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			Patients without Septic Associated Acute Kidney Injury and with Septic Associated Acute Kidney Injury Requiring Continuous Renal Replacement Therapy					
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Background: Material/Methods: Results:			Acute kidney injury (AKI) is one of the most common organ failures. An early diagnosis of AKI using specif- ic biomarkers is essential for effective treatment. This study determined the serum concentrations of select- ed amino acids and amines using targeted liquid chromatography coupled with tandem mass spectrometry (LC-MS/MS) in patients with AKI during sepsis and septic shock treated in the Intensive Care Unit (ICU). A sample of 41 patients was divided into 2 groups: (1) patients with sepsis and septic shock along required continuous renal replacement therapy (CRRT) due to AKI (n=13), and (2) patients with sepsis and septic shock but without AKI (n=28). LC-MS/MS was used to measure a serum concentration of 6 amino acids and amines: arginine, ornithine, asymmetric dimethylarginine (ADMA), symmetric dimethylarginine (SDMA), dimethylamine (DMA), and citrulline. There was a statistically significantly higher median DMA level in AKI patients compared to those without AKI (8.1 vs 5.2 umol/L; <i>P</i> =0.022). The results for the remaining molecules showed no significant differences (<i>P</i> >0.05).					
Conclusions:			Patients with DMA \geq 14.95 umol/L (n=5; 100%) and treated with CRRT presented DMA level below the cut-off point (n=7; 20%). Subjects with creatinine levels \geq 1.19 mg/dL (n=11; 28%) and treated with CRRT presented creatinine levels below the cut-off point (n=1; 3%). In patients with sepsis, increased serum levels of DMA were significantly associated with AKI requiring CRRT. It remains unclear whether increased DMA concentrations are secondary to sepsis-induced AKI or are a cause.					
	I	Keywords:	Acute Kidney Injury • Biomarkers • Dimethylamine • Intensive Care Units • Renal Insufficiency • Sepsis					
	Ful	ll-text PDF:	https://www.medscimonit.com/abstract/index/idArt/937784					

Measurement of Serum Levels of 5 Amino

Chromatography-Tandem Mass Spectrometry in

Acids and Dimethylamine Using Liquid



e937784-1

Background

Acute kidney injury (AKI) is one of the most common organ failures in patients hospitalized in intensive care units (ICUs) [1]. The incidence varies from 5% to 50% depending on the literature and is associated with a mortality rate of up to 50% [2-4]. Additionally, 20-50% of patients who survive later develop chronic kidney disease (CKD), and about 5% develop end-stage renal disease [5-9]. Renal tubular ischemia can be caused by sepsis, cardiogenic shock, low cardiac output, drug toxicity, surgery, and hypovolemia [10].

An early diagnosis of AKI is essential for effective treatment. According to the Kidney Disease: Improving Global Outcomes (KDIGO) guidelines, the 2 basic diagnostic criteria in the diagnosis of AKI are the dynamics of the increase in serum creatinine concentration and urine output [11,12]. CRRT is the method of choice for AKI treatment in patients treated in the ICU. Due to the high hemodynamic instability of patients, it has an advantage over IHD. The main indications for starting CRRT are life-threating electrolyte, fluid, and acid-based changes [13].

In kidney injuries, when the glomerular filtration rate (GFR) decreases, the half-life of creatinine increases from 4 to 24-72 h, meaning that change in creatinine level is delayed by up to 24-36 h [14,15]. New biomarkers, such as neutrophil-gelatinase associated lipocalin (NGAL), cystatin C (CC), kidney injury molecule-1 (KIM-1), interleukin-18 (IL18), liver-type fatty acid-binding protein (L-FABP), N-acetyl- β -D-glucosaminidase (NAG), α -glutathione transferase (alpha-GST), metalloproteinase tissue inhibitor type 2 (TIMP-2), and insulin-like growth factor binding protein 7 (IGFBP-7), might be more beneficial for the early detection of acute kidney injuries [16-18].

In the diagnosis of AKI in the course of sepsis and septic shock, metabolic and cytokine factors associated with mortality and organ dysfunction, such as IL-6, IL-10, IL8, IL9, TNF β , and DMA, may also be useful [19]. A clinically relevant biomarker should have high sensitivity and specificity, and the ability to predict clinical outcomes through the use of a rapid, low-cost measurement technique [20].

