

Association between Metabolic Syndrome and Erythrocyte Fatty Acid Profile in Mexican Adolescents: A *Trans* Fatty Acid Approach

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ABSTRACT

The type of fat consumed in the Mexican diet could predispose to the development of Metabolic Syndrome (MS) which has been associated with an increased risk to develop cardiovascular disease and type 2 diabetes mellitus. Our study included adolescents between 12 and 16 years of age, divided in two groups: Control Group (n = 31) and MS Group (n = 44). Waist circumference, blood pressure, fasting glucose, triglycerides, and HDL-cholesterol were determined. Erythrocytes' fatty acids methyl esters were quantified using gas chromatography with ionized flame detector. We identified 16 fatty acids (FA) with chain lengths from C12 to C24, with emphasis in four *trans* FA (TFA) isomers: vaccenic (C18:1n7t), elaidic (C18:1n9t), linoelaidic (C18:2n6t), and conjugated linoelaidic acids (C18:2n7t). MS Group had a less proportion of: myristic (C14), palmitoleic (C16:1), C18:1n7t, and linoleic acids (C18:2); and a higher one of C18:1n9t, C18:2n7t, and nervonic acids (C24:1) when compared to the control group. C24:1 and C18:1n9t had a significant positive association with MS (OR = 14.17 and OR = 12.94, respectively); whereas C14 (OR = 0.14), C18:1n7t (OR = 0.14), and C18:2 (OR = 0.22) appear to have a protective effect against the disease. The proportion of specific FAs in erythrocytes' membranes differs between adolescents with MS and healthy controls; these FA not only showed a strong association with MS, but also correlated with most of its individual components. Interestingly, TFA displayed an antagonistic behavior; while C18:1n9t had a strong association with MS, apparently C18:1n7t confers a protective effect; these results suggest that analyzing each TFA separately will constitute a more accurate approach to determine the role of TFAs in the pathogenesis of MS or other related metabolic disorders.

Keywords: Metabolic Syndrome; Fatty Acid Profile; Erythrocyte Membrane

1. Introduction

Metabolic Syndrome (MS) in children is defined as the presence of three or more of the following features: obesity, dyslipidemia (increased triglycerides, TG; and/or decreased high density lipoprotein cholesterol, HDL-C), hypertension, and/or impaired glucose metabolism [1].

The growth in obesity, which has reached alarming proportions, has contributed to increase the prevalence of MS in children [2-4]. One of the main concerns about MS in pediatric population is that after due consideration of age, puberty, growth, gender, and ethnic-specific influences; the gathering of these metabolic detriments

may predict the early onset of adulthood cardiovascular disease and type 2 diabetes mellitus [5].

It is well known that diet varies markedly across cultures, however in the last few decades, major changes concerning the quality and quantity of dietary fat have occurred worldwide, changes that may be directly related with the progression of this metabolic impairment [6,7].

Both, observational studies [8] and clinical trials [9] have revealed that fatty acid (FA) composition in human tissues varies widely as a result of fat intake [10]; whereas plasma lipid profile mirrors only dietary fat, within the previous few days [11]; the measurement of FA profile in erythrocytes' membranes is a better long-term dietary fat biomarker, which also reflects endoge-

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nous FA metabolism [12] and constitutes a fair approach to FA composition in muscle cells [13].

FA profile has a marked clinical importance, because it can interfere adversely with the physicochemical properties of the cell membrane, such as protein binding affinity and eicosanoids activation and synthesis [7,14].

Although to our knowledge, there is no evidence concerning changes in the FA profile of erythrocytes' membranes in adolescents with MS, cutting-edge studies in adult population have revealed that a lower proportion of polyunsaturated and monounsaturated fatty acids (PUFA and MUFA) in erythrocytes' membranes is inversely associated with MS, while a high content in saturated fatty acids (SFAs) is positively associated [14-16].

Trans Fatty Acids (TFAs), unsaturated FA with at least one double bond in *trans* configuration, may also provide a good MS-risk biomarker given the fact that they rely almost exclusively on dietary intake [17-19] and are known to have adverse effects on serum lipids [20]. TFAs are mostly produced during the partial hydrogenation of vegetable oils into semisolid fats in the food industry [8,20,21]; however TFAs may also be formed naturally throughout the rumination process, so small amounts of TFAs are present in dairy products and meat [22]. Vaccenic acid, from which conjugated linoleic acid is formed, is the predominant *trans*-isomer in ruminants [17]. Recent studies in adults and in pediatric populations have shown that a high dietary intake of TFAs increases serum low-density lipoprotein cholesterol (LDL-C) and TG, and decreases serum HDL-C [20, 23-28]; besides, the human lipase enzyme is specific for *cis* configuration and is ineffective with *trans* configuration, so *trans* fat remains in the bloodstream for a much longer period of time [26].

