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# Guanidinoacetic Acid and Creatine are Associated with Cardiometabolic Risk Factors in Healthy Men and Women: A Cross-Sectional Study

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**Abstract:** Guanidinoacetic acid (GAA) conversion to creatine is thought to be involved in cardiometabolic disturbances through its role in biological methylation and insulin secretion. We evaluated the association of serum GAA and creatine with cardiometabolic risk factors in a cohort of 151 apparently healthy adults (82 women and 69 men) aged 18–63 years. Serum levels of GAA and creatine were measured with liquid chromatography-tandem mass spectrometry. A multiple linear regression model adjusted for age and sex was employed to examine the relationship of serum GAA and creatine with cardiometabolic risk factors. Higher GAA levels were associated with an unfavorable cardiometabolic risk profile (higher insulin, higher total homocysteine, and higher body fat percentage), while having elevated serum creatine levels ( $\geq 31.1$   $\mu\text{mol/L}$ ) was associated with being overweight (body mass index  $\geq 25.0$  kg/m). The results from our study suggest a possible role of the GAA–creatine axis in the pathogenesis of cardiovascular and metabolic diseases.

**Keywords:** guanidinoacetic acid; creatine; cardiometabolic risk; homocysteine; overweight

## 1. Introduction

Guanidinoacetic acid (GAA) and creatine are natural amino acid derivatives that are heavily involved in cellular energy metabolism [1–3]. Much attention has been given recently to the potential role of these energy-related compounds in cardiometabolic diseases. For example, the high creatine kinase (CK) phenotype has been found in hypertension- and obesity-prone patients [4–6], and high CK activity has been reported to lead to dysfunctional vascular contractility and insulin resistance [6]. In addition, the biosynthesis of creatine from GAA is considered critical for methyl group consumption in humans [7], with altered methylation and consequent homocysteine production suggested as additional factors for cardiovascular disease risk [8,9]. Although the main focus has been on homocysteine metabolism, there have also been links between the GAA–creatine axis and classical cardiometabolic biomarkers, including insulin sensitivity and blood cholesterol [10]. In this study, we evaluated the associations of serum GAA and creatine with cardiometabolic risk factors—including body mass index (BMI), lipid profiles, high-sensitive C-reactive protein (CRP), glucose, and insulin—in apparently healthy men and women. A prespecified hypothesis suggested a negative correlation between an unfavorable cardiometabolic risk profile (including higher fasting insulin, higher total homocysteine, and higher body fat percentage) and serum levels of GAA and creatine.

## 2. Materials and Methods

### 2.1. Participants

This study involved a secondary analysis of a subset of the Diet and Physical Activity for Health Initiative (DiPAH) cohort (ClinicalTrials.gov ID: NCT01958333) that underwent cardiometabolic risk assessment and had archived serum. The DiPAH was started in 2008 as a long-term, nationally recognized health study that is focused on strategies for the prevention of chronic diseases in the Serbian population. The present study was a cross-sectional study designed to identify relationships between energy-related biomarkers of micronutritional status and cardiometabolic disease risk factors in healthy adults. Participants were selected to join the study on the basis of the following criteria: (a) previous participation in DiPAH trials; (b) age between 18 and 65 years; (c) no medical conditions that would limit the successful completion of the protocol; and (d) current residence in Serbia. Pregnant women, individuals taking dietary supplements, and individuals engaged in a programmed exercise regimen were excluded from the study. Of 11069 DiPAH subjects initially considered, 151 participants (82 women and 69 men) were eligible and consented to participate in the current study. The minimal sample size ( $n = 120$ ) was calculated according to power analysis for correlation point biserial model, with the effects size set at 0.25, a two-tail alpha level of 0.05, and a study power of 0.80 (G-Power 3, Heinrich Heine University Düsseldorf, Düsseldorf, Germany). The study was conducted according to the guidelines laid down in the Declaration of Helsinki, and all procedures were approved by the local IRB at the University of Novi Sad.

