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Serum amyloid a protein as a potential biomarker in predicting acute onset and association with in-hospital death in acute aortic dissection



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Abstract

Background: Acute aortic dissection (AAD) is a life-threatening disorder in vascular surgery with a high early mortality. Serum amyloid A (SAA) is a kind of acute-phase protein with a rapid diagnostic value in other diseases. However, the researches on the performance of SAA for the diagnosis of AAD is still lacking. This retrospective study aimed to evaluate the SAA levels and further explore its potential diagnostic role in AAD patients.

Methods: SAA levels were measured by enzyme-linked immunosorbent assay (ELISA) in 63 controls and 87 AAD patients. Laboratory examinations were also performed. And relative clinical information was collected from participants included in this study.

Results: SAA levels were significantly higher in AAD patients than those in healthy controls. SAA levels were independently associated with the risk of AAD. There was a positive significant correlation between SAA and C reactive protein (R = 0.442, and P = 0.001). Based on receiver-operating characteristic (ROC) analysis, the area under the curve (AUC) of SAA for the diagnosis of AAD were 0.942 with optimal cut-off points of 0.427 mg/L. For in-hospital mortality, the AUC of SAA were 0.732 with optimal cut-off points of 0.500 mg/L. According to logistic regression analysis, higher SAA levels represent a higher risk of in-hospital mortality (OR = 1.25; 95%CI: 1.07–1.47; P = 0.005).

Conclusion: Our findings demonstrated that SAA levels were significantly enhanced in AAD. SAA was closely correlated with inflammatory parameters and coagulation-related parameters in AAD. Furthermore, SAA could be a potential bio-marker for identifying AAD in the early diagnosis. Finally, SAA > 5.0 mg/L are independently related to AAD in-hospital mortality.

Keywords: Serum amyloid A, Acute aortic dissection, Biomarker, Inflammation

Background

Acute aortic dissection (AAD) is a life-threatening disorder in vascular surgery defined as the separation of aortic wall layer [1, 2]. In line with the Stanford classification, AAD is commonly divided into type A aortic

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dissection (TAAD) (involving ascending aorta) as well as type B aortic dissection (TBAD) (ascending aorta not affected) by considering both dissection extent and lesion site [3, 4].

AAD is characterized by acute onset, rapid progression as well as high morbidity and mortality in early stage. It has been previously reported that approximately 48.6% of untreated AAD patients die pre-hospitally [5]. Meanwhile, both TAAD and TBAD have high short-term inhospital mortality. Medical imaging, including computed tomography (CT) and magnetic resonance imaging (MRI), is a reliable approach for AAD diagnosis [2, 6]. However, due to the mimic clinical manifestations between AAD



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In addition, the clinical values of several serum biomarkers in the early diagnosis of AAD are increasingly recognized [8, 9]. For instance, C reactive protein (CRP) and D-dimer have been included in the ESC guideline for aortic diseases for the assessment of AAD patient conditions [10–12]. An ideal biomarker should be generally characterized by a rapid diagnostic value, a low cost, noninvasive and easy to perform with its potential sensitivity and specificity.

Serum amyloid A (SAA), is a specific apolipoprotein of high-density lipoprotein cholesterol (HDL-C), involved in the acute phase response. Diverse functions of SAA have been proved in cardiovascular diseases (CVD), including regulating matrix metalloproteinase-2 activity in aorta [13], modifying the vascular functionality of HDL-C [14], and regulation of systemic inflammatory reaction [15]. Recent research efforts have focused on the potential role of SAA as a biomarker in clinical disorders [16, 17]. However, there has been no study concerning the value of SAA in the early diagnosis and prognostic prediction in AAD.

In this study, the levels of SAA were assessed in AAD subjects as well as normal controls, followed by analysis of the potential correlation of SAA with AAD. We also investigated the diagnostic performance of SAA as a novel clinical biomarker in AAD prediction. Moreover, we also aimed to analyze the association between SAA and these classical biomarkers, including CRP as well as D-dimer level, followed by analysis on the possible combined diagnostic efficacy to detect AAD. Finally, the possible correlations between AAD prognosis and SAA level were identified.

Methods

Study population

In this retrospective study, eligible aortic dissection patients were selected from two Affiliated Hospital of China Medical University (Shenyang, China). In brief, 131 consecutive AAD patients with corresponding peripheral blood samples and clinical data were collected from the Department of Vascular Surgery of the First Hospital of China Medical University (CMU) immediately after admission between May 2014 and February 2019. In addition, another 93 AAD patients were collected from the Department of Clinical Laboratory, Shengjing Hospital of CMU between January 2016 and January 2018. Among these 93 patients, 50 and 43 patients were diagnosed with TAAD and TBAD, respectively, during hospitalization (Fig. 1). The diagnosis of AD was based on imaging outcomes (CT, MRI and echocardiography). The AD was categorized in line with Stanford classification [18].

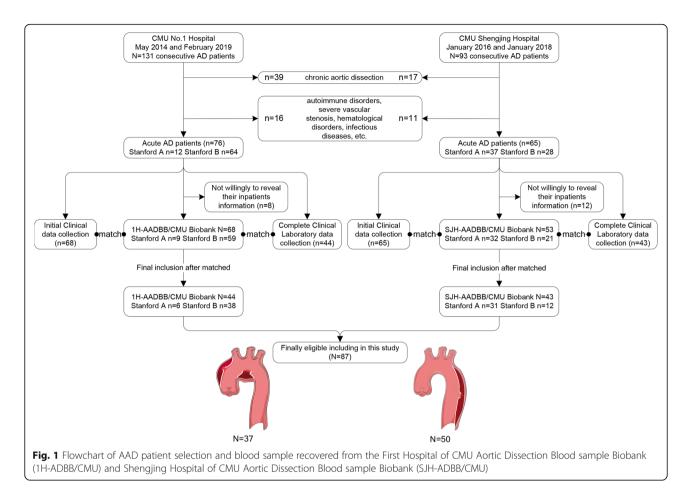
Definitions

AD could be divided into acute as well as chronic phases. AAD is diagnosed if a patient is admitted to hospitalization within 14 days of symptom onset, otherwise, chronic AD is considered [5]. Moreover, acute AD patients admitted to the department of emergency of our two clinical centers were clearly diagnosed after careful assessment within 24 h following symptom onset. The diagnostic criteria of hypertension included a clinical record of systolic blood pressure (BP) ≥140 mmHg and/or diastolic BP \geq 90 mmHg, and the administration of anti-hypertensive drugs. The definition of diabetes mellitus (DM) included fasting glucose level \geq 7.0 mmol/L, glycosylated haemoglobin $A1c \ge 6.5\%$ and the administration of oral hypoglycaemic drugs or insulin. In addition, the smoking status was judged in accordance with selfreport current smokers.

The definitions of terms utilized in this research were listed in the following. The definition of a communicating false lumen in AAD is the opacification of at least partial false lumen with contrast media except ulcer-like projection [19]. On the contrary, a non-communicating false lumen in AAD is defined by the completely occlusive false lumen by a thrombus, as well as ulcer-like projection. Moreover, the definition of upper and lower strata SAA levels were set in line with cut-off value of receiver operating curve (ROC), which was 0.427 mg/L as cut-off value in the present study.

Exclusion criteria

The AD patients with the following disorders were eliminated: chronic aortic dissection, malignancy, autoimmune disorders, severe aortic stenosis which was defined by an aortic valve area less than 1.0cm² or less than $0.6 \text{ cm}^2/\text{m}^2$ if indexed to body surface area [20], hematological disorders, infectious diseases, coronary artery disease which was defined as the presence of at least one≥50% stenosis in a coronary artery≥2.0 mm in diameter based on either coronary computed tomography angiography or invasive coronary angiography examination [21], severe organ failure, congenital heart disorders, previous aortic operation, Marfan syndrome, Ehlers-Danlos syndrome, other types of combined connective tissue or vascular disorders, and those receiving non-steroidal anti-inflammatory drugs or steroids. These disorders were ruled out using angiographic diagnosis, imaging examinations, laboratory tests, echocardiography, and other medical examinations based on the clinical presentations and medical history of patients. Eventually, there were 55 from First Hospital and 28 from Shengjing



Hospital incongruent patients eliminated in this study (Fig. 1).

Aortic dissection blood sample biobank

After excluding incongruent participates, on February 2019, eight and twelve patients were unwillingly to provide their inpatient record for publication, including their clinical data or blood samples, respectively (Fig. 1). Finally, aortic dissection blood samples from 68 patients were eligible for further analyses as Biobank, which registered in the First Hospital of CMU Aortic Dissection Blood sample Biobank (1H-ADBB/CMU) were selected to identity AAD. As a result, a total of 53 blood samples from AAD patients registered in Shengjing Hospital of CMU Aortic Dissection Blood sample Biobank (SJH-ADBB/CMU) were eligible and further enrolled in this study.

