

Metabolite Profile Analysis Reveals Functional Effects of 28-Day Vitamin B-6 Restriction on One-Carbon Metabolism and Tryptophan Catabolic Pathways in Healthy Men and Women^{1–3}

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Abstract

Suboptimal vitamin B-6 status, as reflected by low plasma pyridoxal 5'-phosphate (PLP) concentration, is associated with increased risk of vascular disease. PLP plays many roles, including in one-carbon metabolism for the acquisition and transfer of carbon units and in the transsulfuration pathway. PLP also serves as a coenzyme in the catabolism of tryptophan. We hypothesize that the pattern of these metabolites can provide information reflecting the functional impact of marginal vitamin B-6 deficiency. We report here the concentration of major constituents of one-carbon metabolic processes and the tryptophan catabolic pathway in plasma from 23 healthy men and women before and after a 28-d controlled dietary vitamin B-6 restriction (<0.35 mg/d). Liquid chromatography-tandem mass spectrometry analysis of the compounds relevant to one-carbon metabolism showed that vitamin B-6 restriction yielded increased cystathionine (53% pre- and 76% postprandial; $P < 0.0001$) and serine (12% preprandial; $P < 0.05$), and lower creatine (40% pre- and postprandial; $P < 0.0001$), creatinine (9% postprandial; $P < 0.05$), and dimethylglycine (16% postprandial; $P < 0.05$) relative to the vitamin B-6-adequate state. In the tryptophan pathway, vitamin B-6 restriction yielded lower kynurenic acid (22% pre- and 20% postprandial; $P < 0.01$) and higher 3-hydroxykynurenine (39% pre- and 34% postprandial; $P < 0.01$). Multivariate ANOVA analysis showed a significant global effect of vitamin B-6 restriction and multilevel partial least squares-discriminant analysis supported this conclusion. Thus, plasma concentrations of creatine, cystathionine, kynurenic acid, and 3-hydroxykynurenine jointly reveal effects of vitamin B-6 restriction on the profiles of one-carbon and tryptophan metabolites and serve as biomarkers of functional effects of marginal vitamin B-6 deficiency. *J. Nutr.* 143: 1719–1727, 2013.

Introduction

Vitamin B-6, as pyridoxal 5'-phosphate (PLP)¹⁵, serves a coenzymatic function in many phases of metabolism. Marginal

vitamin B-6 status, defined as plasma PLP of 20–30 nmol/L, is common in the American population (1,2) and has been associated with increased risk of coronary artery disease and stroke (3–6). The mechanism responsible for these disease associations has not been determined but frequently is not associated with homocysteine elevation.

Vitamin B-6 plays an integral role in one-carbon metabolism, because PLP serves as a coenzyme for glycine decarboxylase of the glycine cleavage system, cytoplasmic and mitochondrial forms of serine hydroxymethyltransferase (SHMT), and cystathionine β -synthase (CBS) and cystathionine γ -lyase (CSE) in the transsulfuration pathway. Inadequate vitamin B-6 status has been associated with cardiovascular disease risk (3–6), possibly due to derangement of one-carbon metabolism. Whether any of these specific metabolic processes are involved in mediating disease risk remains unclear. *In vivo* kinetic studies and mathematical modeling analysis suggest that most aspects of one-carbon metabolism are sufficiently adaptive to maintain function over the range of

¹ Supported by NIH grant R01 DK072398 and NIH National Center for Research Resources CTSA grant 1UL1RR029890.

² Author disclosures: V. R. da Silva, L. Rios-Avila, Y. Lamers, M. A. Ralat, Ø. Midttun, E. P. Quinlivan, T. J. Garrett, B. Coats, M. N. Shankar, S. S. Percival, Y.-Y. Chi, K. E. Muller, P. M. Ueland, P. W. Stacpoole, and J. F. Gregory III, no conflicts of interest.

³ Supplemental Table 1 is available from the "Online Supporting Material" link in the online posting of the article and from the same link in the online table of contents at <http://jn.nutrition.org>.

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¹⁵ Abbreviations used: ADMA, asymmetric dimethylarginine; CBS, cystathionine β -synthase; CSE, cystathionine γ -lyase; DMG, dimethylglycine; IDO, indoleamine 2,3-dioxygenase; MS/MS, tandem MS; PLP, pyridoxal 5'-phosphate; SAM, S-adenosylmethionine; SHMT, serine hydroxymethyltransferase; TDO, tryptophan 2,3-dioxygenase; VIP, Variable Influence on Projection.

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mild to moderate deficiency of vitamin B-6 (7–12). However, our recent metabolomic investigation showed extended metabolic changes associated with marginal vitamin B-6 deficiency in controlled nutritional studies, including perturbations of amino acid, lipid, and organic acid profiles (13).

Severe vitamin B-6 deficiency has substantial effects on amino acid metabolism as reflected by changes in the concentration of several amino acids in plasma, tissue, and urine (14–16). Concentrations of glycine and, to a lesser extent, serine have been shown to increase in plasma, urine, and liver after vitamin B-6 depletion (14,15,17,18). A marginal vitamin B-6 deficiency, consistent with plasma PLP concentration of 20–30 nmol/L, (19,20) has been shown to affect one-carbon metabolism. Previous studies reported significant elevations in plasma glycine and cystathionine after a 28-d dietary vitamin B-6 restriction (8,9) and an inverse association of cystathionine with plasma B-6 vitamers in a large-scale population study (21). Little is known about the influence of vitamin B-6 deficiency on the pattern of other constituents of one-carbon metabolism and related pathways, especially during marginal deficiency. Mathematical modeling suggested that the PLP-dependent reactions in one-carbon metabolism exhibited little change of flux in vitamin B-6 deficiency (11), but the pattern of the various metabolites was not systematically investigated and is addressed herein.

