Relationship between *Helicobacter pylori* babA2 status with gastric epithelial cell turnover and premalignant gastric lesions

J Yu, W K Leung, M Y Y Go, M C W Chan, K F To, E K W Ng, F K L Chan, T K W Ling, S C S Chung, J J Y Sung

**Background:** *Helicobacter pylori* blood group antigen binding adhesin (BabA) mediates bacterial adherence to human blood group antigens on gastric epithelium. Although strains harbouring babA2 were recently found to be associated with peptic ulcer and gastric cancer, the role of babA2 in cellular turnover, severity of gastritis, and premalignant changes is poorly understood.

**Aim:** We correlated *H pylori* babA2, vacuolating toxin (vacA), and cytotoxic associated gene A (cagA) genotypes with the severity of gastric inflammation and epithelial cell turnover in a group of Chinese patients from an area with a high incidence of gastric cancer.

**Patients and methods:** *H pylori* isolates were obtained from 104 Chinese patients who participated in a gastric cancer prevention programme. Genotype variants of babA2, vacA, and cagA were determined by polymerase chain reaction. Antrum and corpus histopathology was examined according to the updated Sydney classification. Apoptosis was scored by terminal uridine deoxynucleotidyl nick end labeling (TUNEL) and proliferation by Ki-67 immunostaining.

**Results:** Of the 104 patients, 102 (98.1%) harboured cagA+ strains and all had vacA s1 genotype. The babA2+ strains were found in 83 (79.8%) patients and were associated with higher lymphocytic infiltration (p=0.028), presence of glandular atrophy (odds ratio (OR) 7.5, 95% confidence interval (CI) 2.3–24.3), and intestinal metaplasia (OR 7.4, 95% CI 2.2–25.3) in the antrum. Increased epithelial proliferation was also noted in individuals infected with babA2+ strains (p=0.025). Strains harbouring cagA+/vacA s1 genotypes lacked this association in the absence of babA2.

**Conclusions:** The presence of babA2: *H pylori* strains alone or in combination with cagA+ and vacA s1 was associated with the presence of premalignant gastric lesions.

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**MATERIALS AND METHODS**

**Study design and subjects**

This was a cross sectional study that examined the correlation between various *H pylori* genotypes, gastric inflammation, and cellular turnover. Subjects were previously recruited in a randomised control trial of *H pylori* eradication in the prevention of gastric cancer in Yantai County, Shandong province of China, where the incidence of gastric cancer is among the highest in China. Endoscopy was performed in all subjects, with gastric biopsies obtained. Two antral and two corpus biopsies were fixed in 10% buffered formalin and processed for histological examination. One extra antral biopsy was used for bacterial culture for *H pylori*.

In this study, 104 *H pylori* positive subjects (median age 50 years; range 39–75) were studied. These patients were randomly selected from those with persistent active *H pylori* infection at first follow up, as detected by bacterial culture. None of these patients had previously received anti-*Helicobacter* therapy.

**H pylori culture**

One antral biopsy was placed in selective transport medium and cultured on horse blood agar at 37°C under micro-

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**Abbreviations:** IM, intestinal metaplasia; cagA, cytotoxin associated gene A; VacA, vacuolating toxin; AT, atrophy; TdT, terminal deoxynucleotidyl transferase; TUNEL, triphosphate nick end labeling; AI, apoptosis index; PI, proliferation index; OR, odds ratio; BabA, blood group antigen binding adhesin; PCR, polymerase chain reaction.
aerophilic conditions, as previously described.11 *Helicobacter pylori* was identified by typical colony morphology, Gram stain, and positive biochemical testing for urease, catalase, and oxidase. Bacteria were harvested from the plates using sterile cotton swabs and stored at −70°C in brucella broth plus 30% (vol/vol) fetal calf serum and 20% (vol/vol) glycerol. All frozen isolates were controlled for contamination.

**Detection of *H pylori* genotypes and sequencing**

*Helicobacter pylori* cagA, vacA, and babA2 genotypes were determined by polymerase chain reaction (PCR). Genomic DNA was extracted from *H pylori* using the High Pure PCR Template Preparation Kit (Roche, Germany). The integrity of the DNA was assessed by 1.2% agarose gels stained with ethidium bromide. PCR reactions were performed in a total volume of 50 μl which contained 50 pmol of primers, 100 ng of genomic DNA, 1.0 mM of each of the four dNTPs, and 2 units of AmpliTaq DNA polymerase (Perkin-Elmer, Norwalk, Connecticut, USA). Primer sequences were described previously12−17 and are listed in table 1 for reference. The amplified PCR products were resolved in 1.5% agarose gels, stained with ethidium bromide, and visualised under a short wavelength ultraviolet light source. The sequences of the PCR products were confirmed by automated sequencing (ABI Prism 310 Genetic Analyzer; Perkin Elmer, Branchburg, New Jersey, USA) using the same primer pairs.

