

# Expression Analysis of p53, Ki-67 and bcl-2 in Pre-Malignant Lesions of the Cervix

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## Abstract

*Objectives*: The present study has tried to analyze the individual and combined expression of the three markers—p53, Ki-67 and bel-2-in HPV associated pre-malignant lesions of the cervix. *Methods*: Thirty-four cases of high-grade lesions, 21 low-grade cases and 18 normal cases were histologically assessed. HPV-DNA presence and the viral types were determined through PCR and RFLP, respectively. Marker expression was performed by immuno-histochemistry. *Results*: With regards to viral types, HPV with high oncogenic potential represented the majority, with just one case reporting the low oncogenic potential for viral infection (HPV 6). HPV was the most frequent and could be associated with the high-grade lesion group, however, no significant associations were found between the viral ones and the marker expression. When the expression location, beyond the basal layer, was taken into account, the three markers proved to be good indicators of the lesion grade. The markers' combined expression: p53(-), Ki-67(+), bcl-2(-) and p53(-), Ki-67(-), bcl-2(-) was statistically significant when associated with the lesion grade. *Conclusions*: The results suggest that the p53, Ki-67 and bel-2 markers can contribute to differentiate the lesion grades, whether in its isolate form or in combined form.

# Keywords

p53, Ki-67, bcl-2, Pre-Malignant Lesions of the Cervix, HPV

# **1. Introduction**

Cervical cancer is one of the most frequent neoplasias in women, with approximately 530 new cases worldwide per year; cervical cancer is the third most common kind of cancer among women, being responsible for 274

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thousand women's deaths per year [1]. It is the third most frequent tumor in the female population, after breast and colorectal cancer, and the fourth most common cause of female death by cancer in Brazil, with 15,590 new cases expected in 2014. [2] A epidemiological and experimental evidence indicates that certain kinds of HPV are directly involved in cervical cancer pathogenicity.

Although it is present in about 100% of cervical carcinoma, isolatedly, HPV cannot be considered solely responsible for the malignant transformation [3]. It is believed that HPV as an initiator factor, as the progression of the HPV infection into cancer also requires a number of other co-factors to be taken into consideration: environmental factors, such as physical and chemical carcinogens, and host restricted factors, such as immune rsponse, genetic susceptibility, among others [4]-[7].

Taking into consideration that most viral infections, including high risk HPV, recede spontaneously and that just a small fraction of infected women will progress into cervical cancer, it is necessary to identify early in which, among the infected patients, it will really evolve into cancer [5] [8] [9].

The cell protein p53 suppresses the transcription of some cells. However, its greatest tumor suppressor effect is due to the genes transcriptional activation that maintains genomic stability [10]-[13]. Under normal conditions, protein p53 acts as an *emergency brake* in hypoxia or DNA lesion situations due to irradiation, ultra-violet light or mutagenic agents. Through p53 action, repair genes are induced and the cell cycle is interrupted in G1 phase. When repair becomes impossible, the injured cell is induced to cell death by p53 (apoptosis) [13] [14]. The inactivity or loss of p53 causes the loss of control in cell proliferation which results in genetic instability and accumulation of oncogenic mutations responsible for the development of malignant tumors [15] [16].

The p53 gene can also be inactivated through other mechanisms besides somatic and hereditary mutations. The HPV E6 transformed protein can bind with p53 protein and induce its degradation. The affinity of the viral onco-proteins with p53 protein varies according to the HPV type. The binding affinity of the E6 low risk genital HPV onco-protein is lower than the high oncogenic risk HPV [15] [17]. The importance of binding between E6-p53 in cervical cancer pathogenicity has been supported by recent molecular studies [17]-[25].

Ki-67 is characterized as a cell proliferation marker in phases G1, S, M and G2 of the cell cycle, able to identify a nuclear protein present in the interphase proliferative cells, as well as in the mitotic cells, thus, it has been widely used in the growth stage investigation and in several tumor kinetic activities, such as breast cancer, colon and rectum cancer, and lymphomas [26] [27].

The bcl-2 protein is characterised for being an intracellular membrane protein that does not allow cell death (apoptosis) in several situations. An inadequate bcl-2 expression might extend the life-time of the damaged cells, thus increasing significantly the probability of malignant transformation. In addition, the bcl-2 super-expression can interrupt the arrest in the G1 stage of the cell cycle, mediated by p53, therefore inhibiting apoptosis [28]-[30].