The involvement of PLP in tryptophan catabolism represents another important role of vitamin B-6 in amino acid metabolism by regulating tryptophan turnover and providing NAD synthesis (22). This pathway is influenced by inflammatory conditions (23–25), cancer (26), pregnancy (27,28), and oral contraceptive usage (28–32) as well as insufficiency of cellular PLP (33–35). The major pathway of tryptophan oxidation to *N*-formylkynurenine exists in liver and is catalyzed by tryptophan 2,3-dioxygenase (TDO) (22,36). TDO is induced by tryptophan (37–39) and hormones such as glucocorticoids (38,40,41), glucagon (42), and estrogens (43). Tryptophan oxidation is catalyzed in nonhepatic tissues by indoleamine 2,3-dioxygenase (IDO) (44,45). In contrast to TDO, IDO is induced primarily by IFN γ (3,12); thus, IDO is associated with the immune response (24,44). Various inflammatory conditions frequently are associated with low plasma PLP (23–25), although vitamin B-6 deficiency in otherwise healthy humans does not induce inflammation (8). Despite having similar roles in supplying the tryptophan catabolic pathway, TDO and IDO have distinct functions in regulating tryptophan metabolism (39). Both hepatic and nonhepatic pathways of tryptophan catabolism involve the bifunctional PLP-dependent enzymes kynureninase and kynurenine aminotransferase (46,47) and are subject to effects of vitamin B-6 deficiency. Kynureninase catalyzes both the conversion of kynurenine to anthranilic acid and 3-hydroxykynurenine to 3-hydroxyanthranilic acid. Kynurenine aminotransferase catalyzes the conversion of kynurenine to kynurenic acid and 3-hydroxykynurenine to xanthurenic acid (48–50). Kynureninase appears to be more sensitive to the effects of vitamin B-6 insufficiency than kynurenine aminotransferase (47), as confirmed by recent mathematical modeling of tryptophan metabolism (51). It has been proposed that induction of the IDO-dependent nonhepatic pathway of tryptophan catabolism and other PLP-dependent enzymes involved in the inflammatory response can account for the association of low plasma PLP and markers of systemic inflammation (23).

Relatively little information is available regarding the response of tryptophan catabolites in blood, tissue, and urine to various levels of vitamin B-6 status. The relationship between circulating B-6 vitamers and plasma tryptophan metabolites has been examined in large-scale studies of individuals with systemic

inflammation and showed that elevation of 3-hydroxykynurenine was a biomarker of vitamin B-6 deficiency (52,53), which complements similar studies of the relationship of plasma B-6 vitamers and trans-sulfuration metabolites (21), although the extent to which inflammation affected the latter relationship is unclear. The sensitivity of tryptophan catabolites to nutritional effects (i.e., controlled vitamin B-6 restriction) is addressed in the present study.

We report here metabolic profile analyses intended to extend our understanding of the functional consequences of marginal vitamin B-6 deficiency in healthy men and women. These investigations involved analysis of the constituents of one-carbon metabolism and related pathways and of tryptophan catabolism of plasma samples from healthy men and women obtained before and after a dietary vitamin B-6 restriction protocol (9,12). We also interpret findings in relation to simulations using mathematical models of one-carbon metabolism (11) and tryptophan catabolism (51). These studies broaden our understanding of the metabolic effects of vitamin B-6 deficiency and extend recent metabolomic findings (54).

Materials and Methods

Materials

All standards were purchased from Sigma-Aldrich except symmetric dimethylarginine and asymmetric dimethylarginine (ADMA), which were purchased from Axxora (formerly Alexis Biochemicals). Internal standards [$^{13}\text{C}_2$]-glycine; [$^{13}\text{C}_5$]-methionine; [$^{15}\text{N}_2$]-arginine; [$^2\text{H}_3$]-serine; [$^2\text{H}_3$]-sarcosine; [$^2\text{H}_6$]-dimethylglycine; [$^2\text{H}_9$]-choline; [$^2\text{H}_{11}$]-betaine; [$^2\text{H}_3$]-creatine; [$^2\text{H}_3$]-creatinine; [$^2\text{H}_2$]-threonine; [$^2\text{H}_4$]-cystathionine; [$^2\text{H}_2$]-guanidinoacetic acid] were purchased from Cambridge Isotopes. [$^{13}\text{C}_5$] Methionine sulfoxide was synthesized from [$^{13}\text{C}_5$]methionine by oxidizing with 5% hydrogen peroxide.

Human vitamin B-6 restriction protocols

Participants. The plasma samples analyzed in this study were obtained from 23 healthy men and women participants in 2 identical dietary vitamin B-6 restriction studies previously reported (9,12). Nutritional adequacy was indicated by normal concentrations under preprandial conditions of serum folate (>7 nmol/L), serum vitamin B-12 (>200 pmol/L), plasma PLP (>30 nmol/L), and plasma total homocysteine (<12 $\mu\text{mol/L}$). All participants gave written informed consent. The University of Florida Institutional Review Board and the University of Florida Clinical Research Center Scientific Advisory Committee reviewed and approved this protocol (9,12).

Dietary protocol. All meals were prepared by the Clinical Research Center Bionutrition Unit at the University of Florida. Prior to the first blood sample, participants consumed nutritionally adequate meals with standardized composition for 2 d, after which they consumed a low-vitamin B-6 diet (0.37 ± 0.04 mg/d) for 28-d, intended to induce a state of marginal vitamin B-6 deficiency as reflected by plasma PLP between 20 and 30 nmol/L (9,12). All participants received daily a custom vitamin-mineral supplement containing no vitamin B-6 to maintain adequate status of other micronutrients.

