

Using Parameters of Protein Oxidative Modifications for Differential Diagnosis of Progressive Ovarian Cancer

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Abstract

Background: The aim of the study was to elaborate a method of specific diagnostics of progressive ovarian cancer.

Methods and findings: We observed 21 parameters in plasma of 300 primary ovarian cancer patients, stage III and IV. Using the least squares method, approximation of the relative frequencies by the Rayleigh distribution density was conducted for each parameter. The likelihood functions were constructed and intervals for each measurement on each stage were determined. Approximation accuracy was identified while checking the hypothesis of correspondence of parameter practical value to the theoretical law of density distribution by Pearson criterion.

Conclusions: The obtained results suggest that the level of the oxidative modification of proteins may serve as a reliable indicator for the differential diagnosis of stages III and IV ovarian cancer.

Keywords: Ovarian cancer; Protein oxidative modification; Cytokines

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Introduction

Ovarian cancer (OC) remains the leading cause of death from cancer among gynecologic tumors [1]. According to EuroCare-5 Study statistics, the 5-year survival rate for OC is 30% [2]. 11,700 ovarian malignant tumors and 7,300 deaths from them are registered annually in Russia [3]. Low survival rate of ovarian cancer patients results from late diagnosis. At present, 70% of cases are diagnosed advanced disease [4]. The observed increase in 5-year survival of patients with malignant ovarian tumors in Europe by 3% and in the U.S. by 4% over the past decade can be explained not so much by diagnostics improvement, as by effective application of platinum-based chemotherapy in a treatment of disseminated forms of ovarian cancer and germ cell tumors.

Economic loss resulting from the consequences of malignant tumors incidence is about 90 billion rubles per year and, according to some authors, is going to increase [5]. Therefore it is necessary to correlate the response to treatment with economic factors to determine the cost efficiency of the management of patients with OC. At present, there is a view of chemotherapy futility for cancer patients in terminal stages [6-9]. In this regard, the urgency of clarifying diagnostics methods of advanced (III-IV) disease stage is enhancing.

Ovarian malignant tumors are characterized by absence of symptoms. There are no pathognomonic symptoms of gonadal neoplasia. According to National guidelines [10], diagnosis of OC includes not only precise history taking, full physical, bimanual, recto-vaginal examination, abdominal and transvaginal ultrasound and CT scan, but also chemistry panel and detailed biochemical and clinical blood tests, and determination of the level of tumor markers. For final diagnosis appropriate surgical staging is performed. Due to the complexity of diagnosis and staging of ovarian cancer, the search is conducted for markers that would help to clarify the stage of the tumor process.

Protein oxidative modification is considered one of the earliest and reliable indicators of redox homeostasis disruption [11]. The aim of our research was to develop a clarifying diagnosis method of progressive OC forms on the basis of determining the blood plasma POMP level.

Methods

We studied 300 patients with primary OC stage III-IV according to the classification of the International Federation of Obstetrics and Gynecology (FIGO) and the TNM (7th edition 2010), who were treated the first time during the period from January 2010 till August 2013 in Ulyanovsk Regional Clinical Oncology Center. FIGO staging of ovarian cancer changed from 1 January 2014. However, a study was carried out on the basis of the previous FIGO classification, which has been used on the territory of the Russian Federation until 2015. FIGO stage based on surgical data and TNM categories based on clinical and/or pathological classification. Using the standardized methods, 21 indicators were defined in blood plasma of each patient: leukocyte count, neutrophil count, levels of metalloproteinases MMP-2 and MMP-9 in neutrophils and blood serum, levels of cytokines IL-1 β , IL-6, IL-10, IFN- γ , TNF- α , levels of Her-2/neu and CA125, 4 indicators of POMP in plasma: the level of malondialdehyde (MDA), the activity of glutathione reductase (GR), glutathione S-transferases (GSTs), and catalase. The parameters are shown in **Tables 1-4** and distributed randomly.