Venipuncture was conducted on eligible patients after their admissions, followed by sample collection of in EDTA plastic tubes (BD Vacutainer^{\circ} lavender, 5.0 mL) and anticoagulant and silica/gel plastic tubes (SST BD Vacutainer^{\circ} gold, 5.0 mL). Moreover, blood sample was centrifuged to collect plasma, which was reserved at – 80 °C for further test (up to 1 year). Peripheral blood mononuclear cells (PBMCs) of these AD patients were extracted by Ficoll-sodium diatrizoate density gradient centrifugation as described previously [22].

The present AD Biobank research was conducted in line with the Guidelines of the World Medical Association Declaration of Helsinki, and was approved by the Ethics Committee of Shengjing Hospital (Ethics Approval No. 2016PS085K) of China Medical University. All 121 subjects signed written informed consent.

Serum measurements

ELISA kits (SAA: Wuhan Boster Biotechnology Company, China) were purchased to determine the SAA levels according to the manufacturer's protocol.

Laboratory examinations

Lipid panel

The plasma levels of low-density lipid cholesterol (LDL-C) and high-density lipid cholesterol (HDL-C) were directly detected using selective solubilization method (LDL-C test Kit or Determiner L HDL, Kyowa Medex, Tokyo). Additionally, levels of total cholesterol (TC) as well as TG were measured by enzymatic methods. Automatic biochemistry analyzer (ARCHIRECT ci16200, Abbott Laboratories, USA) was utilized to produce lipid profiles.

Additional biochemistry

Alanine aminotransferase (ALT) and glutamic oxalacetic transaminase (AST) were determined utilizing International Federation of Clinical Chemistry approach (Abbott Laboratories, USA). The plasma concentration of total protein (TP) was determined using biuret method (FUJIFILM Wako Pure Chemical industries Ltd., Japan). The fasting plasma glucose (FPG) levels were determined by urease GLDH and glucose oxidase methods (DiaSys Diagnostic Systems GmbH, Germany).

D-Dimer and CRP

Immunoturbidimetry was used to assess plasma D-dimer levels (Diagnostica Stago, France, normal limit $\leq 0.5 \,\mu$ g/mL). High-sensitivity assay with BN II nephelometer (Dade Behring, Germany) was used to detect CRP levels (normal limit $\leq 0.17 \text{ mg/L}$).

All blood analyses were carried out by Department of Clinical Laboratory at CMU Shengjing Hospital and Department of Clinical Laboratory of First Hospital of China Medical University for SJH-AADBB/CMU Biobank and 1H-AADBB/CMU Biobank, respectively. Follow by clinical data collection, blood Biobank establishment, and laboratory examinations, total of 87 eligible participates (TAAD = 37 and TBAD = 50) were finally identified to meet the inclusion criteria. A summary of the flow of participants' selection and inclusion process is illustrated in Fig. 1.

Control group (ctrl)

The blood sample from 87 patients with AAD were available from the venipuncture together with the blood samples from 63 matched controls. Shengjing Hospital of China Medical University Hospital medical examination database was used to identify healthy control by thoroughly searching all patients admitted to the emergency department diagnosed with trauma or motor vehicle accident during the 2014-2018 period (with imaging examinations, CT and/or MRI, at admission). All participants that were hemodynamicly stable and had non-typical symptoms, were considered as candidates of the control group. The control group was selected because these patients would not be expected to present any inherent bias favoring AAD. We also selected 20 matched patients diagnosed with stable angina as another control group. Stable angina was defined to chest discomfort that is classically retrosternal, triggered by exertion, and relieved by rest or nitrates within minutes [23]. Subjects diagnosed with vascular or connective tissue disorders using imaging examinations, CT and/or MRI, at admission were excluded from the control group. Other exclusion criteria included malignancy, infection, drug history, or any other immunerelated disorders. Eventually, 63 participants in the control group and 20 patients in the angina group were included in this study. And all 83 subjects as control also signed written informed consent and conducted in line with the Guidelines of the World Medical Association Declaration of Helsinki.

Statistical analysis

SPSS 22.0 (SPSS Inc., Chicago, IL, USA) was utilized for statistical analysis. Data were shown as medians with upper or lower quartiles for continuous variables due to the non-Gaussian data distribution. The difference between two groups was measured by non-parametric Mann-Whitney test, respectively. Comparisons among three groups were performed by one-way ANOVA, followed by Tukey's post-hoc test, or nonparametric tests, followed by Kruskal-Wallis 1-way ANOVA test, according to the normality of the values. Categorical variables were shown as numbers with percentages, and the differences between two groups (for both biochemical and clinical parameters) was determined by Chi-square test. Correlations between continuous variables were analyzed through partial correlation analysis accounting for age, gender, and smoking. Furthermore, multiple logistic regression analysis was conducted to evaluate the correlation of serum SAA and D-Dimer or CRP with AAD risk following the adjustment of possible confounding factors. For instance, demographic characteristics and comorbidities were adjusted as confounding factors in all multivariable logistic regression models. Receiver operator characteristic (ROC) curves with area under the curve (AUC) along with logistic models were employed to determine the corresponding cut-off points and to assess the diagnostic performance of serum SAA and D-Dimer or CRP individually, and combined for AAD detection. P values < 0.05 were considered as statistical significance.

Results

Basic clinical characteristics of patients

The detailed clinical features of all subjects were shown in Table 1. Patients had significantly higher levels of heart rate, higher ratio of hypertension and smoker, higher white blood cell (WBC), platelet (PLT), and fast plasma glucose (FPG) (P < 0.001, P = 0.001, P = 0.048, P < 0.001, P = 0.033 and P < 0.001 respectively), but lower level of hemoglobin (Hb) in overall AAD, type A and type B groups in comparison to the control group (P < 0.001). Additionally, TAAD was not significantly different from TBAD in other comparisons. Nevertheless, these features were not statistically different between TAAD group and TBAD group. Moreover, in terms of other clinical features, including age, gender, BMI and rate of diabetes mellitus, healthy controls were not significantly different from

Table 1 Demographic and	laboratory parameters of	participants includ	led in this study

	Control (<i>n</i> = 63)	Angina (<i>n</i> = 20)	AAD (n = 87)	P-value	TAAD (n = 37)	TBAD (n = 50)	P-value
				AAD vs Control			
				AAD vs Angina			
Male, n (%)	43 (68.25%)	15(75.00%)	65 (74.71%)	0.657	27 (72.97%)	38 (76.00%)	0.75
Hypertension, n (%)	21 (33.33%)	8(40.00%)	54 (62.69%)	0.002*	27 (72.97%)	27 (54.00%)	0.073
Smoking, n (%)	19 (30.16%)	7(35.00%)	31 (35.63)	0.774	12 (32.43%)	19 (38.00%)	0.594
DM, n (%)	10 (15.87%)	6(30.00%)	26 (29.89%)	0.123	11 (29.73%)	15 (30.00%)	0.978
Age, Yr	53.44 ± 11.52	57.80 ± 2.11	53.87±11.42	0.957	53.32±11.12	54.1 ± 11.73	0.848
				0.355			
BMI	21.8 ± 3.68	23.70 ± 0.73	22.87 ± 5.14	0.357	22.24 ± 3.92	24.02 ± 5.99	0.149
				0.702			
HR, bmp	78.06 ± 9.6	79.45 ± 1.68	89.46 ± 12.68	< 0.001***	90.49 ± 11.86	88.21 ± 13.28	0.273
				0.001***			
WBC, \times 10 ⁹ /L	5.83 ± 0.77	6.39 ± 0.40	10.74 ± 4.24	< 0.001***	11.54 ± 4.50	10.06 ± 3.92	0.198
				< 0.001***			
HGB, g/L	150.98±17.68	135.50 ± 4.08	132.39 ± 20.97	< 0.001***	128.63 ± 17.86	135.61 ± 23.03	0.052
				0.297			
PLT, $\times 10^9$ /L	173.3 ± 43.14	201.35 ± 13.27	190.59 ± 68.15	0.194	199.34 ± 67.05	183.12 ± 69.00	0.196
				0.744			
FPG, mmol/L	4.97 ± 4.87	5.54 ± 0.24	7.21 ± 2.42	0.001***	7.61 ± 2.84	6.92 ± 2.05	0.355
				0.155			
SAA, mg/L	0.36 ± 0.08	0.72 ± 0.40	4.43 ± 0.31	< 0.001***	4.22 ± 0.33	4.58 ± 0.48	0.595
				0.029*			

Note, AAD acute aortic dissection, TAAD type A aortic dissection, TBAD type B aortic dissection, BMI body mass index, HR heart rate, DM diabetes mellitus, WBC white blood cell, HGB hemoglobin, PLT platelet, FPG fast plasma glucose, SAA serum amyloid A. *P<0.05; **P<0.01; ***P<0.001

AAD groups. Expectedly, the level of SAA protein was significantly enhanced in AAD group in comparison to the healthy control group (AAD vs Ctrl, P < 0.001; TAAD vs Ctrl, P < 0.001; TAAD vs Ctrl, P < 0.001; Table 1 and Fig. 2a). The SAA level was not significantly different between TAAD group and TBAD group (P = 0.595; Table 1 and Fig. 2a).

