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Blood biomarker panel recommended for personalized prediction, prognosis, and prevention of complications associated with abdominal aortic aneurysm

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Abstract

The aim of the study was to evaluate the ability of following biomarkers as diagnostic tools and risk predictors of AAA: C-reactive protein, interleukin-6, pentraxin-3, galectin-3, procollagen type III N-terminal peptide, C-terminal telopeptide of type I collagen, high-sensitive troponin I, and brain natriuretic peptide. Seventy-two patients with an AAA and 100 healthy individuals were enrolled into the study. We assessed individual biomarker performance and correlation between the AAA diameter and biomarker levels, and also, a multivariate logistic regression was used to design a possible predictive model of AAA growth and rupture risk. We identified following four parameters with the highest potential to find a useful place in AAA diagnostics: galectin-3, pentraxin-3, interleukin-6, and C-terminal telopeptide of type I. The best biomarkers in our evaluation (galectin-3 and pentraxin-3) were AAA diameter-independent. With the high AUC and AAA diameter correlations in AAA patients. Authors recommend to add biomarkers as additional parameters to the current AAA patient management. Main addition value of biomarkers is in the assessment of the AAA with the smaller diameter. Elevated biomarkers can change the treatment decision, which would be done only based on AAA diameter size. The best way how to manage the AAA patients is to create a reliable predictive model of AAA growth and rupture risk. A created multiparameter model gives very promising results with the significantly higher efficiency compared with the use of the individual biomarkers.

Keywords Abdominal aortic aneurism \cdot Biomarker panel \cdot Pentraxin-3 \cdot Galectin-3 \cdot Interleukin-6 \cdot Procollagen type III N-terminal peptide \cdot C-Terminal telopeptide of type I collagen \cdot High-sensitive troponin I \cdot Brain natriuretic peptide \cdot Multivariate stepwise logistic regression \cdot Multivariate model \cdot Predictive preventive personalized medicine \cdot Patient stratification

Abbreviations

AAA	abdominal aortic aneurysm
AUC	area under the curve
BNP	brain natriuretic peptide
CRP	C-reactive protein
hsTnI	high-sensitive troponin I

ICTPC-terminal telopeptide of type I collagenIL-6interleukin-6PIIINPprocollagen type III N-terminal peptidePTX-3pentraxin-3ROCreceiver operating characteristicSASStatistical Analysis Software

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Introduction

Abdominal aortic aneurysm (AAA), the enlargement of the abdominal aorta to a diameter at least 3 cm, is a common disease in the western part of the world. The prevalence in overall population ranges between 3 and 7% and is higher in White men and current smokers with the increasing risk with the age [1]. Smoking, White race, male gender, genetic factors, presence of other aneurysms in large vessels, and hypercholesterolemia are named as main risk factors of AAA. Also, modifiable lifestylerelated factors are significantly associated with AAA. Men with active lifestyle (walking or bicycling) have three times lower risk of AAA than men who never do walking or bicycling [2]. The growth rate of AAA is individual. It is estimated that the time to develop AAA of 3 cm or larger is approximately 5 years [3]. The most risky complication is AAA rupture. The mortality associated with ruptured AAA is still high. When including operative mortality, 80-90% of patients with ruptured AAA died [4]. The pathogenesis of AAA is a multifactorial process and is characterized by chronic inflammatory reaction in the aortic wall, particularly in infiltration of tunica media and tunica adventitia by inflammatory cells and extracellular matrix degradation [5]. AAA is usually diagnosed by physical examination, abdominal ultrasound, or computer tomography (CT). However, the speed of growing is individual, and also, the risk of rupture is not just dependent on the size of the aneurysm [6, 7]. Rupture even in small AAA was reported, while large aneurysms remain stable. Stability of the AAA wall depends on its size and the degree of the proteolytic enzyme activity on the enhanced degradation of the extracellular matrix and the related degree of inflammation occurring in the AAA wall [8]. Individual growth rate and multifactorial pathogenesis of AAA do not allow standardized approach to the patients. Individualization of the diagnostic and treatment process is one of the main conditions of future success [9]. The pathological processes in the aortic wall are connected with the production and involvement of various biomarkers. The research of these biomarkers is very extensive worldwide [10]. The aim of our study was to evaluate the individual biomarkers' performance in AAA diagnostics to correlate the AAA diameter with the age and selected biomarkers, and finally, using multivariate analysis creates the multiparameter model to AAA patient stratification. This model could be used as diagnostic and prognostic tool to identify potentially risky aneurysms, which allow to apply a personalized approach in treatment [11]. The biomarkers were selected as the following-the biomarkers regulating inflammatory response: C-reactive protein (CRP), interleukin-6 (IL-6), pentraxin-3 (PTX-3), and galectin-3; the degradation