In the Mexican Health and Nutrition 2006 National Survey (ENSANUT 2006), a sample of 8690 children and adolescents, underwent a food-frequency questionnaire. According to the World's Health Organization's recommendations, the data obtained from the Survey revealed that Mexican children and adolescents had an insufficient intake of PUFAs, and an excessive intake of SFAs and TFAs [29]. These findings suggest that the quality and quantity of the fat consumed in the Mexican diet could predispose youngsters to the development of metabolic disorders and cardiovascular complications in early adulthood; therefore, the aim of this study was to evaluate if the presence of MS is associated with the FA profile in erythrocytes' membranes, emphasizing TFA contribution.

2. Methods

The study was conducted in the Research Unit in Medical Nutrition, of the Mexican Institute of Social Security

in Mexico City. The protocol was approved by the ethics committee of this institute (R-2010-3603-35). The legal guardian of all study subjects provided written informed consent, and every child gave his approval to participate in the study.

2.1. MS Definition

MS was defined according to the following criteria: waist circumference (WC) > 90th percentile and at least three of the following; HDL-C < 50 mg/dL, TG > 110 mg/dL, diastolic and/or systolic blood pressure (BP) > 90th percentile adjusted by age, gender and height, and/or fasting glucose (FG) > 100 mg/dL [30,31].

2.2. Clinical and Anthropometric Determinations

WC was measured at the narrowest point between the lowest rib and the uppermost lateral border of the right iliac crest. BP was measured three times and the mean was obtained; the measurement was done in the right arm after a 5 minute rest, in a seated position. Weight was assessed with a clinical scale (Tanita, Arlington Heights, IL, TBF-300A) and height was obtained using a wall stadiometer, body mass index (BMI) was computed as [weight (kg)/height (m)²].

2.3. Biochemical Data

After a ten hour overnight fast venous blood samples were collected (10 ml). FG, TG, and HDL-C were measured in serum by enzymatic colorimetric method (SPIN 120 automatic analyzer Shenzhen, Mindray) with commercially available kits.

2.4. Erythrocyte Fatty Acid Profile

Immediately after the blood sample was obtained it was centrifuged at 3000 rpm, for 10min at 4°C. Plasma was discarded and the cells were suspended in an isotonic sodium chloride solution (0.09%). After mixing by inversion, the samples were centrifuged again, and this washing procedure was repeated twice.

FA extraction was performed according to Folch's modified method, with an isopropanol-hexane (6.5:4.0) solution. The solvent solution was added slowly to the red blood cells under vortex mixing conditions, in order to prevent the cells from caking. To separate the lipid fraction the solvent solution was evaporated under a nitrogen stream. The fatty acid portion was methylated with methanolic sodium methoxide (0.5 N) at 50°C for 20min followed by an acid-catalyzed methylation with boron trifluoride in methanol (14%, BF₃) at the same time and temperature conditions.

Erythrocytes' FA methyl esters (FAMES) were separated and measured on a gas chromatographer (Hewlett Packard™ 5890 Series II, Palo Alto, US) coupled to an hydrogen flame detector (FID). A fused silica capillary column of 100 m, 0.25 mm internal diameter, and 0.20 μm film thickness (J&W DB-225) was used. The volume of injection was of 1.0 μL. Helium was used as carrier gas at a flow rate of 1.2 ml/min. Flame ionization temperature was fixed at 270°C and the injector temperature at 250°C. The oven temperature was programmed initially at 70°C with a progressive rate of 30°C/min until 175°C, and then at 1.2°C/min until 230°C, once this temperature was achieved, it was held constant for 5 min.

We identified FAs ranging from C12 to C24 chain lengths; peaks were identified by comparing their retention times against those of high purity (<99%) standard mixtures (Sigma-Aldrich Chemie GmbH, 37 FAs mixture). In addition, TFAs: vaccenic (C18:1n7t), elaidic (C18:1n9t), linoelaidic (C18:2n6t), and conjugated linoleic acid (C18:2 n7t) were verified with their high purity authentic standards (Nu-Check Prep, Inc.).

