

P R A C E O R Y G I N A L N E  
*ginekologia*

# COX-2 expression pattern is related to ovarian cancer differentiation and prognosis, but is not consistent with new model of pathogenesis

Ekspresja COX-2 jest związana z różnicowaniem raka jajnika i prognozą ale nie wiąże się z nowym modelem patogenezy

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## Abstract

**Objective:** Numerous studies suggest that cyclooxygenase-2 (COX-2) is overexpressed in cancer. Our objective was to investigate the relationship between COX-2 expression in ovarian carcinoma and clinicopathological factors. An emphasis was put on the association with the new pattern of tumorigenesis that divides tumors into type I – less aggressive, and type II – more aggressive one. The prognostic significance of COX-2 expression was evaluated.

**Methods:** Ovarian cancer tissues were obtained from 65 patients in FIGO III stage (23 with type I and 42 with type II ovarian cancer). COX-2 expression was evaluated by immunohistochemistry. The statistical analysis was performed in order to assess the connection between COX-2 expression and characteristic factors of ovarian cancer patients as well as the new division for type I and type II ovarian cancer.

**Results:** COX-2 expression was detected in 91% of tissue samples. It was markedly elevated in well differentiated tumors ( $p=0.0041$ ). The platinum - resistant tumors had significantly higher expression of COX-2 ( $p=0.0337$ ). There was no difference between COX-2 expression in type I and type II ovarian cancer ( $p=0.6720$ ). The COX-2 staining was not associated to age, CA125 level, the presence of ascites or any special histological type. An increased expression of COX-2 was an unfavorable prognostic factor for overall survival ( $p=0.0369$ ) and progression-free survival ( $p=0.0218$ ). Multivariate analysis confirmed that COX-2 overexpression is an independent unfavorable prognostic factor of shorter progression-free survival ( $p=0.048$ ).

**Conclusions:** COX-2 expression is an unfavorable prognostic factor for progression-free survival and overall survival in ovarian cancer. There is no relationship between COX-2 expression in ovarian cancer tissue and the examined model of ovarian cancer pathogenesis.

Key words: **ovarian cancer / pathogenesis / COX-2 / immunohistochemistry /**

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## Streszczenie

**Cel:** Liczne badania pokazują, że cyclooxygenaza 2 ulega nadekspresji w nowotworach złośliwych. Celem tego opracowania była analiza związku pomiędzy ekspresją COX-2 w raku jajnika a czynnikami kliniczno-patologicznymi. Szczególny nacisk położono na zbadanie związku z nowym modelem patogenezy raka jajnika, który dzieli nowotwór na typ I – mniej agresywny i typ II – bardziej agresywny. Oceniono prognostyczne znaczenie ekspresji COX-2.

**Metoda:** Tkankę raka jajnika uzyskano od 65 pacjentek w stopniu FIGO III (23 z typem I i 42 z typem II raka jajnika). Ekspresję COX-2 oceniono przy pomocy immunohistochemii. W analizie statystycznej zbadano związek między ekspresją COX-2 a charakterystycznymi cechami kliniczno-patologicznymi oraz nowy podział raka jajnika na typ I i II.

**Wyniki:** Ekspresję COX-2 wykazano w 91% tkanek raka jajnika i była ona istotnie zwiększona w guzach dobrze zróżnicowanych ( $p=0,0041$ ). Platynoooporne guzy miały istotnie wyższą ekspresję COX-2 ( $p=0,0337$ ). Nie znaleziono różnicy pomiędzy ekspresją COX-2 w typie I i II raka jajnika ( $p=0,6720$ ). Ekspresja COX-2 nie była związana z wiekiem, poziomem CA125, obecnością wodobrzusza czy typem histopatologicznym. Zwiększona ekspresja COX-2 okazała się niekorzystnym czynnikiem rokowniczym dla całkowitego czasu przeżycia ( $p=0,0369$ ) oraz czasu wolnego od wznowy ( $p=0,0218$ ). Analiza wieloczynnikowa potwierdziła, że zwiększona ekspresja COX-2 jest niezależnym niekorzystnym czynnikiem prognostycznym czasu wolnego od wznowy ( $p=0,048$ ).