The diet was planned using the following criteria. Actual intake varied depending on participants' daily intake patterns, with food components adjusted according to food preferences and energy needs. The diets provided: 50–60 g/d protein for females or 60–70 g/d males; 1.0 mg/d vitamin B-6 in the 2-d control diet period and <0.5 mg/d in the 28-d vitamin B-6 restriction period; 400 $\mu\text{g/d}$ dietary folate equivalents; energy adjusted to maintain weight using the Mifflin St.-Jeor equation and individualized activity factors; and macronutrient composition varied depending on individual food preferences. Actual intakes for the 2-d control period were: energy, $10,900 \pm 1850$ kJ/d; energy from fat, $28.4 \pm 3.8\%$; energy from carbohydrate, $62.0 \pm 4.0\%$; energy from protein, 9.4 ± 1.3 ; vitamin B-12, 1.7 ± 0.7 $\mu\text{g/d}$; naturally occurring folate, 150 ± 36 $\mu\text{g/d}$; folic acid, 240 ± 36 $\mu\text{g/d}$; and vitamin B-6, 1.02 ± 0.11 $\mu\text{g/d}$. The actual intakes for the 28-d restriction

period were: energy, $11,700 \pm 25,10$ kJ/d; energy from fat, $29.1 \pm 2.6\%$; energy from carbohydrate, $60.4 \pm 2.9\%$; energy from protein, 10.4 ± 1.0 ; vitamin B-12, 1.33 ± 0.34 $\mu\text{g}/\text{d}$; naturally occurring folate, 92 ± 29 $\mu\text{g}/\text{d}$; folic acid, 265 ± 28 $\mu\text{g}/\text{d}$, and vitamin B-6, 0.37 ± 0.04 $\mu\text{g}/\text{d}$. Dietary intake data were collected using Nutrition Data System for Research software versions 2005 and 2007, developed by the Nutrition Coordinating Center, University of Minnesota, Minneapolis, MN. Final calculations were completed using NDSR version 2012. The NDSR time-related database updates analytic data while maintaining nutrient profiles true to the version used for data collection.

Postprandial plasma was obtained during steady-state tracer infusion protocols after 8 administrations of hourly portions of a nutritive formula containing protein, carbohydrate, and fat (9,12). As previously reported (9), hourly administration of this formula beginning 2 h before the infusion maintained a postprandial state. The formula provided a balanced pattern of amino acids at a rate based on requirements of 0.8 g protein \cdot $\text{kg}^{-1} \cdot \text{d}^{-1}$, equivalent to an hourly protein dose of 0.03 g/kg with 5.23 and 5.44 kJ \cdot $\text{kg}^{-1} \cdot \text{d}^{-1}$ for women and men, respectively. The formula also provided an adequate energy intake according to the requirements of 126 and 130 kJ \cdot $\text{kg}^{-1} \cdot \text{d}^{-1}$ for women and men, respectively.

Blood sample collection. Preprandial and postprandial blood samples were collected in EDTA tubes at the end of the 2-d controlled diet and after the 28-d vitamin B-6-restricted diet. Within 2 h of collection, plasma was separated by centrifuging whole blood at $1650 \times g$ for 15 min at 4°C and then stored at -80°C for later analysis.

Analytical methods

Plasma PLP was determined as the semicarbazone-derivative by reverse-phase HPLC with fluorescence detection (55). Aminothiols were measured as ammonium 7-fluorobenzo-2-oxa-1,3-diazole-4-sulfonate derivatives by reverse-phase HPLC with fluorescence detection (56).

A panel of 16 one-carbon metabolites and related compounds, including betaine, dimethylglycine (DMG), serine, glycine, methionine, methionine sulfoxide, threonine, choline, guanidinoacetate, creatine, creatinine, cystathionine, arginine, leucine, symmetrical dimethylarginine, and symmetrical dimethylarginine, was quantified by LC-tandem MS (MS/MS) at the Biomedical Mass Spectrometry Laboratory, Clinical and Translational Sciences Institute, University of Florida, which was a modification of the methods of Ueland et al. (57). In this analysis, the internal standard mixture (20 μL) was added to 60 μL of plasma and mixed by vortexing. Proteins were precipitated by adding 3 volumes of methanol, mixing, and then storing at 4°C for 30 min, followed by centrifuging at $2000 \times g$ for 10 min at 4°C . The supernatant was removed and analyzed (10 μL sample injection) by LC-MS/MS. Samples were chromatographed on a Luna C18 150×4.6 -mm column (Phenomenex) and eluted in 0.02% heptafluorobutyric acid using a methanol gradient at 0.6 mL/min. To achieve retention of the early eluting compounds, the column was equilibrated for 3 min with 0.1% heptafluorobutyric acid between injections. Samples were analyzed in ESI mode using a TSQ Quantum triple quadrupole mass spectrometer (Thermo-Finnigan). Analysis was performed in positive ionization mode (capillary temperature, 300°C ; vaporizer temperature, 250°C ; sheath gas pressure, 50 arbitrary units; auxiliary gas pressure, 20 arbitrary units; collision gas pressure, 1.5 mTorr) using selected reaction monitoring with the m/z transitions shown (Supplemental Table 1). Analyte concentrations were determined by calculating the analyte:internal standard peak area ratios and comparing them with standard concentration curves prepared using authentic standards.

Quantitative analysis of constituents of tryptophan catabolism, including tryptophan, kynurenine, kynurenic acid, anthranilic acid, 3-hydroxykynurenine, xanthurenic acid, 3-hydroxyanthranilic acid, quinolinic acid, N^1 -methyl-nicotinamide, nicotinamide, and nicotinic acid, was performed by LC-MS/MS (58) at Bevitall.

To assess any inflammatory response in this dietary vitamin B-6 restriction protocol, 3 markers of inflammation were evaluated. Neopterin was determined concurrently with the tryptophan metabolites, and the kynurenine:tryptophan ratio also was derived from results of that analysis (58). We also measured C-reactive protein by ELISA in preprandial plasma samples (High Sensitivity C-Reactive Protein kit no. CR120C, Calbiotech).

