

Co-overexpression of bcl-2 and c-myc in uterine cervix carcinomas and premalignant lesions

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Abstract

To establish the role of co-overexpression of bcl-2 and c-myc protooncogenes in uterine cervix carcinogenesis, we examined 138 tissue samples of low grade cervical squamous intraepithelial lesions (SIL), high grade SIL, portio vaginalis uteri (PVU) carcinoma *in situ* and PVU *invasive* carcinoma, stage IA-IIA (study group) and 36 samples without SIL or malignancy (control group). The expression of bcl-2 and c-myc was detected immunohistochemically using a monoclonal antibody. Fisher's exact test ($P < 0.05$) was used to assess statistical significance. Overexpression of bcl-2 was found to increase in direct relation to the grade of the cervical lesions. High sensitivity was of great diagnostic significance for the detection of these types of changes in the uterine cervix. On the basis of high predictive values it can be said that in patients with bcl-2 overexpression there is a great possibility that they have premalignant or malignant changes in the uterine cervix. Co-overexpression of bcl-2 and c-myc oncogenes was found only in patients with PVU *invasive* carcinoma (6/26-23.0%). Statistically significant difference was not found in the frequency of co-overexpression in patients with PVU *invasive* carcinoma in relation to the control group (Fisher's test; $P = 0.064$). The method's sensitivity of determining these oncogenes with the aim of detecting PVU *invasive* carcinoma was 23%, while specificity was 72.2%. On the basis of high predictive values (100%), speaking in statistical terms, it can be concluded that all patients with co-overexpression of bcl-2 and c-myc oncogenes will have PVU *invasive* carcinoma. We confirmed in our research that co-overexpression of bcl-2 and c-myc oncogenes was increased only in PVU *invasive* carcinoma. However, a more extensive series of samples and additional tests are required to establish the prognostic significance of bcl-2 and c-myc co-overexpression in cervical carcinogenesis.

Introduction

Squamous cell carcinoma of the uterine cervix is currently one of the most common malignancies in women worldwide. In searching ways of prevention, early diagnosis and effective treatment of premalignant lesions and malignant tumors of the uterine cervix, molecular-genetic researches played a significant role in the last few years.

Alterations at the genetic or protein level of the protooncogenes result in oncogenic conversion and impaired cellular growth control mechanisms, causing tumor development.¹

Analysis of oncogenes expression in human cancer is increasingly important to gain a better insight in the process of tumorigenesis and identify new markers for early diagnosis of malignant transformation. Abnormal expression of different cellular oncogenes in various cancers assessed by hybridization and immunological techniques has been previously reported.² The results of these analyses do not appear to be of any early diagnostic value since oncogene expression is, in general, only demonstrable in tumors that can already be classified as malignant.

The key difference between normal and malignant cell is in a subtle change of specific genes that control, by their products, the processes of growth, division and differentiation of a cell and are called protooncogenes. These genes that are present in the genotype of every normal cell possess the oncogenic potential because their disturbed expression may lead to malignant transformation.

The bcl-2 family of related proteins is one of the key regulators of the apoptotic process. It consists of two opposing groups of proteins: death antagonists (bcl-2, bcl-XL, Mcl-1) and death agonists (bax, bak, bcl-XS).³ Apoptosis occurs through competing dimerization between the two protein groups. The relative proportions of which ultimately control the sensitivity or resistance of cells to apoptotic stimuli. Overexpression of bcl-2 and bax and their prognostic significance have been reported in several epithelial cancers.⁴ However, the data on squamous cell carcinoma of the uterine cervix are limited and the results are conflicting.^{2,5} *In vitro* studies have also shown that bcl-2 overexpression prolonged cell survival in cell culture studies.⁶ Therefore, an *in vivo* study on the association between these apoptosis regulatory proteins may be a prerequisite for the complete understanding of the molecular pathogenesis underlying cervical cancer development.

