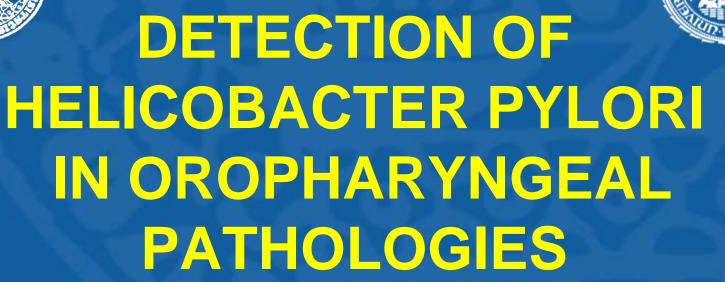
UNIVERSITAS CAROLINA PRAGENSIS



First Faculty of Medicine, Prague, Czech Republic

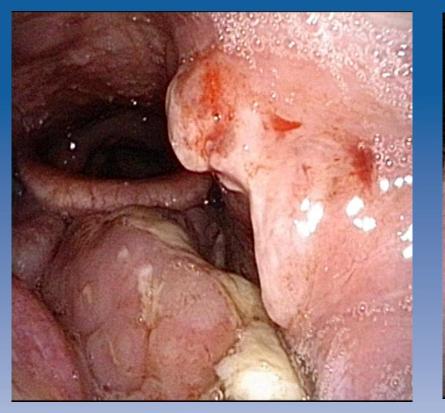


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INTRODUCTION

Increasing rate of oropharyngeal malignancies in recent years in certain countries forced investigators to study possible role of chronic infections in development and progression of cancer in oropharyngeal area.





Tonsillar cancer, male patient , age37

AIMS OF THE STUDY

- To confirm presence of *H. pylori* in the lymphatic tissue of oropharyngeal area in patients with different oropharyngeal pathology requiring surgical intervention, using different techniques for both direct and indirect diagnostics
- To genotype DNA of H. pylori isolated from peroperative biopsies for presence of *cagA* and variability of *vacA* genes using synthetised TaqMan hybridisation probes
- To confirm infection of each patient by at least one of different diagnostic tests: UBT, gastroscopy, specific antibody detection in blood (ELISA, WB)
- To compare, in cooperative patients undergoing gastroscopy with positive result, genotypes of isolates from both the gastric and oropharyngeal samples 3

MATERIAL

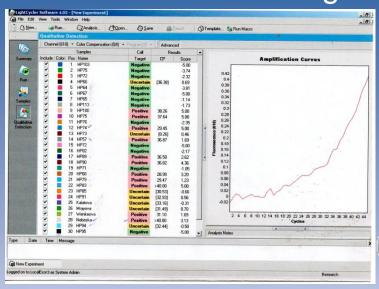
Peroperative biopsies taken from patients with following diagnoses: Tonsillar cancer - 42 Oropharyngeal cancer - 10 Chronic tonsillitis – 78 SAS



METHODS

- Sample collection: biopsies placed directly to Remel M4RT sample collection and transport media. Pre-analytic storage at +4°C (up to 5 days), long-term storage at -20°C
- DNA Isolation: Roche MagNA Pure Compact Extractor: MagNA Pure Compact Nucleic Acid Isolation Kit I, Protocol 400-100
- Real-time PCR: Instrument Roche LightCycler v.





METHODS

- Primers and TaqMan Probes cagA, vacA s1a, vacA s1b, vacA s2, vacA m1 vacA m2 (vanDoorn et al. 1999) synthetised and optimised by TIB MolBiol Berlin, Germany
- Urea Breath Test: in order to confirm H. pylori gastric infection, performed according manufacturer's instruction in package insert
- Detection of specific IgM, IgA and IgG: by ELISA AND IMMUNOBLOT (WB) techniques based on commercial diagnostic tests

Tonsillar Cancer Patients:

Total: 42

| HP not detected | HP detected | cagA detected | cagA not detected | |
|--|-------------|-------------------------|-------------------------|--|
| 8 | 34 | 4 | 30 | |
| 19.04 % | 80.95 % | 0.095 %/11.76 % | 71.43 %/ <u>88.23 %</u> | |
| Most frequent genotype: cagA- vacA s1b m2 (8 - 23.53 %) | | | | |
| Oropharyngeal cancer patients: | | Total: 10 | | |
| HP not detected | HP detected | cagA detected | cagA not detected | |
| 4 | 6 | 2 | 4 | |
| 40 % | 60 % | 20 % / 33.33 % | 40 %/ 66.66 % | |
| Most frequent genotype: cagA- vacAs1b m2 (5 - 83.33 %) | | | | |
| Summarised Data: Total: 52 cagA- vacAs1bm2 (13 - 32.5 %) | | | | |
| | | | | |
| HP not detected | HP detected | cagA detected | cagA not detected | |
| 12 | 40 | 6 | 34 | |
| 23.07% | 76,93 % | 11.54 %/ <i>15.00 %</i> | 65.38 %/ <i>85.00 %</i> | |

Chronic Tonsillitis Patients:

Total: 78

| HP not detected | HP detected | cagA detected | cagA not detected |
|-----------------|-------------|-----------------|-------------------------|
| 13 | 65 | 14 | 51 |
| 16.66 % | 83.34 % | 17.95 %/21.54 % | 65.39 %/ 78.46 % |

Most frequent genotype: cagA- vacAs1b m1 (20), cagA- vacAs2 m1 (7)

SAS – originaly chosen as Control Group Total: 29

| HP not detected | HP detected | cagA detected | cagA not detected |
|-----------------|-------------|-------------------------|-------------------------|
| 5 | 24 | 19 | 5 |
| 17.24 % | 82.76 % | 65.57 %/ 79.16 % | 17,24 %/ <i>20.84 %</i> |

Most frequent genotype: cagA – vacAs1a m2 (4), cagA+vacAs1a m2 (4)

- In summer 2012 a new commercial test for detection of Helicobacter pylori – Bioron RealLine Fla-Format was launched on the market. We have used it to confirm our results
- Testing of all our genotyped isolates showed a 100% agreement in results
- Total samples: 130 + 29

n= 159

| GENOTYPING | BIORON Fla FORMAT rt-PCR | | |
|------------|--------------------------|----------|--|
| Rt- PCR | POSITIVE | NEGATIVE | |
| POSITIVE | 129 | 0 | |
| NEGATIVE | 0 | 30 | |

 This supports evidence of H. pylori presence in tissue of oropharyngeal malignancies

- Genotyping H. pylori strains obtained from different body systems – stomach and oropharynx:
- First 6 patients have completed both the oropharynx peroperative biopsy and GIT biopsy
- Hypothesis to be confirmed: Individuals are colonised by the same bacteria strain
- Result: in 3 patients, genotyping of cagA and vacA genes showed identical result.
- In 3 patients comparison showed difference in both genes tested
- Eventhough no strong data on the issue exist yet, we are able to conclude, H. pylori strains may be different.

DISCUSSION

- H. pylori is a declared type I carcinogen (IARC 1994), Supposed 3 ways of its carcinogenic action are:
- <u>Direct mutagen action</u> (interaction of intracellular signaling molecules and CagA may predispose cells to accumulate genetic changes promoting multistep carcinogenesis – Hatakeyama 2006)
- <u>VacA production</u> causes immunosupression by blocking T cell proliferation (Boncristiano et al. 2003)
- <u>Cell proliferation induction by cytokins and</u> <u>regulatory molecules</u> involved in cell transformation and tumor formation – TGF, EGF and NOS (Konturek 1997, Gallo 1998, Sakaguchi 1999, Keates 2001, Gobert 2002, Schiemann 2002, Wang 2002)

| Di Bonaventura | 2001 | 75 tonsillar swabs and biopsies | PCR | 0 (0 %) |
|-------------------|------|--|-------------------------------|--|
| Cirak | 2003 | 23 tonsillar and adenoid tissue | PCR (16S ribosomal RNA, CagA) | 7 (30 %) H. pylori positive, 5 out of (71 %) cagA+ |
| Yilmaz et. al. | 2005 | 38 adenoid tissue, mesotitis secretion | PCR (23S ribosomal RNA) | 12 (67 %)positives in mesotitis , 1 (5 %) in adenoid tissue |
| Bitar | 2005 | 25 adenoid tissue | nested PCR (UreA) | 0 (0 %) positives in nested PCR |
| Bulut | 2006 | 71 tonsillar and adenoid tissue | PCR (CagA - glmM gene) | 29 (24,6 %) H. pylori positive, 17 out of (58,6 %) CagA + |
| Bitar | 2006 | 28 adenoid tissue | PCR (ureC-gene) | 0 (0 %)PCR positive |
| Yilmaz et. al. | 2006 | 22 adenoid and tonsillar tissue, mesotitis secretion, promotorium mucose | PCR (16S RNA) | mesotitis secretion 7 PCR+, promotorium 7 PCR+, adenoid tissue 14 (64 %) PCR+, tonsilar tissue 14 (64 %) PCR+ |
| Eyigor et. al. | 2009 | 55 35 adenoid tissues, 20 tonsills | PCR (glmM gene) | 0 % PCR positives |
| Vilarinho et. al. | 2010 | 62 tonsillar and adenoid tissue | PCR - DNA (vacA gene) | 0 PCR positives |
| Abdel-Monem | 2011 | 20 tonsillar and adenoid tissue | PCR (ureC-gene) | 5 (16,6 %) PCR positives |

CONCLUSIONS

- Our study provides data supporting hypothesis, that Helicobacter pylori participates in oropharyngeal pathology more than it has been supposed
- Long-term colonisation of tonsillar tissue by H. pylori lacking CagA protein, concidered to be responsible for GIT pathology, may lead to development of chronic inflammation and alteration of immune system mechanisms resulting in uncontrolled formation of tonsillar neoplasias
- Eventhough detailed recognition of this process requires further investigation, prevention through eradication of so called non.pathogenic strains of the bacteria by antibiotics may help to reduce incidence of oropharyngeal neoplasias



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THANK YOU

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