Statistical analysis

All data are presented as means \pm SDs. MANOVA was used to evaluate significance of the overall change across all targeted metabolites from the LC-MS/MS analyses of one-carbon metabolites and tryptophan metabolites for the fasting and fed data separately (59). Differences in concentration after vitamin B-6 restriction were determined after log base 2 transformation to meet the Gaussian requirement. If a significant overall effect was found, stepdown tests with adjustment for multiple testing were performed. For one-carbon metabolites, the resulting adjusted P values were computed with methods that have control of the false discovery rate (60). For the tryptophan metabolites, the adjusted P values were calculated with Sidak adjustment (61). The main effect of the vitamin B-6 restriction protocol on the plasma ratios of cystathionine:creatinine, 3-hydroxykynurenine:kynurenic acid (potential biomarkers of vitamin B-6 deficiency), and kynurenine:tryptophan (an indicator of inflammatory response) were evaluated using the 2-way repeated-measures ANOVA. A 2-sample t test was used to assess differences between men and women on the change of plasma PLP from the restriction protocol. The target level of significance was set to 0.05 or lower. All data analyses were performed using SAS 9.3 software.

Multivariate evaluation of overall effects of the vitamin B-6 restriction protocol on the combined profiles of constituents of one-carbon metabolism and tryptophan catabolism pathways was conducted and the results were visualized using multilevel partial least squares-discriminant analysis, which accounts for the paired structure of the data (62). For this multilevel analysis, we processed the aggregate metabolite profile data according to Westerhuis et al. (62) using Microsoft Excel and then conducted partial least squares-discriminant analysis using SIMCA version 13 (MKS Umetrics). The effect of vitamin B-6 restriction on overall metabolite profiles was visualized with score plots and discriminatory variables (i. e., responding metabolites) were identified using Variable Influence on Projection (VIP) plots in SIMCA version 13 (63). The VIP values in these plots represent weighted sums of squares of the PLS weights of the various metabolites and take into account the proportion of Y-variance in each dimension (64). In the VIP plots, each variable is provided with a VIP value and a 95% CI derived from jack-knifing (64–66). Variables for which the $\text{VIP} \pm 95\%$ CI exceeds 1 are designated as significant biomarkers in this analysis.

Results

Summary of participants and effect of vitamin B-6 restriction protocol. All participants had serum folate, vitamin B-12, and plasma PLP and total homocysteine in the normal range at the beginning of the study (9,12) (Table 1). The dietary vitamin B-6 restriction protocol lowered the plasma PLP concentration from 52 ± 14 to 21 ± 5 nmol/L ($P < 0.05$), consistent with induction of marginal deficiency (9,12,54). Following restriction, the PLP values ranged from 12.3 to 29.3 nmol/L. Plasma total homocysteine and cysteine concentrations were not changed by the 28-d vitamin B-6 restriction (9,12,54).

The 2-sample t test on the change of PLP revealed no difference between male and female participants ($P = 0.11$). The average decrease for males was 35.8 nmol/L with a SD of 12.8 nmol/L; the average decrease for females was 25.9 nmol/L with a SD of 15.4 nmol/L.

One-carbon metabolism. Targeted analysis of one-carbon metabolites in both preprandial and postprandial plasma successfully quantified a panel of 16 compounds (Table 2). MANOVA testing showed an effect of vitamin B-6 deficiency on the overall profile in both preprandial and postprandial states ($P < 0.05$). MANOVA also indicated that the pattern of changes in metabolite concentration in preprandial was different from that of postprandial conditions ($P < 0.01$), but the differences were not due to changes in concentration of any individual metabolite. After adjusting for multiple comparisons using the false discovery

TABLE 1 Baseline characteristics of adults who completed the 28-d dietary vitamin B-6 restriction¹

	Men	Women
<i>n</i>	12	11
Age, <i>y</i>	24 ± 5 ²	25 ± 6
BMI, <i>kg/m</i> ²	24.5 ± 2.7	23.6 ± 2.3
Plasma PLP, <i>nmol/L</i>	59 ± 13	45 ± 13*
Serum folate, <i>nmol/L</i>	34 ± 8	30 ± 7
Serum vitamin B-12, <i>pmol/L</i>	349 ± 115	388 ± 129
Plasma homocysteine, <i>μmol/L</i>	7.5 ± 1.0	6.4 ± 1.3

¹ Data previously reported (9,12,13,80). *Different from men, *P* < 0.05 (2-sample *t* test). PLP, pyridoxal 5'-phosphate.

² Mean ± SD (all such values).

rate method, preprandial plasma concentrations of serine (12.4% increase, adjusted *P* < 0.05), creatine (40.2% decrease, adjusted *P* < 0.0001), and cystathionine (53.3% increase, adjusted *P* < 0.0001) were significantly different after the 28-d vitamin B-6 restriction compared with baseline (Table 2), whereas the changes in preprandial concentrations of creatinine (7.7% decrease, adjusted *P* = 0.09), glycine (14.8% increase, adjusted *P* = 0.09), threonine (15.5% increase, adjusted *P* = 0.09), and DMG (11.3% decrease, adjusted *P* = 0.09) approached significance. In the postprandial state, DMG (16% decrease, adjusted *P* < 0.05) and creatinine (8.4% decrease, adjusted *P* < 0.05), and creatine (40.1% decrease, adjusted *P* < 0.05) and cystathionine (76.4% increase, adjusted *P* < 0.001) plasma concentrations significantly differed in the vitamin B-6–restricted state compared with adequate status (Table 2).