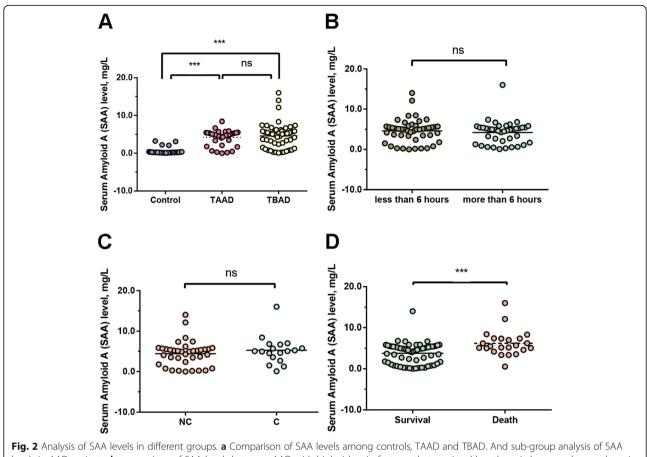
Additionally, we compared SAA levels between control subjects and AAD, angina patients and AAD patients stratified by cardiovascular risk factors (Table 2). SAA levels of AAD patients significantly elevated, compared with the control group and angina patients in each stratified analysis.

Sub-group analysis of AAD patient group was summarized in Table 3, without any significant differences between high-risk pain features that persisted less than 6 hours and more than 6 hours (P = 0.583; Fig. 2b). Additionally, there was no significant differences between and communicating and non-communicating false lumen (C and NC) for these clinical characteristics (P = 0.416; Fig. 2c). Thus, both survival and death cases were enrolled in the study subjects. As a result, WBC count (12.78 ± 4.05

vs $9.96 \pm 4.08 \times 10^9$ /L, P = 0.004), percent of neutrophil (77.88 ± 10.9 vs 70.15 ± 14.72, P = 0.017), lactate dehydrogenase (826.64 ± 1148.14 vs 359 ± 255.33, P = 0.036), FPG (7.90 ± 2.25 vs 6.92 ± 2.46 mmol/L, P = 0.032), plasma D-dimer (6.20 ± 6.91 vs 1.83 ± 3.05 µg/mL, P = 0.006), CRP (138.21 ± 68.8 vs 48.78 ± 50.86 mg/L, P < 0.001) concentrations and SAA (6.214 ± 0.651 vs 3.783 ± 0.314 mg/L, P = 0.001) were significantly increased in death cases compared to those in survival cases (Fig. 2d).

Association of SAA levels with clinical features

The correlation of SAA levels with clinical features was assessed in AAD subjects. As a result, SAA levels were positively correlated with heart rate (R = 0.333, and P = 0.001; Fig. 3a), FPG level (R = 0.654, and P < 0.001; Fig. 3b), WBC count (R = 0.257, and P = 0.042; Fig. 3c), lactate dehydrogenase (R = 0.357, P = 0.006), neutrophil count (R = 0.257, P = 0.025), Cystatin C level (CYSC, R = 0.248, P = 0.046) and plasma CRP concentration (R = 0.442, and P = 0.001; Fig. 3d). Correlations between SAA and other clinical characteristics were shown in Table 4.



levels in AAD patients: **b** comparison of SAA levels between AAD with high-risk pain features that persisted less than six hours and more than six hours, and **c** comparison of SAA levels between AAD with communicating and non-communicating false lumen (C and NC), and **d** comparison of SAA levels between death cases and survival cases. *P < 0.05, **P < 0.01, ***P < 0.001, ns, no significance

 Table 2 Comparison of SAA levels between AAD and control group, AAD and angina group stratified by cardiovascular risk factors

Variables		Control	Angina	AAD	P Value	
		SAA (mg/L)	SAA (mg/L)	SAA (mg/L)	AAD vs Control	AAD vs angina
Age	<60Yr	0.418 ± 0.113	0.661 ± 0.333	4.436 ± 0.308	< 0.001	0.033
	≥60Yr	0.263 ± 0.084	0.632 ± 0.323	6.850 ± 2.535	< 0.001	0.023
Gender	male	0.375 ± 0.093	0.697 ± 0.295	5.542 ± 1.456	< 0.001	0.014
	female	0.332 ± 0.146	0.446 ± 0.244	5.227 ± 0.444	< 0.001	0.032
BMI	< 25	0.300 ± 0.067	0.652 ± 0.301	5.879 ± 1.496	< 0.001	0.017
	≥25	0.652 ± 0.316	0.630 ± 0.364	4.369 ± 0.494	< 0.001	0.016
Hypertension	Yes	0.445 ± 0.164	0.808 ± 0.204	4.892 ± 0.517	< 0.001	0.024
	No	0.319 ± 0.085	0.535 ± 0.342	6.397 ± 2.771	< 0.001	0.046
Smoking history	Yes	0.510 ± 0.215	0.728 ± 0.326	7.698 ± 2.990	< 0.001	0.006
	No	0.297 ± 0.063	0.516 ± 0.278	4.225 ± 0.344	< 0.001	0.030
DM	Yes	0.232 ± 0.093	0.566 ± 0.356	8.157 ± 3.569	< 0.001	0.007
	No	0.385 ± 0.093	0.677 ± 0.311	4.314 ± 0.317	< 0.001	0.047

Note, AAD acute aortic dissection, BMI body mass index, DM diabetes mellitus, SAA serum amyloid A

