

Immunohistochemical assessment of the MIB, CK-14, p63, p16, Cal A, and Cys A markers in spindle cell squamous cell carcinomas of the lip

Avaliação imuno-histoquímica dos marcadores MIB, CK-14, p63, p16, Cal A e Cys em carcinomas espinocelulares de lábio

Marília Gerhardt de Oliveira*

Patrícia W. Fregapani Worm**

Juliana Dreyer da Silva de Menezes***

Luciana Maria Pereira Ramalho****

João Feliz Duarte de Moraes*****

Ferdinando De Conto*****

Abstract

Objective: with the objective of testing the expression of the protein markers MIB, CK14, p63, p16, Cal A, and Cys A in the pathogenesis of oral spindle cell carcinoma, we conducted an immunohistochemical study of the expression of the protein markers MIB, CK14, p63, p16, Cal A, and Cys A in human biopsy specimens of these lesions. **Methods:** fifteen histological specimens of spindle cell squamous cell carcinoma of the lower lip were obtained from the Department of Oral Pathology, Bahia Federal University. Immunohistochemical analyses were performed at the Molecular Biology Laboratory of the Department of Otorhinolaryngology, Heidelberg University, Germany. **Results:** statistical analysis revealed no association between markers. There was strong positive staining for Ck14, MIB, and Cal A in 93.3% of cases, thus establishing a strong association. **Conclusion:** p63, p16, MIB, Cal A, Cys A are markedly expressed and p16 is strongly suppressed in oral cavity tumors, which suggests that the latter protein may play a role in negative regulation of cell cycle progression.

Keywords: Diagnostic. Oral cancer.

Introduction

Immunohistochemical identification of molecular genetic events in the progression of preneoplastic lesions to spindle cell squamous-cell carcinoma enables early detection of lesions with the potential for malignant progression, thus permitting timely intervention^{1,2}.

The various markers that enable assessment of the progression of preneoplastic lesions to spindle cell carcinoma include the p16 protein, which halts the cell cycle and induces apoptosis by pRb-mediated phosphorylation of cyclin-dependent kinase 4 (CDK4). Functional loss of p16 may lead to uncontrolled cell proliferation^{3,4}. Another protein, calgranulin A (Cal A), is involved in the regulation of several cell processes, including the cell cycle and cell differentiation. Its expression has recently been associated with the development of severe dysplasias⁵⁻⁸. Cystatin A (Cys A), a cysteine protease inhibitor, is a precursor of proteins involved in keratinocyte keratinization, and is expressed during the late phase of differentiation of these cells. Studies suggest that expression of cystatin A is inversely associated with malignant progression of cancer⁹.

Other markers, such as retinoblastoma and p53, may be related with early steps of carcinogenesis in

* Full Professor, Pontifícia Universidade Católica do Rio Grande do Sul (PUCRS), Porto Alegre, Brazil.

** Oral and Maxillofacial Surgeon, PUCRS - Porto Alegre - Brazil

*** Dentist, PUCRS, Porto Alegre, Brazil

**** Professor, School of Dentistry, UFBA, Salvador, Brazil

***** Professor, School of Mathematics, UFRGS, Porto Alegre, Brazil

***** Professor, School of Dentistry, UPF, Passo Fundo, Brazil

oral cavity squamous cell carcinoma. Furthermore, higher Rb expression has also been observed in malignant lesions¹⁰. The p63 protein, a homologue of p53, may be associated with tumor formation in the epithelial tissue, acting as an oncogene^{11,12}. Expression of p63 is almost exclusively restricted to epithelial cells, mutations in this gene are infrequent, and its expression is increased in a variety of solid tumors, particularly those of the head and neck area^{12,13}. MIB (Ki-67) is one of the cell cycle regulator proteins that is found during duplication, being higher in carcinomas than in hyperplasias, indicating poor prognosis^{14,15}. CK14 is positive in basaloid squamous cell carcinoma, and is mostly distributed diffusely in the basaloid cells; it is a parameter of proliferative activity and metastatic potential of squamous cell carcinoma of the lung^{16,17}.

Methods

Sample selection

Fifteen histological specimens of spindle cell squamous cell carcinoma of the lower lip were obtained from the Department of Oral Pathology, Bahia Federal University. Preliminary histological analysis with H&E staining was performed to grade lesions according to degree of keratinization, nuclear polymorphism, pattern of invasion, and lymphoplasmacytic infiltrate, using the system developed by Byrne et al. (1992)² (Table 1). Changes were scored on a scale of 1 to 4 points according to intensity, and scores were added to yield a mean lesion grade for each slide.