2.5. Statistical Analysis

Statistical analysis was performed with SPSS software (SPSS Inc., version 19, Chicago, IL, USA). All data are expressed as mean ± standard deviation (SD). FAs on erythrocytes' membranes are expressed as percentage of total FAs. Differences between groups were assessed with Student's T-test and were considered significant at P < 0.05. Pearson's correlation coefficient was performed to evaluate the association between FA profile and each component of the MS. Odds ratio (OR) with a 95% confidence interval (95% CI) was obtained; in order to compute OR with an appropriate cut-off point, a receiver operating characteristic (ROC) curve analysis was performed.

3. Results

We recruited 177 Mexican adolescents between 12 and 16 years of age. By convenience analysis we selected 75 participants (37 females and 38 males) in order to form two opposite study groups: 1) MS Group (n = 44), and 2) Control Group (n = 31). Adolescents who didn't met the criteria for MS, and had only 1 or 2 components of the syndrome were excluded (n = 102) from the study protocol. As expected, significant differences in all anthropometric, biochemical, and clinical parameters were found when groups were compared. Data is summarized in **Table 1**.

FA profile in erythrocytes' membranes is displayed in **Table 2**; we can notice that MS Group had a less proportion of myristic (C14), palmitoleic (C16:1), vaccenic

Table 1. Clinical and biochemical characteristics of study subjects.

Parameters	Control Group (n = 31)	Metabolic Syndrome Group (n = 44)
Gender (F/M)	18/13	19/25
Age	14.0 ± 1.1	13.3 ± 1.4*
Body mass index	19.8 ± 2.1	29.6 ± 4.2*
Waist circumference	77.4 ± 6.6	101.4 ± 11.6 ^o
Systolic blood pressure	99.4 ± 6.3	114.3 ± 10.2*
Diastolic blood pressure	61.8 ± 4.0	71.9 ± 9.4*
Fasting glucose	82.9 ± 7.8	92.1 ± 10.2*
Triglycerides	69.4 ± 20.8	178.8 ± 82.4*
HDL-cholesterol	60.7 ± 7.8	43.2 ± 6.0*

Data is expressed as mean ± SD. *P < 0.05; ^oP < 0.01; *P < 0.001.

Table 2. Erythrocyte FA profile.

Fatty acids	Control Group (n = 31)	MS Group (n = 44)	P
C12:0	0.23 ± 0.20	0.15 ± 0.14	0.058
C14:0	0.78 ± 0.36	0.51 ± 0.29	0.001*
C16:0	26.91 ± 2.83	26.30 ± 3.33	0.411
C16:1	0.83 ± 0.57	0.59 ± 0.24	0.020*
C18:0	19.29 ± 2.60	19.78 ± 2.97	0.467
C18:1	16.49 ± 2.51	15.62 ± 2.29	0.124
C18:1n7t	0.11 ± 0.035	0.066 ± 0.034	0.000*
C18:1n9t	0.33 ± 0.17	0.57 ± 0.20	0.000*
C18:2n6	14.27 ± 1.55	13.03 ± 1.83	0.003*
C18:2n6t	0.59 ± 0.55	0.58 ± 0.57	0.968
C18:2n7t	0.065 ± 0.066	0.16 ± 0.14	0.000*
C18:3n3	0.24 ± 0.16	0.22 ± 0.086	0.359
C20:4n6	13.80 ± 3.66	15.16 ± 4.18	0.148
C20:5n3	0.39 ± 0.14	0.43 ± 0.16	0.329
C22:6n	3.03 ± 0.84	3.06 ± 1.16	0.908
C24:1	2.66 ± 0.93	3.75 ± 1.21	0.000*
ΣSat	47.21 ± 4.56	46.74 ± 5.57	0.700
ΣMUFA	20.34 ± 2.40	19.96 ± 1.86	0.441
ΣPUFA	31.36 ± 4.93	31.88 ± 6.19	0.695
n6/n3	9.16 ± 2.46	8.79 ± 4.81	0.698
Σtrans	1.09 ± 0.61	1.38 ± 0.63	0.051

Data are expressed as mean ± SD (% of FA content). *P < 0.05.

(C18:1n7*t*), and linoleic acids (C18:2); and a higher one of elaidic (C18:1n9*t*), conjugated linoelaidic (C18:2n7*t*), and nervonic acids (C24:1) when compared to the Control Group; the sum of saturated, monounsaturated and polyunsaturated fats showed no significant difference between groups; however, the sum of total *trans* fat showed a marginal significant difference.