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Table 3

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Parameters	TAAD	TBAD	<i>P</i> value	less than 6 h	more than 6 h	P value	communicating false lumen	non-communicating false lumen	<i>P</i> value	Death	Survival	P value
WBC, $\times 10^{9}$ /L	11.54 ± 4.5	10.06 ± 3.92	0.198	10.73 ± 4.75	10.75 ± 3.78	0.917	12.1 ± 5.8	10.41 ± 3.74	0.522	12.78 ± 4.05	9.96 ± 4.08	0.004
BA/WBC, %	0.20 ± 0.23	0.28 ± 0.34	0.646	0.24 ± 0.3	0.25 ± 0.30	0.723	0.3 ± 0.28	0.23 ± 0.30	0.231	0.22 ± 0.21	0.26 ± 0.32	0.954
EO/WBC, %	1.37 ± 1.65	0.89 ± 1.51	0.139	0.95 ± 1.37	1.26 ± 1.76	0.409	1.04 ± 1.34	1.13 ± 1.65	0.783	0.66 ± 1.06	1.28 ± 1.72	0.039*
MO/WBC, %	13.29 ± 14.10	8.48 ± 9.03	0.002**	9.29 ± 7.37	11.96 ± 14.7	0.909	15.02 ± 19.28	9.63 ± 9.04	0.648	9.21 ± 9.16	11.26 ± 12.71	0.227
LY/WBC, %	14.87 ± 9.26	15.98 ± 9.1	0.462	14.76 ± 8.5	16.1 ± 9.72	0.596	14.24 ± 8.78	15.77 ± 9.26	0.514	11.62 ± 5.68	16.94 ± 9.79	0.043*
NE/WBC, %	69.92 ± 15.24	74.31 ± 12.97	0.198	74.49 ± 12.41	70.31 ± 15.41	0.275	69.4 ± 18.71	73 ± 12.86	0.638	77.88 ± 10.9	70.15 ± 14.72	0.017*
RBC, $\times 10^{12}$ /L	4.26 ± 0.64	4.52 ± 0.76	0.014*	4.36 ± 0.65	4.43 ± 0.77	0.730	4.32 ± 0.80	4.42 ± 0.69	0.523	4.46 ± 0.82	4.38 ± 0.68	0.582
HGB, g/L	128.63 ± 17.86	135.61 ± 23.03	0.052	134.92 ± 18.44	130.13 ± 23	0.328	126.73 ± 27.9	133.79 ± 18.92	0.497	131.19 ± 25.29	132.85 ± 19.31	0.968
MCH, pg	30.45 ± 3.12	30.49 ± 1.94	0.373	31.06 ± 2.15	29.94 ± 2.75	0.015*	29.89 ± 4.59	30.61 ± 1.73	0.958	29.68 ± 3.41	30.77 ± 2.07	0.104
MCV, fL	91.66 ± 8.00	92.11 ± 5.52	0.696	93.2 ± 5.67	90.74 ± 7.43	0.181	90.10 ± 11.44	92.35 ± 5.01	0.681	89.83 ± 9.28	92.69 ± 5.36	0.168
MCHC, g/L	331.77 ± 11.11	331.12 ± 11.83	0.731	333.17 ± 9.78	329.85 ± 12.65	0.348	330.8 ± 17.09	331.57 ± 9.75	0.445	329.81 ± 11.29	332.04 ± 11.53	0.183
HCT, L/L	0.38 ± 0.06	0.41 ± 0.07	0.083	0.41 ± 0.06	0.39 ± 0.08	0.428	0.38 ± 0.09	0.40 ± 0.06	0.567	0.39 ± 0.08	0.40 ± 0.06	0.802
RDW, %	13.73 ± 2.43	13.25 ± 1.3	0.864	13.38 ± 1.47	13.51 ± 2.17	0.753	14.26 ± 2.64	13.25 ± 1.59	0.253	13.74 ± 1.78	13.33 ± 1.89	0.089
PLT, ×10 ⁹ /L	199.34 ± 67.05	183.12 ± 69	0.196	177.75 ± 64.26	202.15 ± 70.25	0.218	166.07 ± 62.69	196.62 ± 68.56	0.168	182.33 ± 59.6	193.75 ± 71.4	0.597
MPV, fL	9.48 ± 1.13	9.93 ± 1.67	0.158	9.95 ± 1.43	9.52 ± 1.46	0.380	9.51 ± 1.77	9.77 ± 1.39	0.675	9.91 ± 1.06	9.65 ± 1.58	0.480
PCT, L/L	1.02 ± 2.26	0.36 ± 1.11	0.076	0.80 ± 1.99	0.56 ± 1.54	0.264	1.22 ± 2.57	0.55 ± 1.52	0.670	0.87 ± 2.10	0.59 ± 1.62	0.472
PDW, 10GSD	14.51 ± 2.72	14.4 ± 2.90	0.657	14.9 ± 3.00	14.06 ± 2.59	0.260	15.26 ± 2.12	14.27 ± 2.91	0.308	14.51 ± 2.53	14.42 ± 2.92	0.489
TP, g/L	63.93 ± 7.68	63.78 ± 10.34	0.540	64.76 ± 8.76	63.05 ± 9.56	0.410	61.72 ± 9.1	64.38 ± 9.19	0.277	62.27 ± 10.19	64.46 ± 8.77	0.591
PA, mg/dL	15.96 ± 6.67	17.71 ± 6.29	0.231	14.18 ± 6.55	18.88 ± 5.74	0.005**	16.94 ± 6.67	16.78 ± 6.52	0.868	14.75 ± 5.06	17.56 ± 6.83	0.177
ALB, g/L	35.75 ± 6.12	36.84 ± 6.93	0.341	37.03 ± 6.36	35.75 ± 6.74	0.470	34.45 ± 7.3	36.82 ± 6.34	0.185	36.63 ± 7.29	36.23 ± 6.32	0.813
ALP, U/L	106.79 ± 81.75	99.61 ± 43.41	0.583	104.84 ± 59.66	100.79 ± 66.18	0.793	102.42 ± 62.94	102.83 ± 63.19	0.925	103 ± 59.7	102.66 ± 64.29	0.773
ALT, U/L	140.94 ± 407.11	57.24 ± 61.02	0.151	91.74 ± 203.78	98.2 ± 333.43	0.210	119.07 ± 305.66	89.22 ± 274.01	0.408	147.86 ± 454.6	74.7 ± 170.15	0.224
AST, U/L	94 ± 364.89	60.56 ± 101.49	0.249	59.69 ± 101.85	91.36 ± 346.71	0.502	43.86 ± 45.18	83.97 ± 288.02	0.709	170.45 ± 488.11	41.54 ± 56.58	0.123
GGT, U/L	128.38 ± 212.94	↓ 94.02 ± 122.55	0.766	139.77 ± 181.81	83.2 ± 154.87	0.057	127.2 ± 192.56	105.2 ± 164.37	0.942	99.95 ± 158.57	113.35 ± 174.47	0.402
CHE, 1000 U/L	6.02 ± 1.65	5.96 ± 1.71	0.666	5.70 ± 1.54	6.22 ± 1.74	0.247	5.48 ± 1.68	6.13 ± 1.65	0.273	5.83 ± 1.54	6.05 ± 1.72	0.599
TBA, µmol/L	4.76 ± 4.87	4.16 ± 5.86	0.312	5.84 ± 7.11	3.36 ± 2.95	0.170	4.19 ± 3.29	4.54 ± 5.79	0.873	3.15 ± 2.67	4.95 ± 5.97	0.415
DBIL, µmol/L	8.93 ± 9.78	5.77 ± 2.84	0.493	9.7 ± 10.4	5.66 ± 3.36	0.122	8.72 ± 7.26	7.31 ± 8.06	0.463	6.58 ± 3.14	7.95 ± 8.77	0.815
TBIL, µmol/L	17.95 ± 14.17	16.75 ± 8.70	0.568	20.2 ± 14.67	14.81 ± 6.88	0.103	18.37 ± 10.52	17.01 ± 11.68	0.389	16.06 ± 8.68	17.74 ± 12.29	0.742
CK, U/L	186.1 ± 220.13	382 ± 1184.96	0.865	217.87 ± 314.36	360.08 ± 1186.69	0.940	287.88 ± 338.95	295.68 ± 974.92	0.122	562.31 ± 1711.62	203.15 ± 288.52	0.645
CKMB, U/L	17.95 ± 9.85	22.39 ± 22.55	0.343	18.16 ± 8.53	21.32 ± 21.17	0:930	18.23 ± 9.23	20.25 ± 17.83	0.914	26.84 ± 28.62	17.31 ± 8.18	0.607
Urea, mmol/L	7.56 ± 3.99	7.94 ± 8.41	0.127	7.84 ± 6.95	7.7 ± 6.56	0.640	8.29 ± 5.08	7.64 ± 7.08	0.296	7.87 ± 4.51	7.73 ± 7.41	0.646
Uric, mmol/L	273.5 ± 159.03	352.75 ± 214.01	0.280	332.93 ± 185.1	277.47 ± 184.43	0.213	314.57 ± 211.96	302.35 ± 179.47	0.623	376.29 ± 204.76	283.57 ± 175.97	0.177

