

Detection of *Helicobacter pylori* in Saliva of Patient with Oral Lichen Planus

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Lichen planus is a common, chronic inflammatory disease of the skin and mucous membrane for which no precise causes have been confirmed. But it is often connected with infections.

Helicobacter pylori (*H. pylori*) among various bacteria has been associated with the cause of gastritis, peptic ulcer and gastric cancer. Considering the similarities of histological features between gastric ulcer and oral ulcers, it is reasonable to assume that *H. pylori* might also be involved in the development oral mucosal ulceration. So we employed this study to investigate the possible involvement of *H. pylori* in the aetiology of erosive oral lichen planus.

We analyzed detection rate of *H. pylori* in saliva of patients with erosive oral lichen planus by nested PCR. As a result, it revealed a significant difference statistically by showing positivity in 16 to 21 (76.2%) saliva samples of patients group and in 11 of 44 (25%) saliva samples of control group ($P > 0.001$). We were able to suppose that *H. pylori* in saliva can be related to cause of erosive oral lichen planus.

Key wards: Erosive oral lichen planus, *Helicobacter pylori*, saliva

I. INTRODUCTION

Helicobacter pylori (*H. pylori*) has been associated with development of chronic gastritis, peptic ulcer and gastric cancer.^{1,2)} *H. pylori*, microaerophilous and gram-negative bacteria, exist mainly stomach in human and act important role of

development of gastrointestinal disease. But finding of *H. pylori* in saliva, plaque and periodontal pockets mean that oral cavity can be another reservoir of *H. pylori*.³⁾

The potential ability of *H. pylori* to colonize in oral cavity is related with selective and specific adhesion between *H. pylori*, *Fusobacterium nucleatum* and *Fusobacterium periodontium* in human dental plaque. Such selective coaggregations show that dental plaque is may be provided as a reservoir for pathogens outside of human stomach.⁴⁾ Also, additional characteristic of *H. pylori* is that it combine with salivary mucins which cover the oral epithelium. This is because sulfated oligosaccharide on salivary mucins provides the receptor structure for adhesion of *H. pylori* on oral surface.⁵⁾

Li et al⁶⁾ reported that they had found a specific DNA of *H. pylori* by PCR in 30 of 40 (75%) saliva samples of adults infected *H. pylori*, and

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Czésnikiewicz-Guzik et al⁷⁾ reported that as results of research about existence of *H. pylori*, they had found *H. pylori* in 51 (51%) gastric mucosa sample and in 54 (54%) saliva sample.

Because *H. pylori* exist in stomach and oral cavity like this, a causative role for *H. pylori* in the pathogenesis of oral mucosal ulcers has been suggested. Birek et al⁸⁾ proved that it is relation of oral aphthous ulcer and *H. pylori*, Sanli et al⁹⁾ reported results of endoscopy and histopathological research in upper gastrointestinal tract of patients with oral lichen planus.

But, Riggio et al¹⁰⁾ reported that results of research do not support a definitive aetiological role for *H. pylori* in recurrent aphthous ulcer, Shimoyama et al¹¹⁾ asserted that *H. pylori* might not have a direct association with oral ulcerations.

The purpose of our study was to investigate detection rate of *H. pylori* in saliva of patient with oral lichen planus, oral ulcerative disease, because of the similarities of histological feature between gastric ulcer and oral ulcers.

II MATERIALS AND METHODS

1. MATERIALS

21 Patients with oral lichen planus of erosive type diagnosed in Department of Oral Medicine, Dental Hospital, Chosun University were selected as the patients group, and 44 individuals whose oral condition was good clinically were selected as control group. The age of patients group was from 42 to 81 years (8 males and 13 females). The age of control group was 40 and over years (19 males and 25 females).

For the study, we collected saliva from them into 1.5ml microcentrifuge tube and the secreted saliva was stored frozen -20°C prior to experiments.

2. METHODS

1) DNA extraction

From collected saliva, we extracted the DNA

using AccuPrep[®] Genomic DNA Extraction Kit(Bioneer, Daejeon, Korea). First, we mixed 200 μ l of phosphate buffered saline(PBS), 20 μ l of Proteinase K(20mg/ml), and 200 μ l of Binding Buffer(GC) into 1.5ml microcentrifuge which contains saliva, then executed vortexing. After heating it for 10 minutes in 60°C, we added 100 μ l of isopropanol into a tube then vortexed it for 5 seconds. Such mixed solution was poured into prepared binding column tube to conduct centrifuge in 8,000rpm for one minute, then threw away the filtered solution.