### 2.2. Experimental Procedures

Blood samples after an overnight fast of 10 h and anthropometric measures were obtained by trained research staff. A fasting sample of venous blood was immediately centrifuged, with serum stored in a freezer at  $-80\text{ }^{\circ}\text{C}$ , and the sample was analyzed for specific biomarkers after study completion. Serum GAA and creatine were measured by liquid chromatography-tandem mass spectrometry (SCIEX LC-MS/MS 5500QTRAP, AB Sciex Ltd., Concord, ON, Canada). Lipid profiles and glucose were analyzed by standard enzymatic methods with an automatic analyzer (Hitachi 704, Tokyo, Japan), while C-reactive protein was measured with a high sensitivity immunoturbidimetric assay (Hitachi 912, Tokyo, Japan). Insulin was evaluated with the CLIA method by automated chemiluminescence. Total serum homocysteine was determined with a chemiluminescent immuno-assay method using a chemistry analyzer (DPC Immulite 2000, Siemens, Berlin, Germany). Height was measured by a stadiometer (Seca 213, Hamburg, Germany), while weight and body fat percentage were measured by a bioelectrical impedance analyzer (Omron BF 511, Kyoto, Japan). Body mass index (BMI) was calculated as weight in kilograms divided by the square of height in meters, and participants were categorized as non-overweight ( $\text{BMI} < 25.0\text{ kg/m}^2$ ) or overweight ( $\text{BMI} \geq 25.0\text{ kg/m}^2$ ). All participants were assessed in underwear after voiding, and the same trained technician did the anthropometric assessment in aim to minimize the testing error.

### 2.3. Statistical Analyses

Data are presented as the means  $\pm$  standard deviation (SD) or frequencies (%). A multiple linear regression model with a stepwise method (adjusted for age and sex) was used to examine the relationships between serum GAA and serum creatine levels, and cardiometabolic risk factors. Multivariate-adjusted odds ratios for overweight participants were calculated across quartile categories of GAA and creatine, with a 95% confidence interval (CI) presented and a linear trend assessed across quartiles. Serum GAA and creatine were compared across sex and BMI categories using a two-tailed independent *t*-test. The significance level was set at  $p \leq 0.05$ . The data were analyzed using the statistical package SPSS, version 21.0 (SPSS Inc., Chicago, IL, USA).

### 3. Results

Baseline characteristics of the study sample are depicted in Table 1. The mean  $\pm$  SD age was  $25.5 \pm 6.9$  years, with both genders were almost equally represented (54.3% women), and ~1 in 4 participants was overweight. The mean serum creatine levels were greater by  $5.1 \mu\text{mol/L}$  in men than in women (95% CI 2.2–8.0  $\mu\text{mol/L}$ ;  $p = 0.001$ ), whereas creatine concentrations were lower by  $3.5 \mu\text{mol/L}$  in normal weight vs. overweight participants (95% CI 0.1–6.9  $\mu\text{mol/L}$ ;  $p = 0.05$ ) (Table 2). A positive correlation has been found between serum creatine and BMI ( $r = 0.22$ ,  $p = 0.01$ ) (Figure 1).