For each indicator $\chi_i, i = 1, \dots, N, i \in \mathbb{N}$, functions $\{\rho_i^1(x)\}_{i=1, \dots, N}$ and $\{\rho_i^2(x)\}_{i=1, \dots, N}$.

were constructed, where each function describes probability density of reaching each stage out of two under consideration.

On the basis of these functions, likelihood ratio was constructed for each indicator (P.410) [3].

$$\left\{ z^i(x) = \frac{\rho_i^2(x)}{\rho_i^1(x)} \right\}, i = 1, \dots, N \quad (1)$$

Table 1 Optimal values σ (Le, PMN, Her-2/neu, MMP-2, 9 in PMN) for stages III and IV ovarian cancer.

Indicator Stage	Le	PMN	Her-2/neu	MMP-2 PMN	MMP-9 PMN
III stage	0,79675	0,596502	1,96315	0,650714	1,380432
IV stage	1,86213	0,8433268	1,94747	0,63294	2,034398

Table 2 Optimal values σ (POMP, MDA, Catalase) for stages III and IV ovarian cancer.

Indicator Stage	POMP 356 nm	POMP 370 nm	POMP 430 nm	POMP 530nm	MDA	Catalase
III stage	5,25719	6,46599	3,93999	2,17439	0,996194	3,54199
IV stage	2,29001	2,63601	1,53241	0,53801	0,617518	0,28201

Table 3 Optimal values σ (IL-1 β , IL-6, IL-10, IFN- γ , CA-125) for stages III and IV ovarian cancer.

Indicator Stage	IL-6	IL-10	IL-1 β	IFN- γ	CA-125
III stage	3,69995	0,88705	2,4446012	1,237234	1,20299
IV stage	1,35715	1,25999	2,304716	1,155144	1,18701

Table 4 Optimal values σ (GR, GST, TNF- α , MMP-2, 9 in serum) for stages III and IV ovarian cancer.

Indicator Stage	GR	GST	TNF- α	MMP-2 serum	MMP-9 serum
III stage	1,89599	1,57761	1,044316	2,0501928	3,932618
IV stage	4,01999	0,67055	0,9374848	1,780624	3,879402

Thus, for each indicator $\chi_i, i = 1, \dots, N, i \in \mathbb{N}$, intervals $(a_{ik}; b_{ik})$ and $(c_{ik}; d_{ik}), k \geq 1$ were defined, where $z_i \geq D$ and $z_i \leq \frac{1}{D}$, correspondingly (D-threshold level). We united intervals $(a_{ik}; b_{ik}), k \geq 1$ into the multitude $K_2^i = U_{k \geq 1}(a_{ik}; b_{ik})$, and $(c_{ik}; d_{ik}), k \geq 1$ into the multitude $K_1^i = U_{k \geq 1}(c_{ik}; d_{ik})$, correspondingly.

For each indicator $\chi_i, i = 1, \dots, 21$ characteristic values were simply ordered by a tenfold-hundredfold reduction in each disease stage.

$$K_1^i = U_{k \geq 1}(c_{ik}; d_{ik}),$$

Consider three situations:

1) if $\exists i_0$ is with the property that $y_{i_0} \in K_j^{i_0}, j = 1, 2$, then it is necessary to check that $\forall i \neq i_0, y_i \in K_j^i$. If there is no contradiction, then disease stage j can be diagnosed.

2) if $\exists i_0$ is with the property that $y_{i_0} \in K_j^{i_0}, j = 1, 2$, then it is necessary to check that $\forall i \neq i_0, y_i \in K_j^i$. If there is a contradiction, then D should be increased by 3%-10%.

3) if there is no i_0 with the property that $y_{i_0} \in K_j^{i_0}$, then one should calculate the value

$$\bar{z} = \prod_{i=1}^N z^i \quad (2)$$

If $\bar{z} \leq \frac{1}{D}$ or $\bar{z} \geq D$, then a corresponding disease stage can be

diagnosed exactly. But if, $\frac{1}{D} < \bar{z} < D$ then a patient's disease stage can't be determined exactly.