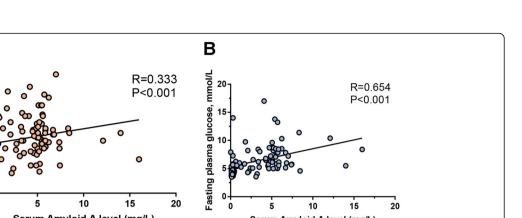
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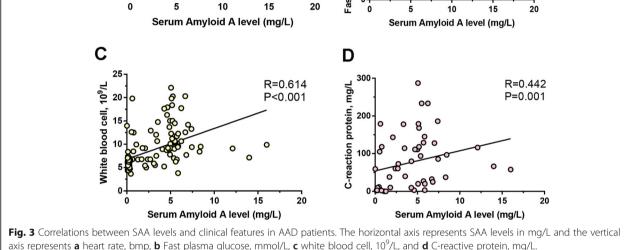
Crea mmol/L 91 CYSC, mg/L 1 CYSC, mmol/L 18 Na, mmol/L 13 K, mmol/L 3.4 CL, mmol/L 10						value	communicating false lumen	false lumen	value	Deall	JULVIVAI	P value
	91.22 ± 43.83	111.17 ± 82.2	0.789	110.42 ± 66.54	88.13 ± 55.74	0.083	99.33 ± 70.69	98.91 ± 59.94	0.614	100.3 ± 22.1	98.58 ± 69.99	0.054
<u> </u>	1.30 ± 0.53	1.42 ± 0.95	066.0	1.38 ± 0.79	1.34 ± 0.73	0.654	1.54 ± 0.84	1.32 ± 0.74	0.412	1.35 ± 0.63	1.36 ± 0.81	0.837
	182.2 ± 35.56	200.05 ± 38.89	0.172	177.55 ± 36.91	198.33 ± 35.83	0.074	181.63 ± 33.57	191.32 ± 38.81	0.763	176.91 ± 31.95	191.81 ± 38.46	0.563
	138.48 ± 4.85	141.52 ± 6.29	0.037*	140.08 ± 7.12	140.15 土 4.48	0.731	139.35 ± 8.79	140.3 ± 4.94	0.293	140.66 ± 6.7	139.91 ± 5.53	0.991
	3.63 ± 0.87	4.27 ± 0.66	0.015*	3.92 ± 0.62	3.84 ± 1.02	0.724	4.18 ± 0.42	3.79 ± 0.91	0.095	3.7 ± 1.14	3.94 ± 0.74	0.274
	103.36 ± 5.92	111.18 ± 7.39	0.000***	104.75 ± 8.74	108 ± 5.91	0.052	105.88 ± 11.64	106.56 ± 6.14	0.875	108.57 ± 8.95	105.71 ± 7.01	0.704
HCO ₃ , mmol/L 24	24.15 ± 2.41	23.98 ± 2.95	0.767	24.37 ± 2.47	23.76 ± 2.92	0.236	23.71 ± 2.39	24.14 ± 2.79	0.341	22.42 ± 2.71	24.69 ± 2.45	0.011*
PH value 6.9	6.93 ± 0.69	6.49 ± 1.08	0.106	7.06 ± 0.6	6.44 ± 1.02	0.021*	6.5 ± 1.04	6.81 ± 0.85	0.915	6.8 ± 0.81	6.73 ± 0.93	0.649
TC, mmol/L 4.	4.19 ± 0.83	4.59 ± 1.17	0.255	4.62 ± 1.32	4.3 ± 0.84	0.589	4.92 ± 1.47	4.32 ± 0.93	0.361	4.51 ± 1.13	4.39 ± 1.03	0.333
	1.34 ± 0.56	1.34 ± 0.67	0.834	1.24 ± 0.47	1.41 ± 0.71	0.677	1.5 ± 0.57	1.31 ± 0.63	0.283	1.23 ± 0.44	1.39 ± 0.69	0.664
ApoA1, mmol/ 1. L	1.14 ± 0.18	1.21 ± 0.22	0.469	1.18 ± 0.17	1.18 ± 0.22	0.938	1.14 ± 0.31	1.19 ± 0.18	0.776	1.24 ± 0.16	1.14 ± 0.22	0.218
ApoB, mmol/L 0.	0.85 ± 0.22	0.76 ± 0.13	0.674	0.75 ± 0.22	0.83 ± 0.13	0.277	0.85 ± 0.13	0.79 ± 0.18	0.506	0.76 ± 0.2	0.82 ± 0.16	0.425
LDL-C, mmol/L 2.	2.58 ± 0.78	2.87 ± 1.06	0.246	2.87 ± 1.04	2.68 ± 0.92	0.620	3.23 ± 1.00	2.66 ± 0.94	0.133	2.65 ± 1.05	2.81 ± 0.93	0.899
HDL-C, mmol/ 1.(L	1.08 ± 0.34	1.22 ± 0.31	0.139	1.18 ± 0.25	1.16 ± 0.37	0.649	1.09 ± 0.38	1.19 ± 0.31	0.570	1.25 ± 0.25	1.13 ± 0.35	0.150
LDH, U/L 45	451.88 ± 779.15	487.03 ± 489.75	0.138	505.85 ± 538.16	444.28 ± 694.87	0.171	410.67 ± 242.15	487.85 ± 692.2	0.645	826.64 ± 1148.14	359 ± 255.33	0.036*
INR 1.	1.14 ± 0.22	1.08 ± 0.11	0.446	1.13 ± 0.21	1.09 ± 0.13	0.172	1.09 ± 0.12	1.11 ± 0.18	0.788	1.18 ± 0.26	1.08 ± 0.10	0.152
PT, s 13	13.11 ± 2.41	13.28 ± 1.33	0.149	13.51 ± 2.27	12.96 ± 1.51	0.119	12.9 ± 1.57	13.28 ± 1.96	0.634	13.91 ± 2.65	12.89 ± 1.36	0.131
PTA, % 85	85.87 ± 17.42	89.28 ± 12.04	0.615	85.62 ± 16.02	89.38 ± 13.59	0.198	88.54 ± 13.99	87.53 ± 15.01	0.974	83.4 ± 19.2	89.61 ± 12.04	0.279
АРТТ, s 32	32.28 ± 5.28	35.79 ± 5.86	0.008**	34.39 ± 6.11	34.04 ± 5.69	0.948	34.42 ± 6.43	34.14 ± 5.75	0.834	35.4 ± 6.4	33.67 ± 5.57	0.339
FPG, mmol/L 7.	7.61 ± 2.84	6.92 ± 2.05	0.355	7.2 ± 1.92	7.22 ± 2.81	0.281	6.38 ± 1.63	7.45 ± 2.57	0.069	7.90 ± 2.25	6.92 ± 2.46	0.032*
CRP, mg/L 95	95.73 ± 60.92	70.53 ± 76.14	0.115	90.4 ± 83.01	71.92 ± 61.85	0.597	79.43 ± 84.49	79.81 ± 69.09	0.889	138.21 ± 68.8	48.78 ± 50.86	0.000***
D-Dimer, µg/ 2.(mL	2.02 ± 3.57	4.10 ± 5.73	0.187	3.25 ± 4.94	3.09 ± 5.02	0.234	1.81 ± 1.69	3.49 ± 5.42	0.759	6.20 ± 6.91	1.83 ± 3.05	0.006**
SAA, mg/L 4.	4.22 ± 0.33	4.58 ± 0.48	0.595	4.61 ± 0.42	4.21 ± 0.45	0.582	5.28 ± 0.81	4.44 ± 0.51	0.416	6.21 ± 0.65	3.78 ± 0.31	0.001***
Note, BMI Body Mass Index, HR heart rate, WBC white blood cells, RBC hemoglobin, MCHC mean corpuscular hemoglobin concentration, MCI cell volume distribution width, APT activated partial thromboplastin t CRP C-reactive protein, ALB albumin, ALP alkaline phosphatase, ALT ala transpeptidase, TBA total bile acid, TP total protein, PA pre-albumin, LL apoliporotein B, FPG fasting plasma quccese, HDL-C high-density libo,	s Index, <i>HR</i> hear mean corpusculi tion width, <i>APTT</i> in, <i>ALB</i> albumin, total bile acid, <i>T</i> , G fasting plasma	t rate, <i>WBC</i> white t ar hemoglobin con activated partial th <i>ALP</i> alkaline phosy <i>P</i> total protein, <i>PA</i> o glucose, <i>HDL</i> -C hi	olood cells, / ncentration, / nromboplast phatase, ALT pre-albumin gh-density II	<i>RBC</i> red blood cells. <i>MCV</i> mean corpusc in time, <i>PT</i> prothro ² alanine aminotran ¹ , <i>LDH</i> lactate dehyv i poprotein-choleste	, <i>HGB</i> hemoglobin, <i>H</i> ular Volume, <i>MPV</i> m mbin time, <i>PTA</i> prot sferase, <i>AST</i> aspartat drogenase, <i>Na</i> serum rrol, <i>TC</i> total choleste	2LT Platele hean platel thrombin ε te aminotr n sodium, erol, TG tri	t, BA basophil, EO (et volume, HCT hel activity, INR interna ansferase, DBIL dirr CL serum chlorine, glycerides, LDL-C Ic	Note, <i>BM</i> Body Mass Index, <i>HR</i> heart rate, <i>WBC</i> white blood cells, <i>HBC</i> hemoglobin, <i>PLT</i> Platelet, <i>BA</i> basophil, <i>EO</i> eosinophil, <i>LY</i> lymphocyte, <i>MD</i> monocyte, <i>NE</i> neutrophil, <i>MCH</i> mean corpuscular hemoglobin, <i>MCH</i> can be an evolved of the mean corpuscular hemoglobin and the mean corpuscular hemoglobin and the mean corpuscular hemoglobin. <i>MCHC</i> mean corpuscular hemoglobin and the mean corpuscular hemoglobin and the mean corpuscular hemoglobin. <i>MCH</i> mean corpuscular hemoglobin and the mean corpuscular hemoglobin and the mean corpuscular hemoglobin. <i>MCHC</i> mean corpuscular hemoglobin concentration, <i>MCV</i> mean corpuscular volume, <i>MPV</i> mean platelet volume, <i>HCT</i> hematocrit, <i>PCT</i> platelet hematocrit, <i>PDW</i> platelet distribution width, <i>APTT</i> activated partial thrombophastin time, <i>PT</i> prothrombin activity. <i>INR</i> international normalized ratio, <i>CK</i> creatine kinase. <i>CMB</i> creatine kinase isoenzyme MB, calvate protein, <i>ALP</i> alanine annotransferase, <i>AST</i> apartate aminotransferase, <i>DBIL</i> direct dilirubin, <i>CHE</i> cholinesterase, <i>GST</i> gamma glutamyl cranscipated at the state activates and continue and the corpuscient. <i>PER</i> the advantation and the corpuscient and the advantation and the corpuscient activity and the corpuscient and the corpuscient and the corpuscient and the cor	e, MO mo matocrit, . CK creatin CK creatin C lirubin, Cl ystatin C,	nocyte, NE neutroph PDW platelet distribu e kinase, <i>CKMB</i> creat <i>H</i> E cholinesterase, <i>G</i> C K serum kalium, <i>Apo</i> .	iil, <i>MCH</i> mean corp ttion width, <i>RDW</i> π ine kinase isoenzyr 37 gamma glutamy A1 apolipoprotein .	uscular td blood ne MB, l A-1, <i>ApoB</i>