**Table 1.** Sample characteristics ( $n = 151$ ). Values are mean  $\pm$  standard deviation (SD).

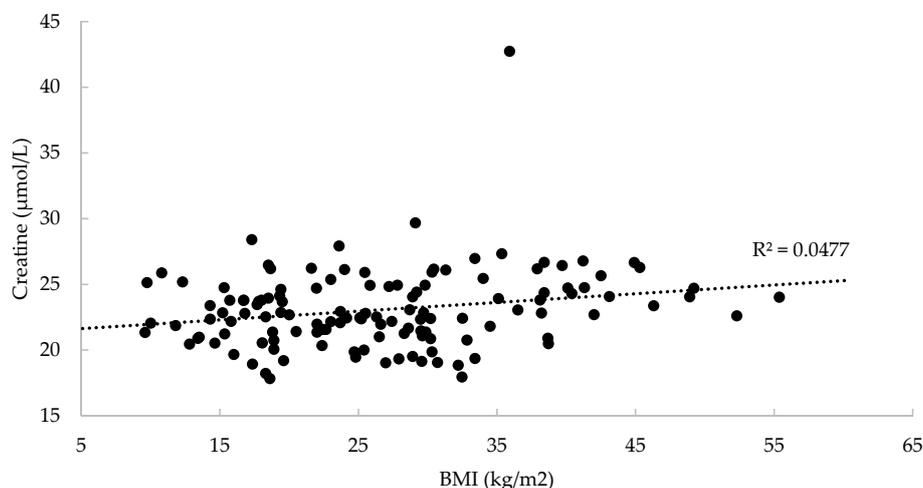
Variable	Value	Range
Age (years)	$25.5 \pm 6.9$	18.0–63.0
Sex, F (%)	54.3	-
BMI ( $\text{kg}/\text{m}^2$ )	$22.9 \pm 2.9$	17.6–42.7
Overweight <sup>1</sup> (%)	25.8	-
Body fat (%)	$21.3 \pm 7.5$	7.4–47.6
Total cholesterol (mmol/L)	$4.7 \pm 0.7$	3.4–7.2
LDL cholesterol (mmol/L)	$3.1 \pm 0.7$	1.4–5.1
HDL cholesterol (mmol/L)	$1.4 \pm 0.3$	0.6–2.7
Triglycerides (mmol/L)	$1.1 \pm 0.6$	0.4–5.8
Glucose (mmol/L)	$4.7 \pm 0.8$	2.2–6.4
Insulin (IU/L)	$8.8 \pm 6.2$	3.9–29.7
tHcy ( $\mu\text{mol/L}$ )	$8.5 \pm 1.9$	4.4–12.5
CRP (mmol/L)	$0.002 \pm 0.001$	0.001–0.004
GAA ( $\mu\text{mol/L}$ )	$2.6 \pm 0.7$	0.9–4.5
Creatine ( $\mu\text{mol/L}$ )	$26.6 \pm 9.3$	9.6–55.4

BMI—body mass index; tHcy—total homocysteine; CRP—C-reactive protein; GAA—guanidinoacetic acid; F—female. <sup>1</sup> BMI  $\geq 25.0 \text{ kg}/\text{m}^2$ .

**Table 2.** Serum levels of guanidinoacetic acid (GAA) and creatine. Values are mean  $\pm$  SD.

	GAA ( $\mu\text{mol/L}$ )	Creatine ( $\mu\text{mol/L}$ )
Sex categories		
Men ( $n = 69$ )	$2.6 \pm 0.7$	$29.4 \pm 10.4$
Women ( $n = 82$ )	$2.6 \pm 0.7$	$24.3 \pm 7.7^*$
BMI categories		
Normal weight ( $n = 112$ )	$2.6 \pm 0.7$	$25.7 \pm 8.9$
Overweight ( $n = 39$ )	$2.6 \pm 0.8$	$29.2 \pm 10.1^*$

An asterisk (\*) indicates a significant difference between categories at  $p < 0.05$  (2-tailed independent  $t$ -test).



**Figure 1.** Correlation between serum creatine and body mass index (BMI) ( $n = 151$ ).

The odds of being overweight were similar across quartiles of serum GAA concentrations after controlling for age and sex (Table 3). A significant trend for greater odds of being overweight has been reported in participants with serum creatine levels in the fourth quartile than in the first three quartiles ( $p = 0.04$ ).

**Table 3.** Multivariate-adjusted odds ratios (95% CI in parentheses) for overweight (BMI  $\geq 25.0$  kg/m<sup>2</sup>) across quartile categories of serum guanidinoacetic acid (GAA) and creatine.

GAA					Creatine				
Quartile with $\mu\text{mol/L}$ Range					Quartile with $\mu\text{mol/L}$ Range				
I	II	III	IV	$p$ trend	I	II	III	IV	$p$ trend
0.90–2.19	2.20–2.79	2.80–2.99	3.00–4.50		9.6–18.9	19.0–26.2	26.3–31.0	31.1–54.4	
1.00	0.57	0.68	0.84	0.66	1.00	1.18	0.97	3.26	0.04
(–)	(0.19 to 1.69)	(0.24 to 1.94)	(0.31 to 2.31)		(–)	(0.38 to 3.68)	(0.30 to 3.09)	(1.14 to 9.32)	

