BRIEF PAPER

Choline metabolite, trimethylamine N-oxide (TMAO), is associated with inflammation in psoriatic arthritis

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ABSTRACT

Objective. Dietary intake of choline has been linked to systemic inflammation through the microbial production of two metabolites, trimethylamine (TMA) and trimethylamine-N-oxide (TMAO). Herein we explore the association between choline metabolites and inflammation in psoriatic arthritis (PsA) patients.

Methods. Thirty-eight patients with PsA, all of whom satisfied the CAS-PAR classification criteria for PsA, were studied. Outcomes reflecting the activity of peripheral arthritis as well as skin psoriasis, Disease Activity Score (DAS)28, Clinical Disease Index (CDAI) and Body Surface Area (BSA) were assessed. Serum concentration of choline metabolites (choline, TMA, TMAO, betaine and carnitine) were determined by LC-MS, and metabolite levels associated with disease scores.

Results. Among the 38 PsA patients included, the mean DAS28PCR was 2.74 ± 1.29 . Twenty-seven patients had active skin disease, with an average BSA of 7.2 ± 16.22 . TMAO, but not TMA or choline, significantly correlated with measures of disease activity for both skin and peripheral joints.

Conclusion. In our cohort, only TMAO, but not TMA, choline, betaine or carnitine, was associated with inflammation in PsA patients, establishing a mechanistic link between TMAO and PsA phenotypes. Future studies will explore the modulation of TMAO and disease severity in PsA.

Introduction

Choline metabolism has been recently strongly related to inflammation in relation to the pathogenesis of atherosclerosis (1). Choline is a semi-essential nutrient found in a variety of foods, but it is particularly abundant in egg yolk, meats, liver, fish, dairy products, nuts, and soybean (2). Choline is a component of phosphatidylcholine, and following oxidation, precursor to betaine (3). Choline and other trimethylamine (TMA) containing species like carnitine are metabolised to TMA by the gut microbiota (4). TMA is subsequently oxidised by at least one member of the flavin-containing monooxygenases, FMO3, forming trimethylamine-N-

oxide (TMAO), which is then released into circulation (4). Several studies have shown that an elevated level of choline, betaine and TMAO increases the risk of atherosclerosis and subsequently of cardiovascular disease (CVD) (5). Also, a recent study revealed a link between the TMAO-producing enzyme FMO3 and obesity (6, 7).

Psoriatic arthritis (PsA) is an inflammatory disease affecting the joints and connective tissue and is associated with psoriasis of the skin and nails. Among the risk factors for psoriasis, evidence is accumulating that nutrition plays a major role in psoriasis pathogenesis. In particular, body weight, nutrition, and diet may exacerbate the clinical manifestations (8). Obesity is a known risk factor for developing PsA in patients with psoriasis and several studies have described that a successful weight loss might be associated with a higher rate of achievement of minimal disease activity in overweight/obese patients with PsA who started treatment with TNF- α blockers (9). Metabolic syndrome and high CVD risk are also more frequent in PsA than other rheumatic diseases such as rheumatoid arthritis (10). As choline metabolism is associated to all these comorbidities, we examined the relationship between circulating levels of choline, betaine, carnitine, TMA and TMAO, and skin and joint clinical scores in a cohort of PsA patients.

Methods

Patients

A cross-sectional cohort of 38 adult patients with PsA fulfilling the classification for PsA (CASPAR) criteria was recruited from the University of California San Diego (UCSD) Arthritis Clinics. The study was approved by the UCSD Institutional Review Board (no. 150272) and obtained the patient's written informed consent to publish the material. Clinical assessment included: evaluation of the number of tender (TJC) and swollen joints (SJC) (out of 28), the number of tender entheseal sites, the percentage of the body surface affected by psoriasis, functional status as assessed by Health Assessment Questionnaire (HAQ), and assessments of pain, fatigue, global disease severity