Dimethylamine (DMA) is an amine circulating in human blood that is excreted in the urine. The occurrence of DMA in human urine was described in 1935 and 1938 [21,22]. A major precursor for endogenous DMA is asymmetric dimethylarginine (ADMA), an endogenous amino acid and inhibitor of nitric oxide (NO) synthesis. ADMA is hydrolyzed to DMA and l-citrulline by dimethylarginine dimethylaminohydrolase (DDAH). The concentration of circulating DMA in healthy young women (n=18) was determined to be 1.43±0.23 μ M in serum, 1.73±0.17 μ M in lithium heparin plasma, and 9.84±1.43 μ M in EDTA plasma.

Serum is recommended as the most appropriate matrix for measuring the DMA level in human blood [23].

In adults, about 90% of daily produced ADMA is estimated to be hydrolyzed to DMA, with the remaining 10% being excreted unchanged in the urine [24]. DMA circulates in the blood in the lower μ M range and is excreted in the urine in the upper μ M range. The diurnal variation of DMA excretion is fairly constant, and drugs such as the diuretic acetazolamide do not affect its urinary excretion rate in healthy humans [25].

The mass spectrometry (MS) technique is based on measurement of the charge-to-mass ratio (m/z) of ions of the tested substances present in the gas phase. Mass spectrometry is a highly specific and sensitive technique; coupled with an appropriate separation technique (liquid chromatography, gas chromatography, capillary electrophoresis), it allows for characterization of many groups of compounds and their quantitative analysis at very low levels, even in such complex biological matrices as serum, plasma, urine, tissues, and cell lines. MS technique is considered the criterion standard in gualitative and quantitative analysis. Due to its potential, it is often used in metabolomics studies, both targeted and untargeted. The method used in the present study is efficient and sensitive, and it allows for simultaneous analysis of L-arginine, L-citrulline, L-ornithine, ADMA, SDMA, and DMA in human serum using benzoyl chloride as a derivatization reagent, analyzed by reversed-phase chromatography, with full separation of ADMA and SDMA in a relatively short time. In contrast to methods using classical detectors, the method using mass spectrometry is highly specific, which generates fewer false high results [26].

Therefore, this study aimed to investigate serum levels of 5 amino acids – , arginine, ornithine, ADMA, SDMA, L-citrulline – and the amine, DMA, measured using LC-MS/MS in 41 patients with and without AKI admitted to the ICU of a single center who required CRRT.

Material and Methods

Ethics

The study protocol was approved by the Bioethics Committee of the Wrocław Medical University, Poland (approval no. KB-166/2018). Confidentiality and privacy were observed for all personal, laboratory, and clinical data. The study was carried out in accordance with the guidelines of the Declaration of Helsinki and Good Clinical Practice. Written informed consent was obtained from all the participants prior to the study.

Table 1. Study group characteristics.

			44					
		n=	11-71		n=29)	No (p-value*	
		n	%	n	%	n	%	
Sex	Male	26	63.4	18	62.1	8	66.7	0.78
	Female	15	36.6	11	37.9	4	33.3	
Sepsis	No	23	56.1	14	48.3	9	75.0	0.12
	Yes	18	43.9	15	51.7	3	25.0	
Septic shock	No	18	43.9	15	51.7	3	25.0	0.12
	Yes	23	56.1	14	48.3	9	75.0	
Source of sepsis	IAI	17	41.5	14	48.3	3	25.0	0.13
	CNS	3	7.3	1	3.5	2	16.7	
	RESP	17	41.5	11	37.9	6	50.0	
	UTI	3	7.3	3	10.3	0	0.0	
	SSTI	1	2.4	0	0.0	1	8.3	
CRRT	No	28	68.3	21	72.4	7	58.3	0.38
	Yes	13	31.7	8	27.6	5	41.7	

n – number of patients; IAI – intrabdominal infection; CNS – central nervous system; RESP – respiratory system; UTI – urinary tract infection; SSTI – soft skin tissue infection; CRRT – continuous renal replacement therapy. * U Mann-Whitney test.

