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► To cite this version:

Sarah Atallah, Beatrice Berçot, Valérie Laurence, Caroline Hoffmann. Association of Mycoplasma Hominis and head and neck cancer with unknown primary. European Annals of Otorhinolaryngology, Head and Neck Diseases, 2020, 137 (1), pp.69 - 71. 10.1016/j.anorl.2019.05.020 . hal-03490046

HAL Id: hal-03490046

<https://hal.science/hal-03490046>

Submitted on 21 Jul 2022

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ASSOCIATION OF MYCOPLASMA HOMINIS AND HEAD AND NECK CANCER WITH UNKNOWN PRIMARY

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Abstract

Introduction

Beside *HPV* infection, there is currently no evidence of association between head and neck squamous cell carcinomas and microbial infections. We report the case of a cervical squamous cell carcinoma by *Mycoplasma hominis*.

Case summary

A 20-years-old woman, consulted for a swelling on the left cervical side. Clinical examination found a large fixed mass. Biological tests found no evidence of infection. Biopsies of the cervical lesion diagnosed an HPV negative squamous cell carcinoma. Microbiological tests of 16sRNA identification showed the presence of *Mycoplasma hominis* in the 3 specimens. The patient was treated by induction chemotherapy associated to antibiotherapy, followed by chemo-radiotherapy.

Discussion

The present case suggests that oropharyngeal infection by *Mycoplasma hominis* might be more frequent than expected, that 16sRNA is an efficient technic to isolate this pathogen and finally that further studies are required to document its potential oncogenic role in head and neck squamous cell carcinomas.

Keywords:

Mycoplasma, infection, squamous cell carcinoma, cervical lymphadenopathy, 16S RNA

Introduction

Beside Human papilloma virus (HPV) infection implicated as an oncogenic factor in oropharyngeal head and neck squamous cell carcinomas (HNSCC) [1], there is currently no evidence of association between HNSCC and microbial infections. To our knowledge, we report the first case of a cervical SCC node with unknown primary, which was HPV negative, but colonized by *Mycoplasma hominis* (*Mh*).

Case report

A 19-years-old woman, with no history of smoking or alcohol use, consulted for a swelling of the left side of her neck. Clinical examination found a large painless fixed mass and no fever. Biological tests found no evidence of infection. Computerized tomography scan (CT scan) revealed a 10cm mass, with a hypodense core and a peripheral enhancement, invading the cervical muscles and surrounding the primary carotid artery. Ultrasound-guided biopsies (3 samples) of the unresectable cervical lesion diagnosed an HPV negative (p16 and Polymerase Chain Reaction (PCR)) squamous cell carcinoma. Because of the young age of the patient, an infection was still suspected and microbial tests were performed on the same biopsies. The 16S ribosomal RNA (16s rRNA) microbiological testing showed the presence of *Mh* in all the specimens: PCR product's sequence showed 100% homology with the gene coding for *Mh* 16S rRNA (GenBank accession number AF443616.3, BLAST search). The serologies performed, HIV, B and C hepatitis were negative. PCR technique associated to mass spectrometry (ESI-MS) found no *Mh* in an oropharyngeal biopsy performed secondarily. Cultures from genital tract samples were negative for *Mh*.

An MRI, a TEP-CT and a left tonsillectomy were performed, but were unable to identify the primary carcinoma (Fig. 1).

The patient was treated by induction chemotherapy (Docetaxel-Cisplatin-5FU) associated to fifteen days of antibiotherapy (josamycin) and followed by chemo-radiotherapy. After 13 months of follow-up, there is no evidence of recurrence (Fig. 1). The diagnosis of Fanconi anemia was also excluded by clinical examination, blood test and skin biopsy.

Discussion

Carcinoma of unknown primary involving cervical nodes in level II to IV are usually related to an oropharyngeal primary tumor [2]. When imaging is not able to identify the primary tumor, tonsillectomy is recommended in order to identify small tumors hidden inside the tonsil, together with HPV testing. In the present case involving a 19-year old patient, additionally to the histological analyses, microbial tests were also performed because the first hypothesis after clinical examination was an infection and not a cancer.

These microbial tests identified *Mh*, which is a common commensal bacterium of the genitourinary tract. Sackel et al. reported that *Mh*, *Ureaplasma urealyticum* and *Neisseria gonorrhoeae* were found in 14.3%, 14.8% and 4,9% of the pharyngeal samples obtained from 1044 men and women attending clinics in Boston, USA, respectively [3]. These results were obtained by cultures. Additionally, they showed that colonization alone was not necessarily associated with sore throat or other upper respiratory diseases [3]. Beyond colonization of the oropharyngeal tract by this genital pathogen, two reports described an association with infection. One study showed that the rate of mycoplasma isolation was significantly higher in children undergoing tonsillectomy for recurrent adenotonsillitis (34.5%) than in the cases of tonsillectomy for sleep apnea (20%) [4]. *Mh* infections, particularly extra-genital infections, are probably underdiagnosed for 3 main reasons: (i) the test is not performed, (ii) the culture technique is long and difficult, and (iii) 16sRNA is not systematically used.