Multiple regression analysis revealed that our model as a whole explained 60.3% of the variance in cardiometabolic risk biomarkers when serum GAA was included as a predictor variable ( $R = 0.78$ ; adjusted  $R^2 = 0.39$ , standard error of the estimate = 0.89) and 18.8% of the variance when serum creatine was included as a predictor variable ( $R = 0.43$ ; adjusted  $R^2 = 1.84$ , standard error of the estimate = 16.6), with gender and age accounted as control variables. There were no significant relationships between blood lipids and serum GAA or serum creatine concentrations when adjusted for age and sex (Table 4). Glucose was inversely associated with serum GAA and serum creatine levels, while insulin was positively associated with GAA levels only. Serum total homocysteine and body fat percentage were positively associated with serum GAA and serum creatine concentrations. Neither C-reactive protein nor body mass index were associated with serum GAA and serum creatine levels. In addition, serum creatine was positively associated with serum GAA concentrations when adjusted for age and sex ( $\beta = 0.32$ ;  $p < 0.0001$ ).

**Table 4.** Multiple linear regression coefficients for cardiometabolic risk factors in relation to serum levels of guanidinoacetic acid (GAA) and creatine ( $n = 151$ ).

	GAA		Creatine	
	$\beta$	$p$	$\beta$	$p$
Blood lipids				
Total cholesterol	−0.11	0.20	0.07	0.41
LDL cholesterol	−0.05	0.54	0.05	0.54
HDL cholesterol	0.08	0.35	0.13	0.12
Triglycerides	0.16	0.08	0.11	0.19
Glucose	−0.28	0.00	−0.26	0.00
Insulin	0.53	0.03	−0.17	0.57
tHcy	0.30	0.00	0.17	0.05
CRP	0.36	0.16	−0.26	0.38
Body mass index	0.02	0.87	0.11	0.23
BFP	0.32	0.03	0.37	0.01

The multiple linear regression model was adjusted for age and sex. tHcy—total homocysteine; CRP—C-reactive protein; BFP—body fat percentage.

#### 4. Discussion

In a cohort of apparently healthy adults, serum GAA, and creatine concentrations had significant associations with several cardiometabolic risk factors. Higher GAA levels were associated with an unfavorable cardiometabolic risk profile (including higher fasting insulin, higher total homocysteine, and higher body fat percentage), while participants with elevated serum creatine levels ( $\geq 31.1$   $\mu\text{mol/L}$ ) had greater odds of being overweight. This suggests a possible role of GAA–creatin axis in the pathogenesis of cardiovascular and metabolic disease.

Our findings were consistent with previous studies reporting a link between GAA and cardiometabolic disorders. Elevated serum GAA levels were associated with higher tHcy levels [11] and insulin hypersecretion [12], unfavorable risk factors for cardiometabolic diseases. We found that tHcy is a very strong predictor ( $\beta = 0.30$ ,  $p < 0.001$ ) and so it could be that tHcys is the root issue, either as an independent factor or through affecting arginine availability, since GAA is synthesized from arginine [1]. Hypothetically, enhanced GAA synthesis might restrain the availability of arginine, an amino acid that has been shown to strongly affect the risk factors of cardiovascular diseases in humans [13]. However, previous reports about the associations between serum creatine and cardiovascular and metabolic risk factors are not in line with our results. In several small-scale studies, higher serum creatine levels (as provoked by oral intake) were associated with a favorable risk-factor profile, including lower tHcy [14], reduced total cholesterol [15], or improved insulin sensitivity [16]. In contrast, a number of studies found no significant relationships between creatine alone and different cardiometabolic markers (for review see [10]), and a recent large study ( $n = 622$ ) suggested no association between serum creatine and tHcy levels in apparently healthy men and women [17]. The conflict between our study and results from previous creatine studies may have been partially due to whether the analyses were based on serum levels or dietary intakes of GAA and creatine. The above studies typically linked creatine levels with cardiometabolic risk factors during an exercise program, an intervention known to reduce cardiometabolic risks per se [18], while we evaluated a diverse population not currently involved in an exercise program or dietary intervention. Our findings are in accordance with previous trials reporting an association between higher BMI and higher serum creatinine, an end-product of creatine metabolism [19], suggesting a link between the GAA-creatinine axis and risk of being overweight. Hypothetically, GAA overload (as indicated by higher levels of serum GAA) may perturb creatine metabolism, thereby resulting in enhanced creatine synthesis that accounts for an equivalent proportion of tHcy production and cardiometabolic burden. We found that serum GAA is positively associated with insulin levels ( $\beta = 0.53$ ), while no such relationship was found between serum creatine and insulin levels. Having higher GAA concentrations appears to be accompanied by elevated levels of insulin circulating in the blood, suggesting a possible link between excess GAA and hyperinsulinemia, a well-known cardiometabolic risk factor. Our results corroborate a previous *in vitro* study showing that GAA stimulates insulin release more potently than creatine, by triggering insulin secretion via kinase-sensitive mechanisms in pancreatic islets cells [12]. However, before putting forward serum GAA as a proxy for insulin release, additional studies are highly warranted to analyze GAA dynamics in populations with normal and impaired insulin secretion and action. In addition, creatine synthesis from GAA might affect cardiovascular risk through depletion of arginine, a precursor of creatine and a key source of nitric oxide (NO) [20]. NO plays an important role in the protection against the onset and progression of cardiovascular disease [21], and creatine overload may induce disturbances in NO bioavailability leading to a loss of the cardioprotective actions and in some populations may increase disease progression [22]. Therefore, synchronized monitoring of GAA-creatinine and arginine-NO axes in future studies should provide more mechanistic evidence by which creatine synthesis affects cardiovascular risk.