Morphological parameters

Table 1 - Morphological parameters assessed by Byrne et al. (1992)

	Degree of keratinization	Nuclear polymorphism	Pattern of invasion	Lymphoplasmacytic infiltration
Grade 1	No keratinization (0-5% of cells)	Extreme nuclear polymorphism (0-25% mature cells)	Marked and widespread cellular dissociation in small groups and/or in single cells (n < 15)	None
Grade 2	Minimal keratinization (5-20% of cells)	Abundant nuclear polymorphism (25-50% mature cells)	Small groups or cords of infiltrating cells (n > 15)	Slight
Grade 3	Moderately keratinized (20-50% of cells)	Moderately abundant nuclear polymorphism (50-75% mature cells)	Infiltrating, solid cords, bands and/or strands	Moderate
Grade 4	Highly keratinized (> 50% of cells)	Little nuclear polymorphism (> 75% mature cells)	Pushing, well delineated infiltrating borders	Marked

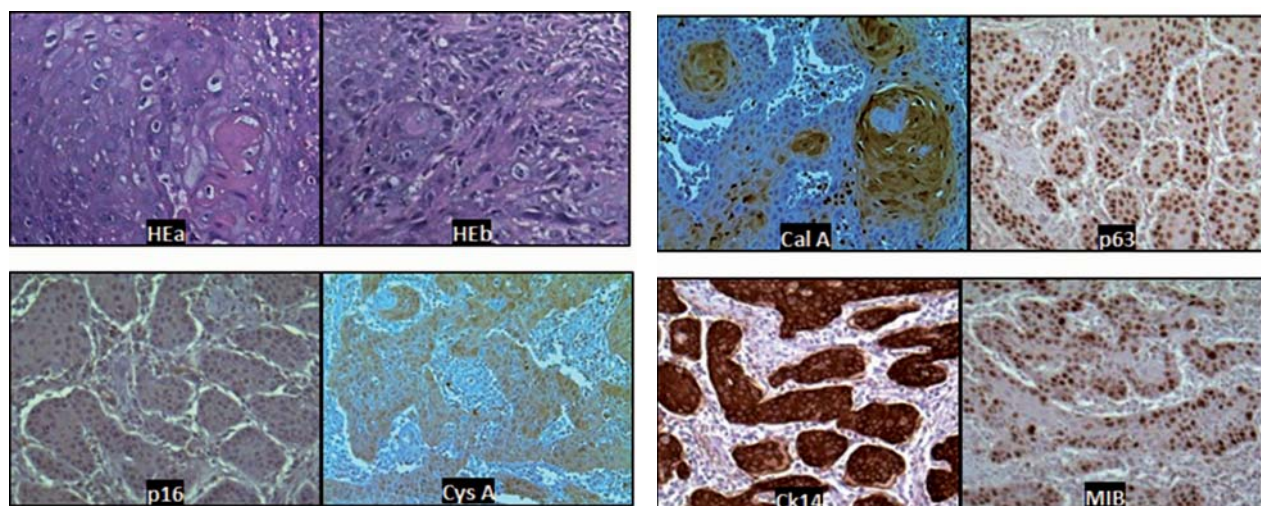


Figure 1 - (A,B). Oral lip cancer. Atypical epithelial cells showing dyskeratosis (A) pleomorphic and hyperchromatic nuclei (B). (Haematoxylin and eosin, x 20)
p16 Immunohistochemical staining of p16 protein in oral lip cancer. Negative staining for p16 in tumor cell islands (Original magnification, x200).
B570E Cys A Immunohistochemical staining of CysA protein in oral lip cancer. Weak and diffuse cytoplasmic and nuclear staining of tumor cell islands. (Original magnification, x200)
B5708E Cal-A 20X2 Immunohistochemical staining of Cal-A protein in oral lip cancer. Overlapping expression pattern observed. Diffuse cytoplasmic staining in central tumor island areas and nuclear staining in scattered tumor cells and distinct stromal cells. (Original magnification, x200)
B570896E p63 20x Immunohistochemical staining of p63 protein in oral lip cancer. Strong and diffuse nuclear staining of tumor cell islands. (Original magnification, x200)
B570896E ck14 Immunohistochemical staining of ck14 protein in oral lip cancer. Strong and uniform cytoplasmic staining of tumor cell islands. (Original magnification, x200)
B570896E MIB 20x Immunohistochemical staining of MIB protein in oral lip cancer. Strong and diffuse nuclear staining of tumor cell islands. (Original magnification, x200)

Immunohistochemistry

After biopsy selection, immunohistochemical analyses were performed at the Molecular Biology Laboratory of the Department of Otorhinolaryngology, Heidelberg University, Germany.

