

Frequency of H.Pylory BabA2 and Hpa Genes in Patients with Dysphagia

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ABSTRACT

Background and Objective: BabA2 and Hpa genes are involved in adherence of *Helicobacter pylori* (H.pylori) to gastric mucosal tissue. This study aimed to investigate the frequency of these genes in isolates of H. pylori from gastric biopsies and their relationship with gastritis, peptic ulcer and gastric cancer.

Methods: Gastric biopsy samples were obtained from patients with gastritis, peptic ulcer and gastric cancer. A sample was sent to the laboratory for urease test and histopathology study, and another sample for DNA extraction. The frequency of BabA2 and Hpa genes was investigated using their specific primers by PCR.

Results: Among the 80 analyzed biopsy samples, 51 (63%) were BabA2 positive, and the frequency of this gene in the samples of gastric cancer, gastritis and peptic ulcer was 61.1, 58.3 and 73.3%, respectively. In addition, 57 samples (71%) were Hpa positive, and the frequency of this gene in the samples of gastric cancer, gastritis and peptic ulcer was 55.5, 69.4 and 84.6%, respectively. There was no significant correlation between the presence of these genes and the type of H.pylori-related diseases.

Conclusion: Frequency of BabA2 and Hpa genes is higher in the samples of peptic ulcer but there was no significant relationship between these genes and H.pylori-related diseases.

Keywords: BabA2, Hpa, Gastric Cancer, Gastritis, Peptic Ulcer.

INTRODUCTION

The colonization of *Helicobacter pylori* (*H.pylori*) in human stomach leads to gastrointestinal diseases such as chronic gastritis, peptic ulcer, lymphoma and gastric adenocarcinoma (1). The outer membrane proteins (OMP), glycoproteins and bacterial lipids on the surface of the bacteria are involved in communication between host cells and bacteria. Thus, the adhesins and OMP are considered as pathogenic factors. The binding factors have the most important role in inflammatory responses among all bacterial factors. *H.pylori* bacterium was isolated and cultured from biopsy samples of patients with chronic gastritis and cultured by Warren and Marshall in 1982. *H.Pylori* is one of the most common bacterial infections in the world (1). BabA2 protein is an important binding factor in *H.pylori*, which causes the binding to B-blood type Lewis antigen on the stomach epithelial cells (2,3). BabA2 protein is the first *H. pylori* adhesin for Lewis B antigen, identified as BabA that is of OMP origin. This adhesion existed in some *H. pylori* enables them to attach to Lewis b antigen of epithelial cells. The encoding gene of this adhesion (BabA2) has two alleles (babA1 babA2), babA1 is the inactive allele and babA2 is the active form of the gene (3-5). The detection of BabA2 gene by Polymerase chain reaction (PCR) indicates the activity of this gene. Several studies on the pathogenesis of *H.pylori*-related diseases have demonstrated the presence of BabA2 gene and investigated the relationship between them (6). The Hpa protein (*H.Pylori* agglutinin), one of the binding factors of *H.pylori*, is coded by Hpa genes and causes the binding of *H.pylori* to the mucosal cells of the stomach. The Hpa is the main sheath flagellar protein with 29 kDa weight and is of *H.pylori* adhesins that plays an important role in the attachment of bacteria (7,8). The Hpa genes code 2 BabA proteins that facilitate bacterial virulence by increasing the production of cytotoxin and cell adhesion to the host cell. The presence of these genes have severe clinical consequences in gastroduodenal patients and brings about dyspepsia (9-11).

MATERIAL AND METHODS

This cross-sectional study was conducted on 80 patients, with gastrointestinal disorders, referring to the endoscopy unit of 5th Azar

medicine ward of the hospital were undergone endoscopy according to diagnosis of gastroenterologist. A sample was taken for histopathology and urease tests and another sample was transferred into sterile vials containing saline to be used for DNA extraction and molecular methods. The urease test was performed using a kit (Baharafshan co.) and then gastric biopsy sample was sent to a pathology lab in 10% formalin. DNA extraction was performed according to the manufacturer's instructions. The extracted DNA was then stored at -20 °C until PCR amplification. In the PCR process, a volume of 25 µl containing 5µl TBE buffer, 2.5 ml MgCl₂, 2µl primer (Forward and Reverse), 0.2 ml dNTP, 2 ml Taq DNA polymerase and 5 µl of extracted DNA was used for babA2 gene. For the Hpa gene, a 25 µl volume consisting of 5 µl TBE buffer, 3 µl MgCl₂, 2 µl primer (Forward and Reverse), 3 µl dNTP, 2.5 µl Taq DNA polymerase and 1ul extracted DNA was used. The thermocycler temperature was programmed for the gene BabA2 as: primary denaturation temperature of 95 °C for 5 minutes and then 35 temperature cycles including: secondary denaturation temperature of 94 °C for 1 min, Annealing temperature of 60 °C for 1 min and Elongation at 72 °C for 5 min. PCR process for the Hpa gene was programmed as the following. Primary denaturation temperature of 95° C for 5 minutes, 35 cycles of temperature including secondary denaturation temperature of 95 °C for 1 min, Annealing temperature of 54 °C for 1 min and Elongation at 72 °C for 1 min. The products of each PCR were stained with ethidium bromide in 1.5% agarose gel and then electrophoresed. Then, using a UV transilluminator and in the presence of a DNA marker (100 bp Fermentase), bands were observed and compared with the marker's pattern. The voltage for electrophoresis was 70V for 40-50 minutes.