Mitochondrial and cell-surface death receptor-mediated apoptosis are the two principal pathways leading to programmed cell death. The mitochondrial pathway is thought to play a

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major role in response to cancer treatments and is mediated by the bcl-2 family proteins.⁷ More than 20 members of this family have been described so far in humans. A positive ratio between pro- and antiapoptotic bcl-2 family members leads to cytochrome-C release from mitochondria, which triggers the final execution of cell death by the caspase cascade.

The c-myc gene belongs to the myc family that includes b-myc, l-myc, n-myc and s-myc. However, only c-myc, l-myc and n-myc have neoplastic potential.⁸ The ability of myc to promote cell proliferation indicates that its deregulation leads to deregulated DNA synthesis and genomic instability.⁹ Deregulated myc expression is linked to increasing in both cyclin A and cyclin E levels.¹⁰ *In vitro* and *in vivo* data indicate that increased expression of c-myc blocks differentiation and enhances proliferative and apoptotic activity.¹¹ Chlamydia trachomatis and HPV high-risk types may contribute to neoplastic changes in the transformation of the uterine cervix and also might modulate expression of c-myc oncogene.¹²

Overexpression of c-myc was frequently found in advanced stages of cervical cancers and was shown to be associated with tumor progression.^{13,14} Moreover, it has been shown in experimental tumor systems that increased expression of bcl-2 and c-myc can cooperate in tumorigenesis.^{14,15}

Materials and Methods

This prospective study was carried out during 2008 and 2009, at the Department of Obstetrics and Gynecology, Faculty of Medicine and the experimental part was performed at the Laboratory for Experimental and Clinical Immunology of the Faculty of Medicine and at

the Immunological Laboratory of the Public Health Institute, Kragujevac, Serbia. From patients operated at the Department of Obstetrics and Gynecology because of premalignant and/or malignant changes of the uterine cervix, some tissue sections were taken from the operative material (hysterectomy, punch biopsy or conization) for pathological verification and used for this research. The control group consisted of 36 females in whom ambulatory biopsy of the uterine cervix was performed (Papanicolaou test was indicated) and where malignant changes or squamous intraepithelial lesions (SIL) were not found by histopathology (cervicitis chronica of mild to moderate degree). The study group consisted of 32 patients with the pathological diagnosis of low grade SIL, 22 with high grade SIL, 22 with PVU carcinoma *in situ* and 26 with PVU invasive carcinoma, stage IA-IIA. Bcl-2 and c-myc oncogene expression followed in patients in the study and the control group. Co-overexpression of this oncogene was observed as the way of detection of premalignant and malignant changes in the uterine cervix. Cryostat sections were sent for intraoperative diagnosis. Extra sections of cervical lesions were snap-frozen in liquid nitrogen and stored at -70°C until used for immunohistochemistry.

Bcl-2 (Anti-Bcl-2 α (Ab-1) Mouse mAb (100/D5), Calbiochem, Oncogene Research, Cambridge, MA, USA) and c-myc (Anti-c-Myc (Ab-1) Mouse mAb (9E10), Calbiochem) monoclonal antibody was used, which was diluted with phosphate buffered saline (PBS, pH 7.2).

Four-micrometer frozen sections were fixed in 100% acetone for 5 min and the endogenous peroxidase activity was quenched by 10 min incubation in 0.5% hydrogen peroxide. For monoclonal antibody bcl-2 and c-myc (concentration of 1 μ g/mL) incubation with primary antibody was carried out overnight at 4°C. Sections of bcl-2 positive lymphoid tissue were used as positive controls. Sections of c-myc positive lung carcinoma were used as positive controls. For negative controls, the samples were taken through the procedure with omission of the primary antibody. Slides were evaluated by two of the authors, unaware of immunohistochemical or clinical data using a semiquantitative method on a Zeiss AXIOSKOP 2 light microscope. The percentage of immunopositive cells in representative areas of the sections was assessed. The intensity of immunostaining was divided into 4 categories, namely i) negative (0-5%) and ii) 1+ (5-25%); iii) positive 2+ (25-50%) and iv) 3+ (50-100%). Tissue sections from the operative material or material taken by biopsy were taken after obtaining informed consent of patients in accordance with the Declaration of Helsinki and recommendations of the World Health Organization (WHO) for experiments

on human material and after getting approval of the Ethics Committee.

