of normal mouse gastric mucosa (fig 1A). These fibroblasts subjacent to the epithelium were easily distinguished from the rest of the mesenchymal elements of the lamina propria, which consist of a loose meshwork of collagen, fibroblasts, haematogenous cells, and capillaries.

 α -SMA is present in pericryptal fibroblasts and is used as a marker for pericryptal fibroblast cells. Pericryptal fibroblasts labelled by α -SMA formed single cell layers that embraced the whole length of the crypts in the normal small and large intestinal mucosa of the mouse (fig 2) and humans (fig 3). α -SMA positive PCFS was seen around many glands. Areas of abutment were broad and the sheath appeared continuous.

We previously generated Cdx2 transgenic mice expressing intestine specific transcription factor *Cdx2* gene exclusively in the gastric epithelium under the control of the β -subunit gene promoter of rat H⁺/K⁺-ATPase.⁷ Cdx2 transgenic mice developed normally into superficially healthy adults and showed intestinal metaplasia in the stomach up to 12 weeks of age. The gastric mucosa of Cdx2 transgenic mouse was completely replaced by intestinal metaplastic mucosa. Cdx2 induced intestinal metaplastic mucosa consisted of terminally differentiated intestinal epithelial cells, including absorptive enterocytes, goblet cells, and enteroendocrine cells.⁷ To clarify whether Cdx2 expression in gastric epithelium affects the underlying mesenchyme in vivo, the intestinal metaplastic mucosa of Cdx2 transgenic mouse stomach was stained for α -SMA. The PCFS expressing α -SMA was not seen around the glands of the normal gastric mucosa (fig 4A, B) whereas PCFS was easily recognised around the crypts of intestinal metaplastic mucosa, PCFS was also recognised around the glands of human intestinal metaplastic mucosa (fig 5C, D) while it was not seen around normal human gastric glands (fig 5A, B).

Electron microscopy revealed an even closer association between pericryptal fibroblasts and the epithelium than was revealed by light microscopy. In intestinal metaplastic mucosa of Cdx2 transgenic mouse stomach, the PCFS was in intimate contact with the epithelial basal lamina (fig 6). Fibroblasts were seen surrounding the base of the crypts in the intestinal metaplastic mucosa. These cells had large areas of contact with the epithelial basal lamina and had a plum fusiform shape (fig 6B, C).

As it is reported that the PCFS is significantly reduced in colorectal epithelial neoplasms, we examined the relationship between PCFS and gastric adenocarcinoma. We observed Cdx2 transgenic mice periodically without carcinogens or H pylori infection. Cdx2 transgenic mice at 50 weeks of age indicated preservation of intestinal metaplasia and no gastric polyp formation, similar to those at 12 weeks (unpublished data). Cdx2 transgenic mice developed gastric polyps in the intestinal metaplastic lesion at two years after birth. Gastric polyps developed from intestinal metaplastic mucosa in all stomachs of 10 Cdx2 transgenic mice examined. The polyps consisted of adenocarcinoma that invaded the submucosa or beyond (unpublished data). Using the adenocarcinoma, we examined the relationship between PCFS and adenocarcinoma. PCFS detected by immunohistochemical stain for α-SMA was virtually absent in the area of gastric



Figure 4 Normal and intestinal metaplastic mucosa of the mouse stomach. Immunohistochemical stain for stomach. Immunohistochemical stain ta α -smooth muscle actin. Pericryptal fibroblast sheath (PCFS) formed by pericryptal fibroblasts is closely embracing the epithelial cells of the intestinal metaplastic mucosa (C, D) while PCFS is not seen in the normal gastric mucosa (A, B). Magnification $\times 100$ (A, C); $\times 400$ (B, D).

adenocarcinoma of both humans and mice (fig 7C, G) while PCFS was easily recognised in the intestinal metaplastic areas (fig 7A, E). PCFS was absent in all 10 Cdx2 transgenic mouse and 10 human intestinal-type adenocarcinomas. Both human and murine gastric carcinomas were classified as intestinaltype according to the criteria of Lauren¹⁴ and as category 5.2 according to the five categories of the Vienna classification.^{15 16} We examined expression of Cdx2 in intestinal metaplasia and adenocarcinoma. Cdx2 staining for the adenocarcinoma lesion (fig 7D, H) was extremely weak

compared with the intestinal metaplastic lesion (fig 7, B and F). The decrease in Cdx2 may explain, in part, the cause of the disappearance of PCFS in the gastric adenocarcinoma. There was no difference in immunoreactivities for Cdx2 and α -SMA in intestinal metaplastic mucosa between 12 and 50 week old Cdx2 transgenic mice (data not shown).

DISCUSSION

Our results demonstrate that intestinal metaplasia is not merely epithelial metaplasia but also affects the underlying