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Heart rate, bmp

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Diagnostic performance of SAA, CRP and their combination for AAD

ROC analysis was conducted to determine the cut-off value of SAA level for the evaluation of TAAD, TBAD, and AAD. The AUCs for TAAD, TBAD, and AAD alone were 0.939, 0.937 and 0.942 with optimal cut-off points of 0.427 mg/L, 0.462 mg/L, and 0.427 mg/L, respectively, associated with sensitivity of 91.9, 88.1 and 90.8%, respectively, and specificity of 93.7, 93.7 and 93.7%, respectively (Table 5 and Fig. 4). Furthermore, the diagnostic performance of SAA, CRP and their combination to discriminate AAD according to ROC analysis, were also shown in Table 5. The AUC (0.977) of combined model (SAA + CRP) was significantly greater than that of SAA (AUC, 0.942), but not combined model (SAA + D-dimer, AUC = 0.900). Furthermore, SAA +CRP yielded sensitivity of 94.4% and specificity of 93.8%. ROC analysis was subsequently conducted to examine the cut-off values of SAA and other prognostic biomarkers for evaluation of in-hospital mortality (Table 5). The cut-off values were 4.998 mg/L for SAA, 82.55 mg/L for CRP, 1.945 µg/mL for D-dimer, and 72.976% for neutrophil to white blood cell ratio (NE%) with their higher sensitivity and specificity, respectively. The AUCs were 0.732 for SAA, 0.826 for CRP, 0.715 for D-dimer and 0.678 for NE%, respectively. In the case of SAA ≥4.998 mg/L, the sensitivity and specificity for inhospital mortality prediction were 73.9 and 62.5% (95% CI 0.61 to 0.85; P = 0.001) for AAD patients. When CRP was ≥82.55 mg/L, the sensitivity and specificity were 0.789 and 0.771 (95% CI 0.709 to 0.942; P < 0.001). When D-dimer was ≥1.945 µg/mL, the sensitivity and specificity in predicting in-hospital death were 0.700 and 0.739 (95% CI 0.562 to 0.869; P = 0.006). When NE% was ≥72.976%, the sensitivity and specificity in predicting in-hospital death were 0.857 and 0.491 (95% CI 0.549 to 0.808; P = 0.017).

The stratum of SAA according to gender and clinical biomarkers

The upper and lower strata of SAA levels were defined according to the cut-off values for the ROC, which in the present study were 0.427 mg/L for AAD patients. The prevalence of upper-stratum SAA levels was (78/87) 89.7% in AAD patients, 87.7% (57/65) in the male AAD group, and 95.5% (21/22) in the female AAD group. There were no significant gender differences of prevalence of upper stratum SAA levels (P = 0.438). Moreover, a multiple logistic regression analysis revealed that the plasma CRP, WBC, neutrophil, and HDL-C levels were positively associated with the upper stratum of SAA levels when compared with the lower stratum (OR = 1.05, 1.74, 1.97, 1.06, respectively, all *P*-values< 0.05). Furthermore, plasma

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Characterization	Indices	R	P value
Baseline Data			
	Age, Yr	- 0.045	0.584
	BMI	0.126	0.123
	HR, bmp	0.333	< 0.001***
Blood routine			
	WBC, ×10 ⁹ /L	0.614	< 0.001***
	RBC, ×10 ¹² /L	-0.111	0.354
	HGB, g/L	-0.395	< 0.001***
	PLT, ×10 ⁹ /L	0.091	0.285
	BA, ×10 ⁹ /L	-0.101	0.385
	EO, ×10 ⁹ /L	-0.243	0.035*
	LY, ×10 ⁹ /L	-0.168	0.147
	MO, ×10 ⁹ /L	0.085	0.468
	NE, ×10 ⁹ /L	0.257	0.025*
	MCH, pg	-0.065	0.574
	MCHC, g/L	-0.202	0.080
	MCV, fL	0.015	0.898
	MPV, fL	-0.098	0.405
	HCT, L/L	-0.198	0.148
	PCT, L/L	-0.011	0.930
	PDW, 10GSD	0.158	0.189
	RDW, %	0.171	0.166
Blood coagulation	function		
	APTT, s	-0.241	0.051
	PT, s	0.03	0.811
	PTA, %	-0.258	0.037*
	INR	0.251	0.042*
	D-Dimer, ug/ml	0.206	0.097
Cardiovascular injur	ry-related parameters		
	CK, U/L	-0.039	0.756
	CKMB, U/L	0.213	0.198
Inflammatory respo	onse		
	CRP, mg/L	0.442	0.001***
Liver function			
	ALB, g/L	-0.237	0.040*
	ALP, U/L	0.124	0.330
	ALT, U/L	0.121	0.300
	AST, U/L	0.163	0.166
	DBIL, umol/L	-0.098	0.553
	TBIL, umol/L	0.012	0.922
	CHE, 1000 U/L	-0.258	0.053
	GGT, U/L	0.028	0.811
	TBA, umol/L	0.138	0.310

Table 4 The correlations between SAA levels and clinical	
features in AAD patients	

Table 4 The correlations between SAA levels and clinical features in AAD patients (Continued)

Characterization	Indices	R	P value
	TP, g/L	-0.177	0.128
	PA, mg/dL	-0.402	0.002**
	LDH, U/L	0.357	0.006**
Re	enal function and serum	electrolyte	
	Na, mmol/L	-0.141	0.224
	C1q, mmol/L	0.026	0.883
	HCO ₃ , mmol/L	-0.282	0.020*
	CL, mmol/L	0.175	0.274
	Crea, mmol/L	-0.116	0.469
	CYSC, mg/L	0.248	0.046*
	PH value	0.113	0.523
	K, mmol/L	0.074	0.644
	Urea, mmol/L	-0.049	0.672
	Uric, mmol/L	0.182	0.336
	Serum lipid prof	ile	
	ApoA1, mmol/L	0.117	0.623
	ApoB, mmol/L	-0.117	0.623
	FBG, mmol/L	0.654	< 0.001***
	HDL-C, mmol/L	0.195	0.175
	TC, mmol/L	0.063	0.678
	TG, mmol/L	-0.223	0.120
	LDL-C, mmol/L	0.026	0.859

Note bloc lymphocyte, MO monocyte, NE neutrophil, MCH mean corpuscular hemoglobin, MCHC mean corpuscular hemoglobin concentration, MCV mean corpuscular Volume, MPV mean platelet volume, HCT hematocrit, PCT platelet hematocrit, PDW platelet distribution width, RDW red blood cell volume distribution width, APTT activated partial thromboplastin time, PT prothrombin time, PTA prothrombin activity, INR international normalized ratio, CK creatine kinase, CKMB creatine kinase isoenzyme MB, CRP C-reactive protein, ALB albumin, ALP alkaline phosphatase, ALT alanine aminotransferase, AST aspartate aminotransferase, DBIL direct dilirubin, TBIL total bilirubin, CHE cholinesterase, GGT gamma glutamyl transpeptidase, TBA total bile acid, TP total protein, PA pre-albumin, LDH lactate dehydrogenase, Na serum sodium, CL serum chlorine, Crea Creatinine, CYSC Cystatin C, K serum kalium, ApoA1 apolipoprotein A-1, ApoB apolipoprotein B, FPG fasting plasma glucose, HDL-C high-density lipoprotein-cholesterol, TC total cholesterol, TG triglycerides, LDL-C low-density lipoprotein-cholesterol

P* < 0.05; *P* < 0.01; ****P* < 0.001

pre-albumin and lymphocyte levels showed negative associations with the upper stratum of the SAA levels when compared with the lower stratum (Table 6).