Statistics

Based on the frequency of the lesions found, patients were split into 4 subgroups and 2x2 contingency tables were formed and specificity and sensitivity were calculated. Also, based on the receiver operating characteristic (ROC) curve, which represents the relation between specificity and sensitivity, the discrimination power of the test was determined. Differences between groups were considered significant (Fisher's exact test) at $P < 0.05$. By establishing sensitivity and specificity of the test, the reliability of these analyses, could be the subject of scientific investigation as one of the factors in the prognosis of premalignant lesions of the uterine cervix, in the course of future researches.

Results

Table 1 and Figure 1 show the expression levels of bcl-2 oncogene in the study and control group. Overexpression of bcl-2 was present in 5.6% (mean \pm SD = 9.61 \pm 13.27) of women in the control group and in 25.0% (mean \pm SD = 20.44 \pm 27.69) (of patients with low grade SIL. Analysis of frequency of patients with positive findings of bcl-2 oncogene revealed no statistical differences when control and low grade SIL groups were evaluated (Fisher's exact test, $P = 0.164$). On the basis of the frequency of patients with bcl-2 overexpression, a 2x2 table of contingency was formed (Table 2), in which sensitivity and specificity were calculated. The sensitivity of determining bcl-2 as a method

for detection of low grade SIL was 25.0% and specificity was 94.4% (Figure 2). The positive predictive value was 80.27%, which means that this percentage of patients with bcl-2 overexpression may be expected to harbor low grade SIL in the cervix. The negative predictive value was 94.8%, meaning that this percentage of

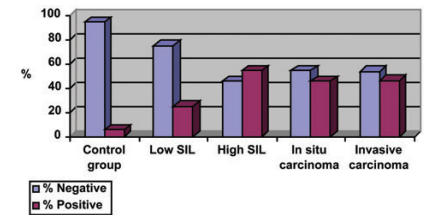


Figure 1. Percentage of positive and negative bcl-2 expression.

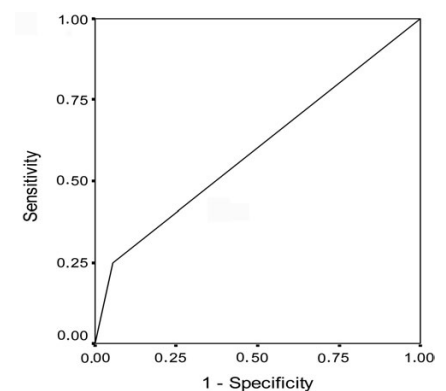


Figure 2. Receiver operating characteristic curve for bcl-2 overexpression in patients with low grade squamous intraepithelial lesions.

Table 1. Expression of bcl-2 in the control and study groups.

bcl-expression expression	Control group, n (%)	Low grade SIL, n (%)	High grade SIL, n (%)	PVU <i>in situ</i> carcinoma, n (%)	PVU <i>invasive</i> carcinoma, n (%)
Negative (-)	20/36 (55.5)	16/32 (50)	8/22 (36.3)	8/22 (36.3)	8/26 (30.8)
Negative (1+)	14/36 (38.9)	8/32 (25.0)	2/22 (9.1)	4/22 (18.2)	6/26 (23.1)
Positive (2+)	2/36 (5.6)	2/32 (6.2)	8/22 (36.3)	6/22 (27.3)	8/26 (30.8)
Positive (3+)	0/36 (0)	6/32 (18.8)	4/22 (18.2)	4/22 (18.2)	4/26 (15.3)

SIL, squamous intraepithelial lesions; PVU, portio vaginalis uteri.