Therefore, this study tried to verify the p53, Ki-67 and bcl-2 marker expression in cervical pre-malignant lesions, HPV associated, as well as relate these marker expressions with viral types and with the lesion grade, trying to propose a combined analysis of the apoptosis and cell proliferation markers for cervical pre-malignant lesions.

#### 2. Material and Methods

This retrospective study has been approved by the University of Caxias do Sul Research and Ethics Committee with no conflicts of interest.

Seventy-three cases of cervix material biopsy were used. These cases were in the archives of the Diagnose-Laboratório de Patologia e Citologia, from January 2003 to December 2004.

The study groups were divided into: 18 cases with normal histology (control-group), 21 cases with low-grade intraepithelial squamous lesions (LEIBG), and 34 cases with high-grade intraepithelial squamous lesions (LEIAG).

The biopsies were submitted to paraffin removal and human DNA extraction by using the methodology suggested by Banerjee *et al.* [31]. They were then sent to generic PCR in order to identify the HPV-DNA using PGMY 09/11(HPV) and GH20/PCO4 (human  $\beta$ -globin) initiators. Viral typing was carried out in 64% of the samples positive for HPV by using RFLP technique, according to the methodology proposed by Bernard *et al.* (1994) [32].

The markers (p53, Ki-67 and bcl-2) expression was analyzed by means of immunohystochemistry (LSAB-

HRP+). The reading criteria were: presence or absence of expression, and the location of the expression in the different epithelial layers. The cases that presented nuclear brown coloration for p53 and Ki-67 or cytoplasmic brown coloration for bcl-2, were classified as positive. For all cases, positive and negative coloration controls were used. Monoclonal anti-bodies used for the reaction were: 1) Anti-human p53 Protein, clone DO-7, dilution 1:50, DAKO<sup>®</sup>, 2) Anti-human Ki-67, clone Ki-S5, dilution 1:50, DAKO<sup>®</sup> e 3) Anti-human bcl-2 Oncoprotein, clone 124, dilution 1:80, DAKO<sup>®</sup>.

The data were analyzed and processed by the SPSS program, version 12.0. For statistical analysis both the non-parametric tests of Pearson's Chi-square  $x^2$ , and Fisher's exact test, were used. P < 0.05 values were considered statistically significant.

#### 3. Results

In the 73 cases studied, protein p53 presented expression in 28 (38.4%) and absence of expression in 45 cases (61.6%).

Out of the 28 cases that presented protein p53 expression, 10 cases (35.7%) were LEIBG distributed in the following way: 5 cases (50%) in the basal layer, 1 case (10%) in the intermediate layer, 1 case (10%) in the surface layer, 1 case (10%) in the intermediate and basal layers and 2 cases (20%) in all layers. Twelve cases (42.8%) were LEIAG, distributed as follows: 4 cases (24%) in the basal and intermediate layers and 5 cases (41.7%) in all layers. Six cases (21.4%) were from the normal group (control), with expression in the basal layer only.

Ki-67 marker showed expression in 53 cases (72.6%) and absence of expression in 16 cases (21.9%). Four cases could not be assessed due to lack of biopsy material along the study.

From the 53 cases that showed expression for Ki-67, 19 cases (35.8%) were LEIBG, distributed as follows: 12 cases (63.1%) in the basal layer, 2 cases (10.5%) in the surface layer, 3 cases (15.8%) in the basal an intermediate layers and 2 cases (10.5%) in all layers. Twenty-three cases (43.4%) were LEIAG distributed as follows: 3 cases (13.4%) in the basal layer, 1 case (4.35%) in the surface layer, 7 cases (30.4%) in the basal and intermediate layers, 11 cases (48%) in all layers, and one case (4.35%) in the intermediate and surface layers. Eleven cases (21%) were from the normal group (control), with (82%) in the basal layer and 2 cases (18%) in the basal and intermediate layers.

Marker bcl-2 showed expression in 38 cases (52.1%) and absence of expression in 35 cases (47.9%). From the 38 cases that showed expression for protein bcl-2, 11 cases (29%) were LEIBG distributed as follows: 10 cases (91%) in the basal layer, 1 case (9%) in all layers. Sixteen cases (42%) were LEIGA distributed as follows: 8 cases (50%) in the basal layer, 6 cases (37.5%) in the basal and intermediate layers and 2 cases (12.5%) in all layers.

Table 1 shows the expression of the three markers and the lesion grade, when the epithelium stratification into upper and basal layers is taken into account.