Specimens were sliced into 4 µm sections, yielding one slide for each tested antibody. After removal of paraffin by xylol immersion and rehydration in a graded alcohol series and deionized water, antigen recovery was performed for the MIB, CK14, p63, p16, Cal A, and Cys A proteins. Sections were placed in citrate buffer (10 mmol/L), microwaved for 3 minutes at 700W and 10 minutes at 200W, left to

cool at room temperature, and rinsed with deionized water and PBS. After inactivation of endogenous peroxidase and soaking in horse serum for 1 hour, sections were incubated in primary anti-MIB, anti-CK14, anti-p63, anti-p16, anti-Cal A, and anti-Cys A antibodies.

Criteria for positive immunohistochemical staining

Immunohistochemistry findings were analyzed semiquantitatively. Expression was considered absent (-) in slides with ≤ 10% positive cells; moderate (+) in those with 10-50% positive cells; and marked (++) in those with ≥ 50% positive cells. Preliminary findings showed differences in expression of the MIB, CK-14, p63, p16, Cal A, and Cys A markers.

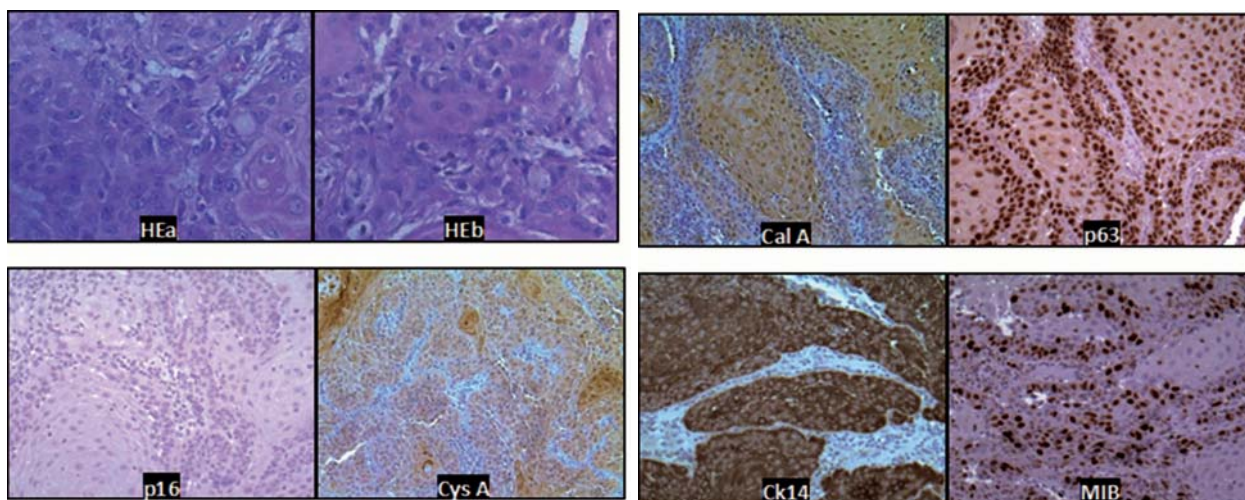


Figure 2 - (A,B). Oral lip cancer. Epithelial tumor cells showing dyskeratosis and atypical mitosis (A) pleomorphic and hyperchromatic nuclei (B). (Haematoxylin and eosin, Original magnification, x200).

p16 20x Immunohistochemical staining of p16 protein in oral lip cancer. Negative staining for p16 in tumor cell islands. (Original magnification, x200)

Cys A Immunohistochemical staining of CysA protein in oral lip cancer. Moderate and diffuse cytoplasmic and nuclear staining of tumor cell islands showing increased reactivity in central areas. (Original magnification, x200)