Table 2. Table of contingency (low grade squamous intraepithelial lesions).

Test	Disease present (low grade SIL)	Disease absent (control group)	Total
bcl-2 overexpression	8 ($P = 0.164$)	2	10
bcl-2 negative	24	34	58
Total	32	36	68

SIL, squamous intraepithelial lesions.

patients with bcl-2 negative expression existence of this type of change is not expected.

Table 1 and Figure 1 show that bcl-2 overexpression was present in 5.6% of women in the control group and in 54.6% (mean±SD=27.27±21.89) of patients with high grade SIL (Fisher's exact test, P=0.006; Figure 3). The sensitivity of determining bcl-2 for detecting high grade SIL changes was 54.6% and specificity was 94.4% (Table 3 and Figure 4).

Regarding the obtained values of sensitivity and specificity, the discrimination power of overexpression of this oncogene with the aim of determining the existence of changes of this type was greater than the discrimination power of negative expression in the control group. The positive predictive value in high grade SIL changes was 85.89% and the negative predictive value was 77.21%.

Table 1 and Figure 1 show that bcl-2 overexpression was present in 5.6% of patients in the control group and in 45.5% (mean±SD=6.27±7.04) of patients with PVU carcinoma *in situ* (Fisher's exact test, P=0.018; Figure 5). The sensitivity of this oncogene for diagnosing this type of lesions was 45.5% and specificity was 94.2% (Table 4 and Figure 6). The positive predictive value was 74.64% and the negative

predictive value was 67.98%. Table 1 and Figure 1 show that bcl-2 overexpression was present in 5.6% of patients in the control group and in 46.1% (mean±SD=22.15±23.99) of patients with PVU invasive carcinoma (Fisher's exact test, P=0.012; Figure 7). The sensitivity and specificity of the test were 46.2% and 94.4%, respectively (Table 5 and Figure 8), and the positive and negative predictive values were 85.84% and 70.80%, respectively. Table 6 and Figure 9 show co-expression levels of bcl-2 and c-myc oncogenes in the control and in the experimental group (6/26-

23.0%; (bcl-2 – mean±SD=51.67±20.17; c-myc - mean±SD=43.33±22.29). Co-overexpression of c-myc was only found in patients with PVU invasive carcinoma (Figure 10).

Analysis of frequency of patients with positive findings of bcl-2 and c-myc oncogenes, revealed no statistical differences (P=0.064) when control and PVU invasive carcinoma groups were evaluated. None of the control patients exhibit co-overexpression of bcl-2 and c-myc, while 23% of the patients with PVU invasive carcinoma showed co-overexpression of bcl-2 and c-myc oncogenes. The method's

Table 3. Table of contingency (high grade squamous intraepithelial lesions).

Test	Disease present (high grade SIL)	Disease absent (control group)	Total
bcl-2 overexpression	12 (P=0.006)	2	14
bcl-2 negative	10	34	44
Total	22	36	58

SIL, squamous intraepithelial lesions.

Table 4. Table of contingency (portio vaginalis uteri carcinoma *in situ*).

Test	Disease present (PVU carcinoma <i>in situ</i>)	Disease absent (control group)	Total
bcl-2 overexpression	10 (P=0.018)	2	12
bcl-2 negative	12	34	46
Total	22	36	58

PVU, portio vaginalis uteri.

Table 5. Table of contingency (portio vaginalis uteri *invasive* carcinoma).

Test	Disease present (PVU <i>invasive</i> carcinoma)	Disease absent (control group)	Total
bcl-2 overexpression	12 (P=0.012)	2	14
bcl-2 negative	14	34	48
Total	26	36	62

PVU, portio vaginalis uteri.