Tryptophan catabolism. LC-MS/MS provided a measurement of 11 constituents of the tryptophan catabolic pathway. MANOVA analysis of data from both preprandial and postprandial states showed effects of vitamin B-6 restriction on the tryptophan metabolite profile (*P* < 0.0002). In this analysis, the magnitude of change of plasma PLP did not have an overall effect (preprandial,

P = 0.149; postprandial, *P* = 0.596) (Table 3). Vitamin B-6 restriction induced the following changes in tryptophan catabolites when assessed under preprandial conditions: kynurenic acid (25.7% decrease, adjusted *P* = 0.0019), 3-hydroxykynurenine (32.2% increase, adjusted *P* = 0.0031), and nicotinic acid (15.8% decrease, adjusted *P* = 0.017). When the effects of vitamin B-6 restriction were evaluated in postprandial plasma, kynurenine acid (22.6% decrease, adjusted *P* = 0.0013) and 3-hydroxykynurenine (28.9% increase, adjusted *P* = 0.0088) changed significantly, but the change in nicotinic acid (10.3% decrease, adjusted *P* = 0.546) was not significant (Table 3).

Vitamin B-6 effects on biomarkers of inflammation. Three indicators of inflammatory response were examined in plasma obtained under both preprandial and postprandial conditions before and after the 28-d vitamin B-6 restriction (Table 4). C-reactive protein, which is an acute-phase protein indicative of inflammation, exhibited a small decrease (*P* = 0.002). Neopterin, which is a pterin indicative of immune activation, exhibited a small increase due to the restriction protocol (*P* = 0.012). The kynurenine:tryptophan ratio, a marker of cellular immune response, exhibited a small decrease due to the B-6 restriction (*P* = 0.013) with a small but positive effect of postprandial status (*P* = 0.008).

Ratios of one-carbon and tryptophan metabolites as biomarkers of vitamin B-6 deficiency. We investigated several plasma product:precursor ratios to probe potential effects of vitamin B-6 restriction on cellular processes (Table 5). The ratio of glycine:serine, reflecting relative rates of glycine cleavage and SHMT reactions, increased in the postprandial state but was not affected by vitamin B-6 status. Vitamin B-6 restriction did not affect the betaine:choline or DMG:betaine ratios in either preprandial or postprandial conditions. The ratio of guanidinoacetic acid:creatine, which reflects the choline synthesis process catalyzed by guanidinoacetate *N*-methyltransferase, increased in vitamin B-6 restriction in both preprandial and postprandial samples and paralleled creatine concentration. The ratio of creatine:cystathionine, which are not directly related

TABLE 2 Concentrations of one-carbon metabolites quantified in plasma of human participants in the preprandial and postprandial states before and after a 28-d dietary vitamin B-6 restriction¹

	Preprandial		Postprandial	
	Baseline	B-6 restricted	Baseline	B-6 restricted
	<i>μmol/L</i>		<i>μmol/L</i>	
Betaine	44.3 ± 15.3	39.0 ± 10.6	39.2 ± 12.2	34.3 ± 9.50
Choline	9.14 ± 2.48	8.57 ± 1.8	8.00 ± 1.70	7.10 ± 1.30
DMG	2.65 ± 0.65	2.35 ± 0.62	2.50 ± 0.60	†2.10 ± 0.40
Serine	113 ± 23.5	†127 ± 38.0	110 ± 24.0	116 ± 27.1
Glycine	277 ± 78.8	318 ± 99.6	283 ± 80.3	308 ± 70.5
Threonine	155 ± 34.2	179 ± 51.1	155 ± 35.7	172 ± 42.7
Guanidinoacetic acid	3.18 ± 1.23	3.00 ± 1.00	2.60 ± 0.60	2.50 ± 0.60
Creatine	31.1 ± 13.7	†18.6 ± 6.36	32.2 ± 12.1	†19.3 ± 6.70
Creatinine	80.2 ± 15.0	74.0 ± 16.9	69.0 ± 10.0	†63.2 ± 9.20
Cystathionine	0.15 ± 0.06	†0.23 ± 0.08	0.17 ± 0.09	†0.30 ± 0.12
Methionine	28.9 ± 6.32	28.1 ± 5.57	30.0 ± 5.40	29.9 ± 5.80
Methionine sulfoxide	0.95 ± 0.31	0.98 ± 0.28	1.00 ± 0.30	1.00 ± 0.30
Leucine	144 ± 36.8	132 ± 27.0	128 ± 23.9	123 ± 18.7
Arginine	105 ± 28.5	96.3 ± 24.7	95.1 ± 17.1	88.8 ± 13.3
Symmetric dimethyl arginine	0.66 ± 0.12	0.64 ± 0.12	0.59 ± 0.06	0.57 ± 0.08
ADMA	0.75 ± 0.15	0.69 ± 0.15	0.66 ± 0.11	0.62 ± 0.10

¹ Values are means ± SDs, *n* = 23. Significant difference from baseline designated by: †adjusted *P* < 0.05, ‡adjusted *P* < 0.001. The adjustment was based on the false discovery rate method. ADMA, asymmetric dimethylarginine; DMG, dimethylglycine.

TABLE 3 Concentrations of tryptophan metabolites quantified in plasma of human participants in the preprandial and postprandial states before and after a 28-d dietary vitamin B-6 restriction¹

	Preprandial		Postprandial	
	Baseline	B-6 restricted	Baseline	B-6 restricted
Tryptophan, $\mu\text{mol/L}$	61.1 \pm 14.1	69.3 \pm 17.8 [#]	62.3 \pm 11.4	66.5 \pm 14.1
Kynurenine, $\mu\text{mol/L}$	1.15 \pm 0.244	1.18 \pm 0.267	1.25 \pm 0.237	1.22 \pm 0.196
Kynurenic acid, nmol/L	30.7 \pm 10.1	22.8 \pm 7.68 [†]	32.8 \pm 8.10	25.4 \pm 7.48 [†]
Anthranilic acid, nmol/L	14.5 \pm 4.32	13.5 \pm 3.44	13.4 \pm 3.36	13.6 \pm 4.87
3-Hydroxykynurenine, nmol/L	24.5 \pm 9.46	32.4 \pm 11.0 [‡]	27.7 \pm 11.8	35.7 \pm 15.9 [‡]
Xanthurenic acid, nmol/L	8.44 \pm 4.43	7.05 \pm 2.43	12.9 \pm 7.02	11.7 \pm 6.00
3-Hydroxyanthranilic acid, nmol/L	19.9 \pm 6.83	18.1 \pm 6.46	24.0 \pm 8.47	22.3 \pm 6.85
Quinolinic acid, nmol/L	253 \pm 59.6	274 \pm 55.2 [@]	264 \pm 63.8	250 \pm 46.4
Nicotinic acid, nmol/L	77.6 \pm 16.4	65.3 \pm 19.1 [‡]	64.8 \pm 14.5	58.1 \pm 8.59
Nicotinamide, nmol/L	292 \pm 184	303 \pm 238	292 \pm 152	272 \pm 147
N ¹ -methylnicotinamide, nmol/L	157 \pm 84.5	146 \pm 79.2	159 \pm 99.6	124 \pm 70.3