Design and settings

This single-center retrospective observational study was conducted at the Department of Anaesthesiology and Intensive Therapy, 4th Military Clinical Hospital, Poland. The standards of Strengthening the Reporting of Observational Studies in Epidemiology (STROBE) were followed, and the relevant checklist was used for enrollment and allocation of participants [27].

Participants

The study group consisted of 41 patients treated in an ICU. This study used a prospective blood sampling and retrospective data analysis of a cohort of patients admitted to the Department of Anesthesiology and Intensive Therapy at Wrocław Medical University, Poland. Patients with sepsis and septic shock within the first 24 h of admittance were enrolled. The inclusion criteria were sepsis or septic shock according to the Sepsis-3 criteria, age >18, while the exclusion criteria disqualified patients with severe immunosuppression (cancer, chronic steroid therapy, chemotherapy) or patients aged <18 years old. All septic patients were treated according to the Surviving Sepsis Campaign guidelines. The detailed characteristics of the patients are presented in **Table 1**.

After their admission to the ICU, 2 groups of patients were distinguished. One group included patients with AKI in the course of sepsis and septic shock, which required the use of CRRT; the second group included patients with sepsis and septic shock without AKI and CRRT. The criteria were established based on the Kidney Disease: Improving Global Outcomes (KDIGO) guidelines [28]. AKI was defined as an increase in creatinine serum (SCr) by \geq 0.3 mg/dL (\geq 26.5 µmol/L) within 48 h, an increase in SCr to \geq 1.5 times baseline, which is known or presumed to have occurred within the prior 7 days, or a urine volume of less than 0.5 mL/kg/h for 6 h. If there was no improvement in urine output or a decrease in creatinine, renal replacement therapy was started in patients with corrected volemia and a MAP \geq 65 mmHg or with life-threatening metabolic and electrolyte disorders. All CRRT patients were treated in CVVHDF mode with citrate anticoagulation. The applied ultrafiltrate dose varied in the range of 20-25 mL/kg/bw.

Laboratory and Clinical Parameters

The patients' demographic, clinical, and biochemical data at the time of ICU admission were recorded. The Acute Physiology and Chronic Health Evaluation (APACHE) II [29] and Sequential Organ Failure Assessment (SOFA) [30] scores were evaluated at the initial time of ICU admission. The SOFA scores were also recorded on days 1, 3, 5, and 7. Data were collected for the following parameters: age, sex, total fluid input and output, serum creatinine, urea, procalcitonin, CRP, Hb, blood lactate, arterial and venous blood gas analysis, parameters of coagulation system and liver, serum electrolytes, and arginine, ornithine, ADMA, SDMA, DMA, and citrulline. Additionally, patients were assessed with respect to their need for CRRT [31].

Serum Sample Collection

Blood samples were collected into serum separator tubes in the morning of the first, third, fifth and seventh days. Blood samples were allowed to clot at room temperature (22-24°C for 15 min and then centrifuged at 720 g for 10 min. For amino acid analysis, the serum was divided into aliquots and stored at -80°C [32].

Preparation of Calibration Standards and Urine and Serum Samples – LC-MS Sample Analysis

Analysis of Arg/NO pathway metabolites was performed using a Xevo G2 Q-TOF MS mass spectrometer coupled with the Acquity UPLC system (Waters, Milford, MA, USA). Calibration standards and plasma samples were conducted in the same manner according to the method described by Fleszar et al [26]. Briefly, 100 µl aliquots was pipetted into 2.0 mL polypropylene tubes with 10 µl of internal standard solution [IS D6-DMA (50 µM); D7-ADMA (20 µM); D7-Arg (100 µM)] and 50 µl of borate buffer. Derivatization was performed by adding the 10% benzoyl chloride (BC)l in acetonitrile and 400 µl of acetonitrile as precipitation solution. After centrifuging, 100 µl of the supernatant was transferred into autosampler glass vials with 400 µl of water for the LC-MS analysis. Samples were separated on a Waters HSS T3 column (1.8 µm, 1.0×50 mm), using the following gradient: 0.0 min – 5% B hold for 1 min, 3.5 min - 14% B, 5.0 min - 60% B, 5.5 min - 90% B hold for 1.1 min, 6.60 min – 5% B, where eluent A was water with 0.1% formic acid (FA) and eluent B was methanol with 0.1% FA. The temperature of the column was 60°C and total flow rate was 0.220 mL/min.