**Wnioski:** Ekspresja COX-2 jest niekorzystnym czynnikiem prognostycznym dla czasu wolnego od wznowy i całkowitego przeżycia w raku jajnika. Nie ma związku pomiędzy ekspresją COX-2 w tkance raka jajnika a badanym modelem patogenezy raka jajnika.

Słowa kluczowe: rak jajnika / patogeneza / COX-2 / immunohistochemia /

## Introduction

A proposed model of ovarian cancer tumorigenesis distinguishes two groups of tumors that differ in the mechanisms leading to malignancy. Briefly, low grade serous, endometrioid, mucinous, clear cell ovarian cancer are considered less aggressive and classified as type I. It is hypothesized that they develop from precursors such as borderline tumors, cystadenomas and endometriosis. On the other hand, high-grade serous and undifferentiated ovarian cancer are assigned to type II subgroup with a more aggressive course of disease. Precursors of type II cancers remain unspecified [1,2]. Cyclooxygenases are the enzymes, which take part in the conversion of arachidonic acid to prostaglandins (PGs) and are responsible for inflammatory reactions. There are two COX isoenzymes: COX-1 gene is constitutively expressed in many types of cells, while COX-2 gene is mainly induced by inflammatory stimuli and mitogens, such as growth factors and cytokines. COX-2 expression is very low and almost undetectable in physiologic conditions. However it is overexpressed in inflamed and neoplastic tissues [3].

The role of COX-2 pathway has been underscored in many neoplasms. Prostaglandins have been reported to stimulate an angiogenesis and suppress cytotoxic immune response [4]. The increased COX-2 expression has been detected in many malignant tumors, among others in: colon cancer, gastric, esophageal, pancreatic, bladder, cervix and endometrial cancer [4-8]. The aim of this research is to analyze how cyclooxygenase-2 expression is related to the hypothesis of ovarian cancer pathogenesis and other clinicopathological factors.

## Materials and methods

### Patients

Immunohistochemical examination was performed retrospectively on tissue samples taken during the first look laparotomy for routine diagnostic purposes. Sixty-five patients operated in 2000-2004 due to ovarian cancer in Department of Obstetrics

and Gynecology (Poznań University of Medical Sciences, Poland) were enrolled in this study. The study was approved by the Institutional Review Board and the patients gave their informed consent before enrollment into the study. The cases were not stratified for any known prognostic factors. The patients distribution in each clinicopathological feature in connection with COX-2 expression level is summarized in table 1. In order to investigate the overall survival (OS) and the progression free survival (PFS) rate, a follow-up examination was performed every 2 months. The mean follow-up period was 37.2 months (range 24-74). The progression free survival time was defined as the time between the end of first line chemotherapy and the diagnosis of the recurrence. Type I and II ovarian cancer groups have been designated on the basis of histopathology and grading, as defined by Kurman and Shih [1, 2]. The type I group includes low-grade (G1) serous, low-grade endometrioid, clear cell and mucinous carcinomas, whereas high-grade (G2 and G3) serous, mesodermal and undifferentiated carcinomas belong to the type II group.

### Tissue

Tissue samples were fixed in 10% buffered formalin and embedded in paraffin. In each case, hematoxylin and eosin stained preparations were subjected to histopathological evaluation by two pathologists. The stage of the tumors was assessed according to the FIGO criteria [9]. Tumors were graded according to the Silverberg grading system [10].