Statistical analysis:

Data were analyzed using SPSS software (version 18) and described using percentages and ratios. The chi-square test was used to compare groups and P-value of <0.05 was considered as statistical significance level.

Table1- Used primers sequences and specific markers in this study

| Gene | Primer |
|--------------|---|
| <i>BabA2</i> | F 5'-AATCC AAAAA GGA GAAAA GTATGAAA-3' |
| | R 5'-TGTTAGTGATTTTCGGTGTAGGACA-3' |
| <i>Hpa</i> | F 5'-ATAAAGCTT TCG GTGGTG GAACG ATG-3' |
| | R 5'-TATCTC GAGTTG TCG GTT TCT TTTGC-3' |

Table 2- Frequency of hpa and babA2 genes in H.pylori sample The

| Peptic ulcer | Gastritis | Gastric cancer | Gene |
|--------------|-----------|----------------|----------------|
| 19(73.7%) | 21(58.3%) | 11(61.1%) | Positive babA2 |
| 7(26.9%) | 15(41.6%) | 7(38.8%) | Negative babA2 |
| 26 | 36 | 18 | Total babA2 |
| 22(84.6%) | 25(69.4%) | 10(55.5%) | Positive hpa |
| 4(15.4%) | 11(30.5%) | 8(44.4%) | Negative hpa |
| 26 | 36 | 18 | Total hpa |

RESULTS

The presence of BabA2 and Hpa genes in the clinical samples of 80 patients with gastrointestinal disorders caused by H.pylori (18 samples of gastric cancer, 36 samples of gastritis, 26 samples of peptic ulcer) were studied and analyzed using the PCR molecular method (Table 1 and 2). The frequency of positive cases for the BabA2 and Hpa genes was 51 cases (63.8%) and 57 cases (71.2%), respectively. In this study, the frequency of positive BabA2 and Hpa genes in peptic ulcer patients was higher than patients with gastritis and gastric cancer. However, there was no significant relationship observed between Hpa and BabA2 genes in gastric cancer, peptic ulcers and gastritis (P-value > 0.05).

DISCUSSION

Binding of H.pylori to gastric mucous and epithelial cells is an important step in the pathogenesis of this bacterium. Following the binding of H.pylori to the mucosal surfaces of the gastrointestinal tract, the host tissue become exposed to the bacterial enterotoxin. Adhesins, proteins, glycoconjugates and bacterial lipids are involved in the early stages of bacterial colonization (3,12). In Shirazi et al. study which was conducted in Tehran, although BabA2 gene frequency in gastric cancer samples was higher than other gastrointestinal diseases, no significant relationship was observed between this gene

and gastric cancer (3). The inconsistency between their findings and this study could be due to geographical and genetic diversity. In this study, the frequency of OMP of H. Pylori and their adhesion ability were studied. BabA2 protein is functionally active and can bind to blood group antigens (6,7). Ishaqi et al. reported no significant relationship between BabA2 genotype, gastritis and peptic ulcer but BabA2 association with gastric cancer was observed. Whereas in our study, no significant association was found between the presence of BabA2 gene, gastric cancer, peptic ulcer and gastritis. Although a large number of articles has indicated the association of BabA2 gene with various H.pylori-related diseases (6), this study similar to some others, observed no significant relationship. In Ghasemian, Safae and colleagues study in 1387, the frequency of BabA2 gene in 81 patients with gastritis, peptic ulcer and gastric cancer was reported as 68.2, 74.1 and 80%, respectively while the overall frequency of BabA2 gene was 71.6% (7). Hpa protein is another H.pylori OMP which is involved in binding and adhesion of H.Pylori to the gastric mucosa, as well as providing the opportunity for gastrointestinal inflammation (10). In a study conducted in the U.S by Carlson, Hpa gene frequency in gastritis, duodenal ulcer and gastric cancer patients was reported as 66%, 44% and 70%,

respectively. However, no significant association was reported between the presence of Hpa and gastric cancer (13).

CONCLUSION

The most H. pylori strains have these genes which increase the incidence of peptic ulcer in patients, harboring the bacteria and also increase the risk of other diseases such as gastritis and gastric cancer. BabA2 and Hpa genes were more frequent in patients with peptic ulcer and less frequent in

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patients with gastric cancer compared with other diseases.

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CONFLICT OF INTEREST

There are no conflicts of interest.

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