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A	B	5			
Characteristic N/Mean (%/SE	D)	Metabolite		Mean (SD)	
Male 22 (58%)		Choline		47.45 (35.96)	
Age 50.8 (10.89)		TMA		45.08 (37.87)	
Duration of PsA in years 10 (9.41)		TMAO		15.96 (15.51)	
Duration of psoriasis in years 16.9 (13.94)		Betaine		792.83 (295.38)	
Number of tender joints 1.9 (range: 0-10)		Carnitine		163.06 (34.71)	
Number of swollen joints 2 (range: 0-10)					
Pain 3.12 (2.84)					
HAQ (Health Assessment Questionnaire) 0.47 (0.57))			
DAS28CRP 2.74 (1.29) CDAI 9.91 (10.04)		Metabolite	Male Mean (SD)	Female Mean(SD)	р
SDAI 11.93 (11.73))	Choline	43.5 (30.1)	52.89 (43.23)	0.46
Enthesitis 8 (21%)	·	TMA	41.9 (32.5)	49.35 (45.01)	0.58
BMI (Body Mass Index) 27.1 (5.42)		TMAO	13.2 (9.7)	19.76 (20.84)	0.25
Body surface affected by psoriasis 7.2 (16.22)		Betaine	887 (314.9)	663.3 (213.21)	0.01
		Carnitine	172.1 (33.3)	150.63 (33.63)	0.06
Carnitine	014	Metabolite	Age ≤ 51 Mean (SD)	Age >51 Mean (SD)	р
	Carnitine	Choline	41.4 (23.6)	53.4 (44.9)	0.31
Betaine		TMA	39.2 (23.7)	50.9 (48)	0.35
Betalle	Betaine	TMAO	12.2 (8)	19.6 (20)	0.14
		Betaine	900.7 (234.5)	684.8 (315.7)	0.02
TMAO	TMAO	Carnitine	167.2 (27.3)	158.8 (41)	0.46
TMA	тма	Metabolite	BMI < 25 Mean (SD)	BMI > 25 Mean (SD)	p
		Choline	55.4 (50.31)	45.3 (31.9)	0.71
Choline	Choline	TMA	53.8 (51.4)	42.7 (35.6)	0.65
amitine setaine TMAO TMA choline	10 0	TMAO	13.6 (15.4)	17.9 (17.3)	0.31
Carnitine Betaine TMAO TMA Choine Carnitine Betaine TMAO	rMA choline	Betaine	697.1 (307.4)	826.6 (320.3)	0.48
Color Key Color Key Color Key		Carnitine	143 (30.2)	164.6 (31.2)	0.12

0 0.02 0.04 0.06 0.08

Fig. 1. Cohort demographics and metabolites clustering.

A: Demographic and disease characteristics of the patients (n=38). B: The average value of the metabolites in our cohort of 38 patients is displayed in Fig. 1B. C: Linear regression was performed between each metabolite – metabolite pair, controlling for age and gender.

Left: The regression coefficients for each pair were used to form a clustered heatmap. TMA: trimethylamine; TMAO: trimethylamine N-oxide. Right: Metabolite regression *p*-values are displayed, where the row and column order are preserved from 1C left.

D: The average value of metabolites per age, gender and body max index (BMI) are displayed. T-test p-values are also displayed. Concentrations are provided as ng/ml.

by patients, and a global assessment of disease by physicians, using a Visual Analogue Scale ranging from 0 to 10. Composite measures of peripheral arthritis were calculated using the above measures: Disease Assessment Score using a 28 joint count and C reactive protein (DAS28-CRP), Clinical Disease Activity Index (CDAI) and Simple Disease Activity Index (SDAI). Blood samples were processed immediately, and sera aliquots were stored at -80°C until analysis.

Metabolite measure

Choline, TMA, TMAO, betaine and carnitine were assayed in serum samples using liquid chromatography coupled to tandem mass spectrometry (LC-MS/MS) as detailed in supplementary methods.

Data analysis

The data, consisting of 38 patient samples measured across 5 choline metabolites and 14 clinical outcomes were processed using R, v. 3.4.1. (www.r-project.org) as detailed in Supplementary methods.