### Immunohistochemistry

Formalin-fixed, paraffin embedded tissue was freshly cut (4  $\mu$ m). The sections were mounted on Superfrost slides (Menzel Gläser, Germany), de-waxed with xylene, and gradually hydrated. Activity of endogenous peroxidase was blocked by 5 min exposure to 3% H<sub>2</sub>O<sub>2</sub>. All the studied sections were boiled for 15 min at 250W in the Antigen Retrieval Solution (DakoCytomation, Denmark). Then, immunohistochemical reactions were

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performed with the use of the following antibody: monoclonal anti-COX-2 (mouse), (Santa Cruz Biotechnology, Santa Cruz, CA), dilution 1:2000 in Antibody Diluent, Background Reducing (DakoCytomation, Denmark) as described previously [5-8]. Tested sections were incubated with antibody for 1h at room temperature. Subsequent incubations involved biotinylated antibodies (15 min, room temperature) and streptavidin-biotinylated peroxidase complex (15 min, room temperature) (LSAB+, HRP, DakoCytomation, Denmark). DAB (DakoCytomation, Denmark) was used as a chromogen (7 min, at room temperature). All the sections were counterstained with Meyer's hematoxylin (DakoCytomation, Denmark). In every case, control reactions, in which specific antibody was substituted by the Primary Mouse Negative Control (DakoCytomation, Denmark), were included.

### Control reactions

In our previous study we have described the control reactions performed in order to evaluate specificity of the COX-2 antibody (Cayman Chemical Company, Ann Arbor, MI, USA) [6].

### Evaluation of reaction intensity

Intensity of immunohistochemical reactions was estimated independently by two pathologists. When in doubt, a re-evaluation was performed using a double-headed microscope and staining was discussed until a consensus was achieved. In order to evaluate the COX-2 expression, a semi-quantitative scale of ImmunoReactive Score (IRS) was applied. It took into account intensity of the colour reaction that was scored 0 to 4 as well as proportion of positive cells that was scored 0 to 3. The final score represented the product of points given for individual characters and ranged between 0 and 12 [11]. Low and high intensity of COX-2 expression was defined as IRS within ranges 0-4 and 6-12, respectively.

### Statistical analysis

For statistical analyses, GraphPad Prism 6 for Windows from GraphPad Software and Microsoft Excel were used. The Mann-Whitney test was used for comparing the intensity of immunoreaction between studied subgroups. To estimate the significance of differences in overall survival time (OS) and progression-free survival time (PFS), Kaplan-Meier statistics (log-rank test) and Cox proportional hazard model were used. If applicable, the relationships between COX expression and clinicopathological features was expressed as Spearman's correlation coefficient. The outcome was recognized as statistically significant when  $p < 0.05$ . All p-values are given for two-sided tests.

### Results

The expression of COX-2 was present in 91% of examined tissues, in each histopathologic type (Figure 1). We did not observe the COX-2 expression in 5 cases (2 serous, 2 undifferentiated, 1 mucinous).

Table I presents an overview of median COX-2 expression values depending on clinicopathological features. The expression was not statistically associated to either ascites or histological type. There was no difference between COX-2 expression in type I and type II cancers ( $p=0.6720$ ). The COX-2 expression was higher in patients who developed the recurrence within 6 months since the end of first line chemotherapy ( $p=0.0337$ ).

**Table I.** Correlation between the COX-2 immunostaining intensity and the clinicopathological factors (Mann-Whitney test).

Characteristics (Number of patients)	Median COX-2 expression	p-value
<b>Histologic type</b> serous (34) non-serous (31)	6.0 4.0	0.1832
<b>Grading</b> G1,G2 (34) G3 (22)	6.0 3.0	0.0041*
<b>Ovarian cancer type</b> I (23) II (42)	6.0 4.0	0.6720
<b>Age</b> ≤50 (34) >50 (31)	5.0 5.0	0.9522
<b>Ascites</b> Yes (20) No (45)	6.0 4.0	0.5690
<b>Time to recurrence</b> ≤ 6months (28) > 6months (37)	6.0 4.0	0.0337*

The median COX-2 expression value in grade 1/2 group was 6.0, whereas in grade 3 it was 3.0 ( $p=0.0041$ ). There was no correlation between COX-2 expression and CA125 level (Spearman  $r = -0.156$ ,  $p=0.278$ ).