¹ Values are means \pm SDs, $n = 23$. Significant difference from baseline designated by: [†], adjusted $P < 0.05$; [‡], adjusted $P < 0.001$. The adjustment was based on the Sidak method. Trend toward significant difference from baseline designated by: [#], adjusted $P = 0.073$; [@], adjusted $P = 0.062$.

metabolically, appears to provide a very sensitive combined biomarker of marginal vitamin B-6 deficiency (Table 5).

Two of the metabolite ratios provided evidence of physiological status. The ADMA:arginine inversely reflects NO synthetase substrate function and ADMA as a competitive inhibitor. In this study, vitamin B-6 restriction yielded an increased ADMA:arginine ratio ($P < 0.001$) in both pre- and postprandial conditions. The ratio of methionine sulfoxide:methionine, which may be an indicator of oxidative stress, did not change significantly due to vitamin B-6 restriction in either pre- or postprandial samples.

Graphical multivariate analysis of pooled one-carbon and tryptophan catabolite profile data. The overall effect of vitamin B-6 restriction on this collection of various constituents of one-carbon metabolism and tryptophan catabolic processes was evaluated further by a multilevel PLS-DA approach (62), which is a graphical method for visualizing overall treatment effects with paired data. The overall metabolite profiles effectively grouped the data according to vitamin B-6 status (i.e., basal vs. restricted) when conducted for preprandial and postprandial samples (Fig. 1A,B). In this analysis of samples in the preprandial state, the first principal component accounted for 85% of the variation modeled in Y and the second component accounted for 5.8%, while 88% of the variation could be predicted by the model. Similarly, for samples in the postprandial state, the first principal component accounted for 85% of the overall variation modeled in Y and the second component accounted for 6.9%, with 88% of the variation predicted by the model. The VIP plot served as an effective means of ranking the variables (metabolites) for biomarker identification (Fig. 2A,B). Using VIP analysis for profiles from preprandial samples, we identified creatine, cystathionine, 3-hydroxykynurenine, and kynurenic acid as the most important terms in the model

(i.e., significant discriminating biomarkers of vitamin B-6 restriction). For profiles from postprandial samples, VIP analysis identified creatine, cystathionine, kynurenic acid, 3-hydroxykynurenine, and DMG as significant discriminating biomarkers.

Discussion

Marginal vitamin B-6 deficiency has been associated with an increased risk of cardiovascular disease and certain cancers (67–69). The dietary protocol used in these studies serves as an effective tool in inducing short-term, marginal vitamin B-6 deficiency in healthy young adults (7–9,12,70). Whether the changes observed in studies of this type successfully predict the metabolic effects of chronic marginal vitamin B-6 deficiency remains to be determined. An important strength of these studies is their focus on the specific nutritional effects of controlled vitamin B-6 deficiency without inflammatory conditions, which often are associated with a low plasma PLP concentration (23,25). The current work augments recent studies of metabolite profiling of vitamin B-6 status in population-based studies. The results of this study confirm and extend our previous conclusion (8) that short-term vitamin B-6 deficiency in healthy young adults is not, in itself, an inflammatory condition (Table 4). The extent to which dietary vitamin B-6 insufficiency contributes to inflammation in various population subgroups (23,25,71,72) remains unclear.

Because PLP serves as a coenzyme for many aspects of metabolism, there are many possible functional impacts of vitamin B-6 deficiency. We recently published evidence from a metabolomic analysis of the same samples used in the current study suggesting wide-ranging effects of vitamin B-6 restriction, including perturbations of amino acid, lipid, and organic acid profiles (54). However, prediction of the effect of marginal

TABLE 4 Indicators of inflammatory status in plasma of human participants in the preprandial and postprandial states before and after a 28-d dietary vitamin B-6 restriction¹

	Preprandial		Postprandial		P value		
	Baseline	B-6 restricted	Baseline	B-6 restricted	B-6 restriction	Preprandial status	Interaction
C-reactive protein, mg/L	2.66 \pm 3.56	1.00 \pm 1.74*	2.41 \pm 3.15	0.947 \pm 1.36*	0.002	0.236	0.618
Neopterin, nmol/L	6.15 \pm 1.15	6.95 \pm 1.46*	6.16 \pm 1.28	6.69 \pm 2.08*	0.012	0.357	0.278
Kynurenine/tryptophan	0.0196 \pm 0.0057	0.0175 \pm 0.0040*	0.0207 \pm 0.0057	0.0190 \pm 0.0044*	0.013	0.008	0.640

¹ Values are means \pm SDs, $n = 23$. *Different from baseline, $P < 0.05$. P values are from repeated-measures 2-way ANOVA.