**Histological assessment**

Haematoxylin-eosin stained sections of the gastric antrum and corpus biopsies were examined by a single pathologist (KFT) who was blinded to the other experimental results. Slides were coded and evaluated histologically for the following parameters: (1) activity of gastritis (granulocytic cell infiltration); (2) chronicity of gastritis (lymphocytic and plasma cells infiltration); (3) glandular AT; and (4) IM. Each parameter was graded as none (0), mild (1), moderate (2), or severe (3) according to the updated Sydney classification.18 Glandular atrophy was defined as loss of gastric glands with replacement by metaplastic epithelium or fibrosis. IM was defined as the presence of foci where at least three neighbouring gastric pits contained two or more goblet cells in each pit.

**Assessment of apoptosis and determination of apoptosis index**

Apoptosis was determined by the terminal deoxynucleotidyl transferase (TdT) mediated deoxyuridine triphosphate nick end labeling (TUNEL) technique (ApopTag: Intergen, New York, USA), as described previously.19 In addition, positively stained cells should also bear morphological characteristics of apoptosis, including chromatin condensation, nuclear disintegration, and formation of crescentic caps of condensed chromatin at the nuclear periphery. The apoptosis index (AI) was calculated as the ratio of TUNEL positive nuclei over the total number of cells counted (usually more than 1000).

**Assessment of proliferation and determination of proliferation index**

Proliferation was assayed by immunoperoxidase staining for Ki-67, a proliferation associated antigen that has been used extensively in the assessment of proliferation in the stomach.19−20 In brief, paraffin embedded sections were dewaxed, dehydrated, and then subjected to microwave antigen retrieval. Endogenous peroxidase was blocked by 3% hydrogen peroxide. Normal rabbit serum (5%) was applied to prevent non-specific binding of primary monoclonal antibodies. The tissue was incubated with monocolonal mouse anti-human Ki-67 antibody (Clone: mib 1, isotype: IgG1, without dilution; Zymed, San Francisco, California, USA). Biotinylated rabbit antimouse antibody (dilution 1:200; Dako, Glostrup, Denmark) was applied as secondary antibody and followed by application of avidin-biotinylated peroxidase complex (dilution 1:100; Dako). The slides were developed in 3,3'-diaminobenzidine tetrahydrochloride (DAB, Dako) and counterstained with Mayer haematoxylin. The proliferation index (PI) was expressed as a percentage of the ratio of Ki-67 positive nuclei to the total nuclei counted.11

**Statistical analysis**

Statistical analysis was performed using the Statistical Package for the Social Science program (SPSS, version 10.0) and MINITAB (Release 7.2, Minitab, Inc). As gastric histology, particularly AT and IM, may be confounded by the effect of age, we have adjusted for the effect of age in this study. Mantel-Haenszel statistics were used to assess the relationship between individual genotypes and the presence of activity, chronicity, glandular AT, and IM after adjusting for the effect of age; the results are expressed as common odds ratios (OR) with the corresponding 95% confidence intervals (CI).

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**Table 1** Primer sequences for polymerase chain reaction based genotyping of *Helicobacter pylori* isolates

<table>
<thead>
<tr>
<th>Ref</th>
<th>Gene</th>
<th>Sequence (5' to 3')</th>
<th>Product size</th>
</tr>
</thead>
<tbody>
<tr>
<td>13</td>
<td>vacA</td>
<td>s1/s2</td>
<td>ATGGAATACAAAGACAACAC 259 bp (s1)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Forward</td>
<td>CTGCTGAATGCGCAAAAG 286 bp (s2)</td>
</tr>
<tr>
<td>17</td>
<td>vacA</td>
<td>m1/m2</td>
<td>CAATCTGCAATCAAGGCCGAG 570 bp (m1)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Forward</td>
<td>GCGTGCTAAATATCCAAG 645 bp (m2)</td>
</tr>
<tr>
<td>13</td>
<td>cagA</td>
<td>Forward</td>
<td>ATAAATGCAAATAGCAACCTGAGGCA 297 bp</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Reverse</td>
<td>TGTTAGTGATTTCGGTGTAGGACA 850 bp</td>
</tr>
<tr>
<td>16</td>
<td>babA2</td>
<td>Forward</td>
<td>AATCCAAAAGGAGAAAAGTATGAAA</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Reverse</td>
<td>TGTAGTGTATTGCGTGGAGCAG</td>
</tr>
</tbody>
</table>