We had repeated this process until the mixed solution had finally disappeared. 500 μ l of W1 Buffer was added, then conducted centrifuge in 8,000rpm for one minute, then we poured out the filtered solution, then 500 μ l of W2 Buffer was added, then conducted centrifuge in 8,000rpm for one minute, then poured out the filtered solution, then finally conducted centrifuge in 12,000rpm for one minutes. Binding column tube was moved to new sterilized 1.5ml tube, and 50 μ l EL Buffer was added into Binding column tube. After making it reaction in room temperature for one minute, centrifuge was conducted in 8,000rpm for one minutes. Then, filtered solution was stored in -20°C before being used in polymerase chain reaction(PCR).

2) The amplification by PCR

For the first PCR of *H. pylori* genomic DNA, 10 μ l template DNA, and 1 μ l of each primer (20 μ mole/ μ l) in HEPY Primer set (Bioneer, Daejeon, Korea) were added into AccuPower PCR Premix (Bioneer, Daejeon, Korea) which contains 10mM of Tris-HCl(pH9.0), 40mM of KCl, 1.5mM of MgCl₂, 250 μ m of dNTP, and 1U of Tag DNA Polymerase, and to make its total volume of 20 μ l, we added 8-mop in HEPY Primer set (Bioneer, Daejeon, Korea).

The PCR was performed under conditions of 94°C 5minutes, 1cycle: 94°C for 30 seconds, 62°C for 30 seconds, 72°C for 60 seconds, 33cycles: 72°C for 5 minutes, 1cycle in PCR thermocycler(Minicycler[™], MJ Research, U.S.A).

The mixture for second PCR of *H. pylori* genomic DNA, contained AccuPower PCR Premix (Bioneer, Daejeon, Korea) containing 10mM of Tris-HCl(pH9.0), 40mM of KCl, 1.5mM of MgCl₂, 250 μ m of dNTP, and 1U of Tag DNA Polymerase, with 2 μ l of first PCR product, and 1 μ l of each Primer(20pmole/ μ l) in HEPY Primer set (Bioneer, Daejeon, Korea), and to make its total volume of 20 μ l, we added 8-mop in HEPY Primer set (Bioneer, Daejeon, Korea). The PCR was performed under conditions of 94 $^{\circ}$ C 5minutes, 1cycle: 94 $^{\circ}$ C for 30seconds, 58 $^{\circ}$ C for 30 seconds, 72 $^{\circ}$ C for 60 seconds,



1 2 3 4 5 6 7 8 9 10 11 12
 Fig. 1. Result of nested PCR products to *H. pylori* DNA analyzed on 1.5% agarose gel electrophoresis
 Lane 1 : 100bp DNA Ladder
 Lane 2, 4, 5, 9, 10 : Saliva of patients with oral lichen planus [Positive]
 Lane 3, 6, 7, 8 : Saliva of patients with oral lichen planus [Negative]
 Lane 11 : Positive control (325bp)
 Lane 12 : Negative control

33cycles: 72 $^{\circ}$ C for 5 minutes, 1cycle in PCR thermocycler(MinicyclerTM, MJ Research, U.S.A).

Besides, for every PCR amplification, negative control and positive control in HEPY Primer set (Bioneer, Daejeon, Korea) had also amplified simultaneously. Therefore we excluded the possibility of contamination and false positive.

3) Electrophoresis

We applied on 3 μ l of PCR product on 1.5% agarose gel which includes ethidium bromide(0.5 μ g/ml) and after electrophoresis, we analyzed the size of PCR product using UV Transilluminator(Fig. 1). For Marker, 100 base pairs(bp) Ladder (Bioneer, Daejeon, Korea) was used.

4) Statistical analysis

To compare the detection rate of patients with oral lichen planus and control group, statistical analysis was executed χ^2 test with SPSS Windows 12.0 program.

III. RESULTS

Nested PCR amplification was executed with template DNA, which was extracted from saliva of 21 patients with oral lichen planus and 44 adults as control group.

From 1.5% agarose gel electrophoresis, we were able to observe amplified PCR product of 325base pair(bp) (Fig. 1). 16 patients (76.2%) with oral lichen planus and 11 adults from control group show positivity, showing statistical significance (Table 1).

Table 1. The detection of *H. pylori* in patients with oral lichen planus and control group

Group	Frequency	%	P-value
Patients (N=21)	16	76.2	< 0.001
Control (N=44)	11	25	

N : Number of individuals

IV. DISCUSSION

Lichen planus is a common, chronic inflammatory disease of the skin and mucous membranes.