Results

Cohort demographics and disease characteristics

Patient characteristics are summarised in Figure 1A. Of the 38 PsA patients included in this study, 58% were male and the mean age was 50.8 years ± 10.9 (range 23-75 years). The mean number of tender joints and swollen joints were 1.9±3 (range 0-10) and 2±3.2 (range 0-10) respectively. The average DAS28-CRP score was 2.74±1.29 (range 1-5.18). Twenty-five patients (71%) had active skin disease, with an average BSA of 7.22±16.22. Eight patients had enthesitis. 65% were receiving biological therapy (46% of them in association with a synthetic diseasemodifying drug (sDMARD), mostly methotrexate. 14% received sDMARD as monotherapy, and 21% received no systemic treatment.

Choline metabolite profiling and clustering

Concentration of circulating betaine, carnitine and choline, TMA and TMAO are shown in Figure 1B. We also analysed choline metabolite clustering (Fig. 1C). Of interest, choline strongly correlated with its metabolite TMA, but did not correlate with TMAO. Betaine and carnitine did not correlate with any of the other metabolites. Concentration of betaine and carnitine were higher in males than in females (p=0.01 for betaine and p=0.06 for carnitine). Betaine was lower in patients older than 51 years old (p=0.02). There were no significant differences in the concentrations of choline and its metabolites between patient with normal BMI *versus* overweight and obese patients adjusted by age and gender (Fig. 1D).

Linear and logistic regression analysis between grouped serum choline metabolites and skin and joint clinical parameters.

We then analysed whether choline metabolites differentiate between high and low skin or joint disease activity scores. Figure 2A shows logistic regression analysis adjusted by age and gender by including these factors as covariates in the model, of each choline metabolite of patients classified between mild plus moderate (BSA ≤ 10) versus severe pso-

A				
Metabolite	BSA ≤ 10 Mean (SD)	BSA > 10 Mean (SD)	p	
Choline	49.3 (39)	40.4 (21.4)	0.59	
TMA	46.9 (41.1)	38 (22.3)	0.63	TMAO
TMAO	12.2 (10.3)	29.8 (23.4)	0.03	
Betaine	772.3 (314.3)	869.6 (207.7)	0.19	Betaine
Carnitine	160.7 (30.9)	171.8 (47.9)	0.19	
В				Carnitine
	DAS28CRP ≤ 2.67	DAS28CRP > 2.67	р	
Metabolite	Mean (SD)	Mean (SD)		TMA
Choline	44.1 (31.6)	41.5 (32.9)	0.84	
TMA	40.8 (32)	39.9 (34.2)	0.96	Choline
TMAO	10.9 (6)	19.7 (14.6)	0.11	
Betaine	781.4 (284.5)	949 (273.7)	0.35	ACT AN AN AN AN CAR CAR AN AND CAR AN AND THE SCAR AN
Carnitine	161.5 (27.6)	160 (44.3)	0.11	and the the the offer a with a be a
	DAS28CRP ≤ 4.09	DAS28CRP > 4.09		all of
Metabolite	Mean (SD)	Mean (SD)	р	Color Key
Choline	44.4 (32.2)	28.5 (2.1)	0.49	-0.5 0 0.5
TMA	44.4 (32.2) 41.6 (32.7)	23.7 (5.4)	0.45	
TMAO	12.3 (8.6)	23.7 (17.3)	0.03	
Betaine	800.9 (281.2)	1140.6 (225.7)	0.08	
Carnitine	161.7 (32.3)	153.5 (19)	0.76	TMAO
C	101.1 (02.0)	100.0 (10)	0.70	
Metabolite	CDAI ≤ 10	CDAI >10	р	Betaine
	Mean (SD)	Mean (SD)	r	
Choline	45.8 (32.7)	42 (29.6)	0.49	Carnitine
TMA	42.1 (33.3)	40.8 (30.3)	0.69	
TMAO	11.4 (6.1)	24.3 (22.2)	0.05	TMA
Betaine	790.8 (298.6)	829.4 (281.3)	0.25	
Carnitine	164.9 (27.2)	157.2 (44.9)	0.76	Choline
Metabolite	CDAI ≤ 22	CDAI > 22		
Wetabolite	Mean (SD)	Mean (SD)	р	and and and an and a set and an and an and and and
Choline	42.8 (29.8)	52.8 (39.8)	0.53	ACT ANT AT AND CAR ANT AP CAR ANT AP CAR ANT AP CAR ANT AND CAR ANT
TMA	39.7 (29.9)	51.5 (42)	0.46	all of
TMAO	13.1 (11.4)	32.5 (24.2)	0.04	Color Key
Betaine	776.6 (289.8)	954.1 (254.9)	0.04	0 0.02 0.04 0.06 0.08 0.1
Carnitine	160.8 (33.2)	168.3 (43.9)	0.36	
			0.00	