*There were three strains that could not be typed.*

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**Table 2** Distribution of different *Helicobacter pylori* genotypes in 104 patients

<table>
<thead>
<tr>
<th>Genotype</th>
<th>No (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>babA2</td>
<td>83 (79%)</td>
</tr>
<tr>
<td>cagA*</td>
<td>102 (98.1%)</td>
</tr>
<tr>
<td>vacA</td>
<td>104 (100%)</td>
</tr>
<tr>
<td>s1</td>
<td>0 (0%)</td>
</tr>
<tr>
<td>s2</td>
<td>43 (41.3%)</td>
</tr>
<tr>
<td>m2</td>
<td>58 (55.8%)</td>
</tr>
<tr>
<td>m2*/s1</td>
<td>102 (98%)</td>
</tr>
<tr>
<td>cagA*/vacA s1</td>
<td>43 (41.3%)</td>
</tr>
<tr>
<td>babA2/cagA*/vacA s1</td>
<td>82 (78%)</td>
</tr>
</tbody>
</table>

*There were three strains that could not be typed.*
The age of the subjects was stratified into two groups (>50 or ≤50 years old) according to the median age of the study population. The Mann-Whitney U test was used to compare the severity of gastritis among different genotypes. Spearman's correlation coefficient was used to assess the correlation between (1) proliferation and severity of inflammation; and (2) age and AT/IM scores. The Pearson correlation coefficient was used to examine the correlation between epithelial cell proliferation and apoptosis. The two tailed independent \( t \) test was performed to examine the difference in AI and PI among various genotypes. A two sided \( p \) value of less than 0.05 was considered statistically significant.

**RESULTS**

**Characteristics of *H pylori* isolates**

Of the 104 *H pylori* infected subjects, 102 (98.1%) harboured *cagA*+ strains. All *H pylori* isolates belonged to the *vacA* s1 genotype and thus 98.1% were *cagA*/vacA s1. For the middle (m) region of the *vacA* gene, 101 isolates could be typed and the remaining three strains could not. There were 43 (42.6%) m1 and 58 (57.4%) m2 strains. Eighty three (79.8%) were *babA2* strains and the simultaneous presence of *cagA*, *vacA* s1, and *babA2* genes (triple positive) was found in 82 (78.8%) *H pylori* isolates (table 2). The presence of the *babA2* genotype did not correlate with the presence of *cagA* or various *vacA* genotypes.

**Histological assessment of gastritis**

Most *H pylori* infected patients had evidence of granulocytic and lymphocytic infiltration in the stomach (fig 1). Granulocytic infiltration was more marked in the antrum than in the body (table 2). The presence of the *babA2* genotype was associated with a higher degree of lymphocytic infiltration (chronicity) in the antrum (\( p=0.028 \), table 3).

**Correlation between *H pylori* genotypes and inflammation**

As most *H pylori* infected patients had evidence of acute and chronic inflammatory infiltrates in the stomach, there was no significant correlation between different *H pylori* genotypes and the presence of granulocytic and lymphocytic infiltrates in the antrum or body of the stomach (\( p>0.3 \)). However, the presence of the *babA2* genotype was associated with a higher degree of lymphocytic infiltration (chronicity) in the antrum (\( p=0.028 \), table 3).

**Correlation between *H pylori* genotypes and atrophy/metaplasia**

After adjustment for age, carriage of *babA2* *H pylori* isolates was associated with a higher frequency of glandular AT (OR 7.5, 95% CI 2.3–24.3) and IM (OR 7.4, 95% CI 2.2–25.3) in the antrum (table 4). A similar association was also demonstrated in triple positive (*babA2*/cagA*/vacA s1*) *H pylori* isolates. The corresponding OR values for AT and IM in triple positive strains were 7.1 (95% CI 2.2–23.3) and 7.0 (95% CI 2.1–24.1). None of the *H pylori* genotype or genotype combinations appeared to correlate with preneoplastic lesions in the corpus.