products of the extracellular matrix: procollagen type III N-terminal peptide (PIIINP) and C-terminal telopeptide of type I collagen (ICTP); and biomarkers of myocardial injury and vascular resistance: high-sensitive troponin I (hsTnI) and brain natriuretic peptide (BNP).

Materials and methods

Study population

A total of 172 participants were enrolled into the study. The patient group comprised 72 patients with the AAA.

Inclusion criterion was the diagnosis of AAA and open surgery or endovascular treatment at the university hospital in Pilsen. Exclusion criteria were malignancy in the anamnesis, infection or autoimmune disease, and renal and hepatic dysfunction. The control group comprised 100 healthy individuals who underwent regular preventive examinations at our university hospital. A summary of the description of patient and control group is shown in Table 1.

Serum samples

Blood samples were collected prior to the endovascular treatment or open surgery. Peripheral blood was drawn using VACUETTE® Z Serum Sep tubes (Greiner Bio-One, Kremsmünster, Austria) and allowed to clot. Serum was separated within 3 h of collection by centrifugation at $1700 \times g$ for 10 min, and all samples were immediately aliquoted and frozen to -80 °C. Samples were stored under a 24-h monitoring system. Serum samples were thawed only once, just prior to the analysis. Serum levels of PTX-3, IL-6, galectin-3, hsTnl, CRP, BNP, PIIINP, and ICTP were measured for each sample.

 Table 1
 Characteristics of the study groups

Characteristic	Patient group	Control group
Male	59 (82%)	56 (56%)
Female	13 (18%)	44 (44%)
Total	72 (100%)	100 (100%)
Age, years (min-max)	56 (36-80)	53 (32–74)
Average size of aneurysm (mm)	65	Not applicable
Current smoking	41 (55%)	46 (46%)
Diabetes mellitus	9 (12%)	7 (7%)
Hypertension	45 (61%)	40 (40%)
Ischemic heart disease	21 (29%)	19 (19%)
Peripheral artery disease	11 (15%)	5 (5%)

Parameter	Status	Mean	Median (min-max)	Lower quartile	Upper quartile	Wilcoxon test p value
CRP (mg/l)	AAA Controls	5.79 2.82	1.92 (0.19–100) 1.19 (0.13–44.05)	0.81 0.56	5.16 2.98	0.0248
IL-6 (pg/ml)	AAA Controls	5.09 2.03	3.03 (0.58–76.72) 1.36 (0.18–18.35)	2.16 0.88	5.76 2.42	< 0.0001
PTX-3 (pg/ml)	AAA Controls	7223 3524	6545 (1246–18,916) 2872 (708–12,673)	4010 1727	9453 4813	< 0.0001
Galectin-3 (ng/ml)	AAA Controls	19.93 13.21	18.85 (8.00–54.70) 12.95 (5.6–27.10)	15.30 10.70	22.20 14.70	< 0.0001
PIIINP (µg/l)	AAA Controls	5.26 3.90	4.86 (2.54–10.19) 3.92 (0.12–6.72)	4.16 3.33	6.09 4.68	< 0.0001
ICTP (µg/l)	AAA Controls	5.46 3.55	4.77 (2.39–15.15) 3.37 (1.75–6.73)	3.63 2.88	6.57 4.08	< 0.0001
hsTnl (ng/l)	AAA Controls	16.30 3.25	4.90 (1.70–754) 2.25 (0.23–37.30)	3.50 1.40	7.40 3.35	< 0.0001
BNP (pg/ml)	AAA Controls	26.86 3.55	14.80 (10.00–135) 3.37 (1.75–6.73)	10.0 2.88	27.5 4.08	0.0128