TABLE 5 Ratios of selected one-carbon and tryptophan metabolites quantified in plasma of human participants in the preprandial and postprandial states before and after a 28-d dietary vitamin B-6 restriction¹

	Preprandial		Postprandial		P value		
	Baseline	B-6 restricted	Baseline	B-6 restricted	B-6 restriction	Preprandial status	Interaction
Product:precursor ratios							
Glycine:serine	2.49 ± 0.608	2.54 ± 0.522	2.62 ± 0.571	2.73 ± 0.573	0.153	0.021	0.795
Betaine:choline	5.09 ± 1.99	4.59 ± 0.972	5.13 ± 1.75	4.86 ± 1.26	0.513	0.252	0.575
DMG:betaine	0.0651 ± 0.0216	0.0620 ± 0.0161	0.0674 ± 0.0245	0.0633 ± 0.0151	0.486	0.163	0.765
Creatine:guanidinoacetic acid	11.3 ± 6.65	6.51 ± 2.33*	13.9 ± 6.88	8.28 ± 3.07*	<0.001	<0.001	0.91
ADMA:arginine	0.00822 ± 0.00422	0.00732 ± 0.00118*	0.00761 ± 0.00176	0.00702 ± 0.00117*	0.005	0.22	0.74
Other biomarker ratios							
Creatine:cystathionine	229 ± 94.5	85.6 ± 31.1*	214 ± 91.0	74.9 ± 34.3*	<0.001	0.056	0.26
3-Hydroxykynurenine: kynurenic acid	0.903 ± 0.244	1.56 ± 0.759*	0.900 ± 0.530	1.47 ± 0.735*	<0.001	0.67	0.35

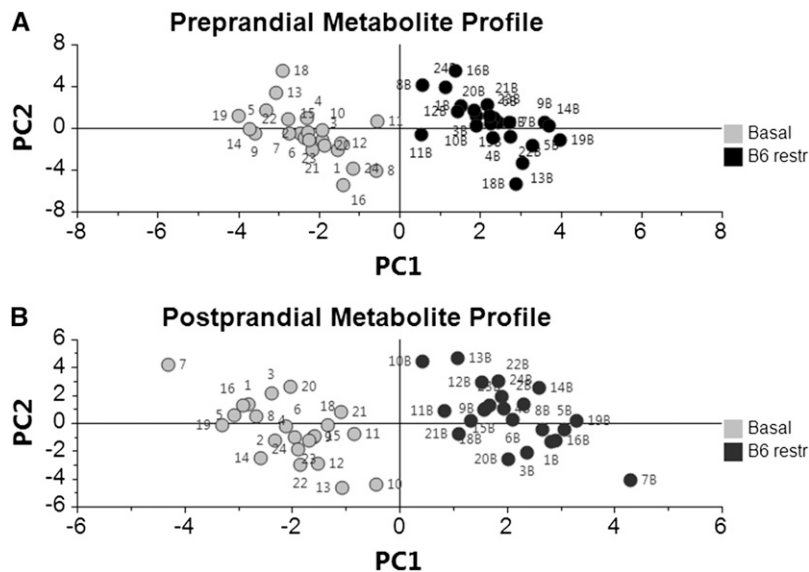
¹ Values are means ± SDs, *n* = 23. *Different from baseline, *P* < 0.05. *P* values are from repeated-measures 2-way ANOVA. ADMA, asymmetric dimethylarginine; DMG, dimethylglycine.

deficiency on specific PLP-dependent enzymes on in vivo fluxes and metabolite concentrations has not been successful. For example, marginal deficiency of vitamin B-6 yields little or no change of in vivo SHMT flux (7,9) and only subtle changes in serine and glycine concentrations (7,9,54), as supported by mathematical modeling (11) and observations of this study (Tables 2 and 4). The trans-sulfuration pathway of homocysteine catabolism, comprised of the PLP-dependent enzymes CBS and CSE, exhibits differential effects of PLP insufficiency. In this case, CBS tends to maintain its activity and in vivo flux (8,73), whereas CSE is sensitive to loss of activity with an increased proportion of apo-CSE and loss of total CSE activity. However, CSE in vivo flux is maintained by the great increase in cystathionine, which, relative to the high *K_m* of this enzyme, allows net in vivo flux of the trans-sulfuration pathway to be maintained (8,12,73). Consequently, as shown here and previously, elevated cystathionine is a sensitive biomarker for marginal vitamin B-6 deficiency, whereas cysteine flux (8,11) and cysteine concentration are depressed only in more severe deficiency (8,11,12,21,73,74).

Several additional findings of interest arose from our metabolite profile analysis. Plasma methionine sulfoxide concentrations and the methionine sulfoxide:methionine ratio (Tables 2 and 5) were not affected by vitamin B-6 restriction, nor were

plasma glutathione concentrations (10). Methionine sulfoxide may be formed from oxidation of methionine by reactive oxygen species and thus may serve as an indicator of oxidative stress damage (75). Therefore, the marginal vitamin B-6 deficiency induced in our human participants did not appear to increase oxidative stress. The ADMA:arginine ratio is another unexpected example. Because ADMA arises from the turnover of methylated proteins, the formation of ADMA is not directly related to PLP-dependent processes. However, vitamin B-6 restriction was associated with a reduction in the ADMA:arginine ratio. A reduction in the ADMA:arginine ratio would be expected to improve endothelial function, because ADMA is a competitive inhibitor of endothelial NO synthase-derived NO production (76). The reduction in the ADMA:arginine ratio appears to be in contrast with the association of marginal vitamin B-6 deficiency with a variety of forms of vascular disease (3–6). Perhaps the most unexpected change in the product:precursor ratios was creatine:guanidinoacetate, which reflects the *S*-adenosylmethionine-dependent guanidinoacetate methyltransferase reaction. Because guanidinoacetate methyltransferase is not PLP dependent and neither guanidinoacetate production nor creatine catabolism involves PLP-dependent steps in metabolic proximity, the mechanism responsible for the association of vitamin B-6 restriction with a lower creatine:guanidinoacetate ratio is

FIGURE 1 Score plots from multilevel partial least squares-discriminant analysis of the overall one-carbon metabolism and tryptophan catabolism data sets from analysis of plasma sampled before and after 28-d vitamin B-6 restriction. (A) Analysis of metabolite profiles of preprandial plasma. (B) Analysis of metabolite profiles of postprandial plasma. Each data point represents a function of the entire metabolite profile of a human participant (numerical designations provided); e.g., 1 and 1b represent data points for participant 1 in basal and vitamin B-6 restricted states. PC1, principal component 1; PC2, principal component 2; restr, restricted.