Univariate and multivariate logistic regression analyses for in-hospital mortality

The following ten variables were indicated to be related with hospital short-term mortality, WBC, eosinophils, lymphocyte, neutrophil, HCO₃ level, lactate dehydrogenase, FPG, CRP, D-Dimer, and SAA concentration (P value

Table 5 Diagnostic value of SAA and its combinations with CRP and D-Dimer for AAD patients

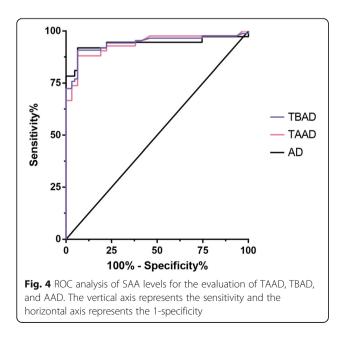
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	AUC	95%CI	P-value	cut-off	Sensitivity (%)	Specificity (%)
AAD	0.942	0.902-0.981	< 0.001	0.427	0.908	0.937
TAAD	0.939	0.873-1.004	< 0.001	0.427	0.919	0.937
TBAD	0.937	0.883-0.990	< 0.001	0.462	0.881	0.937
Combination						
SAA + CRP	0.977	0.955-0.999	< 0.001	-	0.944	0.938
SAA + D-Dimer	0.900	0.838-0.963	< 0.001	-	0.848	0.938
Death						
SAA (death v.s. survival)	0.732	0.612-0.853	0.001	4.998	0.739	0.625
CRP (death v.s. survival)	0.826	0.709-0.942	< 0.001	82.55	0.789	0.771
D-Dimer (death v.s. survival)	0.715	0.562-0.869	0.006	1.945	0.700	0.739
NE% (death v.s. survival)	0.678	0.549-0.808	0.017	72.976	0.857	0.491

Note, AAD acute aortic dissection, TAAD type A aortic dissection, TBAD type B aortic dissection, SAA serum amyloid A, CRP C reactive protein, AUC area under the curve, NE% neutrophil to white blood cell ratio

was approximately 0.05 in univariate analysis) (Table 3). Therefore, the ten variables were incorporated into multivariate logistic regression analysis, revealing that SAA (OR = 1.25; 95%CI: 1.07–1.47; P = 0.005), CRP (OR = 1.03; 95%CI = 1.01–1.10; P < 0.001), WBC count (OR = 1.17; 95%CI: 1.03–1.33; P = 0.015) and eosinophils (OR = 1.06; 95%CI: 1.00–1.11; P = 0.041) were significantly related to hospital short-term mortality (Table 7) adjusted as confounding factors. Nevertheless, in-hospital death was not significantly associated with any of variables enrolled in this research.

Discussion

AAD is a severe aortic disorder associated with inflammation [8, 15, 24]. The inflammatory responses play a



critical role in initiating further necrosis and apoptosis of smooth muscle cells as well as degeneration of elastic tissue, contributing to aorta rupture. SAA, an acutephase protein, reflects the status of inflammation [25]. The early diagnostic value of SAA as a serum biomarker has been increasingly recognized in other diseases. To our knowledge, it was the first study to focus on the association between SAA and AAD, which demonstrated SAA as a potentially useful biomarker for AAD detection.

In this study, SAA levels were significantly higher in AAD patients than healthy controls. However, SAA levels were not significantly different between TAAD and TBAD, indicating that SAA was not associated with the specific subtype of AAD. Moreover, SAA was correlated with several laboratory examination outcomes, which were necessary to evaluate the status and to detect complications. According to the correlation analysis, SAA levels were significantly correlated with inflammation-related parameters, including WBC count, neutrophil count and CRP. CRP, an acute phase reactant, has been widely used as an independent predictor of poor prognosis in AAD patients, which is also related to the in-hospital mortality in AAD patients [12]. Additionally, SAA was also correlated with prothrombin activity and international normalized ratio in this study. AAD is widely known to be influenced by activated coagulation and fibrinolytic systems [26, 27]. Among these, Ddimer shows its important diagnostic and prognostic value in AAD patients [9]. However, SAA is not correlated with myocardial biomarkers in this study. The closely correlation between SAA and admission hematological parameters indicates the potential clinical significance of SAA in the clinical process of AAD.

Additionally, based on the elevated SAA levels in AAD patients, the diagnostic value of SAA alone and its combination with CRP as well as D-dimer was assessed for AAD detection. As a result, SAA was able to provide

Table 6 Associations of parameters with the upper stratum of the SAA levels when compared with the lower stratum

	SAA lower	SAA upper	P
		OR (95%CI)	value
Age, Yr	1 Reference	0.98 (0.92; 1.04)	0.534
Hypertension, n	1 Reference	3.12 (0.76; 12.84)	0.114
Smoker, n	1 Reference	1.64 (0.31; 8.54)	0.558
D-Dimer, µg/mL	1 Reference	1.72 (0.79; 3.75)	0.175
CRP, mg/L	1 Reference	1.05 (1.00; 1.09)	0.039
CAD, n	1 Reference	1.49 (0.17; 13.14)	0.719
Type01, n	1 Reference	1.11 (0.28; 4.50)	0.879
WBC, ×10 ⁹ /L	1 Reference	1.74 (1.18; 2.56)	0.005
RBC, ×10 ¹² /L	1 Reference	0.55 (0.18; 1.66)	0.287
HGB, g/L	1 Reference	0.96 (0.92; 1.00)	0.071
PLT, ×10 ⁹ /L	1 Reference	1.00 (0.99; 1.01)	0.974
FPG, mmol/L	1 Reference	1.17 (0.80; 1.70)	0.419
BA, ×10 ⁹ /L	1 Reference	0.18 (0.03; 1.20)	0.076
EO, ×10 ⁹ /L	1 Reference	0.79 (0.01; 114.81)	0.925
LY, ×10 ⁹ /L	1 Reference	0.85 (0.77; 0.93)	0.000
MO, ×10 ⁹ /L	1 Reference	1.02 (0.94; 1.10)	0.708
NE, ×10 ⁹ /L	1 Reference	1.97 (1.24; 3.11)	0.004
CKMB, u/L	1 Reference	1.01 (0.89; 1.15)	0.836
MCH, pg	1 Reference	0.83 (0.56; 1.21)	0.332
MCHC, g/L	1 Reference	0.97 (0.91; 1.04)	0.432
MCV, fL	1 Reference	0.95 (0.84; 1.08)	0.459
MPV, fL	1 Reference	0.77 (0.47; 1.27)	0.306
PCT, L/L	1 Reference	0.51 (0.01; 3.93)	0.906
PDW, 10GSD	1 Reference	1.03 (0.80; 1.32)	0.829
RDW, %	1 Reference	2.43 (0.99; 5.97)	0.054
_DH, U/L	1 Reference	1.01 (1.00; 1.02)	0.063
ALB, g/L	1 Reference	0.94 (0.84; 1.05)	0.304
ALP, U/L	1 Reference	1.01 (0.99; 1.03)	0.383
ALT, U/L	1 Reference	1.01 (0.99; 1.02)	0.452
AST, U/L	1 Reference	1.00 (0.99; 1.02)	0.567
GGT, U/L	1 Reference	1.00 (1.00; 1.01)	0.529
DBIL, µmol/L	1 Reference	1.77 (0.71; 4.43)	0.223
TBIL, μmol/L	1 Reference	1.02 (0.94; 1.10)	0.617
PA, mg/dL	1 Reference	0.83 (0.71; 0.96)	0.010
TP, g/L	1 Reference	0.97 (0.89; 1.05)	0.396
TBA, µmol/L	1 Reference	0.97 (0.86; 1.10)	0.668
K, mmol/L	1 Reference	3.31 (0.91; 12.06)	0.070
CL, mmol/L	1 Reference	0.98 (0.81; 1.17)	0.801
Na, mmol/L	1 Reference	0.95 (0.86; 1.05)	0.313
HDL-C, mmol/L	1 Reference	1.06 (1.01; 1.10)	0.014
LDL-C, mmol/L	1 Reference	1.27 (0.52; 3.11)	0.605
TC, mmol/L	1 Reference	1.40 (0.56; 3.52)	0.471

 Table 6 Associations of parameters with the upper stratum of the SAA levels when compared with the lower stratum (Continued)

()						
	SAA lower SAA upper		Р			
		OR (95%CI)	value			
TG, mmol/L	1 Reference	0.50 (0.15; 1.60)	0.240			
Urea, mmol/L	1 Reference	1.17 (0.87; 1.55)	0.295			
Uric, mmol/L	1 Reference	1.01 (0.98; 1.05)	0.389			
Crea, mmol/L	1 Reference	1.09 (0.97; 1.23)	0.133			
C1q, mmol/L	1 Reference	1.03 (0.98; 1.08)	0.264			
HCO ₃ , mmol/L	1 Reference	0.77 (0.57; 1.04)	0.086			
PH value	1 Reference	0.08 (0; 921.56)	0.596			
APTT, s	1 Reference	0.90 (0.80; 1.01)	0.079			
PT, s	1 Reference	1.08 (0.71; 1.64)	0.724			
PTA, s	1 Reference	0.96 (0.90; 1.02)	0.164			