The combined expressions of the markers, in the format of an immuno-histochemical panel considering the epithelium stratification in the basal layer and in upper layers with the lesions grade, are represented in Table 2.

Combinations p53(-) Ki-2(n = 19) and p53(-) Ki-67(-) bcl-2(-) (n = 34) were the most frequent among the

Cable 1. Expression of the markers and lesion grade.				
Marker	Expression Location	Normal	LEIBG	LEIAG
p53*	Basal Layer $(n = 15)$	40% (n = 6)	33.3% (n = 5)	26.7% (n = 4)
	Upper Layers (n = 13)	0% (n = 0)	38.5% (n = 5)	61.5% (n = 8)
Ki-67**	Basal Layer ( $n = 24$ )	37.5% (n = 9)	50% (n = 12)	12.5% (n = 3)
	Upper Layers $(n = 29)$	6.9% (n = 2)	24.1% (n = 7)	$69\%^{***}$ (n = 20)
bcl-2****	Basal layer $(n = 29)$	37.9% (n = 11)	34.5% (n = 10)	27.6% (n = 8)
	Upper Layers (n = 9)	0% (n = 0)	11.1% (n = 1)	88.9% ***** (n = 8)

\*Normal × LEIBG: P = 0.027. \*\*Normal × LEIBG × LEIAG: P = 0.001. \*\*\*Expression location × LEIAG: P = 0.001. \*\*\*\*Normal × LEIBG × LEIAG: P = 0.004.

Markers	Group I (normal)	Group II (LEIBG)	Group III (LEIAG)
p53+Ki67+ bcl-2+	0	0	2
p53+Ki67+ bcl-2-	0	3	2
p53+Ki67-bcl-2+	0	0	1
p53+Ki67-bcl-2-	0	2	1
p53-Ki67+ bcl-2+	0	1	4
*p53-Ki67+bcl-2-	2	3	14
p53-Ki67-bcl-2+	0	0	2
**p53-Ki67-bcl-2-	16	12	6
Total	18	21	32

Table 2. Immuno-histochemical panel considering the location of the marker expression and the lesion grade.

-: absence of expression, or expression restricted to the basal layer; +: expression beyond the basal layer; \*Normal × LEIBG × LEIAG: P = 0.013. \*\*Normal × LEIBG × LEIAG: P = 0.001.

groups. Combinations p53(-) Ki-67(+) bcl-2(-) presented significant difference among the others, as well as with the lesions grade (P = 0.013). Combination p53(-) Ki-67(-) also showed significant difference among the others, as well as association with the lesions grade (P = 0.001).

HPV-DNA presence was not found in the control group. However, different kinds of HPV were found in the lesion groups. HPV 16 was the predominant viral type (65.7%). The remaining viral types represented just a small portion (34.3%). The presence of multiple infections was not considered significant within the groups, since just one case of HPV 26 and HPV 18 association was observed. According to the oncogenic potential of the virus, just one sample showed low oncogenic potential (HPV6), the others presented high oncogenic potential HPV (HPV16, 18, 26, 31, 33, 39, 45, 52 and 58).

The association between the type of virus and the lesion grade was significant only for HPV 16 (P = 0.001). The association between the other viral types and the lesions was not significant, such as the association of the viral types and the expression of the markers.

Any positive association between the viral types and the markers used in this study has been found.

#### 4. Discussion

The epithelium division into two levels (basal layer and surface layers) allowed to associate the marker expression with the lesion grade. The criteria used for the expression reading (presence or absence of expression and location on the epithelium layers) enabled a new approach through a differentiated, fast and accurate analysis. Studies have shown, in their findings, the importance of the epithelium stratification when assessing intraepithelial lesions. Other authors have used quantitative analysis of the expression by using distinct classification systems [33] [34]. These parameters have not been considered, especially for presenting great inter-observational variability in slide reading, making the assessment subjective, in addition to being laborious and needing a longer period of study.

The individual expression of the three markers (p53, Ki-67 and bcl-2) has shown to be a good indicator for the lesion grade as the expression passes through the basal layer towards the surface layers.

Although the immuno-histochemical results for p53 protein detection in the cervix are still contradictory [35] [36], the present study shows that a p53 expression is beyond the basal layer in the LEIBG and LEIAG groups, and that the most frequent expression reaches all layers in the latter. The location beyond the basal layer can be related to loss of epithelium maturation in high grade dysplasia as well as to a process of loss of normal prolifer-ative cells regulation [37] [38].