The relationships between COX-2 expression and overall survival time (OS) as well as progression-free survival time (PFS) were assessed by means of Kaplan-Meier analysis (log-rank test). Both OS ( $p=0.0369$ ; HR 2.211; 95%CI 1.049-4.658) (Figure 2A) and PFS ( $p=0.0218$ ; HR 2.293; 95%CI 1.077-4.883) (Figure 2B) were shorter in patients with high expression of COX-2. In patients with high COX-2 expression the median OS and PFS were equal to 28 and 9 months, respectively. Whereas in a group with low COX-2 expression the median OS and PFS were 32 and 11 months.

OS and PFS were also estimated within groups demonstrating one particular clinicopathological factor (Table II). We have found an association between high COX-2 expression and decreased OS as well as PFS in following groups: non-serous histologic type ( $p=0.0475$  for PFS), G1/G2 grade ( $p=0.0417$  for OS), age  $< 50$  ( $p=0.0024$  for OS;  $p=0.0149$  for PFS), optimal cytoreduction ( $p=0.0007$  for OS;  $p=0.0049$  for PFS). The observed relationships are presented at the figures 3-4. We have found no relationships between COX-2 expression and either OS or PFS within groups with serous histologic type, G3 grade, type I, type II ovarian cancer, age  $> 50$ , elevated CA125, the presence of the ascites.

In multivariate analysis COX-2 expression was an independent prognostic factor associated with unfavourable PFS ( $p=0.048$ ; HR 1,632; 95%CI 0.958-2.779), however did not influence OS ( $p=0.691$ ; HR 1.224; 95%CI 0.452-3.316).

### Discussion

Our research has revealed an increased expression of COX-2 in ovarian cancer tissue. The staining was observed in nearly all cancer samples. This finding has been corroborated by a number

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**Table II.** High COX-2 expression as a prognostic factor within subgroups divided according to clinicopathological traits (Cox proportional hazard model).

	High COX-2 expression (IRS 6-12)			
	Overall survival		Progression free survival	
	p	HR (95%CI)	p	HR (95%CI)
<b>Serous histologic type</b>	0.2481	1.815 (0.6598 – 4.995)	0.9476	1.026 (0.4734 – 2.225)
<b>Non-serous histologic type</b>	0.1069	2.527 (0.8188 - 7.797)	0.0475*	2.314 (1.009 - 5.303)
<b>G1/G2 grade</b>	0.0417*	3.032 (1.042 - 8.822)	0.7394	1.135 (0.5391 – 2.388)
<b>G3 grade</b>	0.3819	1.758 (0.4962 - 6.232)	0.2702	1.856 (0.6181 - 5.574)
<b>Type I</b>	0.1121	2.986 (0.7745 to 11.51)	0.5046	1.472 (0.4727 – 4.585)
<b>Type II</b>	0.8489	1.066 (0.5409 - 2.123)	0.5134	1.203 (0.6802 - 2.266)
<b>Age &lt;50</b>	0.0024*	5.030 (1.769 - 14.30)	0.0149*	2.606 (1.206 - 5.634)
<b>CA125 elevated</b>	0.5147	1.431 (0.4871 - 4.203)	0.0868	2.489 (0.8766 - 7.066)
<b>Presence of ascites</b>	0.2294	2.066 (0.6328 - 6.748)	0.2120	1.822 (0.7102 - 4.677)
<b>Optimal cytoreduction</b>	0.0007*	10.92 (2.752 - 43.36)	0.0049*	4.880 (1.618 - 14.72)