The mass spectrometer was operated in full scan mode from 100 to 650 m/z with leucine–enkephalin as lock mass. The spray voltage, source temperature, and the desolvation temperature were set at 0.5 kV, 120°C, and 400°C, respectively. For quantitative analysis, extracted ion chromatograms (EIC) of compounds were used. Data were acquired and quantified using MassLynx software (Waters) [33].

Preparation of Calibration Standards and Urine and Serum Samples – Chemicals and Reagents

The Arg/NO pathway metabolites were of analytical standard: hydrochloride salts of unlabeled dimethylamine (DMA), hexadeutero- dimethylamine (D6-DMA, declared as 99 atom% 2H), L-arginine, SDMA, ADMA, L-citrulline, L-ornithine monohydrochloride, labeled L-ornithine hydrochloride (3,3,4,4,5,5-D6ornithine), benzoyl chloride (BCl) and sodium tetraborate were procured from Sigma-Aldrich (Poznan, Poland). Isotope labeled L-arginine: HCl (D7-arginine, 98%) and asymmetric dimethylarginine (2,3,3,4,4,5,5-D7-ADMA, 98%) were obtained from Cambridge Isotope Laboratories (Tewksbury, MA, USA). Methanol, acetonitrile (ACN), water, and formic acid (FA) were acquired from Merck Millipore (Warsaw, Poland), and leucine– enkephalin was obtained from Waters (Warsaw, Poland) [34,35].

Sample Size

Sample size analysis was performed using Statistica 13 (TIBCO Software, Inc.). Based on the available preliminary results carried out in our center (pilot incidence, n=9), we assessed how the results differ between the 2 groups (group A – without AKI and without CRRT, n=5; group B – with AKI and CRRT, n=4). In the sample size estimation analysis, the means, and standard deviations of the amino acid level scores in both groups were used. The estimated sample size is based on a two-sample *t* test (independent-sample *t* test). The alpha level was set at 0.05 and the test power at 0.8. It was also assumed that the assessed variables were not correlated, and a two-sided null (zero) hypothesis was adopted.

Statistical Analysis

Statistical analysis was performed using the Statistica 13 program (TIBCO, Inc., USA). For measurable variables, the medians, the lower quartile, and the upper quartile were calculated. All investigated quantitative variables were checked with the Shapiro-Wilk test to establish the type of distribution. The results were compared using the Mann-Whitney U test. The comparison of qualitative variables was performed using the chi-square test. The receiver operating characteristic (ROC) curves for selected variables were presented along with the determined optimal cut-off points (Youden's method) and the relationship between sensitivity and specificity are shown. Moreover, to assess the reliability of diagnostic tests, sensitivity and specificity were determined. Additionally, the positive (PPV) and negative (NPV) predictive value was indicated, as well as the likelihood ratio of positive (LR+) and negative (LR-) results. An attempt was made to evaluate the relationship of the results of selected parameters with AKI and for CRRT (logistic regression for each variable separately - one-way model of predictors included in the analysis). Odds ratios and confidence intervals were determined.

Results

AKI Biomarkers and Procalcitonin Level

There was a statistically significantly higher median serum level of procalcitonin obtained in patients with AKI who underwent CRRT (Me=54.7 ng/mL) compared to those without AKI and CRRT (Me=-2.9 ng/mL) (P<0.001). In addition, a statistically significant difference in serum urea results was found. The median in patients with AKI who did not receive CRRT therapy was 66.0 mg/dL and was lower than in patients with AKI who received

	AKI and CRRT									
		Yes (n=13)			p -value*					
	Me	Q1	Q3	Me	Q1	Q3				
Procalcytonin (ng/mL)	54.7	37.5	90.0	2.9	1.3	13.6	<0.001			
Urea (mg/dL)	95.0	60.0	158.0	66.0	42.0	90.0	0.018			
Creatinine (mg/dL)	1.8	1.4	4.0	1.3	0.8	2.1	0.022			
Urine output (mL/24h)	100.0	0.0	330.0	1675.0	1012.5	2825.0	<0.001			
HCO3- (mmol/L)	17.5	15.7	21.1	21.9	18.7	26.5	0.029			
Lactate (mmol/L)	5.0	1.8	6.4	1.7	1.1	2.4	0.012			

Table 2. Comparison of selected laboratory blood tests in patients with and without AKI and CRRT.

n - number of patients; Me - median; Q1 - lower quartile; Q3 - upper quartile. * U Mann-Whitney test.