Fig. 2. Relation of choline metabolites with skin and joint disease activity.

A: Logistic regression was performed for each metabolite in patients with BSA ≤ 10 compared with patient with BSA > 10, adjusting for age and gender. B and C: Logistic regression was performed for each metabolite in patients with low disease activity (measured by either DAS28CRP ≤ 2.67 or CDAI ≤ 10) and patients with moderate and high disease activity (DAS28CRP > 2.67 or CDAI > 10), adjusting for age and gender. Logistic regression was also performed for each metabolite in patients with moderate by either DAS28CRP ≤ 4.09 or CDAI ≤ 22) and patients with high disease activity (DAS28CRP > 4.09 or CDAI ≤ 22) and patients with high disease activity (DAS28CRP > 4.09 or CDAI ≤ 22) and patients with high disease activity (DAS28CRP > 4.09 or CDAI ≤ 22) and patients with high disease activity (DAS28CRP ≥ 4.09 or CDAI ≤ 22) and patients with high disease activity (DAS28CRP ≥ 4.09 or CDAI ≤ 22) and patients with high disease activity (DAS28CRP ≥ 4.09 or CDAI ≤ 22) and patients with high disease activity (DAS28CRP ≥ 4.09 or CDAI ≤ 22) and patients with high disease activity (DAS28CRP ≥ 4.09 or CDAI ≤ 22) and patients with high disease activity (DAS28CRP ≥ 4.09 or CDAI ≤ 22) and patients with high disease activity (DAS28CRP ≥ 4.09 or CDAI ≤ 22) adjusting for age and gender.

D: Linear regression was performed between each clinical score – metabolite pair, controlling for age and gender. In the left panel, the regression coefficients for each pair were used to form a clustered heatmap, to lend insight into which clinical scores were correlated with which metabolite. Row clusters have been identified by cophenetic cutting of the row dendrogram. In the right panel, metabolite regression *p*-values are displayed, where the row and column order are preserved from left panel.

TJC: tender joint count; SJC: swollen joint count; CDAI: Clinical Disease Activity Index; SDAI: Simple Disease Activity Index; HAQ: Health Assessment Questionnaire; PhGA: physician global assessment; PtGA: patient global assessment; BMI: body mass index; BSA: body surface area; DAS28CRP: disease assessment score 28. Concentrations are provided as ng/ml.

riasis (BSA >10). We also conduct this analysis of each choline metabolite of patients classified as low or high joint disease activity (Fig. 2B-C). TMAO was the only metabolite that significantly increased in patients with higher skin and joint scores. We also conducted linear regression that was performed between each choline metabolite-clinical parameter pair, controlling for age and gender. The regression coefficients for each choline metabolite-clinical parameter pair were used to form a clustered heatmap to lend insight into which groups of choline metabolites were correlated with which clinical parameters (Fig. 2D, left panel). Metabolite regression *p*-values are displayed in Figure 2D (right panel), where the row and column order are preserved from left panel. Interestingly, TMAO correlated with parameters of both skin (BSA) and

joint disease activity (TJC, SJC, CDAI, SDAI, DAS28CRP). TMAO also correlated with HAQ and fatigue. Of note, two metabolites, choline and TMA, negatively correlated with fatigue.