**Epithelial cell proliferation and apoptosis**

The association between cellular proliferation and apoptosis was examined. There was a significant correlation between PI and AI (\( r=0.37, p=0.001 \)) but there was no direct correlation between proliferation and severity of inflammation (both activity and chronicity).

**babA2 genotypes and cell proliferation**

Proliferation indexes of different genotypes are shown in fig 2. PI was significantly higher in *babA2* isolates (mean (SD) 21.3 (17.7)) than in *babA2* isolates (12.2 (13.9); mean difference 9.1, 95% CI 1.46–16.70; \( p=0.025 \)). In addition, there was a significant difference in PI between the *babA2*/cagA*/vacA s1* (21.4 (17.9)) and *babA2*/cagA*/vacA s1* (12.9 (13.9)) isolates (mean difference 8.5, 95% CI 0.64–16.29; \( p=0.041 \)). There was however no difference in proliferation between *vacA* m1 (21.6 (18.6)) and *vacA* m2 (16.5 (15.6)) strains (mean difference 5.1, 95% CI –2.73–12.95; \( p=0.19 \)).

**babA2 genotypes and apoptosis**

Cellular apoptotic indexes of different *H pylori* genotypes are demonstrated in fig 3. In contrast with proliferation, there was no difference in AI between *babA2* and *babA2* isolates (mean (SD) 0.67 (0.61) v 0.66 (0.52); mean difference 0.01, 95% CI –0.27–0.29; \( p=0.96 \)). In addition, AI was comparable between *babA2*/cagA*/vacA s1* and *babA2*/cagA*/vacA s1* (0.68 (0.61) v 0.67 (0.53); mean difference 0.00, 95% CI –0.28–0.29; \( p=0.98 \)).

**Table 3** *Helicobacter pylori* genotypes and median (range) gastritis score

<table>
<thead>
<tr>
<th></th>
<th>babA2* (n=83)</th>
<th>babA2+ (n=21)</th>
<th>p Value</th>
<th>vacA m1 (n=43)</th>
<th>vacA m2 (n=58)</th>
<th>p Value</th>
<th>babA2*/cagA*/vacA s1 (n=82)</th>
<th>babA2*/cagA*/ vacA s1 (n=20)</th>
<th>p Value</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Antrum</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Activity</td>
<td>2.00 (0–3)</td>
<td>2.00 (0–3)</td>
<td>0.49</td>
<td>2.00 (0–3)</td>
<td>2.00 (0–3)</td>
<td>0.12</td>
<td>2.00 (0–3)</td>
<td>2.00 (0–3)</td>
<td>0.59</td>
</tr>
<tr>
<td>Chronicity</td>
<td>2.00 (1–3)</td>
<td>2.00 (1–3)</td>
<td>0.028</td>
<td>2.00 (1–3)</td>
<td>2.00 (1–3)</td>
<td>0.34</td>
<td>2.00 (1–3)</td>
<td>2.00 (1–3)</td>
<td>0.065</td>
</tr>
<tr>
<td><strong>Corpus</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Activity</td>
<td>1.00 (0–2)</td>
<td>1.00 (0–2)</td>
<td>0.73</td>
<td>1.00 (0–2)</td>
<td>1.00 (0–2)</td>
<td>0.77</td>
<td>1.00 (0–2)</td>
<td>1.00 (0–2)</td>
<td>0.77</td>
</tr>
<tr>
<td>Chronicity</td>
<td>2.00 (1–2)</td>
<td>2.00 (1–2)</td>
<td>0.82</td>
<td>1.00 (1–2)</td>
<td>2.00 (1–2)</td>
<td>0.59</td>
<td>2.00 (1–2)</td>
<td>2.00 (1–2)</td>
<td>0.99</td>
</tr>
</tbody>
</table>

Comparison between different groups was made using the Mann-Whitney U test. Activity=degree of granulocytic cell infiltration; chronicity=degree of lymphocytic and plasma cell infiltration.
patients were infected with H. pylori in this population with a high background incidence of gastric cancer such as Japan and China tend to have a higher prevalence of babA2 genes. Recently, a similar association between gastric inflammation and preneoplastic lesions was identified factors are also likely to be involved. Otherwise, one might expect to find a wider difference in inflammation scores between infection by babA2+ and babA2− isolates (table 3).

