Helicobacter pylori in both the sinuses and the stomach*

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Abstract

Background: The role played by Helicobacter pylori in the sinuses, and its association with the same organism's gastric infection, are still unclear.

Methods: In order to compare H. pylori colonization patterns in the nose and stomach we conducted a cohort analysis of 14 patients, eligible for sinus surgery due to chronic medically refractory rhinosinusitis, who were tested for simultaneous presence of H. pylori, by histology, culture and polymerase chain reaction, in pathologic sinus tissue collected during surgery and in gastric mucosa obtained through gastroduodenal endoscopy.

Results: H. pylori DNA was found in the sinus mucosa of 15.4% of patients with chronic rhinosinusitis, and all of them showed concurrent H. pylori stomach infection. Sinus colonization was not found without simultaneous gastric colonization, although most patients with gastric infection did not have the bacterial DNA in their sinuses. H. pylori's presence in the nose was not associated with local inflammatory status, and no cultures could be obtained from any of the sinus tissue samples, including those positive for H. pylori DNA.

Conclusions: Only H. pylori DNA, and not the culturable active form of the microorganism, could be found in the sinus mucosa of some patients with H. pylori gastric infection. We could not find evidence, however, that the bacterium's presence in the nose contributes to local mucosal inflammation.

Key Words: chronic rhinosinusitis, Helicobacter pylori, gastric infection

Introduction

Helicobacter pylori DNA has been detected in a number of extra-gastric locations, such as the oral cavity, tonsils and adenoids⁶-⁷, and even the middle ear and paranasal sinuses⁸-¹⁵. However, the significance of the microorganism's presence at these aero-digestive and respiratory sites is still unclear. It is believed that the oral cavity is an important reservoir for H. pylori, that contributes to the oral-oral route of transmission and acts as a source of stomach re-infection⁶. Others suggest that the bacterium may be capable of causing damage to the aero-digestive and respiratory mucosa in the same way it does to the gastric mucosa⁹. However, this capacity for extra-gastric disease remains unproved, and some authors argue that it even seems unlikely.

In order to initiate tissue damage, not only does H. pylori require a set of events unique to the gastric milieu, but also culturable active forms of H. pylori have only been recovered outside the stomach in the aero-digestive tract in the vicinity of the upper esophagus (i.e. in tracheal secretions in intubated patients), and never as far away as the bronchi, lung, or the upper respiratory tract¹⁰. That far from the stomach, only the microorganism's DNA suggests the presence of H. pylori at such locations.

We conducted the present investigation in order to see how H. pylori colonization patterns in the nose and stomach are associated in chronic rhinosinusitis patients, as a way to also provide information on the eventual existence of an interaction between two seemingly unconnected pathologies, one in the digestive
Enrolment required that the patients also agree to undergo a chain reaction (PCR) to amplify genomic DNA. Depending on the inflammatory infiltrate cell density, inflammation was classified as mild, moderate, or severe (1+, mild; 2+, moderate; 3+, severe). All patients were subjected to: 1) histopathological examination, to assess the overall degree of local inflammation (graded as 0+, absent; 1+, mild; 2+, moderate; 3+, severe inflammation, depending on the inflammatory infiltrate cell density); 2) histology to identify *Helicobacter pylori* in Warthin-Starry silver stained gastric mucosal biopsies, also graded as 0+, no bacteria found; 1+, mild; 2+, moderate; 3+, severe density colonization); and 3) microbiological culture of gastric biopsy specimens specific for *H. pylori*.

All sinus and stomach tissue samples were collected aseptically, transported to the laboratory at 4°C in Portagerm pylori (bioMérieux, France), and processed less than four hours after collection. For microbiological culture, a drop of tissue macerate was plated onto *H. pylori*-selective medium (Brucella supplemented with 10% horse blood and Brucella supplemented with 10% horse blood and Oxoid *H. pylori*-selective supplement (Dent)) and incubated at 37°C in a microaerophilic atmosphere (Campygen, Oxoid) for up to 15 days. Colonies were then tested for urease, catalase and oxidase, and motility and were stained with Gram stain.