Table 6. Co-expression of bcl-2 and c-myc in the control and study groups.

bcl-2 and c-myc co-expression	Control group, n (%)	Low grade SIL, n (%)	High grade SIL, n (%)	PVU <i>in situ</i> carcinoma, n (%)	PVU <i>invasive</i> carcinoma, n (%)
Negative (-)	28/36 (77.8)	24/32 (75)	16/22 (72.8)	18/22 (81.9)	12/26 (46.2)
Negative (1+)	8/36 (22.2)	8/32 (25.0)	6/22 (27.2)	4/22 (18.1)	8/26 (30.8)
Positive (2+)	0/36 (0)	0/32 (0)	0/22 (0)	0/22 (0)	2/26 (7.7)
Positive (3+)	0/36 (0)	0/32 (0)	0/22 (0)	0/22 (0)	4/26 (15.3)

SIL, squamous intraepithelial lesions; PVU, portio vaginalis uteri.

Table 7. Table of contingency (portio vaginalis uteri *invasive* carcinoma).

Test	Disease present (PVU <i>invasive</i> carcinoma)	Disease absent (control group)	Total
bcl-2 and c-myc co-overexpression	6 (P=0.064)	0	6
bcl-2 and c-myc negative	20	36	56
Total	26	36	62

PVU, portio vaginalis uteri.

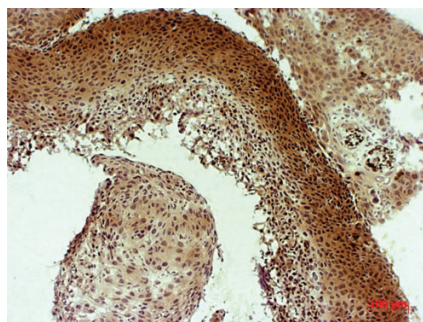


Figure 3. Bcl-2 overexpression (3+) in high grade squamous intraepithelial lesions.

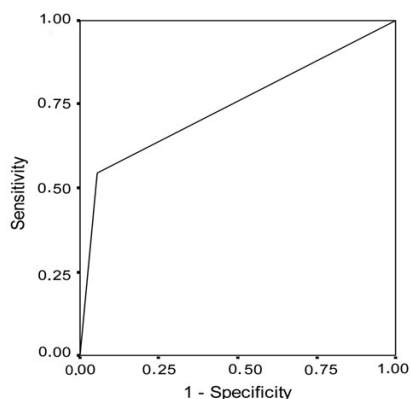


Figure 4. Receiver operating characteristic curve for bcl-2 overexpression in patients with high grade squamous intraepithelial lesions.

sensitivity of determining this oncogene with the aim of detecting PVU invasive carcinoma was 23% while specificity was 72.2%. On the basis of the frequency of patients with bcl-2 and c-myc co-overexpression, a 2x2 table of contingency was formed in which sensitivity and specificity was calculated (Table 7). On the basis of the obtained values of sensitivity and specificity, discrimination power of negative expression in patients with normal findings in the cervix was greater than discrimination power of co-overexpression of this oncogene, with the aim of determining the existence of change of this type (Figure 11). The positive predictive value was 100% which, statistical wise, means that PVU invasive carcinoma may be expected in all patients with co-overexpression of bcl-2 and c-myc. However, the substantiation of the claim requires much more extensive research. The negative predictive value was 64.34%, implying that the existence of this type of change is not expected at this percentage of patients with negative co-expression of bcl-2 and c-myc oncogene.

Discussion

This study evaluated the expression of bcl-2 and c-myc oncogenes in a range of tissues obtained from normal, dysplastic and neoplastic conditions of the cervix, in an attempt to elucidate the expression of this oncoprotein in uterine cervix premalignant and malignant lesions. Bcl-2 is an oncogene that has been investigated for prognostic significance in various malignancies, including carcinoma of the cervix.¹⁶⁻¹⁸ Up to now, conflicting results have been obtained. Bcl-2 has been shown in some studies to be an independent predictor of poor prognosis in carcinoma of the cervix.¹⁹ On the other hand, other studies have revealed no statistically significant correlation with adverse outcome.²⁰

The conflicting results may be due to differences in institutional treatment standards and to varied subjective interpretations of staining intensity and distribution between centers. Since staining is judged on a continuum, differences in institutional cut-off determinations for positive staining may also affect the correlation with clinicopathological results.