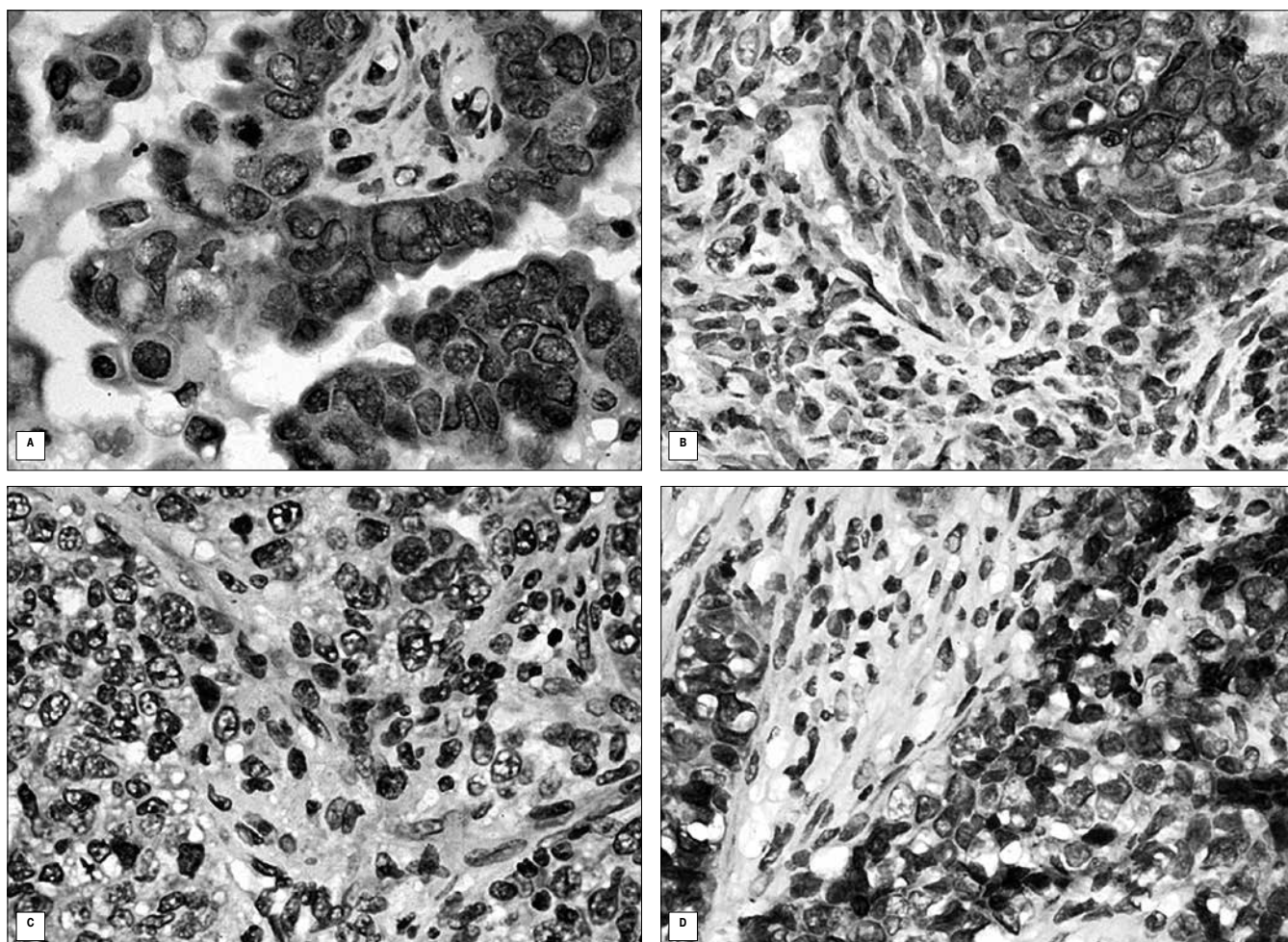
of studies, concerning different types of neoplasms [12, 13]. In the analysis conducted by Denkert on the material from 117 patients, COX-2 expression was revealed in 42% of 86 malignant tumors and 37% of 19 borderline tumors. They did not observe expression of this protein in 12 cystadenomas and in 2 specimen of healthy ovarian tissues [14]. This work pointed that COX-2 expression is present only in certain group of ovarian cancers. Seo et al. also reports that COX-2 expression is markedly increased in non-mucinous histological tumor types compared to mucinous [15]. In our study mucinous ovarian cancer had in two cases high expression and in one case no expression. However this group is too small to draw any conclusions. In one of the previous studies the Northern Blot analysis revealed no mRNA for COX-2 in 12 cases of mucinous ovarian cancer [16]. Moreover, in patient with non-mucinous ovarian cancer, COX-2 expression was related with staging ( $p < 0.001$ ) and the presence of ascites ( $p = 0.036$ ) [15]. Klimp et al. confirmed COX-2 expression in 15 from 18 cases of ovarian cancers and in 10 from 15 borderline tumors [17]. Thus, COX-2 expression in ovarian cancer patients is significantly related to neoplasm histology.