Cal-A 20x2 Immunohistochemical staining of S100A8 protein in oral lip cancer. Diffuse cytoplasmic and scattered nuclear staining in tumor cell islands and nuclear reactivity in a few distinct stromal cells. (Original magnification, x200)

p63 20x Immunohistochemical staining of p63 protein in oral lip cancer. Strong and diffuse nuclear staining of tumor cell islands. (Original magnification, x200)

ck14 Immunohistochemical staining of ck14 protein in oral lip cancer. Strong and uniform cytoplasmic positivity of tumor cell islands. (Original magnification, x200)

MIB 20x Immunohistochemical staining of MIB protein in oral lip cancer. Strong and diffuse nuclear staining of tumor cell islands. (Original magnification, x200)

Results

Descriptive statistics were used to assess the immunohistochemical expression of CK-14, MIB, p63, Cys A, Cal A, and p16 in lower lip spindle cell carcinoma specimens. Fisher's exact test was used to tabulate data.

All patients were male. Mean age was 55 (range, 43-71 years) (Table 2).

Table 2 - Patient age

	N	Minimum	Maximum	Mean	SD
Age	15	43	71	55.2	9.9

Statistical analysis revealed no association between markers. There was strong positive staining for Ck14, MIB, and Cal A in 93.3% of cases, thus establishing a strong association (Table 3):

Table 3 - Frequency of the CK-14, MIB, and Cal A markers (Fisher's exact test)

Staining	MIB		CK-14		Cal A	
	Frequency	Percent	Frequency	Percent	Frequency	Percent
Strong	1	6.7	1	6.7	1	6.7
Very strong	14	93.3	14	93.3	14	93.3
Overall	15	100	15	100	15	100

There was strong, positive tissue staining for the p63 and Cys A genes in 86.7% of cases, as shown in Table 4:

Table 4 - Frequency of the p63 and Cys A markers (Fisher's exact test)

Staining	p63	Percent	Cys A	Percent
	Frequency		Frequency	
None	2	13.3	0	0
Strong	0	0	2	13.3
Very strong	13	86.7	13	86.7
Overall	15	100	15	100

There was no positive association with the p16 gene, as staining for this marker was absent in 93.3% of cases (Table 5):

Table 5 - Frequency of the p16 marker (Fisher's exact test)

p16	Frequency	Percent	Valid percent	Cumulative percent
None	14	93.3	93.3	93.3
Strong	1	6.7	6.7	100
Overall	15	100	100	

Discussion

To overcome this challenge and gain a better understanding of cell survival and apoptosis, cell proliferation and tumor suppression markers have been studied in the search for more reliable indicators for detection and prediction of the progression of preneoplastic or malignant lesions¹⁸⁻²¹. A better understanding of the molecular mechanisms that involve control of neoplasm cell growth may enable identification of the factors involved in tumor regulation and progression. From the simple count of mitoses, which provides generic evidence^{2,22}, to the regulatory mechanisms of the cell cycle, the relationship between growth factors, oncogenes, tumor suppressor genes, and their protein products plays an important role in the detection and quantification of proliferating cells¹. Therefore, biomarkers are relevant to the clinical management of patients with malignant neoplasms, as they assist in diagnosis, staging, assessment of treatment response, detection of recurrence, and prognostication. Despite their major clinical application, there is little scientific evidence in this area^{23,24}, which justifies further research, such as this study.

The possibility of analyzing mRNA from the archives of pathology laboratories is exciting, as it allows for large retrospective studies. Formalin is

the most common fixative used in the surgical pathology routine, and its promotion of nucleic acid degradation is well known. In addition, the proteins extracted from formalin-fixed and paraffin-embedded tissue seem to be the same compared with those extracted from fresh frozen tissue. Additionally, easier-to-use immunohistochemical markers of apoptosis, applicable in archived paraffin-embedded tissue, have been commercially developed.²⁵⁻²⁷

Among the various markers reported to be positive in premalignant and malignant lesions, p16 stands out²¹. This study evaluated the p16 protein due to its important role in the pathogenesis of head and neck cancer²⁸, as cell cycle progression depends on the action of cyclin-CDK complexes. This process is inhibited by a group of proteins that modulate the action of these complexes, known as cyclin-CDK complex inhibitors (CKIs)²⁹, which includes p16. Functional loss of this gene may be associated with development and progression of a variety of malignant neoplasms³⁰. Under normal conditions of controlled cell division, p16 acts as a negative regulator of the cell cycle, which may predispose to occurrence of the initial steps required for malignization²⁸. In this study, there was no expression of p16 in 93% of cases, which confirms the hypothesis that p16 is a tumor suppressor gene.