The p53 expression, restricted to the basal layer in normal epithelium and in LEIBG, also found by Kurvinen *et al.* (1996) [39], might be representative of cells in proliferation.

The viral types of high oncogenic potential have not shown p53 expression beyond the basal layer, in accordance with the Kurvinen *et al.*'s (1996) [24] [39] findings that were found low or absence of p53 expression in

53% of the lesions associated with HPV 16 and 18 types, and according to the hypothesis that the E6 viral oncoprotein from the viral types of high oncogenic potential has more affinity with p53, is capable of inducing fast protein degradation, thus decreasing its expression levels [17] [37].

Marker Ki-67 has been studied as an important tumor marker. Its expression is related to proliferative action, lesion grade and HPV in several studies [40] [43]. In this study, the normal group presented expressions in the basal layer with more frequence. Only one case presented expression as far as the intermediate layer, which corroborated Carrilho *et al.*'s studies (2003) [33] who found Ki-67 expression restricted to the basal layer in normal epithelia.

LEIBG and LEIAG groups had expression beyond the basal layer, for the LEIBG group the most frequent expression reached the basal layer only, and for the LEIAG, the most frequent expression reached all the epithe-lium layers.

The findings have shown that as the Ki-67 expression crosses the basal layer, there is a relation with the presence of lesion (LEIBG and LEIAG), especially with more serious lesions (LEIAG).

There was a strong association between LEIAG and the expression location in the epithelium layers, found to be in accordance with the association found between Ki-67 positivity and the levels of dysplasia observed in Keating *et al.*'s studies (2001) [44], whose highest proliferative activity measured by Ki-67, was found in high grade squamous lesions.

The bcl-2 intense expression in normal epithelia, restricted to the basal layer, was also shown by Kurvinan *et al.* (1996) [45]. Hockenbery *et al.* (1991) [46] suggested the existence of a basal layer natural protection system against apoptosis in order to explain the finding. However, as the epithelial cells present dysplastic alterations, bcl-2 expression expands to the upper layers of the epithelium, especially in LEIAG group, proving bcl-2 marker to be a good indicator for lesion grade.

Few studies assess the marker expression in cervix pré-malignant lesions in a combined way. Queiroz *et al.* (2006) [47] assessed the expression of p16, cyclin D1, p53 and Ki-67 proteins, and found greater lesion progression correlation for p16 and Ki-67 than for p53 and cyclin D1. Taking into account the expression results for the three markers p53, Ki-67 and bcl-2 in the present study, the combination p53(-) Ki-67(+) bcl-2(-) has shown a significant difference among the others, as it can also be associated to the lesions grade, suggesting that the p53 and bcl-2 protein absence of expression, plus Ki-67 super-expression beyond the basal layer, indicates high-grade lesion. Similarly, the combination p53,(-) Ki-67(-) bcl-2(-) shows a difference among the others, as it can also be associated to the lesion grade. However, it suggests that the absence of p-53, Ki-67, bcl-2 Te expressions in the upper layers, indicates the opposite: control-group and low grade lesion (Figure 1).

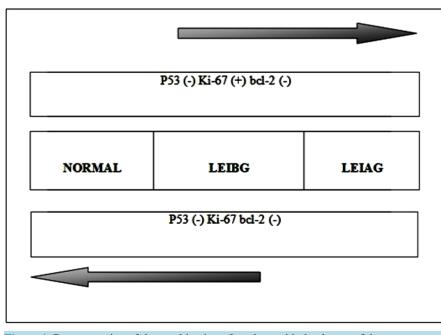


Figure 1. Representation of the combination of markers with the degree of damage.

As expected, different types of HPV have been found in the lesion groups, being HPV 16 the predominant viral type. Data found in the literature refer to HPV 16 as being the most frequent one, both in the intra-epithelial squamous lesions and in invasive squamous carcinomas [6] [36] [48] [49]. Although a association has been found between HPV 16 and the lesion grade, especially in LEIAGs, the association among the other viral types and the lesion grade has not been significant, probably because of the low frequency of the other viral types in this study.

The absence of association among the types in question and the marker expression, both in individual form and combined form, has probably occurred due to the limited number of viral types, therefore, demanding a larger number of samples.

The search for biological markers has been the focus of many research centers, and several molecules involved in cell growth control have been studied singly. Considering that few studies have assessed several markers simultaneously, the findings in this study have shown that p-53, Ki-67 and bcl-2 markers, when assessed in combination, can contribute for differentiating the cervix pre-malignant lesion grade, working as an auxiliary tool in lesion prognosis.

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