COX-2 expression appeared to be strongly related to the grading of a neoplasm. Well differentiated (G1,G2) tumors have higher COX-2 concentration than those described as poorly differentiated (G3). Such a pattern has been observed in colorectal, lung and hepatocellular cancer [18-20]. What is more, COX-2 expression was detected in precancerous conditions such as colon adenomas [18]. It has been thus suggested that COX-2 is involved in early stages of tumorigenesis. It is worth to stress, however, that the models of neoplastic development in these types of cancer are based on the transition from low grade to high

grade lesions. For this reason it is hardly reproducible to ovarian cancer biology, as current data imply an existence of two distinct pathophysiological pathways leading to ovarian cancer emergence [1, 2, 21]. The genetic profiling suggests that low-grade cancers (mainly type I) develop in a completely different mechanism from high grade tumors (mainly type II) [1]. These studies imply that the transition of low-grade into high grade lesion is not probable. The type I and type II tumors in our investigation, however, do not differ in terms of COX-2 expression.

The assessment of COX-2 expression turned out useful in the division of tumors with respect to response to chemotherapy. Platinum refractory and resistant ovarian cancers (recurrence before 6 months) have enhanced COX-2 expression compared to partly sensitive and sensitive tumors (recurrence after 6 months). This finding has potential clinical implications. A more intense surveillance over the group of patients with increased COX-2 expression in tumor tissue seems therefore reasonable. These patients could especially benefit from the maintenance therapy, as well. Given the link between COX-2 and angiogenesis an antiangiogenic therapy could be considered. Ferrandina et al. was the first who mentioned about relationship between increased COX-2 expression and resistance to chemotherapy in ovarian cancer. The two patient groups were compared, after optimal cytoreduction and after explorative laparotomy. It was proved that in both groups the percentage of COX-2 positive cases was significantly higher among the patients resistant for platinum than among patients sensitive to treatment. From the analyzed clinic-pathological factors, only increased COX-2 expression and the age above 60 were independent prognostic factors of worse treatment response [22]. It has been suggested that COX-2 may

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**Figure 1 A-D.** Immunohistochemical localization of COX-2 expression in ovarian carcinoma cells. Cytoplasmic reaction. A – IRS 9, B – IRS 6, C – IRS 4, D – IRS 3 (hematoxylin, x400).

increase the expression of proteins responsible for multidrug resistance phenomenon, especially the resistance to cisplatin [23].