Table 3. Comparison of the level of amine DMA and selected amino acids in the group with and without AKI.

		Yes (n=13)			p-value*		
	Me	Q1	Q3	Me	Q1	Q3	·
Arg [umol/L]	49.0	34.3	55.0	52.8	39.6	78.0	0.189
Orn [umol/L]	10.3	3.7	32.0	7.5	5.6	16.1	0.965
ADMA [umol/L]	0.58	0.53	0.62	0.64	0.49	0.76	0.507
SDMA [umol/L]	1.11	0.75	1.69	1.06	0.51	1.46	0.434
DMA [umol/L]	8.1	5.4	18.7	5.2	4.0	7.4	0.022
Cit [umol/L]	22.0	10.7	32.1	18.5	13.6	26.2	0.734

n – number of patients; Me – median; Q1 – lower quartile; Q3 – upper quartile; Arg – L-arginine; Orn – L-Ornithine; ADMA – asymmetric dimethylarginine; SDMA – symmetric dimethylarginine; DMA – dimethylamine; Cit – L-citrulline. * U Mann-Whitney test.

CRRT (Me=95.0 mg/dL) (P=0.018). A statistically significant difference in serum creatinine levels was also demonstrated. A higher median of creatinine was achieved by patients with AKI who received CRRT (Me=1.8 mg/dL) compared to those without AKI who did not receive CRRT (Me=1.3 mg/dL) (P=0.022).

It was found that statistically significant higher levels of urine output were obtained by patients who did not undergo CRRT (Me=1675.0 mL/day) compared to those who received CRRT (Me=100.0 mL/day) (P<0.001). In addition, there was a statistically significant difference in blood bicarbonate (HCO3-) results: the median in patients without CRRT was 21.9 mmol/L and was higher than in those with CRRT (Me=17.5 mmol/L) (P=0.029). A statistically significant difference in lactate levels was also shown, with a higher median for the group of patients undergoing CRRT (Me=5.0) compared to the patients not undergoing CRRT (Me=1.7) (P=0.012). **Table 2** compares

the results of selected blood tests in patients with and without AKI and CRRT.

Selected Amino Acids and DMA

Table 3 compares the concentrations of amino acids (arginine, ornithine, ADMA, SDMA, and citrulline) and DMA in the serum (expressed as μ mol/L) between the septic group with and without AKI.

There was a statistically significantly higher median level of DMA obtained from the AKI patients (Me=8.1 umol/L) compared to the patients without AKI (Me=5.2 umol/L) (P=0.022). The results for the remaining molecules, as well as for the values of selected amino acids and DMA in patients with sepsis and septic shock, revealed no statistically significant differences (P>0.05) (**Table 4**).

	Sepsis							Septic shock						
	Yes (n=18)			No (n=22)		P- value*	Yes (n=22)			No (n=18)			p- value*	
	Me	Q1	Q3	Me	Q1	Q3		Me	Q1	Q3	Me	Q1	Q3	
Arg [umol/L]	44.8	37.5	58.7	55.0	40.0	74.5	0.407	55.0	40.0	74.5	44.8	37.5	58.7	0.407
Orn [umol/L]	7.5	5.8	15.2	8.5	4.0	25.5	0.989	8.5	4.0	25.5	7.5	5.8	15.2	0.989
ADMA [umol/L]	0.6	0.6	0.8	0.6	0.4	0.7	0.161	0.6	0.4	0.7	0.6	0.6	0.8	0.161
SDMA [umol/L]	1.2	0.9	1.7	1.0	0.6	1.5	0.471	1.0	0.6	1.5	1.2	0.9	1.7	0.471
DMA [umol/L]	5.4	4.0	7.6	5.8	4.6	11.0	0.438	5.8	4.6	11.0	5.4	4.0	7.6	0.438
Cit [umol/L]	20.2	11.1	35.4	18.0	10.9	28.3	0.596	18.0	10.9	28.3	20.2	11.1	35.4	0.596

Table 4. Comparison of the level of amines and amino acids in the group with and without sepsis and septic shock.

n – number of patients; Me – median; Q1 – lower quartile; Q3 – upper quartile; Arg – L-arginine; Orn – L-Ornithine; ADMA – asymmetric dimethylarginine; SDMA – symmetric dimethylarginine; DMA – dimethylamine; Cit – L-citrulline.