Polymerase chain reaction

For PCR testing, two sets of primers targeting the 16S rRNA and 23S rRNA genes of *H. pylori* were used, chosen to maximize detection as together they target sequences common to virtually all *H. pylori* strains. The first primer pair, PCR-G, amplifies a 780bp fragment from the 16S rRNA gene specific for the *Helicobacter* genus, Helico F – 5’ CTATGACGGGATCCGGC 3’ and Helico R – 5’ CTCAGGACACGCTGAC 3’. The second primer pair, PCR-S, amplifies a 267 bp fragment from the 23S rRNA gene of the *H. pylori*, HPYS - 5’ CGCATGATATTCCCATTAGCAGT 3’ and HPYA - 5’ AGGTAAAGGATGCGTGCT 3’. Ethidium bromide-stained agarose gel electrophoresis was employed for separating the PCR products according to their size. When interpreting the results, we considered a positive genus-specific PCR test a sign of presence of DNA for one of the various *Helicobacter* species, whilst the combination of a positive genus-specific with a positive species-specific PCR test was taken as a clear sign of presence of the actual *H. pylori* species.
Table 1. Status of gastric and nasal inflammation and *H. pylori* infection for all patients.

<table>
<thead>
<tr>
<th>Patient</th>
<th>Breath test</th>
<th>Stomach</th>
<th>Sinus</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Inflammation</td>
<td><em>H. pylori</em> histology</td>
<td><em>H. pylori</em> culture</td>
</tr>
<tr>
<td>1</td>
<td>Pos</td>
<td>Antrum: 2+</td>
<td>Antrum: 3+</td>
</tr>
<tr>
<td>2</td>
<td>Neg</td>
<td>Antrum: NA</td>
<td>Antrum: NA</td>
</tr>
<tr>
<td>3</td>
<td>Pos</td>
<td>Antrum: NA</td>
<td>Antrum: NA</td>
</tr>
<tr>
<td>4</td>
<td>Pos</td>
<td>Antrum: 2+</td>
<td>Antrum: 0+</td>
</tr>
<tr>
<td>5</td>
<td>Neg</td>
<td>Antrum: 1+</td>
<td>Antrum: 0+</td>
</tr>
<tr>
<td>6</td>
<td>Pos</td>
<td>Antrum: 2+</td>
<td>Antrum: 3+</td>
</tr>
<tr>
<td>7</td>
<td>Pos</td>
<td>Antrum: NA</td>
<td>Antrum: NA</td>
</tr>
<tr>
<td>8</td>
<td>Pos</td>
<td>Antrum: 2+</td>
<td>Antrum: 3+</td>
</tr>
<tr>
<td>9</td>
<td>Pos</td>
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<td>10</td>
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<tr>
<td>11</td>
<td>Pos</td>
<td>Antrum: 2+</td>
<td>Antrum: 0+</td>
</tr>
<tr>
<td>12</td>
<td>Neg</td>
<td>Antrum: 1+</td>
<td>Antrum: 0+</td>
</tr>
<tr>
<td>13</td>
<td>Pos</td>
<td>Antrum: 1+</td>
<td>Antrum: 0+</td>
</tr>
<tr>
<td>14</td>
<td>Pos</td>
<td>Antrum: NA</td>
<td>Antrum: NA</td>
</tr>
</tbody>
</table>

Neg = negative result. Pos = positive result. Inflammation grade: 0+= no inflammation; 1+= mild inflammation; 2+= moderate inflammation; 3+= severe inflammation. *H. pylori* histology: 0 = no bacteria found; 1+ = low density of bacteria; 2+ = moderate density of bacteria; 3+ = high density of bacteria. *H. pylori* PCR: G - genus pair of primers; S - species pair of primers. NA = data not available.