Cheung *et al.*, in their analysis of 44 patients with cervical adenocarcinoma, found that increased expression of bcl-2 was correlated with a poorer prognosis.²¹ The absence of correlation between clinical prognosis and bcl-2 expression was instead described by Jain *et al.*²² In our study, no statistically significant difference was found in the frequency of overexpression of bcl-2 between the control group and patients with low grade SIL. However, sta-

tistically significant difference was found in the frequency of overexpression in patients with high grade SIL, PVU carcinoma *in situ* and PVU carcinoma *invasive* in relation to the control group. High sensitivity values speak in favor of great diagnostic significance for the detection of these types of changes in the uterine cervix. On the basis of high predictive values, it can be said that, in patients with overexpression of bcl-2 oncogene, there is a great possibility that they have premalignant or malignant changes in the uterine cervix.

The c-myc oncogene is known to regulate

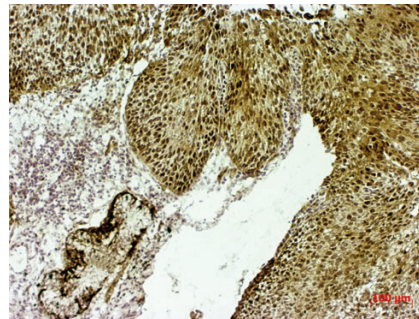


Figure 5. Bcl-2 overexpression (3+) in portio vaginalis uteri carcinoma *in situ*.

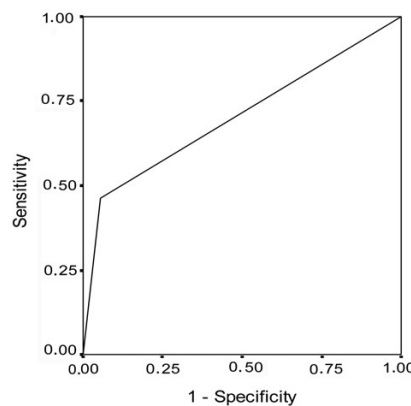


Figure 6. Receiver operating characteristic curve for bcl-2 overexpression in patients with portio vaginalis uteri carcinoma *in situ*.

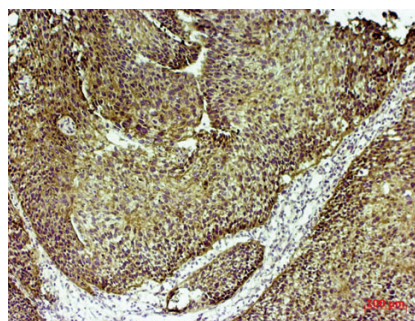


Figure 7. bcl-2 overexpression (3+) in portio vaginalis uteri *invasive* carcinoma.

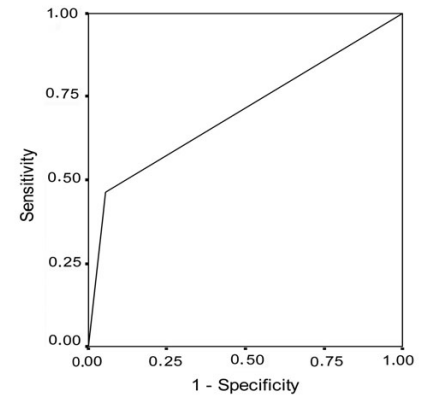


Figure 8. Receiver operating characteristic curve for bcl-2 overexpression in patients with portio vaginalis uteri *invasive* carcinoma.