* U Mann-Whitney test.

Table 5. The AUC of biomarkers for detection of AKI.

	AUC	SE	AUC lower 95%	AUC upper 95%	z	p-value*
Arg [μmol/L]	0.366	0.09	0.19	0.54	-1.49	0.137
Orn [μmol/L]	0.494	0.12	0.26	0.73	-0.05	0.960
ADMA [µmol/L]	0.430	0.10	0.23	0.63	-0.70	0.486
SDMA [µmol/L]	0.580	0.10	0.39	0.77	0.84	0.403
DMA [µmol/L]	0.732	0.09	0.55	0.91	2.55	0.011
Cit [µmol/L]	0.536	0.105	0.33	0.741	0.341	0.733

Orn – L-Ornithine; ADMA – asymmetric dimethylarginine; SDMA – symmetric dimethylarginine; DMA – dimethylamine; Cit – L-citrulline; AUC – area under the curve; SE – standard error. * U Mann-Whitney test.

AUC/ROC Analysis of the Concentration of Selected Amino Acids, Amin DMA, and Serum Creatinine Relative to the Incidence of AKI in Sepsis and Septic Shock

The AUC analysis of the selected amino acids, arginine, ornithine, ADMA, SDMA, citrulline, and DMA, as predictors of AKI, is presented in **Table 5**.

Only DMA was selected for ROC analysis as its concentrations were shown to be significantly different between septic patients with and without AKI and CRRT (**Table 5, Figure 1**). The ROC curves of DMA and creatinine were plotted to identify a cut-off value that could differentiate patients with and without AKI and CRRT (**Figures 1, 2**). For DMA and creatinine, a measure of relationship, such as sensitivity, specificity, PPV, NPV, ACC, LR+, and LR-, is shown. **Table 6** summarizes the AUC results for the variables DMA and creatinine.

Table 7 shows the number of people who obtained a result above or below the established cut-off point for DMA and creatinine, according to whether they were treated with CRRT. In subjects with DMA greater than or equal to 14.95 umol/L, 100% (n=5) were treated with CRRT, and in the group of subjects with a DMA score below the cut-off point, 20% were treated with CRTT (n=7). It was also observed that in subjects with creatinine levels greater than or equal to 1.19 mg/dL, 28% of the subjects (n=11) were treated with CRRT, and in the group of subjects with creatinine scores below the cut-off point, only 3% were so treated (n=1). The table also includes the results of sensitivity, specificity, PPV, NPV, AVV, and LR-/+. Additionally, a positive, statistically significant correlation between creatinine and DMA was observed (**Figure 3**).

Survival Analysis

A comparison of APACHE II and SOFA scores in the groups of patients with and without AKI is presented in **Table 8**. Patients

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Figure 1. DMA ROC curve in AKI and CRRT patient group. DMA – dimethylamine; ROC – receiver operating characteristic; AKI – acute kidney injury; CRRT – continuous renal replacement therapy.



Figure 2. Creatinine ROC curve in AKI and CRRT patient group. ROC – receiver operating characteristic; AKI – acute kidney injury; CRRT – continuous renal replacement therapy.

Table 6.	The AUC	of DMA	biomarker	and	creatinine	for	detection	of CRRT	patients.
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	AUC	SE	AUC lower 95%	AUC upper 95%	z	p-value
DMA [umol/L]	0.732	0.09	0.55	0.91	2.55	0.011
Creatinine[mg/dL]	0.702	0.09	0.53	0.88	2.30	0.022

AUC - area under the curve; SE - standard error.