According to statistics, each parameter was divided into 5 groups. Statistics approximation of the distribution density

$$\rho(x) = \frac{x}{\sigma^2} \exp\left(-\frac{x^2}{2\sigma^2}\right).$$

was made by the least square method.

Optimal values σ are presented in **Tables 1-4** Suppose threshold level

$D=1$, then indicators intervals are defined as follows (**Tables 5-8**).

Results

Let's consider a patient in disease stage III (**Table 9**).

With reference to the 13 parameters (№ 1, 2, 4, 5, 6, 8, 10, 12, 13, 14, 18, 20 and 21)-stage III can be diagnosed, with reference to the rest 8 parameters (№ 3, 7, 9, 11, 15, 16, 17 and 19)-stage IV.

As it is impossible to diagnose the stage exactly in this case, then using formula (2) we get $\bar{z} = 2,74796E-13$. Suppose $D=1$, then $\bar{z} < \bar{D}$

, consequently the patient has stage III of disease.

Let's consider a patient in disease stage IV (**Table 10**).

Table 5 Indicator intervals for stages III and IV ovarian cancer: Le, PMN, Her-2/neu, MMP-2,9 in PMN.

Indicator Stage	Le	PMN	Her-2/neu	MMP-2 PMN	MMP-9 PMN
III stage	[0;1,278]	[0;0,993]	(2,765;+ ∞)	(0,907;+ ∞)	[0;2,341]
IV stage	(1,278;+ ∞)	(0,993;+ ∞)	[0;2,765]	[0;0,907]	(2,341;+ ∞)

Table 6 Indicator intervals for stages III and IV ovarian cancer: POMP, catalase, MDA.

Indicator Stage	POMP 356 nm	POMP 370 nm	POMP 430 nm	POMP 530 nm	Catalase	MDA
III stage	(4,638;+∞]	(5,469; +∞]	(3,232;+∞]	(1,312; +∞]	(0,900; +∞]	(1,088; +∞]
IV stage	[0;4,638]	[0;5,469]	[0;3,232]	[0;1,312]	[0;0,900]	[0;1,088]

Table 7 Indicator intervals for stages III and IV ovarian cancer: IL-1β, IL-6, IL-10, CA-125 in serum.

Indicator Stage	IL-6	IL-10	IL-1β	IFN-γ	CA-125
III stage	(2,922;+∞]	[0;1,480]	(3,355;+∞]	(1,690;+∞]	(1,690;+∞]
IV stage	[0;2,922]	(1,480;+∞]	[0;3,355]	[0; 1,690]	[0;1,690]

Table 8 Indicator intervals for stages III and IV ovarian cancer: GR, GST, TNF-α, MMP-2,9 in serum.

Indicator Stage	GR	GST	TNF-α	MMP-2 serum	MMP-9 serum
III stage	[0;3,728]	(1,370; +∞]	(1,398; +∞]	(2,697; +∞]	(5,523;+∞]
IV stage	(3,728;+∞]	[0; 1,370]	[0; 1,398]	[0;2,697]	[0;5,523]

With reference to 3 parameters we can diagnose stage III, with reference to 18 parameters-stages IV. As it is impossible to diagnose the stage exactly in this case, then using formula (2), we get $\bar{z} = 280189, 27$. Suppose threshold level $\bar{z} > \bar{D}$, then $\bar{z} > \bar{D}$, consequently the patient has stage IV.

Approximation accuracy

Approximation accuracy was identified while checking the hypothesis of correspondence of parameter practical value to the theoretical law of density distribution by Pearson criterion (P.444) [3]. For checking statistical data is introduced

$$\chi^2 = \sum_{i=1}^N \frac{(f^{theor} - f^{pract})^2}{N \times f^{theor}}, \quad (3)$$

Where N -the number of parameters groups, equals 5. The hypothesis is accepted when $\chi^2 < \chi_{\alpha, N}^2$, where α -the level of significance, equal 0.05. Approximation accuracy is calculated as $\gamma = 1 - \alpha$, equal 0.95.