Although some other studies have reported a relatively low correlation between items assessing joint symptoms with items assessing skin symptoms (11), in our series, these parameters showed high correlation between them (Fig. 3A-B). We then conducted linear regression between each choline metabolite-joint disease scores (TJC, SJC, DAS28, CDAI and SDAI) adjusted for BSA. We also conducted linear regression between each choline metabolite and BSA adjusted for CDAI. Figure 3C shows regression coefficients and Figure 3D metabolite regression *p*-values. Of note, TMAO still significantly correlates with both skin and joint clinical scores.

Discussion

In the present study, we show that circulating levels of TMAO correlates with skin and joint severity in a cohort of PsA patients. Previous studies showed that choline, betaine and TMAO were all associated with atherosclerosis and higher risk of CVD risk (5). Also, TMAO concentration was dependent on the dietary intake of choline (1), hence it was suggested that changes in diet could be a therapeutic approach for decreasing CVD risk. In our patients, however, there is no correlation between the concentration of choline and TMAO, and, furthermore, only TMAO and not choline and betaine correlated with skin and joint disease activity. This suggests that TMAO concentration does not depend on choline concentration in PsA patients, but more likely on the activity of FMO3,

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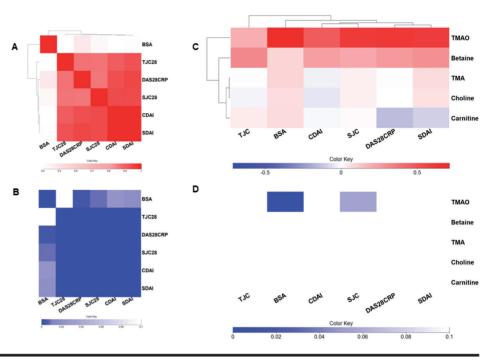
Fig. 3. Choline metabolite regression after BSA and CDAI.

A: The Pearson correlation coefficients for each pair of clinical scores were used to form a clustered heatmap, to evaluate the relationship between joint and skin disease scores.

B: The *p*-values are displayed in Fig. 3B where the row and columns are preserved from Fig. 3A.

C: Linear regression was performed between each clinical score – metabolite pair, controlling for age and gender. TJC, SJC, DAS28CRP, CDAI and SDAI were also adjusted by BSA, and BSA was also adjusted by CDAI. The regression coefficients for each pair were used to form a clustered heatmap.

D: Metabolite regression *p*-values are displayed in Figure 3D, where the row and column order are preserved from Fig. 3C. TJC: tender joint count; SJC: swollen joint count; CDAI: clinical Disease Activity Index; SDAI: Simple Disease Activity Index; BSA: body surface area; DAS28CRP: disease assessment score 28.



which might be elevated in patients with inflammation and obesity.

A recent study showed that expression of FMO3 in adipose tissue in overweight patients positively correlated with BMI and waist-to-hip ratio, and negatively correlated with insulin sensitivity (6, 7), suggesting a link between TMAO-producing enzyme FMO3 and obesity. In our cohort, although TMAO levels did not reach significance in overweight patients, the ratio TMAO/TMA was higher in the obese group (0.3 ± 0.2) in BMI ≤ 30 vs. 0.7 ± 0.5 in BMI > 30, p=0.08) suggesting a higher activity of FMO3 in our cohort as well. Given that there is solid epidemiologic evidence linking psoriasis and PsA to metabolic syndrome (12, 13) an increase of FMO3 activity could explain the elevated TMAO levels in PsA patients.

Very few studies have attempted to study whether or not elevated levels of choline metabolites have an impact on cell function *in vitro* or *in vivo*. Although previous studies have suggested that TMAO promotes vascular inflammation by activating several inflammatory pathways including the NLRP3 inflammasome (14), mitogen-activated protein kinase and nuclear factor-KB pathways (15), further studies are needed to explore the mechanistic links between TMAO and inflammation in PsA. In conclusion, in our cohort, only TMAO, but not TMA, choline, betaine or carnitine, was associated with inflammation in PsA patients. Future studies are needed to explore the mechanistic links between TMAO and skin and joint inflammation.

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