Table 7. The sensitivity, specificity, PPV, NPV, AVV, and LR of DMA and creatinine for detection of CRRT patients.

5144	CR	RT	Curstinius	CRRT		
DMA	No (n) Yes (n)		Creatinine	No (n)	Yes (n)	
DMA <14.95 umol/L	28	7	Creatinine <1.19 mg/dL	13	1	
DMA ≥14.95 umol/L	0	5	Creatinine ≥1.19 mg/dL	15	11	
Sensitivity	1.000		Sensitivity	0.464		
Specificity	0.417		Specificity	0.917		
LR+	1.7	/14	LR+	5.571		
LR-	0.0	000	LR-	0.584		
PPV	0.800		PPV	0.929		
NPV	1.000		NPV	0.423		
ACC	0.825		ACC	0.600		

n – number of patients; DMA – dimethylamine; LR+ – positive likelihood ratio; LR- – negative likelihood ratio; PPV – positive predictive value; NPV – negative predictive value; ACC – accuracy.



Figure 3. Creatinine vs DMA correlation for the whole group (n=41). DMA – dimethylamine.

Table 8. Comparison of APACHE II and SOFA scale results against the occurrence of AKI.

	AKI									
		Yes (n=13)			No (n=28)		p-value*			
	Me	Q1	Q3	Me	Q1	Q3				
APACHE II	25.0	23.0	32.0	21.0	15.5	27.0	0.038			
SOFA	14.0	13.0	15.0	9.0	7.0	11.5	<0.001			

n - number of patients; Me - median; Q1 - lower quartile; Q3 - upper quartile. * U Mann-Whitney test.

with AKI (Me=25.0 points) had statistically significantly higher APACHE II scores than patients without AKI (Me=21.0 points) (P=0.038). In addition, a statistically significant difference in SOFA scores (P<0.001) was demonstrated; patients with AKI (Me=14.0 points vs Me=9.0 points) had higher median SOFA scores.

Discussion

In our study, the usefulness of determining the value of DMA as a marker of AKI in the course of sepsis and septic shock and the need for CRRT was demonstrated. As a biomarker of septic AKI, a very high sensitivity (100%) and a specificity of 42% were demonstrated for and above a 14.95 μ mol/L of DMA serum level. Due to the renal metabolism of dimethylamine and its excretion by urine, it seems to be useful in the diagnosis of kidney damage.

The kidney plays an important role in the metabolism of amino acids and the control of plasma concentrations. Abnormal plasma and muscle amino acid profiles in chronic renal failure first indicates malnutrition, which can be partially corrected by supplementation. Investigations in normal kidneys have shown that glutamine uptake maintains acid-base homeostasis, removes glycine and citrulline, and releases serine and arginine into the circulation. These metabolic processes are impaired in chronic renal failure. Uremia affects most tissues and causes malnutrition, while acidosis activates the catabolism of amino acids and proteins in muscle.

The kidney converts citrulline to arginine. Loss of this function likely contributes to the increasing ratio of citrulline to arginine as the GFR declines below 50 mL/min/1.73 m² [36]. Similarly, reduced renal production of serine from glycine probably underlies the rise in the plasma glycine-to-serine ratio.

In particular, DMA, which circulates in human blood and is excreted in the urine, is a useful biomarker of kidney injury. The major precursor for endogenous DMA is ADMA, an endogenous inhibitor of NO synthesis, which is hydrolyzed to DMA and l-citrulline by DAH. In adults, about 90% of daily produced ADMA is estimated to be hydrolyzed to DMA, with the remaining 10% being excreted unchanged in the urine. DMA circulates in the blood in the lower mM range and is excreted in the urine in the

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upper mM range. The variation of DMA excretion in the urine is fairly constant, and drugs, such as the diuretic acetazolamide, do not affect its urinary excretion rate in healthy humans [25]. ADMA is an endogenous inhibitor of nitric oxide synthase (NOS) [37]. Protein-bound arginine is methylated [38,39] by endothelial cells to form ADMA. Elevated ADMA concentrations are observed during renal failure and have been shown to contribute to cardiovascular mortality in those patients [40,41]. ADMA is converted to arginine and dimethylamine by the enzyme N-G,N-G-dimethylarginine dimethylaminohydrolase in the kidney or in the liver [42-44]. Approximately 4.5% of ADMA is excreted in the urine [45], and the remainder is metabolized to arginine, arginine-derived amino acids, and by-products.