The 16S rRNA technique is a recent, reliable and reproducible tool for bacteria identification [5], now routinely used for pathogen identification, particularly in cases of slow-growing, unusual, fastidious bacteria and for bacteria that are poorly differentiated by conventional methods. This technique consists of amplification and sequencing the *RRS* gene coding for 16S rRNA, a chromosomal gene of about 1500 base pairs, present in all bacterial species (universal gene). The 5' and 3' ends are preserved in all bacteria and are used to link the two complementary universal primers that initiate the PCR reaction. The gene's sequence is specific to each species and allows their precise identification by sequencing the amplification product. For each sequence obtained, homologous sequences are searched in databases via the BLASTN programs of the National Center for Biotechnology Information (NCBI) [6]. Contamination is one limit of this technique. In the present case, this bias has been excluded by using a standard hybridation technique for *Mh* identification, and by the repeated biopsies confirming the colonization.

The role of mycoplasma infections in carcinogenesis remains a matter of debate. Recent epidemiologic, genetic, and molecular studies suggest that mycoplasma infection and inflammation may initiate prostate cancer. Namiki and al. showed that *Mycoplasma* chronic infection (19 weeks of culture with benign human prostate cells) promoted malignant transformation resulting from a dysregulation of genes such as Ras, Myc or P53 [7]. Logunov and al., in a model of rat fibroblasts infected by *Mycoplasma arininis*, showed that the bacteria suppressed the transcriptional activity of p53 and activated the Nf-kB pathway, eventually inducing a Ras-dependent malignant transformation. This finding indicates that mycoplasma infection can act as an equivalent of p53-suppressing oncogene [8] (Fig. 2).

Conclusion

This report describes an original case of association between *Mh* and cervical SCC in a young patient. It may suggest that, besides HPV infections, other genital pathogenic agent may be related to HNSCC. Further studies are required to document this association and the hypothesis of a potential oncogenic role.

Disclosure statement

Dr Caroline Hoffmann: consultant and honorary, Nanobiotix. No disclosure related to this publication.

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Figure 1 – Cervical MRI before treatment (A) and after 13 months of follow-up (B)

Figure 2 – Schematic presentation of the three proposed mechanisms by which *Mycoplasma* sp. may promote tumor progression: epigenetic changes, inhibition of p53 pathway, and activation of NFkb pathway.

Figure 1 – Cervical MRI before treatment (A) and after 13 months of follow-up (B)

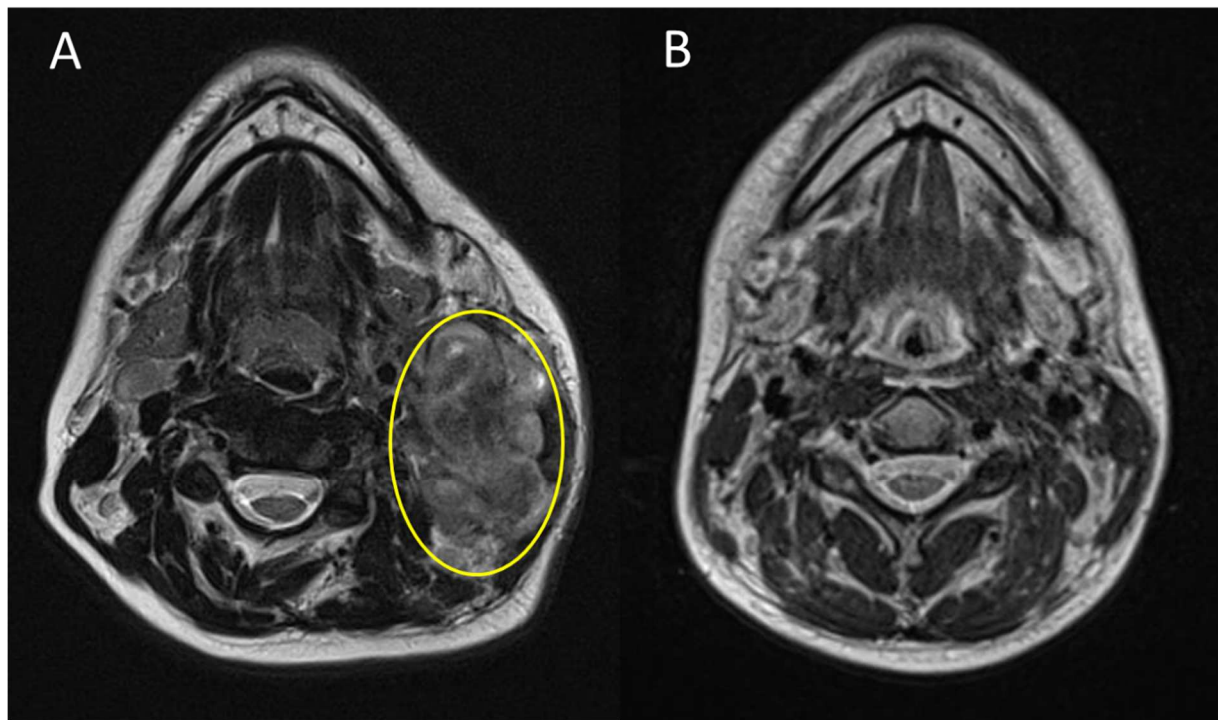


Figure 2 – Schematic presentation of the three proposed mechanisms by which *Mycoplasma sp.* may promote tumor progression: epigenetic changes, inhibition of p53 pathway, and activation of NFkb pathway.