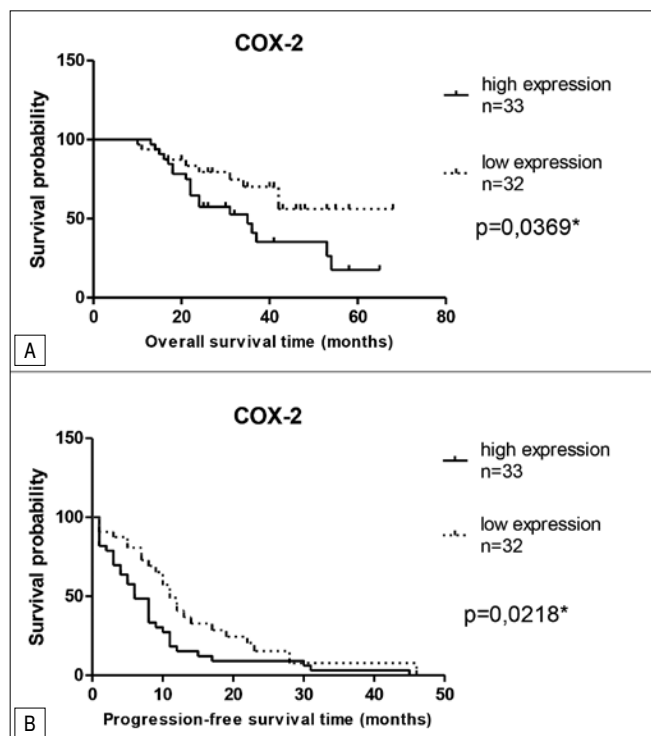
In multivariate analysis the COX-2 expression appeared to be an independent prognostic factor. A high expression is related to low overall survival and progression free survival. Given the heterogeneity of ovarian tumors, an analysis of the prognostic value of COX-2 expression has been conducted in the stratified subgroups. Its usefulness in assessing prognosis has been observed especially in patients with well differentiated tumors, at the age below 50 years and in those who had undergone an optimal cytoreduction. Again, it proves that the tumor differentiation decides about a distinct cancer biology. G1 ovarian cancers are usually related to better survival. An assessment of COX expression may indicate the group of patients vulnerable to recurrence in spite of low grading. It could be especially useful in stage FIGO I when the level of differentiation among others decides about the administration of chemotherapy. The unfavorable prognostic value of COX-2 expression has been corroborated by the meta-analysis [24].

The use of non-steroid anti-inflammatory drugs has appeared effective in prevention of colon cancer [25]. A number of studies has been conducted to confirm this activity with respect to ovarian cancer. Their results, however, remain inconclusive. It has been

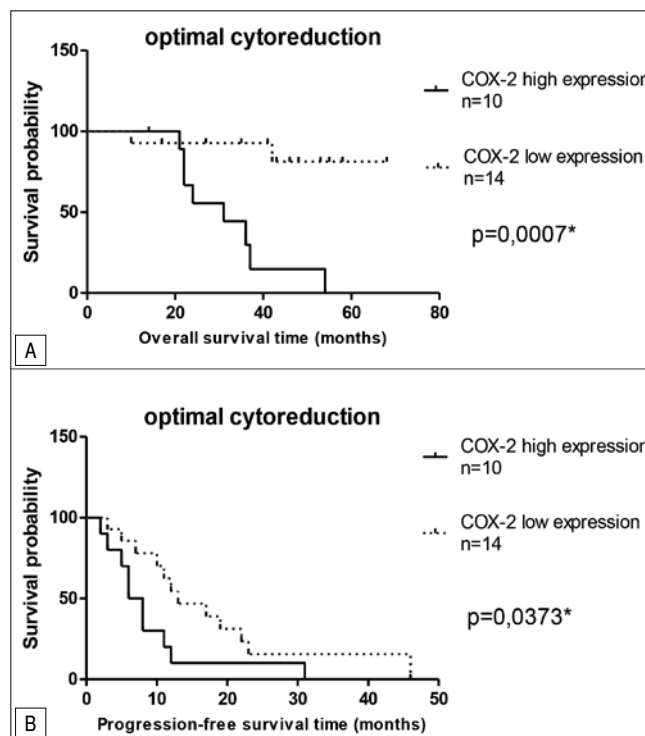
shown by some researchers that regular NSAID intake decrease the risk of ovarian cancer [26-30]. The others suggest that there is no relationship between these drugs administration and ovarian cancer incidence [31].