Figure 2  Proliferation index in different Helicobacter pylori genotypes. Results are plotted as mean and 95% confidence interval. The squares indicate the mean and the horizontal lines the upper and lower values of the 95% confidence interval. Significant differences were noted in the proliferation index between (a) babA2+ and babA2− (p=0.025); and (b) babA2+/cagA+/vacA s1 and babA2−/cagA+/vacA s1 (p=0.041).

and between vacA m1 and vacA m2 (0.62 (0.54) v 0.65 (0.49); mean difference -0.02, 95% CI -0.26-0.21; p=0.86) isolates.

DISCUSSION
In the present study, we have examined three genotypes (cagA+, vacA and babA2) that have been suggested to have higher pathogenic potential in H. pylori isolates from northern China. All H. pylori isolates were of the vacA s1 genotype and 98% were cagA+, indicating that the cagA+/vacA s1 genotypes are highly prevalent in this population with a high background incidence of gastric cancer. Additionally, we found that approximately 80% of patients were infected with babA2+ strains and 41.3% patients were harbouring vacA m1 strains. Carriage of the babA2+ strains was associated with more intense chronic inflammation (table 3), and presence of glandular AT and IM in the gastric antrum (table 4). This association with preneoplastic gastric lesions was also demonstrated in H. pylori isolates that were triple positive (babA2+/cagA+/vacA s1) compared with cagA+/vacA s1 strains that lacked babA2. Recently, a similar association between gastric inflammation and babA2+ genotypes was reported in a German study. The investigators reported that type 1 (cagA+/vacA s1) H. pylori strains that were negative for babA2 lacked the association with gastric inflammation and preneoplastic lesions. In contrast with the German study that reported a 38% prevalence of the babA2 gene, babA2 was present in approximately 80% of our study population. In a recent Japanese study, the prevalence of the babA2 genotype was reported to be 85%. The differential prevalence of babA2+ H. pylori isolates in these three different studies appears to parallel the gastric cancer incidence of the countries. Countries with a high background incidence of gastric cancer such as Japan and China tend to have a higher frequency of babA2+ H. pylori. As attachment to gastric epithelial cells by H. pylori is mediated by BabA, it has been postulated that this attachment process precedes all other putative pathogenic processes. Intuitively, adhesion to epithelial cells via BabA may facilitate the effective delivery of bacterial products such as the CagA protein into the host cells and the subsequent tyrosine phosphorylation process. None the less, babA2 is probably only one of the mediators for the attachment process and other unidentified factors are also likely to be involved. Otherwise, one might expect to find a wider difference in inflammation scores between babA2+ and babA2− isolates (table 3).

Age is one of the factors that may be associated with the presence of AT and IM in H. pylori infected individuals. In our previous study, we showed that older subjects are more likely to demonstrate progression of metaplasia than younger

Table 4  Odds ratio of glandular atrophy (AT) and intestinal metaplasia (IM) in different Helicobacter pylori genotype combinations after adjustment for age