**Histology**

For histological identification of *H. pylori* in paraffin sections of sinus tissue, the modified Warthin-Starry silver staining kit (Merck, Germany) was used. The grading of the histological material, as well as all the molecular biology procedures, were performed after the identification of the samples was masked.

**Statistics**

Statistical analysis employed descriptive statistics and the Spearman’s rank correlation coefficient (Spearman’s rho).

**Results**

The study group included 8 patients with asthma, 6 with allergic rhinitis, and 2 with intolerance to nonsteroidal anti-inflammatory drugs (NSAIDs). Other co-morbidities included: hypertension (n=2), hypothyroidism (n=1), goiter (n=1), angioma of the liver (n=1), ankylosing spondylitis (n=1), fibromyalgia while on NSAID’s (n=1), gastroesophageal reflux disease (GERD) (n=2) and chronic gastritis (n=1). Two patients were undergoing revision sinus surgery, and three admitted having taken an oral antibiotic in the immediate three months prior to surgery. One subject had history of *H. pylori* gastric infection treatment, while five
were taking a PPI until the day before surgery. The patient with ankylosing spondylitis was under treatment with sulfasalazine at the time of the surgical procedure, and the stiffness of his entire spine was so severe that the gastroenterologist was unable to perform the required gastroduodenal endoscopy, even under general anesthesia. Another patient underwent upper gastrointestinal endoscopy outside of the time frame imposed by the strict study criteria (either intra-op or up to two weeks post sinus surgery), so the gastric results were excluded from the analysis. Two patients had histology results for one gastric site only, and in one patient the gastric histological results were altogether absent.

The inflammatory status of the sinus mucosa at the time of surgery for each patient is displayed in Table 1. Severe inflammation was not encountered in any subject; moderate inflammation was found in 50% (n=5) of the asthmatics and in 16.6% (n=1) of the non-asthmatics, whilst mild inflammation was encountered in 50% (n=4) of the asthmatics and in 83.3% (n=5) of the non-asthmatics. A total of 60% of the antrum biopsies showed moderate inflammation, with mild inflammation occurring in 40%, whereas in corpus biopsies 70% had mild and 30% moderate inflammation.

Histological identification of H. pylori in gastric mucosa biopsies of the antrum was negative in 50% of the cases (n=5), revealing mild infection in 10% (n=1), moderate infection in 10% (n=1), and severe infection in 30% of the cases (n=3). For the corpus biopsies, the results were negative in 50% (n=5), with mild infection in 10% (n=1), moderate in 30% (n=3), and severe infection in 10% (n=1).

The results of cultures of gastric mucosa biopsies were positive for H. pylori in 66.7% of the cases (n=8), whilst the PCR-G test was positive in 54.5% (n=6) of the antrum samples and in 50% (n=5) of the corpus samples, and the PCR-S test was positive in 72.7% (n=8) of the antrum samples and in 66.7% (n=8) of the corpus samples.

A total of 66.6% of the patients receiving PPI treatment were H. pylori negative.

Table 2 shows the correlations between the variables found to

<table>
<thead>
<tr>
<th>Variables</th>
<th>Correlation Coefficient</th>
<th>p = value</th>
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</thead>
<tbody>
<tr>
<td>PCR-G results in the nose</td>
<td>PCR-G results in the antrum</td>
<td>0.655</td>
</tr>
<tr>
<td>PCR-S results in the nose</td>
<td>Previous anti-H. pylori treatment</td>
<td>0.677</td>
</tr>
<tr>
<td>Breath test</td>
<td>Cultural identification of H. pylori in the stomach</td>
<td>0.625</td>
</tr>
<tr>
<td>Cultural identification of H. pylori in the stomach</td>
<td>Inflammatory status in the antrum</td>
<td>0.802</td>
</tr>
<tr>
<td>Histological identification of H. pylori in the antrum</td>
<td>Inflammatory status in the antrum</td>
<td>0.769</td>
</tr>
<tr>
<td>Histological identification of H. pylori in the antrum</td>
<td>Histological identification of H. pylori in the corpus</td>
<td>0.732</td>
</tr>
<tr>
<td>Breath test</td>
<td>PCR-G results in the corpus</td>
<td>0.655</td>
</tr>
<tr>
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<td>0.936</td>
</tr>
<tr>
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<td>PCR-G results in the corpus</td>
<td>0.816</td>
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<tr>
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<td>PCR-G results in the antrum</td>
<td>0.828</td>
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<tr>
<td>Histological identification of H. pylori in the corpus</td>
<td>PCR-G results in the antrum</td>
<td>0.763</td>
</tr>
<tr>
<td>PCR-G results in the antrum</td>
<td>PCR-G results in the corpus</td>
<td>1</td>
</tr>
</tbody>
</table>
have statistical significance.