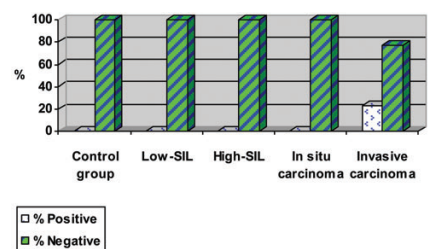


Figure 9. Percentage of positive and negative bcl-2 and c-myc co-expression.

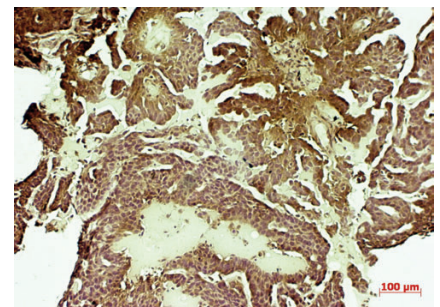


Figure 10. C-myc overexpression (3+) in portio vaginalis uteri *invasive* carcinoma.

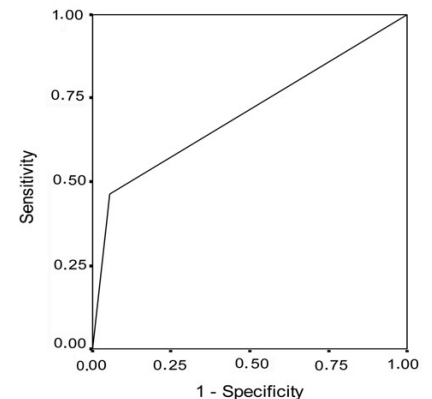


Figure 11. Receiver operating characteristic curve for bcl-2 and c-myc co-overexpression in patients with portio vaginalis uteri *invasive* carcinoma.

neoplastic development and apoptotic cell death. Myc protein is widely distributed in different tissues and is predominantly localized to nuclei of cells where its positivity related to proliferation rate.²³

Bcl-2 at high concentrations protects cells apoptosis induced by c-myc. However, the associations of these proteins and the correlation of their expression and the prognosis of cancer are still controversial. Cooperation between c-myc and bcl-2 in the lymphoid tumors has been demonstrated in double transgenic mice.²⁴

Jinyoung et al. found that overexpression of c-myc and bcl-2 plays a role as prognostic factor in pulmonary adenocarcinomas.²⁵

The positive role of c-myc in promoting cell proliferation places this oncogene at the head of bcl-2 partners for metastatic progression and lymph node involvement of small and well differentiated breast tumors, because bcl-2 may also act as an effective suppressor of the apoptotic action of c-myc, without affecting its ability to promote cell proliferation.²⁶ Liu *et al.* noted a significant positive relationship between bcl-2 and c-myc in female genital tract sarcomas.²⁷

In our study, co-overexpression of bcl-2 and c-myc oncogenes was found only in patients with PVU invasive carcinoma (3/13-23.0%). Analysis of frequency of patients with positive findings of bcl-2 and c-myc, revealed no statistical significant differences ($P=0.064$) when control and PVU invasive carcinoma groups were evaluated. The method's sensitivity of determining this oncogenes with the aim of detecting PVU invasive carcinoma was 23% and specificity was 72.2%. On the basis of the obtained values of sensitivity and specificity, discrimination power of negative expression in patients with normal findings in the cervix was greater than discrimination power of co-overexpression of these oncogenes with the aim of determining the existence of change of this type.

On the basis of high predictive values, it can be concluded that, statistically wise, all patients with co-overexpression of bcl-2 and c-myc oncogenes will have PVU invasive carcinoma.

Conclusions

The findings presented in this study indicate that the evaluation of co-expression of bcl-2 and c-myc may provide additional and independent prognostic information to predict the clinical course of cervical cancer. In our study we confirmed that co-overexpression of these oncogenes suggests with great certainty, at least statistically, that there are malignant changes in the uterine cervix. However, a more extensive series of samples and additional

tests are required to establish the prognostic significance of co-expression of bcl-2 and c-myc in uterine cervix carcinogenesis.

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