The mechanisms responsible for the worse prognosis in cases with increased COX-2 expression ovarian cancer are still unknown. Some functions induced by COX-2 were described in the biology of different neoplasms: increased cells proliferation, apoptosis inhibition, angiogenesis stimulation as well as inhibition of immune control mechanisms [3]. In the colon cancer COX-2 expression was unfavourable prognostic factor and correlated with the level of tumor vascularization [32]. COX-2 expression in stomach cancer was related to the tumor invasion into the lymph vessels as well as metastases to the lymph nodes [33]. COX-2 expression can be related to metastasis formation. In lung adenocarcinoma COX-2 expression was increased in metastatic tumors compared to the primary cancer focus [34]. Therefore it may be hypothesised that ovarian cancer cases with increased COX-2 expression have elevated potential for metastases formation and thus worse prognosis comparing to COX-2 negative tumors. Thus, an overexpression of COX-2 in ovarian cancer cannot be only a coexisting phenomenon but rather a mechanism important in the way the tumor progress.

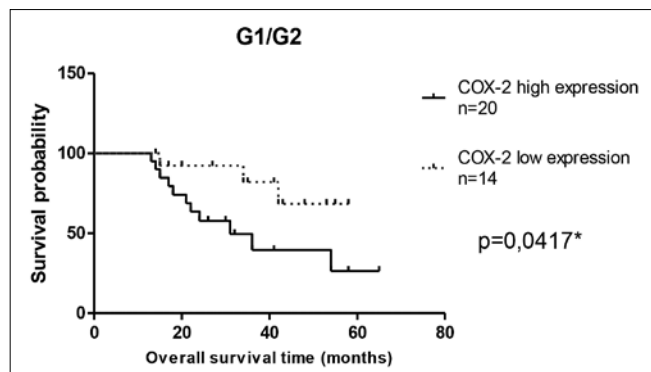
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**Figure 2.** Kaplan Meier curves for A – overall survival time and COX-2 expression; B – progression-free survival time and COX-2 expression.



**Figure 2.** Kaplan Meier curves for A – overall survival time and COX-2 expression; B – progression-free survival time and COX-2 expression.



**Figure 3.** Overall survival time was significantly shorter in cases with high COX-2 expression and G1/G2 grade.

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## Conclusions

COX-2 expression is an unfavorable prognostic factor for progression-free survival and overall survival in ovarian cancer. (Ovarian cancer patients with elevated COX-2 expression may have shorter time to recurrence and shorter survival time). There is no relationship between COX-2 expression in ovarian cancer tissue and the examined model of ovarian cancer pathogenesis.

## Oświadczenie autorów

1. Magdalena Magnowska – zebranie materiału, analiza statystyczna wyników, opracowanie wyników badań, przechowywanie dokumentacji, przygotowanie manuskryptu i piśmiennictwa – autor zgłaszający i odpowiedzialny za manuskrypt.

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**Tematy wykładów:**

1. Obrazowanie wczesnej ciąży (5-10. tygodnie) – nowoczesne metody, a możliwości diagnostyczne.
2. Diagnostyka prenatalna między 11 a 14 tygodniem ciąży – nowości.
3. Praktyczne zasady diagnostyki wad rozwojowych płodu na podstawie przypadków klinicznych, prezentacje multimedialne.
  - układ moczowy
  - układ pokarmowy
  - układ kostny
  - centralny układ nerwowy
- Omówienie rozwoju postnatalnego.  
Follow-up do 10 roku życia.  
Sesje z udziałem neonatologów, chirurgów i pediatrów.
4. Markery ultrasonograficzne aberracji chromosomalnych. Kiedy i na co zwracać uwagę? Odrębności diagnostyczne
5. Diagnostyka aberracji chromosomalnych – ocena DNA płodowego w surowicy matki – nowe możliwości i zagrożenia.
6. Ultrasonografia szyjki macicy i progesteron w leczeniu zagrażającego poronienia i porodu przedwczesnego
7. Diagnostyka ultrasonograficzna nowotworów piersi – nowe wyzwania?
8. Obrazowanie ultrasonograficzne 2D/ 3D/4D w diagnostyce niepłodności i onkologii ginekologicznej.
9. Diagnostyka sonograficzna w uroginekologii
10. Nowości w diagnostyce ultrasonograficznej w ostatnich latach (2013-2014)

Szczegóły na stronie:

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