Note, HR heart rate, WBC white blood cells, RBC red blood cells, HGB hemoglobin, PLT Platelet, BA basophil, EO eosinophil, LY lymphocyte, MO monocyte, NE neutrophil, MCH mean corpuscular hemoglobin, MCHC mean corpuscular hemoglobin concentration, MCV mean corpuscular Volume, MPV mean platelet volume, HCT hematocrit, PCT platelet hematocrit, PDW platelet distribution width, RDW red blood cell volume distribution width, APTT activated partial thromboplastin time, PT prothrombin time, PTA prothrombin activity, CKMB creatine kinase isoenzyme MB, CRP C-reactive protein, ALB albumin, ALP alkaline phosphatase, ALT alanine aminotransferase, AST aspartate aminotransferase, DBIL direct dilirubin, TBIL total bilirubin, CHE cholinesterase, GGT gamma glutamyl transpeptidase, TBA total bile acid, TP total protein, PA pre-albumin, LDH lactate dehydrogenase, Na serum sodium, CL serum chlorine, Crea Creatinine, CYSC Cystatin C, K serum kalium, ApoA1 apolipoprotein A-1, ApoB apolipoprotein B, FPG fasting plasma glucose, HDL-C high-density lipoprotein-cholesterol, TC total cholesterol, TG triglycerides, LDL-C low-density lipoprotein-cholesterol

 Table 7
 Univariate and multivariate logistic regression analyses

 for in-hospital mortality
 Image: Second Second

	Model 1		Model 2	
	OR (95%CI)	P-value	OR (95%CI)	P-value
Type of AAD	0.47 (0.16; 1.32)	0.152	0.48 (0.17; 1.36)	0.167
HR, bmp	1.01 (0.97; 1.05)	0.584	1.01 (0.97; 1.05)	0.536
Hypertension, %	1.22 (0.41; 3.64)	0.724	1.13 (0.37; 3.40)	0.832
Smoker, %	0.63 (0.20; 1.96)	0.421	0.63 (0.20; 2.00)	0.432
D-Dimer, µg/mL	1.19 (0.98; 1.43)	0.074	1.20 (0.99; 1.46)	0.065
CRP, mg/L	1.02 (1.01; 1.03)	0.000	1.03 (1.01; 1.04)	0.000
WBC, ×10 ⁹ /L	1.17 (1.03; 1.32)	0.013	1.17 (1.03; 1.33)	0.015
EO, ×10 ⁹ /L	0.68 (0.40; 1.15)	0.148	0.68 (0.40; 1.16)	0.161
LY, ×10 ⁹ /L	0.92 (0.85; 0.99)	0.030	0.92 (0.85; 0.99)	0.031
NE, ×10 ⁹ /L	1.05 (1.00; 1.11)	0.040	1.06 (1.00; 1.11)	0.041
HCO ₃ , mmol/L	0.69 (0.53; 0.89)	0.004	0.67 (0.51; 0.88)	0.004
LDH, U/L	1.00 (0.99; 1.00)	0.097	1.00 (0.99; 1.00)	0.151
FPG, mmol/L	1.17 (0.95; 1.44)	0.133	1.16 (0.95; 1.43)	0.149
SAA, mg/L	1.22 (1.05; 1.42)	0.011	1.25 (1.07; 1.47)	0.005

Note, AAD acute aortic dissection, HR heart rate, CRP C reactive protein, WBC white blood cell, EO eosinophil, LY lymphocyte, NE neutrophil, LDH lactate dehydrogenase, FPG fasting plasma glucose, SAA serum amyloid A, Cl confidence interval, Model 1 no adjustments, Model 2 adjusted for age, gender, BMI

high sensitivity and specificity for TAAD diagnosis by referring to the ROC curves. However, our data revealed high specificity but relatively weak sensitivity of SAA when used alone for TBAD. Interestingly, compared with SAA alone, CRP-SAA combination led to improved diagnostic accuracy along with the increased sensitivity as well as higher AUC, while D-Dimer-SAA combination failed to show the significant elevation of AUC. Taken together, these findings indicated that SAA may be a potential non-invasive predictor of AAD, especially for AAD patients with normal CRP value.

Furthermore, SAA has been identified as an apolipoprotein of high-density lipoprotein cholesterol (HDL-C). Previous studies have indicated that HDL-C is negatively associated with cardiovascular risks, and low HDL-C levels may represent a high in-hospital mortality in AAD patients [28]. SAA-HDL complex in blood has been demonstrated to have a high affinity for macrophages [14] and to further eliminate them from the blood in patients with sarcoidosis [29]. Thus, increased SAA levels in AAD patients may be caused in response to the systemic inflammation, showing its potential value in predicting AAD characteristics. Additionally, on the one hand, the progressive degradation of extracellular matrix is also considered as a critical pathological feature in the pathogenesis and progression of AAD [30]. On the other hand, SAA has been reported to elicit several kinds of matrix metalloproteinases induction in vascular smooth muscle cells in vitro [15, 31, 32]. Thus, elevated SAA level in AAD is of great significance for the evaluation of AAD development. In this study, AAD patients with low SAA levels are more likely to control disease status than those with high SAA levels. Collectively, these outcomes suggest the potential prognostic value of SAA in the clinical procedure of AAD.

In this study, we observe that high SAA levels are associated with high in-hospital mortality. It has been previously reported that high SAA levels were associated with mortality in other diseases. For example, higher SAA concentration was associated with all-cause mortality in patients with end-stage renal disease [33]. Additionally, as an apolipoprotein of HDL, SAA can transforms HDL from a protective lipoprotein into a pro-atherosclerotic lipoprotein, which contributed to the substantially worsened cardiovascular outcome, which have been reported previously [34]. Our results suggest that SAA levels in AAA patients may be a critical marker of all-cause mortality in AAD. Additionally, it has been reported previously that CRP, D-Dimer [12] and NE% [35] are important risk factors for in-hospital mortality in AAD. In our study, we observed that CRP with AUC of 0.826, the highest sensitivity and specificity, is the best prognostic indicator of the mortality, and the AUC of SAA in predicting in-hospital mortality (AUC = 0.732) is better than that of D-Dimer (AUC = 0.715) and NE% (AUC = 0.678). Therefore, our result revealed the potential value of SAA as a prognostic biomarker of the in-hospital mortality in AAD.

However, there are some limitations in this study. To begin with, the population size was relatively small. Secondly, the specific study population was collected from only two clinical centers. Additionally, the detailed backgrounds of healthy controls were not clearly defined due to the incomplete recording of their previous medical history. Furthermore, a combination of tissue expression and mechanism study would definitely give rise to more detailed information concerning the role of SAA in AAD.

Conclusions

Collectively, our findings demonstrated that SAA levels were significantly enhanced in both TAAD as well as TBAD patients. Then, we suggested the close correlations of SAA with inflammatory parameters and coagulation-related parameters in AAD patients at the level of serology. Furthermore, we provided the proof that SAA may be a useful biomarker for AAD identification, and SAA > 5.0 mg/L is considered as important risk factors, which are independently related to AAD in-hospital mortality. Nevertheless, prospective, large-scale clinical studies are warranted to further validate these outcomes, which would definitely contribute to the prognostic evaluation of AAD.

Abbreviations

1H-ADBB/CMU: The first hospital of CMU aortic dissection blood sample biobank; AAD: Acute aortic dissection; ALT: Alanine aminotransferase; AST: Glutamic oxalacetic transaminase; AUC: The area under the curve; BP: Blood pressure; C and NC: Communicating and non-communicating false lumen; CMU: China Medical University; CRP: C-reactive protein; CT: Computed tomography; Ctrl: Control group; CVD: Cardiovascular diseases; CYSC: Cystatin C level; DM: Diabetes mellitus; FPG: Fasting plasma glucose; HDL-C: High-density lipoprotein cholesterol; LDL-C: Low-density lipid cholesterol; MRI: Magnetic resonance imaging; NE%: Neutrophil to white blood cell ratio; PBMCs: Peripheral blood mononuclear cells; ROC: Receiveroperating characteristic; SAA: Serum amyloid a; SJH-ADBB/CMU: Shengjing hospital of CMU aortic dissection blood sample biobank; TAAD: Type a aortic dissection; TBAD: Type B aortic dissection; TP: Total protein; WBC: White blood cell

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Authors' contributions

YCH and JX performed the experimental work. YSH and YCH analyzed the results and composed the manuscript. SYW, CCM, CJ and YCH participated in the collection of clinical data. JZ, and YSH designed the study and participated in editing the manuscript. All authors read and approved the final manuscript.

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Availability of data and materials

The data used in this study are available from the corresponding author if needed.

Ethics approval and consent to participate

Participants gave written informed consent before the study, and the study protocol was approved by the Ethics Committee of China Medical University (CMU), in accordance with the Declaration of Helsinki.

Consent for publication

All the participants provided written informed consent for the publication of the results of this study.

Competing interests

The authors declare that they have no conflict interests.

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