Sepsis is defined as a life-threating organ dysfunction caused by the dysregulation of the host's response to an infection [39]. Due to abnormal tissue perfusion, over 47% of patients with sepsis and septic shock develop an AKI [46]. Patients with sepsis-associated AKIs have a 62% risk of death compared to 32% in patients without associated renal injury. For both sets of patients, diagnosis using sensitive and specific biomarkers of kidney damage can significantly affect the speed of diagnosis and the prevention of sepsis-associated AKIs. Biomarkers can also help in monitoring kidney function during sepsis treatment, for which drugs and fluids with nephrotoxic potential are often used.

The basic markers of renal failure used today are serum creatinine and the measurement of daily urine output. In the case of creatinine, there is no clear position on the determination of the baseline value in the absence of primary concentration measurements. In addition, changes in creatinine concentration are delayed due to renal reserves and the kinetics of creatinine itself. Since serum creatinine determination remains the standard in defining AKIs, the positive correlation of serum creatinine with DMA confirms the usefulness of DMA determination in the diagnosis of AKIs. At a DMA serum concentration of >14.95 umol/L, the high sensitivity of DMA determination (100%) in patients treated with CRRT was observed. Rodrigues et al [19] described the usefulness of monitoring DMA concentration in urine as a biomarker of kidney function.

Urine output depends mainly on the patient's volemic state before admission to the hospital. Urine output is the 1 of 2 KDIGO criteria of AKI diagnosis, but urine output is dependent upon many factors, including fluid administration, volemic status, and drug administration. Decrease of urine output does not necessarily indicate a decrease of glomerular filtration rate, but can be physiological renal adaptation to the body volume or electrolytes homeostasis. In the case of hypovolemia, urine output may be inadequately low.

A highly specific and sensitive biomarker to assess kidney damage is still being sought. In the course of sepsis and septic

shock, the pathomechanism of kidney damage results from hypoperfusion and ischemia, as well as inflammation and apoptosis [47]. In addition, during sepsis there are immunological disturbances and macrovascular and microvascular organ dysfunctions. In the course of sepsis, there is an increase in the expression of proinflammatory cytokines and the activation of leukocytes, leading to capillary plugging and micro clots. As a result, reactive oxygen species and the induction of NOS production occurs, which can damage the endothelium and glycocalyx [24]. All of this changes during sepsis and septic shock, resulting in structural and functional changes in the kidneys.

In our study, an attempt was made to assess the usefulness of amino acids and DMA, synthesized in the human body, as biomarkers of kidney damage. Only DMA seems to be an independent biomarker of kidney damage caused by sepsis and septic shock.

Our study has some potential limitations to be discussed. It was a single-center study and the number of patients enrolled was relatively small. The method of measurement of amino acids and DMA, as we explained before, is sensitive and highly specific; however, like any quantitative method, it has specific inaccuracies regarding quantification and detection, as well as limitations related to the amount of sample necessary to use, confirmed by validation, which was carried out in accordance with the Food and Drug Administration (FDA) guidelines. Method is characterized by the limits of detection (1.7 μ M L-arginine, 0.03 μ M ADMA, 0.02 μ M SDMA, 0.36 μ M l-citrulline, 0.06 μ M DMA) and quantification (3.2 μ M L-arginine, 0.08 μ M ADMA, 0.05 μ M SDMA, 1.08 μ M l-citrulline, 0.19 μ M DMA). The volume of the sample, depending on the concentration of the tested substances in analyzed matrix, ranges from 50 μ L to 100 μ L [26].

Conclusions

In patients with sepsis, increased serum levels of DMA were significantly associated with AKI requiring CRRT. It remains unclear whether increased DMA concentrations are secondary to sepsis-induced AKI, or are a cause. Serum DMA might be useful as a specific and sensitive biomarker of renal failure in patients with sepsis and septic shock.

Department and Institution Where Work Was Done

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Declaration of Figures' Authenticity

All figures submitted have been created by the authors who confirm that the images are original with no duplication and have not been previously published in whole or in part.

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