 Table 2
 Levels of the selected biomarkers in the AAA group and control group

Legend: AAA, abdominal aortic aneurysm; CRP, C-reactive protein; IL-6, interleukin-6; PTX-3, pentraxin-3; PIIINP, procollagen type III N-terminal peptide; ICTP, C-terminal telopeptide of type I collagen; hsTnI, high-sensitive troponin I; BNP, brain natriuretic peptide

Sample analysis

Serum levels of CRP were measured using the chemiluminescent assay with the Maglumi Instrument (Snibe Co., Shenzhen, China). IL-6 and hsTnI were measured using the chemiluminescent ACCESS assays with the UniCel DxI 800 Instrument (Beckman Coulter, Brea, CA, USA). PTX-3 was measured using Simple Plex assay with the Ella System (ProteinSimple, San Jose, CA, USA). Galectin-3 and BNP were measured using the chemiluminescent assays with the Architect 1000i System (Abbott, Libertyville, IL, USA), and PIIINP and ICTP were measured using the radioimmunoassays UniQ PIIINP RIA and ICTP RIA (Orion Diagnostica, Espoo, Finland) with the STRATEC SR 300 Instrument (Stratec Biomedical AG, Birkenfeld, Germany).

Statistical analysis

Statistical Analysis Software release 9.2 (SAS Institute Inc., Cary, NC, USA) was used for all statistical analyses. Basic descriptive statistic parameters are presented: mean, median, minimum, maximum, lower quartile, and upper quartile. Spearman rank correlation coefficient was used to estimate correlation between AAA diameter, biomarkers' levels, and age. The Wilcoxon test was used to compare distributions of biomarker values between the groups of patients and the group of healthy persons, while a *p* value of less than 0.05 indicated statistical significance. Receiver operating characteristic (ROC) curves and the area under the curve (AUC) are presented to aid an assessment of the usefulness of selected biomarkers for AAA diagnosis. A multivariate stepwise logistic regression was used whereby the biomarkers (as AAA predictors) were added one-by-one until they achieved statistical significance after addition to the model. The previously added predictors were eliminated if, after the addition of other predictors, they had become statistically non-significant. This ensures that all the predictors, which are in the final logistic regression model, remain statistically significant [12].

Results

Table 2 shows the results of the comparison of CRP, IL-6, PTX-3, galectin-3, PIIINP, ICTP, hsTnl, and BNP levels in the AAA patient group and in the group of healthy persons. AAA group had statistically significantly higher levels of all the examined biomarkers compared with the control group. We plotted ROC curves and calculated the AUC for each individual biomarker. The values of the AUC are shown in Table 3. The best values were achieved by galectin-3 (AUC = 0.8383), hsTnl (AUC = 0.8306), PTX-3 (AUC = 0.8029), IL-6 (AUC = 0.7890), and ICTP (AUC = 0.6014). ROC curves are shown in Figs. 1, 2, 3, 4, 5, 6, 7, 8.

Table 4 shows the correlation between the AAA diameter and biomarkers. The correlation with statistical significance was observed in the case of CRP, hsTnl, IL-6, PIIINP, and

 Table 3
 AUC values of all tested biomarkers

Parameter	AUC	95% confidence limit	
Galectin-3	0.8383	0.7744	0.9022
hsTnl	0.8308	0.7701	0.8916
PTX-3	0.8029	0.7374	0.8685
IL-6	0.7890	0.7209	0.8570
ICTP	0.7689	0.6933	0.8446
PIIINP	0.7448	0.6664	0.8232
BNP	0.6084	0.5244	0.6924
CRP	0.6014	0.5149	0.6879
Model	0.9012	0.8499	0.9526

Legend: *AUC*, area under the curve; *hsTnI*, high-sensitive troponin I; *PTX-3*, pentraxin-3; *IL-6*, interleukin-6; *ICTP*, C-terminal telopeptide of type I collagen; *PIIINP*, procollagen type III N-terminal peptide; *BNP*, brain natriuretic peptide; *CRP*, C-reactive protein; *Model*, multivariate stepwise selection model created using galectin-3, PTX-3, IL-6, and ICTP

ICTP. Galectin-3, PTX-3, BNP, and age did not correlate with the AAA diameter.