The p63 protein, a p53 homologue, may be associated with tumor formation in epithelial tissues, thus acting as an oncogene^{11,12}. There has been substantial recent interest in the role of p63 as a regulator of cell proliferation and differentiation in potentially and frankly malignant lesions. As expression of p63 is almost exclusively restricted to epithelial cells, mutations are infrequent, and its expression is increased in a variety of solid tumors, particularly those of the head and neck area^{12,13}. In our sample, approximately 86% of specimens were strongly positive on immunohistochemical staining, which is consistent with the literature. Even more markedly, MIB (Ki-67) and CK-14 were equally expressed (93%) in our specimens, corroborating previous studies that reported a worse prognosis in tumors that exhibit affinity for this marker^{14,15,31}.

Cys A expression has been reported in polymorphonuclear granulocytes, keratinocytes, and liver and spleen tissue³². This protein plays an important role in inhibition of apoptosis caused by caspase-3 inhibition, giving the cell more time to promote DNA repair³³. In this study, Cys A was detected very often (86.7%) in specimens of lower lip cancer, which suggests an association with tumor pathogenesis.

The Cal A protein is closely related to regulation of intracellular activities such as abnormal growth, cell cycle progression, transcription, and cellular differentiation⁶. The Cal A gene has been the object of particular research attention due to its involvement in a variety of human diseases, such as rheumatoid arthritis, acute inflammation, cardiomyopathies, Alzheimer's disease, and cancer¹³. Expression of this protein in normal epithelial tissue, including the oral mucosa, is well established. The role of Cal A in neoplasms is still unclear, but appears to be closely tied to the differentiation potential of epithelial tissue, as it is strongly expressed in keratinocyte hyperproliferation. Dysregulation of this gene has been described in epithelial cell carcinoma of the head and neck, as in a study that revealed strong expression in nasopharyngeal tumors⁵.

Dysregulation of S100 expression may be associated with development of tumors and metastases⁶. Furthermore, protein expression studies have detected increased levels of S100-like proteins, such as Cal A, in a variety of tumor tissues as compared to their corresponding abundance in normal tissue¹³. This is consistent with the data reported in this study, in which approximately 94% of samples exhibited strong immunohistochemical staining for this protein, which suggests it is closely related to spindle cell carcinoma.

Conclusion

The results of this study support investigation of the relationship between cell cycle biomarkers and the pathogenesis of oral cancer. Particularly relevant is the finding that p63, p16, MIB, Cal A, Cys A are markedly expressed and p16 is strongly suppressed in oral cavity tumors, which suggests that the latter protein may play a role in negative regulation of cell cycle progression.

Acknowledgements

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Resumo

Objetivo: O objetivo deste estudo foi avaliar, através de imunohistoquímica, a expressão dos marcadores proteicos MIB, CK14, p63, p16, Cal A e Cys A em lesões bucais com diagnóstico histopatológico de carcinoma espinocelular. **Métodos:** 15 amostras histológicas com diagnóstico confirmado localizado em lábio inferior obtidas do Serviço de Patologia Bucal da Universidade Federal da Bahia foram examinadas no Laboratório de Biologia Molecular do Departamento de Otorrinola-

ringologia da Universidade de Heidelberg, Alemanha. **Resultados:** O estudo estatístico não revelou associação entre os marcadores. Os genes Ck14, MIB e Cal A apresentaram intensa marcação nas células do tecido, em 93,3% dos casos, estabelecendo assim, forte relação. **Conclusão:** Os resultados suportam a investigação revelando que os genes MIB, CK14, p63, Cal A e Cys A se apresentam fortemente evidentes nos tumores de cavidade oral e o p16 suprimido, sugerindo que esta proteína pode exercer um papel de regulador negativo do ciclo celular.

Palavras-chave: Câncer bucal. Diagnóstico. Imuno-histoquímica.

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Address for Correspondence:

Ferdinando De Conto
 Rua Teixeira Soares, 1075 sala 1002
 Edifício Tamandaré
 99010-080 - Passo Fundo-RS
 Fone: (54) 33168402
 E-mail: ferdi@upf.br

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