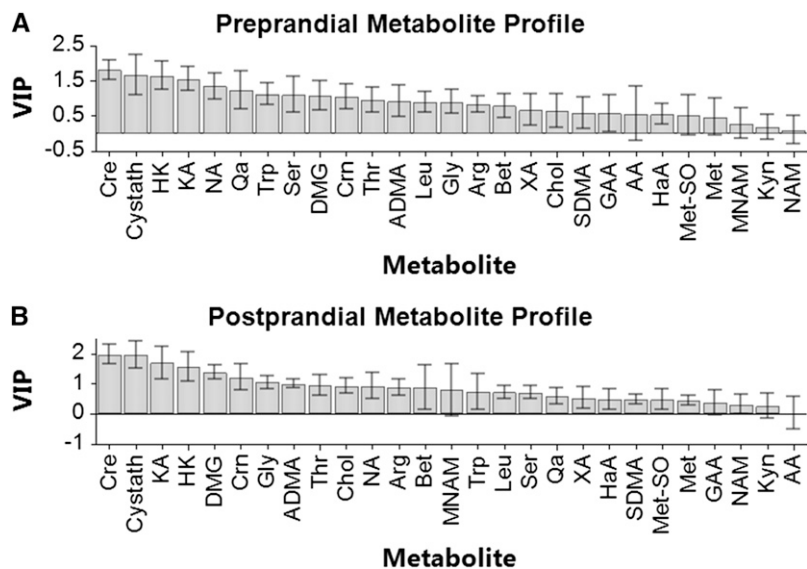


FIGURE 2 VIP plots from multilevel partial least squares-discriminant analysis of the overall one-carbon metabolism and tryptophan catabolism data sets from analysis of plasma sampled before and after 28-d vitamin B-6 restriction. (A) Analysis of metabolite profiles of preprandial plasma. (B) Analysis of metabolite profiles of postprandial plasma. For each metabolite in the VIP analysis, the bar represents a weighted sum of squares with CI. A variable is considered to respond significantly if the VIP value and its 95% CI exceeds 1. AA, anthranilic acid; ADMA, asymmetric dimethylarginine; Arg, arginine; Bet, betaine; Chol, choline; Cre, creatine; Crn, creatinine; Cystath, cystathionine; DMG, dimethylglycine; GAA, Guanidinoacetic acid; Gly, glycine; HaA, 3-hydroxyanthranilic acid; HK, 3-hydroxykynurenine; KA, kynurenic acid; Kyn, kynurenine; Leu, Leucine; Met, methionine; Met-SO, methionine sulfoxide; MNAM, N¹-methylnicotinamide; NA, nicotinic acid; NAM, nicotinamide; Qa, quinolinic acid; SDMA, symmetric dimethylarginine; Ser, serine; Thr, threonine; Trp, tryptophan; VIP, variable influence in the projection (63); XA, xanthurenic acid.

unclear. In fasted participants, creatine is the major product of the methylation cycle (77), and the methylation demand of creatine synthesis is second only to PC synthesis in fed participants. Therefore, decreasing creatine synthesis may spare 1C-groups (or more specifically methyl-groups) in the face of vitamin B-6 restriction. Severe vitamin B-6 deficiency (78) in rats yields a reduction in hepatic SAM concentration and decrease in the SAM:SAH ratio. However, such an effect appears unlikely in this study in view of the lack of effect on plasma methionine, homocysteine, or N¹-methylnicotinamide (which is formed by methylation of nicotinamide by SAM) concentrations. It is possible that differences in the creatine content of the participants' prestudy diet and the controlled vitamin B-6 restriction diet could contribute to the observed results. However, we think that observations demonstrate a physiologically real phenomenon in light of our findings of a similar reduction in creatine in HepG2 cells cultured in low vitamin B-6 conditions (V.R. da Silva and J.F. Gregory III, unpublished results).

Our metabolite profiles demonstrate that the tryptophan catabolic pathway is perturbed during marginal vitamin B-6 deficiency. These results support our mathematical modeling (51) and animal studies (79) as well as a recent large-scale study of low vitamin B-6 status in cardiovascular patients with systemic inflammation (52). Our results of this study and the previous findings of Midttun et al. (52), showing increased 3-hydroxykynurenine and decreased kynurenic acid as well as an increased 3-hydroxykynurenine:kynurenic acid ratio, strongly suggest that kynureninase is more susceptible than kynurenine aminotransferase to PLP insufficiency, as previously proposed (47). The mechanisms and metabolic consequences of the association of systemic inflammation, low plasma PLP, and altered tryptophan catabolism, as recently reviewed (23), require further study.

Our multivariate analysis of the pooled metabolites associated with one-carbon metabolism and tryptophan catabolism has provided important leads in the development of better functional diagnostic criteria for marginal vitamin B-6 deficiency. In view of well-documented associations of marginal vitamin B-6 deficiency with disease risk, we propose that biomarkers such as cystathionine, 3-hydroxykynurenine, kynurenic acid, and creatine, especially in combination, may provide useful new assessment tools.

Acknowledgments

J.F.G. and P.W.S. designed the research; V.R.d.S., L.R.-A., Y.L., M.A.R., E.P.Q., T.J.G., B.C., M.N.S., S.S.P., Ø.M., and P.M.U. conducted the research; Y.-Y.C., K.E.M., and J.F.G. analyzed the data; and V.R.d.S. and J.F.G. wrote the paper. All authors read and approved the final manuscript.

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