When multivariate stepwise logistic regression was applied, the following four biomarkers were selected as the best predictors of AAA: galectin-3, PTX-3, IL-6, and ICTP. We created a multivariate model (AUC = 0.9012, Fig. 9) with the significantly higher efficiency compared with the use of the individual biomarkers only (p < 0.0001).

Fig. 1 ROC curve of galectin-3. ROC analysis of the galectin-3 ability to separate AAA patients and healthy persons. Galectin-3 proved to be a best individual biomarker to this purpose. Model: galectin-3, AUC = 0.8383

Discussion

Recently, a large number of studies have been published that analyze various biomarkers in patients with AAA [2, 8, 13, 14]. We summarized our data, which will be discussed in following text in Tables 2 and 3.

Inflammatory biomarkers

The most common focus is the effort to detect signs of an inflammatory reaction occurring in the wall of the aorta as a potential reason for the aortic wall instability and hence an increased risk of rupture of the aneurysm. Being one of the basic inflammatory markers [15], CRP is a frequently investigated parameter in connection with AAA. CRP is long-established and used as a reliable non-specific inflammatory marker with an important place in inflammatory diagnostics. However, with regard to AAA, the use of CRP is limited. In our study, CRP correlates with the AAA diameter but achieved the worst AUC value (AUC = 0.6014), and we cannot recommend its use as an AAA marker. Inflammation cells, like activated macrophages, monocytes, or eosinophils, produce many kinds of cytokines. There are many studies available on this topic. However, the conclusions regarding the usefulness of individual cytokines are controversial [13]. IL-6 was confirmed as one of the most significant markers. There is a hypothesis that IL-6 is directly produced by an AAA [16]. In our study, IL-6 was higher in the

ROC Curves for Comparisons

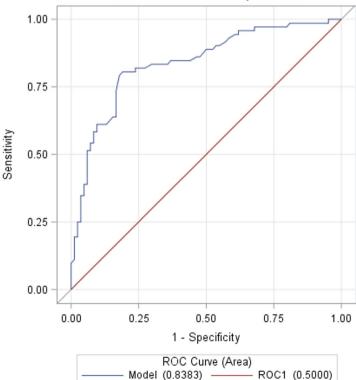


Fig. 2 ROC curve of hsTnI. ROC analysis of the hsTnI ability to separate AAA patients and healthy persons. hsTnI achieved the second best AUC, and it is a candidate to be used as an independent prognostic biomarker of the upcoming heart complications in AAA patients. Model: hsTnI, high-sensitive troponin I, AUC = 0.8308

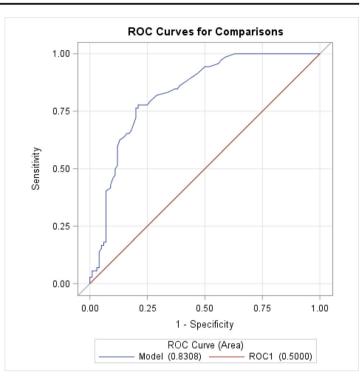


Fig. 3 ROC curve of PTX-3. ROC analysis of the PTX-3 ability to separate AAA patients and healthy persons. PTX-3 proved to be a potential useful part of the multi parameter predictive model of AAA growth and rupture risk. Model: PTX-3, pentraxin-3, AUC = 0.8029

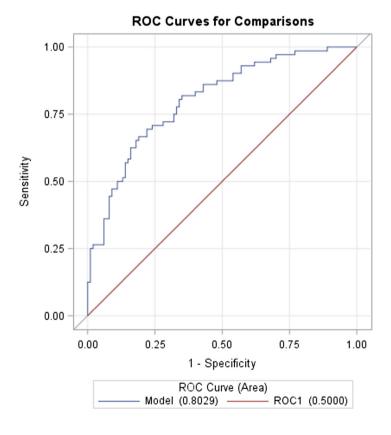


Fig. 4 ROC curve of IL-6. ROC analysis of the IL-6 ability to separate AAA patients and healthy persons. IL-6 is the next candidate to be a potential useful part of the multiparameter predictive model of AAA growth and rupture risk. Model: IL-6, interleukin-6, AUC = 0.7890