Baseline levels of GAA and creatine were in accordance with previous studies reporting reference values for these two compounds [23–25], and gender-related differences for serum creatine found in the present study could be attributed to the effects of testosterone and estrogen on creatine synthesis and transport. The CK activity, which regulates the use and consumption of creatine, is found to be sex-specific [26]. Gender differences in muscle mass and responses to downstream signaling pathways in the skeletal muscle might also affect creatine utilization [27]. In particular, sex hormones appear to differently affect creatine transporter (SLC6A8) expression, with SLC6A8 inhibited by estrogens and stimulated by testosterone [28]. This has been suggested to provoke a higher leakage of body creatine in the urine of healthy women [25], with perhaps less creatine retained in the blood, as we found in the present study. In addition, we found that overweight adults had higher creatine concentrations (by 13.6% on average) compared to adults of a normal weight, and the odds ratio of being overweight

was 3.26-fold higher in participants with a serum creatine  $\geq 31.1$   $\mu\text{mol/L}$ . Previous studies confirm our findings, with a positive association having been reported between serum creatinine and BMI in healthy men and women [19,29]. The origin of creatinine from creatine may explain this correlation and the higher concentrations in men than women [30]. However, the strong positive correlation found between serum creatine levels (also GAA) and body fat percentage in our cohort remains puzzling. This perhaps means that circulating values of these compounds are related not only to body size but also to body composition and other (patho) physiological mechanisms that need to be revealed.

The major limitation of the present study is its cross-sectional design that prevented causal conclusions between the GAA-creatine axis and cardiometabolic risk factors. Second, we recruited a specific cohort of participants (predominantly young normal-weight adults, with no major biochemical disturbances) that limited the interpretation of our results to a healthy population while being unable to generalize the results to patients with cardiometabolic diseases. Third, we used only a finite compendium of biochemical variables to understand a more complete picture of GAA-creatine metabolism. Fourth, no link has been established between GAA and creatine levels and clinical indicators of cardiometabolic diseases, including markers of vascular pathology or metabolic overload. Finally, no skeletal muscle mass was profiled and accounted for in the regression analysis. Since skeletal muscle is a major site of insulin resistance [31], with creatine utilization found to be different among type I and type II muscle fibers [32], controlling for muscle mass in future studies could help to better address the link between biomarkers of GAA-creatine metabolism and cardiometabolic risk.

## 5. Conclusions

In conclusion, our results that GAA and creatine are associated with specific cardiometabolic risk factors (e.g., fasting insulin, total homocysteine, and body fat percentage) suggest a possible role of these energy-related compounds in the pathogenesis of cardiovascular and metabolic disease. The results from our study may help to better understand the metabolic aspects of these disorders, and suggest further evaluation of serum GAA and creatine applicability in the identification of adults with cardiometabolic disease risks.

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## Abbreviations

BFP	Body fat percentage
BMI	Body mass index
CK	Creatine kinase
CRP	C-reactive protein
DiPAH	Diet and Physical Activity for Health Initiative study
GAA	Guanidinoacetic acid
SLC6A8	Creatine transporter
tHcy	Total homocysteine

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