Regarding *H. pylori* presence in the sinonasal mucosa, the species-specific PCR identification was positive in only two cases (15.4%), although 69.2% had sinonasal samples with positive genus-specific PCR results. The first case of positive *H. pylori* DNA in the sinuses had a positive breath urea test and tested positive for *H. pylori* in the stomach by histology, bacterial culture and PCR. The second case, however, had a negative breath test and tested negative for gastric *H. pylori* by histology and bacterial culture, but the species-specific PCR identification test in the stomach was positive (result confirmed at a different laboratory). All attempts to grow *H. pylori* in culture medium from sinonasal samples failed.

**Discussion**

The characteristics of our investigation, with patients undergoing simultaneous extensive sinus and stomach tissue sampling, and with various tests concurrently performed, necessarily restricts the number of patients enrolled in the study, and as a result may not be able to show statistical significance. The available data, however, are sufficient to allow for important conclusions.

Since urease production is a hallmark of active gastric infection, it is perhaps no surprise that most, but not all, patients in our study with a positive urease breath test had indeed *H. pylori* gastric infection, confirmed by histopathology, culture, and/or PCR testing. The breath test was positive in one case where *H. pylori* was detected in the sinuses, but was negative in the other. As the majority of patients in our study had positive breath tests and no evidence of *H. pylori* in their sinuses, this apparently renders the test unsuitable for the specific identification of the bacterium in the nose.

Our data actually reveals that, in spite of the relevant number of patients with positive Helicobacter-genus DNA detected by PCR in the sinus mucosa (which certainly merits separate investigation), specifically *H. pylori* DNA was only found in the sinonasal mucosa of about 15.4% of chronic rhinosinusitis surgical patients. These results are in agreement with previously published data[3-11]. Admittedly, this prevalence could indeed be greater if the bacterium in the nose would follow a similar mosaic pattern of mucosal infection as in the stomach, with patches of diseased mucosa alternating with non-infected normal mucosa, in which case it would require repeated and extensive sampling to allow for a positive *H. pylori* result (five different sites, according to some)[6].

Both cases with *H. pylori* DNA in the sinuses had *H. pylori* DNA simultaneously present in their stomachs, but only about 25-28.6% of the patients with positive gastric species-specific PCR tests had *H. pylori* DNA in their sinuses. This suggests that if the bacterium is to be encountered in the nose, its DNA has to also be present in the stomach, but that not all *H. pylori* gastric infections are necessarily associated with *H. pylori* colonization of the sinuses.

Our data show that while there is a positive correlation between cultural and histological identification of *H. pylori* in the stomach and gastric inflammation, we found no such correlation between *H. pylori* and site inflammation in the nose (Table 2). Therefore, it is perhaps not farfetched to admit that *H. pylori* presence in the nose and sinuses does not contribute to local mucosal inflammation.

The diagnosis of allergic rhinitis, asthma or intolerance to NSAIDs, was not found to statistically relate to either positive or negative sinonasal *H. pylori* results, suggesting that sinonasal *H. pylori* colonization may occur regardless these co-morbidities are present or not.