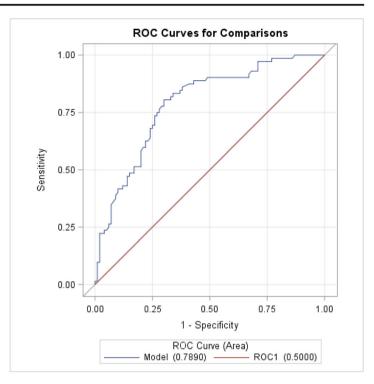


Fig. 5 ROC curve of ICTP. ROC analysis of the ICTP ability to separate AAA patients and healthy persons. ICTP is the last candidate from our biomarker panel to be a part of the multi parameter predictive model of AAA growth and rupture risk. Model: ICTP, C-terminal telopeptide of type I collagen, AUC = 0.7689

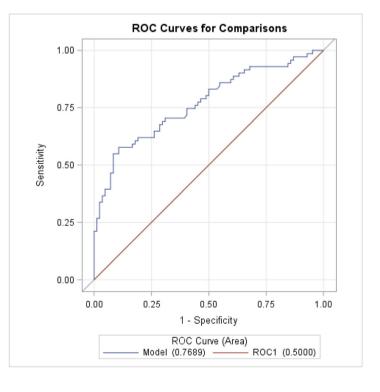
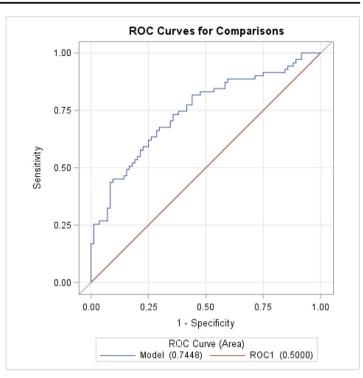
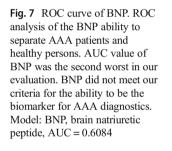


Fig. 6 ROC curve of PIIINP. ROC analysis of the PIIINP ability to separate AAA patients and healthy persons. Despite relatively satisfied AUC value, PIIINP was not chosen by created multi parameter model of AAA growth and rupture risk. Model: PIIINP, procollagen type III Nterminal peptide, AUC = 0.7448





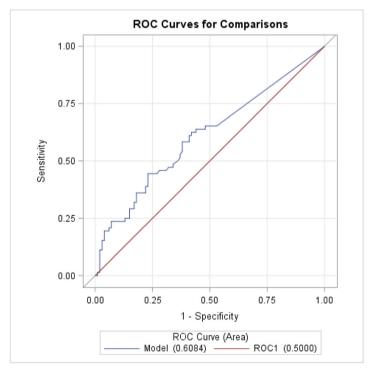


Fig. 8 ROC curve of CRP. ROC analysis of the CRP ability to separate AAA patients and healthy persons. AUC value of CRP was the worst in our evaluation. CRP did not meet our criteria for the ability to be the biomarker for AAA diagnostics. Model: CRP, C-reactive protein, AUC = 0.6014

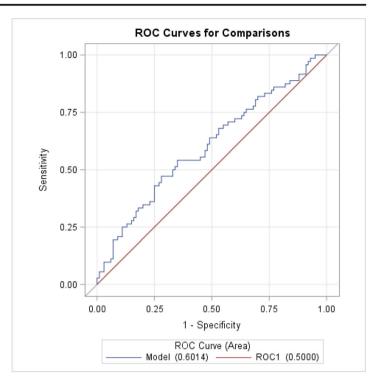


Fig. 9 ROC curve of multi parameter model. ROC analysis of the multi parameter model ability to separate AAA patients and healthy persons. AUC value of this model was statistically significantly higher compared with the individual biomarkers. This result showed that the right way for the future in AAA diagnostics is the shift from single parameter assessment to multi parameter systematic model. Model: multiparameter model, multivariate stepwise selection model created using galectin-3, PTX-3, IL-6, and ICTP, AUC = 0.9012