The disease has been classified into five groups, namely reticular, erosive, atrophic, hypertrophic and bullae form. The reticular form is the most common variant and is characterized by small white papule (Wickham's striae). The erosive form is the second most frequent variant and is characterized by small or extensive painful erosions with isolated papules or lines at the periphery.¹²⁾

The aetiology of lichen planus is unknown, but it is often connected with infections.¹³⁾ *H. pylori* among various bacteria lives beneath the gastric mucous layer, on the surface of epithelial cells and produce various enzymes in abundance—urease is one of the most common enzymes associated with *H. pylori* that has an essential role in the colonization. Urease creates a non-acidic microambiance that allows the bacteria to survive the sterilizing effect of chlorhydric acid. The bacteria travel through the viscous mucus film that protects the gastric epithelium, adhere to the epithelial cells, and begin to replicate. An immune response is subsequently initiated by neutrophils, which results in gastric tissue alteration.¹⁴⁾

Considering the similarities of histological features between gastric ulcer and oral ulcers, it is reasonable to assume that *H. pylori* might also be involved in the development of oral mucosal ulcers.^{10,15)} So a causative role for *H. pylori* in the pathogenesis of oral mucosal ulcers has been suggested previously.⁸⁾ Leimola-Virtanen et al¹⁵⁾ reported *H. pylori* DNA can be found in separate oral mucosal ulcers in apparently immunocompetent adults, Long et al¹⁶⁾ reported that *H. pylori* might in some way associated with recurrent aphthous ulcer, which in turn is associated with an increased incidence of digestive disease, and Sanli et al⁹⁾ asserted that patients with lichen planus should be evaluated for possible gastrointestinal involvement with endoscope. But Vainio et al¹³⁾ reported that the prevalence of *H. pylori* infection in patients with

lichen planus was not significantly different from that in patients with other skin disease, Daudén et al¹⁷⁾ reported that the more virulent cytotoxin-associated gene A(Cag A) positive strains of *H. pylori* are not strongly associated with lichen planus.

Our study aimed to investigate the possible involvement of *H. pylori* in the aetiology of erosive oral lichen planus. In our study, *H. pylori* was detected in 16 of 21 (76.2%) saliva samples of patients with oral lichen planus and in 11 of 44 (25%) saliva samples of control group.

Therefore, we were able to assume that *H. pylori* in saliva can be relate to cause of erosive lichen planus.

Saliva is a relatively simple biological sample for diagnosis of *H. pylori* infection. Recently, many analysis methods using PCR have been developed to detect *H. pylori* in oral cavity. The PCR offers the advantage of detecting *H. pylori* in area outside the stomach where cultures usually fail, or when biopsy material is not available. PCR performed on DNA extracted from bile juice, stools and oral secretions can detect as few as two, or even a single, *H. pylori* organism.^{8,18)}

Most of PCR assays are based on arrangement of urease gene and 16S ribosomal RNA gene.¹⁹⁾

Lamaroon et al²⁰⁾ studied the association *H. pylori* and recurrent aphthous ulcer by a highly sensitive technique, nested PCR and Song et al²¹⁾ reported than nested PCR showed superior detection ability to *H. pylori* than [13C] urea breath test after research in saliva and dental plaque. We also were able to get satisfying result by nested PCR using manufactured HEPY Primer set with control DNA (Bioneer, Daejeon, Korea).

V. CONCLUSION

We analyzed detection rate of *H. pylori* in saliva of patients with oral lichen planus by nested PCR to investigate the possible involvement of *H. pylori* in the aetiology of erosive oral lichen planus. As a result, it revealed a significant difference

statistically by showing positivity in 16 of 21 (76.2%) saliva samples of patients group and in 11 of 44 (25%) saliva samples of control group.

In conclusion, we were able to suppose that *H. pylori* in saliva can be related to cause of erosive oral lichen planus.

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국문요약

구강 편평태선 환자의 타액에서 *Helicobacter pylori*의 검출

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편평태선은 피부와 점막에 발생하는 흔한 만성 염증성 질환으로 정확한 원인은 잘 알려져 있지 않으나 종종 감염과 관련되어 있다. 다양한 박테리아 중 *Helicobacter pylori* (*H. pylori*)는 위염, 위·십이지장 궤양 그리고 위암과 관련되어 있다. 위궤양과 구강 궤양들의 조직학적 특징의 유사성을 고려할 때 *H. pylori*는 구강 점막궤양의 발생과 관련 있음을 추론할 수 있다.

따라서 미란성 구강편평태선의 발생에 *H. pylori*가 관련 있는지를 조사하기위해 이 연구를 수행하였다.

미란성 구강편평태선을 가진 환자의 타액을 중합효소연쇄반응에 의해 분석한 결과 21명의 환자 중 16명 (76.2%)에서 *H. pylori*가 검출되었고, 대조군은 44명 중 11명 (25%)에서 *H. pylori*가 검출되어 통계적 유의성을 나타내었다($P > 0.001$). 이상의 결과를 종합해 볼 때 타액내 *H. pylori*는 미란성 구강편평태선의 발생에 원인이 될 수 있음을 추론할 수 있었다.

주제어: 미란성 구강편평태선, *Helicobacter pylori*, 타액
