

The Value of Genetic Research of Women with Abnormal Perimenopausal Bleeding

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Abstract

This paper makes analyses of the value of genetic research of women with abnormal perimenopausal bleeding. On this case, both methodological and theoretical aspects of the research points were discussed by author. Therefore, genetic research of women with abnormal perimenopausal bleeding were focused in order to identify points as the whole. Finally, research has been pinpointed on research development while accentuating on both outcomes and shortcomings.

IJOBAS
Accepted 13 May 2020
Published 16 May 2020
DOI: 10.5281/zenodo.3831312

Keywords: Value, genetic research, women, abnormal, perimenopausal bleeding,

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Introduction

The period of perimenopause is characterized by a gradual extinction of ovarian function, during which hypoluteinism is replaced by anovulation with relative hyperestrogenia and then hypoestrogenia. At any stage of perimenopause, there is a high probability of the formation of menstrual dysfunction, in particular, abnormal uterine bleeding (AUB). The frequency of AUB in premenopause reaches 60-70% among gynecological diseases [5, 9, 12].

The problem of AUB is studied in various aspects: etiology, pathogenesis, diagnosis, prevention, therapy. However, studies aimed at testing the pathogenesis of proliferative processes in the endometrial mainly affect hormonal disorders in a woman's body and the expression of steroid hormone receptors in the uterine mucosa [12, 14, 15]. Further study requires a comprehensive assessment of risk factors for the occurrence and development of AUB based on molecular genetics and immunological status.

Frequent relapses of AUB and malignancy determine the morphofunctional features of various types of endometrial hyperplasia (EH), as well as epigenetic and genetic disorders leading to inactivation of tumor suppressor genes, increased proliferation, angiogenesis, and decreased apoptosis [7, 8, 11, 12]. In recent years, there has been a tendency to study the molecular genetics foundations of the development of AUB, to search for predictors of its formation and progression in EA. The results of the studies were mixed, which did not contribute to solving the problem. In some studies, the following were selected as prognostically significant markers: loss of PTEN expression, TP53 gene mutations, decrease in the apoptotic BCL-2 / BAX index [1, 14], as well as a low level of expression of both progesterone receptor isoforms (PGR).

Main part

One can also find such a view of the problem in the literature — molecular genetic markers identified for RE (MSI, MSH1, KRAS, (3-catenin and others) can hardly be considered reliable predictors of AUB [15]. Native authors also made a contribution in problem solving [11]. They were able to identify the most informative clinical characteristics and laboratory indicators that reflect the level of methylation of tumor suppressor genes (MLH1, RASSF1, p16, GSTP1, TP53). The reason for these scientific differences is multifunctional and insufficient knowledge of the pathogenesis of AUB. For practical use, it is important to create an integrated prognostic criterion based on the most significant predictors of the formation of ER against the background of AUB.

Many authors evaluate the relationship of molecular genetics indicators of proliferative activity (Ki-67), apoptosis (APAF-1), neoangiogenesis (VEGF), and components of the extracellular matrix (MMP-1, MMP-9, TIMP-1) and the morphological variant of endometrial hyperplasia leading to AUB. [3,5,6].

THE PURPOSE OF THE STUDY:

To determine the role of genetic factors in predicting the course of AUB in perimenopause.

SURVEY RESULTS:

To achieve the above goal, we studied the role of the genetic marker MMP9 and mutation of the TP53 suppressor gene of women, which were divided into 2 groups: the main and the control. The main group consisted of 75 women with AUB in perimenopause. The control group consisted of 25 healthy women.

Matrix metalloproteinase (MMP) are the family of extracellular zinc-dependent endopeptidases that can break down all types of extracellular matrix proteins. They play a

role in tissue remodeling, angiogenesis, cell proliferation, migration and differentiation of cells, apoptosis, and tumor growth inhibition. They involved in the cleavage of membrane receptors, the release of apoptotic ligands, such as FAS, as well as in the activation and deactivation of chemokines and cytokines.

MMP were first described in vertebrates in 1962, and later found in invertebrates and plants. The main differences between MMP and other endopeptidases are their dependence on metal ions, the ability to destroy the structures of the extracellular matrix. [10,12].

MMP9	Gelatinase-B, 92 kDa- Gelatinase	secreted	may play a significant role in the local proteolysis of the extracellular matrix and in the migration of leukocytes. It can participate in proliferation and in endometrial apoptosis. Splits collagen type IV and V into large C-terminal and smaller N-terminal fragments. Cleaves fibronectin, but not laminin.
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In women with various pathological conditions, such as AUB and endometrial cancer, the level of MMP-9 is increased. It was shown that in patients with AUB in serum, the concentration of MMP-9 is significantly higher than in practically healthy individuals.

p53 (p53 protein) is a transcription factor that regulates the cell cycle. P53 acts as a suppressor of the formation of malignant tumors, respectively; the TP53 gene is an anti-oncogen. Mutations of the TP53 gene are found in cells of about 50% of cancers. It is often called the “guardian of the genome” [1,11,14].

For complete impairment of gene function (in particular, anti-oncogen), as a rule, inactivation of both copies of the gene on two chromosomes of the diploid cell genome is necessary. Such a violation may occur as a result of an extended deletion - complete or partial loss of the nucleotide sequence encoding the gene. Substitutions, loss or insertion of single nucleotides in DNA are also possible. Replacing a nucleotide at some point can cause the formation of the so-called termination codon, and protein synthesis at this point will be interrupted. Losses or insertions of single nucleotides lead to a malfunction of the correct framework for the translation of mRNA into protein. After such a mutation, “incorrect” amino acids are included in the growing polypeptide chain and termination codons appear which prematurely terminate the synthesis of the polypeptide chain. Another possibility is the replacement of single nucleotides, which change the meaning of a codon encoding an amino acid residue. This leads to the replacement of one amino acid in the protein by another, and if the replacement occurs on a functionally important part of the protein, then it can change its functionality or even completely lose it. Such substitutions are called missense mutations.