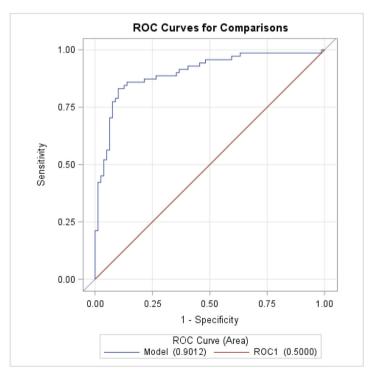


Table 4 Correlation of AAA diameter with the biomarker leve	Table 4	Correlation of AAA	diameter with the	biomarker level
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Item	Spearman correlation coefficient	p value
CRP	0.3210	0.0060
hsTnI	0.3197	0.0062
IL-6	0.3152	0.0070
PIIINP	0.2896	0.0143
ICTP	0.2879	0.0149
BNP	0.1393	0.2429
age	0.1231	0.3028
PTX-3	0.1131	0.3440
Galectin-3	0.0031	0.9791

Legend: *CRP*, C-reactive protein; *PTX-3*, pentraxin-3; *IL-6*, interleukin-6; *PIIINP*, procollagen type III N-terminal peptide; *ICTP*, C-terminal telopeptide of type I collagen; *hsTnI*, high-sensitive troponin I; *BNP*, brain natriuretic peptide

AAA group with a strong statistical significance (p < 0.0001) compared with the healthy group, achieved a good AUC value (AUC = 0.7890), and also correlated with the AAA diameter. This is in concordance with previously published studies our own team's included [15, 17].

In the field of inflammatory markers, many new molecules have recently been introduced. We focused on two of them: PTX-3 and galecin-3. PTX-3 is a glycoprotein, a member of the pentraxin family. It is produced by a variety of cells, such as monocytes, macrophages, dendritic cells, endothelial cells, and fibroblasts in response to an inflammatory stimulus. It is one of the acute phase proteins and is close to CRP. Its role has been studied in connection with acute aortic dissection or acute coronary syndrome. Very little is currently known about the association between the AAA and serum levels of PTX-3 [18, 19]. We proved statistically significantly higher serum levels of PTX-3 in patients with an AAA (p < 0.0001) with a high AUC value (AUC = 0.8029). To our knowledge, only one study reported PTX-3 expression in the tissue of AAA [20]. Based on our results, we can assume that PTX-3 plays an important role in the pathogenesis of AAA and seems to be a potentially new AAA diameter independent biomarker.

Galectin-3 is a member of the family of lectins. It is produced by activated macrophages and is involved in cell adhesion, proliferation, angiogenesis, and apoptosis [21]. Galectin-3 has previously been investigated in connection with inflammation and risk of the cardiovascular diseases and mortality [22]. Data regarding the relation of galetin-3 to AAA are rare. Recent studies show a positive correlation between galectin-3 levels and AAA incidence. In the general population, systemic galectin-3 concentrations are elevated years before the identification of AAA incidence [23]. Increased galectin-3 levels were associated with AAA progression, while galectin-3 inhibition decreased experimental AAA development. In addition, galectin-3 serum levels were associated with the need for surgical repair, independently of other potential diagnostic factors [24]. In our study, we observed statistically significantly higher serum levels of galectin-3 in patients with an AAA (p < 0.0001) with the highest AUC value (AUC = 0.8383). This finding proved an important role of galectin-3 in the AAA pathogenesis and the ability of this molecule to be a good marker of inflammation and increased systemic vascular resistance in AAA. Similarly to PTX-3, galectin-3 seems to be AAA diameter independent. We did not prove any correlation with the AAA diameter in case of both above mentioned biomarkers.

Collagen degradation products

Another group of biochemical markers, which are very often studied in connection with AAA pathogenesis, is the extracellular matrix degradation products, collagen, and elastin, respectively. Collagen, as an important component of this matrix, appears to be a key element in the process of AAA development and progression. If collagen degradation prevails over the synthesis, the irreversible destruction of the aortic wall is started. This situation can directly lead to the expansion and subsequent rupture of an AAA [25]. In our work, we decided to analyze PIIINP and ICTP. We understand these two collagen degradation products to be an indicator of collagenolysis. In the available literature, there are a number of studies of these parameters in patients with AAA [13, 22]. Certain works had shown that PIIINP is able to predict further development and the eventual necessity for intervention in a small AAA [26, 27]. Also, the association of PIIINP with congenital cardiovascular disorders has been proved [24]. In our study, we found statistically significantly elevated PIIINP in AAA patients (p < 0.0001) as well as ICTP (p < 0.0001). The AUC values for PIIINP (AUC = 0.7448) and ICTP (AUC = 0.7689) were still acceptable for possible clinical use. PIIINP and ICTP correlated with the AAA diameter.