Discussion

We have analyzed chemistry and hematology panel values of 300 OC patients and stated the absence of statistically significant

difference between these values in various stages of the disease, as well as the absence of significant correlation relationships between above mentioned values and OC stage [12]. Also, we found no correlation between CA-125 level and cancer stage in primary OC patients.

According to the cited literature, the change in oxygen metabolism plays an important role in carcinogenesis [13]. It is stated that metabolism change from oxidative phosphorylation to glycolysis is related to the higher production of reactive oxygen species (ROS) [14]. Furthermore, electron transfer through pentose phosphate pathway to NADPH and GSH is enhanced. As a result, redox homeostasis in neoplasia changes [15]. There is an opinion stating that high proliferative activity, drug resistance, apoptosis reduction and immortalization of tumor cells can be mediated by increasing of ROS levels, which results in changing of gene expression [16].

Protein oxidative modification products (POMP)-protein carbonyl derivatives-have long-term half-life; their number increases not only due to oxidative posttranslational modification, but also proteolytic destruction.

Applying assumed parameters of follow-up examination, stage clarification was performed in 300 patients with primary OC who were treated at Ulyanovsk Regional Clinical Oncology Center. In 21 patients the stage of cancer was changed: 9 patients had the stage changed from IV to III, 12 patients-from III to IV.

With reference to obtained results, patients with updated stage III received 6 courses of multiagent chemotherapy instead of 4; patients with updated stage IV received immunotherapy.

According to the obtained results, the level of protein oxidative modification products is an accurate indicator for differential diagnostics of ovarian cancer stage III and IV.

The patent was approved for this method of differential diagnostics of ovarian cancer.

Table 9 Likelihood function value Z1 for diagnostics parameters of ovarian cancer (stage III).

Indicator	Value	Likelihood function, Z_i	Stage
Le	0,66	0,678	III
PMN	0,4818	0,588	III
POMP 356	3.5	2,04	IV
POMP 370	7.1	0,292	III
POMP 430	5.5	0,027	III
POMP 530	3.4	1,179E-07	III
Her2	2.05	1,007	IV
CA125	3.1	0,938	`
MMP-2 serum	2.4062	0,0007	IV
MMP-9 serum	6.258	1	III
MMP-2 PMN	0.403	1,06	IV
MMP-9 PMN	1.6141	0,992	III
IL-6	6,27	1,045	III
IL-10	0	0,665	III
IL-1 β	1.9942	1,079	IV
IFN- γ	0.9932	1,094	IV
TNF- α	0.8572	1,144	IV
MDA	1.109	1,083	III
Catalase	0	1	IV
GST	1	1,1111	III
GR	1.65	0,463	III

Table 10 Likelihood function value, Z1 for diagnostics parameters of ovarian cancer (stage IV).

Indicator	Value	Likelihood function, Z_i	Stage
Le	1.28	1,001	IV
PMN	1.0624	1,1051	IV
POMP 356	1,1	4,7999	IV
POMP 370	2,9	3,632	IV
POMP 430	1,9	3,442	IV
POMP 530	1,0	3,227	IV
Her2	0,41	1,015	IV
CA125	1	1,017	IV
MMP-2 serum	2,2355	1,093	IV
MMP-9 serum	6,502	0,989	III
MMP-2 PMN	0,345	1,048	IV
MMP-9 PMN	2,1737	0,859	III
IL-6	2,08	2,689	IV
IL-10	0	1	III
IL-1 β	2,4637	1,055	IV
IFN- γ	1,2837	1,062	IV
TNF- α	0,9715	1,1184	IV
MDA	0,516	2,099	IV
Catalase	0,1	157,65	IV
GST	4	0,2478	IV
GR	0,42	4,713	IV

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