Critically, *H. pylori* could only be recovered from the nose in the DNA form, as all the attempts to culture the bacterium from nasosinusal sites failed. This inability to culture *H. pylori* could be due to the presence of too few microorganisms to be detected, or the simultaneous presence of too many types of other bacteria in the nose that inhibit growth of *H. pylori*. However, it has been shown that the bacterium can be cultured from adverse environments such as the air sampled during vomiting or from a tracheostomy tube[7]. So it is admissible that the reason may not have to do with the method but with the possibility that, in the sinuses, either the organism is represented just by fragments of its DNA and that these are destined to transiently remain there just for a limited time, or the microorganism is, in fact, in a dormant state that precludes culture, and is destined to remain for a long time in the sinuses.

The bacterium is, indeed, known to be able to resist harsh environments by changing to a dormant, inactive state, a non-culturable coccoid form that could still be potentially viable, later on, in the stomach[21,22]. It is therefore possible that *H. pylori* may lay dormant for long periods of time, using the nose and the sinuses as reservoirs, waiting for an eventual return to an active form, either to cause gastric re-infection or to participate in the oral-oral route of transmission. The fact that in a previous study no statistical difference was observed between *H. pylori* nasal colonization in patients with sinusitis when compared to the control group[6], lends credibility to the ‘nose as a reservoir’ hypothesis and reinforces our conviction that the bacterium’s presence in the nose does not contribute to the local inflammatory status.

To account for the bacterium’s presence in the nose, the hypothesis of gastric-nasal transmission seems the most logical explanation since a significant number of patients with GERD and laryngopharyngeal reflux (LPR) also have *H. pylori* gastric colonization. It has been shown that the microorganism has a positive tropism for mucins[23] and mucins cover and protect the sinus and mouth epithelia[24] and is also able to invade epithelial cells[25]. The presence of *H. pylori* in the sinuses could then be regarded as a biomarker of the extent of LPR in the upper
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airway tract. However, at this time, we have no definitive proof of this, and we simply cannot rule out the possibility that the bacterium may use, in alternative or in conjunction, other routes to have its DNA reach the sinuses, for instance, via lymphatic or vascular transmission, from either the stomach or any other extra-digestive site.

Also, the presence of *H. pylori* in the sinuses apparently does not support a local pathogenic role does not entirely rule out the possibility that the bacterium may influence the course of an inflammatory disease of the sinuses. *H. pylori* gastric infection is known to cause a vast array of systemic effects, including a strong immunologic response and gastrin and cytokine release from the stomach mucosa, all of which may indirectly affect chronic inflammation in any part of the respiratory system\(^2\). Definitive proof of the clinical relevance of these *H. pylori*-induced systemic effects is, however, still lacking.

**Conclusion**

Our results suggest that, regardless of how *H. pylori* reaches the sinuses, its presence there does not seem to contribute to the local inflammatory status of the respiratory mucosa. The fact that *H. pylori* could not be cultured from nose samples and is only present in its DNA form, suggests either a transient presence of parts of its genome in the sinuses, or, instead, what could be seen as a defensive adaptive reaction in preparation for a more or less lengthy stay at an inhospitable location, a change to a viable non-culturab form, from which *H. pylori* could hypothetically regain activity, to either play a role in the oral-oral route of transmission or in an eventual gastric re-infection.

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**Authorship contribution**

PBD: Concept and design, data acquisition and analysis, drafting and final approval of the manuscript, accountability for all aspects of the work. TM: Data acquisition and analysis, final approval of the manuscript. MS: Data acquisition and analysis, draft revision, final approval of the manuscript. PLA: Data acquisition and analysis, draft revision, final approval of the manuscript. JV: Data acquisition and analysis, final approval of the manuscript. AMC: Data acquisition and analysis, draft revision, final approval of the manuscript. JV: Concept and design, data acquisition and analysis, drafting, final approval of the manuscript, accountability for all aspects of the work.

**Conflict of interest**

The Jorge Vitor’s lab has received funding from New England Biolabs Inc., USA, since 1995. All the other authors have no conflicts of interest to disclose.

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