Biomarkers of cardiac injury and vascular resistance

The last two biomarkers in our study were hsTnI and BNP, which were primarily connected with cardiac injury and increased vascular resistance [13, 28]. A new generation of high-sensitive troponins (hsTn) opens up a new field of indication for these types of biomarkers [29]. In our work, we used newly introduced hsTnI kit, and to our knowledge, it is the first use in this indication, which was published. We observed statistically significantly elevated hsTnI levels in AAA patients (p < 0.0001), and hsTnI also reached the high AUC value (AUC = 0.8303) when ROC analysis was done and also correlated with the AAA diameter. hsTnI seems to be an independent sign of the cardial risk and can early selected the patients we have to focused on in this regard. In the case of BNP, recent studies describe increased BNP in patients who later developed AAA [30]. In our study, BNP levels were also statistically significantly higher in the group of AAA patients (p < 0.0213); however, its AUC value was the second

worst in our evaluation (AUC = 0.6084), and BNP also did not correlate with the AAA diameter.

Correlation between biomarkers and AAA diameter

Looking at our data, it is visible that the relation between concentration of biomarkers and AAA diameter will not provide reliable information, which can be used in the decision about following treatment procedure. Two biomarkers, which have the best ability to separate AAA patients and healthy persons (PTX-3 and galectin-3), achieved the worst correlation coefficient with the AAA diameter. CRP achieved the best correlation coefficient, but according to the AUC value, it is very unspecific. Age is considered as one of the main risk factor of AAA development. Therefore, we also correlated age of AAA patients with AAA diameter, but in concordance with the literature findings, it is without a positive result [31]. We recommend to assess AAA diameter and biomarker activity as two independent parameters with the addition of the biomarker levels to the current system of AAA evaluation, which is based on AAA diameter.

Multiparameter model

The rapid development of various diagnostic technologies is driving the shift from single parameter model to multiparameter systematical model. This way allows us to better interpretation of the results, which accelerates predictive, preventive, and personalized medicine (PPPM) practice allowing prediction of response with substantially increased accuracy, stratification of particular patients, and eventual personalization of medicine [32]. Using experience from our previous research [11] and other relevant studies [33, 34] as the last step of our eight biomarker panel evaluation, we selected the best candidates for use in clinical practice. For this reason, multivariate stepwise logistic regression was used, and the following four biomarkers were selected as the best predictors of AAA: galectin-3, pentraxin-3, IL-6, and ICTP. We created a multivariate model (AUC = 0.9012) with the significantly higher efficiency compared with the use of the individual biomarkers (p < 0.0001).

Conclusions and expert recommendations

AAA patients are currently followed up using an imaging techniques and AAA diameter scanning. Our data together with the literature evidences show that blood biomarkers have important information, which allows us better stratify patient and individualize the treatment procedures.

 We identified following four parameters with the highest potential to find a useful place in AAA diagnostics: galectin-3, PTX-3, IL-6, and ICTP.

- We created a multivariate model with the significantly higher efficiency compared with use of the individual biomarkers.
- The best biomarkers in our evaluation (galectin-3 and PTX-3) were AAA diameter independent.
- With the high AUC and AAA diameter correlation, the hsTnI can be used as an independent prognostic biomarker of the upcoming heart complications in AAA patients.
- We recommend to add biomarker levels as additional parameters to the current algorithm of AAA patient management.
- The main addition value of biomarkers we see in the assessment of the AAA with the smaller diameter.
 Elevated biomarker levels can work as a warning signal and can changed the treatment decision, which would be done only based on AAA diameter size.
- The best way how to manage the AAA patients is to create a reliable predictive model of AAA growth and rupture risk. Our multiparameter model gives very promising results, and we will focus on the confirmation of current results on the larger group of patients.

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Compliance with the ethical standards

Competing interests The authors declare that they have no competing interests.

Ethical approval All investigations conformed to the principles outlined in the Declaration of Helsinki. Informed consent was obtained from all the participants. The study was approved by the responsible Ethical Committee of the University Hospital in Pilsen on August 12, 2014.

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