<table>
<thead>
<tr>
<th></th>
<th>AT (%)</th>
<th>Non-AT (%)</th>
<th>Odds ratio* (95% CI)</th>
<th>IM (%)</th>
<th>Non-IM (%)</th>
<th>Odds ratio* (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Antrum</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>babA2+</td>
<td>54 (65.1)</td>
<td>29 (34.9)</td>
<td></td>
<td>49 (59.0)</td>
<td>34 (41.0)</td>
<td></td>
</tr>
<tr>
<td>babA2−</td>
<td>3 (23.8)</td>
<td>16 (76.2)</td>
<td>7.5 (2.3–24.3)</td>
<td>4 (19.0)</td>
<td>17 (81.0)</td>
<td>7.4 (2.2–25.3)</td>
</tr>
<tr>
<td>vacA m1</td>
<td>25 (58.1)</td>
<td>18 (41.9)</td>
<td></td>
<td>22 (51.2)</td>
<td>21 (48.8)</td>
<td></td>
</tr>
<tr>
<td>vacA m2</td>
<td>32 (55.2)</td>
<td>26 (44.8)</td>
<td>1.1 (0.5–2.4)</td>
<td>30 (51.7)</td>
<td>28 (48.3)</td>
<td>0.9 (0.4–2.1)</td>
</tr>
<tr>
<td>babA2+/cagA+/vacA s1</td>
<td>32 (55.2)</td>
<td>26 (44.8)</td>
<td></td>
<td>30 (51.7)</td>
<td>28 (48.3)</td>
<td>0.9 (0.4–2.1)</td>
</tr>
</tbody>
</table>
| *Common odds ratio was given after adjustment for the effect of age (>50 or ≤50 years) by the Mantel-Haenszel statistics. 95% CI, 95% confidence interval. As all strains were of the vacA s1 genotype and 98.1% strains were cagA+,
| **Corpus**     |        |            |                      |        |            |                      |
| babA2+         | 8 (9.6) | 75 (90.4) |                      | 5 (6.0) | 78 (94.0) |                      |
| babA2−         | 3 (14.3) | 18 (85.7) | 0.8 (0.2–3.3)        | 1 (4.8) | 20 (95.2) | 1.5 (0.2–14.2)       |
| vacA m1        | 5 (11.6) | 38 (88.4) |                      | 2 (4.7) | 41 (95.3) |                      |
| vacA m2        | 6 (10.3) | 52 (89.7) | 1.0 (0.3–3.6)        | 4 (6.9) | 54 (93.1) | 0.6 (0.1–3.3)        |
| babA2+/cagA+/vacA s1 | 5 (11.6) | 38 (88.4) |                      | 2 (4.7) | 41 (95.3) |                      |
| babA2−/cagA+/vacA s1 | 8 (9.8) | 74 (90.2) |                      | 5 (6.1) | 77 (93.9) |                      |
| babA2−/cagA−/vacA s1 | 3 (15.0) | 17 (85.0) | 0.7 (0.2–3.1)        | 1 (5.0) | 19 (95.0) | 1.4 (0.2–13.2)       |

Figure 3  Apoptosis index in different Helicobacter pylori genotypes. Results are plotted as mean and 95% confidence interval. There was no significant difference in apoptosis index between (a) babA2+ and babA2−; (b) babA2+/cagA+/vacA s1 and babA2−/cagA+/vacA s1; and (c) vacA m1 and vacA m2.

Proliferation index in different Helicobacter pylori genotypes. Results are plotted as mean and 95% confidence interval. The squares indicate the mean and the horizontal lines the upper and lower values of the 95% confidence interval. Significant differences were noted in the proliferation index between (a) babA2+ and babA2− (p=0.025); and (b) babA2+/cagA+/vacA s1 and babA2−/cagA+/vacA s1 (p=0.041).
subjects. However, even after adjustment for age, the frequencies of AT and IM were still significantly higher in individuals harbouring the babA2 strains or the triple positive H pylori strains. In this study, we stratified patients into two groups, with 50 years as the dividing line, because this was the median age of our subjects. Moreover, in a previous decision analysis modelling study of H pylori screening to prevent gastric cancer, the investigators suggested that screening at age 40–50 years is considered to be reasonably cost effective. Therefore, it appears that most gastric changes related to chronic H pylori infection had already developed at 50 years.

Disturbance in the balance between cellular proliferation and apoptosis may predispose to either cell loss with mucosal damage or cell accumulation and cancer development. Recent studies suggested that H pylori infection increases gastric mucosal cell proliferation. However, the effect of H pylori on gastric epithelial cell proliferation is conflicting. One explanation for this discrepancy may be related to the presence of different bacterial genotypes. However, whether H pylori adherence factors, such as BabA, affect cellular kinetics is unclear. Our findings demonstrated that subjects harbouring H pylori babA2 strains had significantly higher gastric proliferation rates than subjects with babA2 strains, which was unrelated to the degree of inflammation. More importantly, this increase in gastric cell proliferation was not accompanied by a parallel increase in apoptosis. Although the triple positive strains (babA2 cagA vacA) had a similar increase in cell proliferation, the conventional type 1 (cagA vacA) H pylori strains which were negative for babA2 lacked such an association. In other words, in subjects infected with H pylori babA2 strains, there was dissociation between gastric epithelial cell proliferation and apoptosis, which may predispose to cancer development.

In conclusion, babA2 H pylori isolates were present in approximately 80% of the study population from Yantai County where the incidence of gastric cancer is among the highest in China. The presence of babA2 H pylori strains alone, or in combination with cagA vacA was associated with the presence of glandular AT and IM, and heightened gastric epithelial proliferation in the antrum. Carriage of babA2 H pylori strains and the associated changes in the gastric epithelium may partly explain the high background incidence of gastric cancer in the study population.

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