Among the tumor suppressor genes, TP53 is the best known. Its product is phosphoprotein p53, which regulates the transcription of several other genes. In a normal cell, p53 is inactive, but in extreme events the genome guard activates and plays the role of performing various anti-cancer functions: it activates the DNA repair system (the Nobel Prize in Chemistry was awarded to study the mechanisms of this system in 2015); if DNA is damaged, p53 delays mitosis of dividing cells, blocking the transition from the G1 phase to the S phase and giving the repair system time to repair the damage; if DNA damage cannot be repaired, p53 includes a cell death program called apoptosis. At the same time, the TP53 gene itself is very often susceptible to mutations in many types of cancerous tumors. For TP53, all of the above types of mutations are known, and usually there is no relationship between the type of cancer and certain types of mutations [10,14,23].

Group I consisted of 75 women with AUB in perimenopause. Exclusion criteria were women with coagulopathic diseases, cancer patients and women in whom AMA was an iatrogenic complication. The age of the women surveyed ranged from 45-51 years. All women underwent the following examination methods:

- anamnesis and assessment of the nature of bleeding
- clinical blood test
- gynecological examination
- Doppler mapping
- sonohysterography
- curettage of the uterine cavity with subsequent histology
- hysteroscopy with targeted biopsy and histology.
- determination of serum levels of MMP3 and mutation of the TP53 gene by PCR.

In 16 (21.2%) of them, endometrial polyps were found, in 17 (22.6%) endometrial hyperplasia, 13 (17.3%) of uterine fibroids were 5.3% of them submucous, adenomyosis was in 6 (8.23%)) women, a combination of endometrial hyperplasia and uterine leiomyoma 12 (16%), a combination of leiomyoma and adenomyosis 11 (14.67%). Pathomorphological causes of abnormal uterine bleeding are shown in the table.

Table 1

Pathomorphological causes of AUB

Pathomorphological causes of AUB	P=75	%
Endometrial polyps	16	21,2
Endometrial hyperplasia	17	22,6
Myoma	5	17,3
Leiomyoma	5	
Submucous node	3	
adenomyosis	6	8,23
Hyperplasia + adenomyosis	11	14,67
Hyperplasia + leiomyoma	12	16
	75	100

The women with AUB, a significant ($p < 0.05$) increase in serum MMP-9 level was observed when comparing the main group and the control group. In the main group, the concentration of MMP-9 was 423.94 ng / ml (SD = 128.9; the range of values was 237.0–740.0 ng / ml), in the control group, 162.7 ng / ml (SD = 23, 7; 120.0–269.0 ng / ml). In the analysis of MMP-9, based on the data obtained, a concentration of 240 ng / ml should be considered a critical value: when it is exceeded, a statistically significant ($p = 0.063$) correlation with the probability of hyperplastic processes in the endometrium is noted. When taking the above specified value of the concentration of MMP-9 as a threshold isolated assessment of the level of MMP-9 in the blood serum, it has a sensitivity of 86%, specificity of 80%.

Table 2
The content of MMP-9 in the blood serum of examined women

The main group of examined MMP-9 (ng/ml)	Control group MMP-9 (ng/ml)
423,94±17,71*	162,7±3,35

* Compared to the control group at $p < 0.05$.

The tumor suppressor gene and its corresponding p53 protein play a central role in the development of apoptosis [22,29]. P53 mutations are the most common genetic disorder in the development of malignant tumors. The appearance of mutant forms of this gene leads to the accumulation in tumor cells of inactive forms of the protein that cannot fulfill the functions of normal p53 [16, 20, 21]. Mutations of this gene can be associated with the aggressive course of the disease and the resistance of tumor cells to the effects of anticancer drugs and radiation therapy. [15; 27]. In RE, according to various researchers, mutant p53 is found in 44–64% of patients [25].

Endometriosis, leiomyoma, polyps can be attributed to multifactorial diseases, in the genesis of which an important role belongs not to one but to many different genes. Given the proximity of these diseases to cancer, it can be assumed that the gene network of these diseases and ER should include identical or similar genes. One of these is TP53, whose mutations naturally occur in a wide variety of tumors. The product of this gene is a key element in controlling the cellular response to various types of stress.

By PCR, the blood serum of the main and control group of patients was examined.

Table 3

Gene polymorphism	Main group		Control group	
	N	M ±M%	N	M ±M, %
TP53 Arg72Pro	75	30,7±4,1	25	4,2±2,4

Of the 75 women in the main group, the level of MMP9 was increased in 69 (92%). In the control group, the level of MMP9 was increased in 3 (12%) women. Mutated p53 was positive in 23 (31%) in the examined women with AUB; in the control group, an increase in the level of mutated p53 was observed in 1 (4%) patient. Thus, in women with AUB, the level of MMP9 is increased in 92% of cases. Based on this result, it can be judged that this genetic marker is specific for AUB.

A positive result for mutated p53 is a criterion for the formation of an increased cancer risk group, which makes it possible to optimize the management of women with this pathology.

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Cite this article:

Author(s), Z.Z. ASKAROVA, M.Z. KURBANIYAZOVA, (2020). "The Value of Genetic Research of Women with Abnormal Perimenopausal Bleeding". Name of the Journal: International Journal of Academic Research in Business, Arts and Science, (IJARBAS.COM), P, 58- 65. DOI: <http://doi.org/10.5281/zenodo.3831312> , Issue: 5, Vol.: 2, Article: 6, Month: May, Year: 2020. Retrieved from <https://www.ijarbas.com/all-issues/>

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