Although we did find a significant difference in conjugated linoelaidic acid's proportion between study groups, data didn't have a normal distribution even after a logarithmic transformation was performed. Moreover, this FA was present in erythrocytes' membranes just as traces and we weren't able to identify it in 23 study subjects (13 of the Control Group, and 10 of MS Group). Because of this, we decided to suppress this FA from all further analyses.

In order to build contingency tables, and to successfully analyze OR with the most accurate data; we pro-

posed different cut-off points according to ROC curve analyses.

Each ROC curve was constructed using the FA's content in erythrocytes' membrane (expressed in %) and the presence or absence of MS (dichotomous variable). We selected the cut-off points according to sensitivity and specificity values (**Table 3**). ROC curve analyses were performed only for the FAs that differed significantly amongst groups.

Table 4 resumes the ORs obtained from the contingency tables; by analyzing data on **table 4**, we can realize that nervonic (24:1) and elaidic acids (C18:1n9*t*) had a strong positive association with MS; whereas myristic (C14), vaccenic (C18:1n7*t*), and linoleic acids (C18:2) appear to have a protective effect against the disease.

Finally, to determine association between those FAs that were different between groups and each component of the MS, we constructed a correlation matrix (**Table 5**).

Table 3. Determination of a cut-off point for selected FAs.

Fatty acid	Cut-off point (%)	Sensitivity (%)	Specificity (%)	AUC	<i>P</i>
C14	0.62	77.3	67.7	0.75	0.000
C16:1	0.55	50	54.8	0.55	0.42
C18:1n9 <i>t</i>	0.39	81.8	74.2	0.81	0.000
C18:1n7 <i>t</i>	0.085	75	71	0.83	0.000
C18:2	13.9	68.2	67.7	0.71	0.002
C24:1	3.31	77.3	77.4	0.81	0.000

Table 4. Odds ratios (95% CI) of selected FA against MS.

Fatty Acid	Control Group		MS Group		Odds Ratio (CI) 95%
	Exposed	Not Exposed	Exposed	Not Exposed	
C14	>0.62%	<0.62%	>0.62	<0.62	0.14 (0.05 - 0.39)
Cases (n)	21	10	10	34	
C16:1	>0.55%	<0.55%	>0.55%	<0.55%	0.82 (0.33 - 2.05)
Cases (n)	17	14	22	22	
C18:1n7 <i>t</i>	>0.085%	<0.085%	>0.085%	<0.085%	0.14 (0.05 - 0.39)
Cases (n)	22	9	11	33	
C18:1n9 <i>t</i>	>0.35%	<0.35%	>0.35%	<0.35%	12.94 (4.26 - 39.25)
Cases (n)	8	23	36	8	
C18:2	>13.9%	<13.9%	>13.9%	<13.9%	0.22 (0.082 - 0.58)
Cases (n)	21	10	14	30	
C24:1	>3.31%	<3.31%	>3.31%	<3.31%	14.17 (4.52 - 44.26)
Cases (n)	6	25	34	10	

Table 5. Correlation matrix between MS components and certain fas.

	WC	FG	TAG	HDL	SBP	DBP
C14	-0.32 [◊]	-0.28*	-0.29*	0.38*	-0.23*	NS
C16:1	NS	NS	NS*	NS	NS	NS
C18:1n7t	-0.488*	-0.47*	-0.49*	0.39*	-0.28*	-0.3 [◊]
C18:1n9t	0.53*	0.27*	0.35 [◊]	-0.4*	0.22*	0.34 [◊]
C18:2	-0.35 [◊]	NS	NS	NS	-0.39*	-0.27*
C24:1	0.31 [◊]	NS	0.39*	-0.31 [◊]	NS	NS

WC = Waist Circumference; FG = Fasting Glucose; TAG = Triglycerides; HDL = High Density Lipoprotein; SBP = Systolic Blood Pressure; DBP = Diastolic Blood Pressure. *P < 0.05; [◊]P < 0.01; [◊]P < 0.001. NS = No significance reported.

A positive association was observed against 24:1 and C18:1n9t and most of the MS components. In contrast, C18:2, C18:1n7t, C14 showed and inverse association.

4. Discussion

Our results demonstrate that the proportion of certain FAs in erythrocytes' membranes differs between adolescents with MS and healthy controls. By analyzing the ORs obtained, we can infer that C24:1 and C18:1n9t play a role in the development of MS; whereas C14, C18:1n7t, and C18:2 apparently provide a protective effect against the disease.

When we analyzed the correlation coefficients between each FA and each component of the MS, we realized that the FAs with the highest ORs, also have the strongest associations; for instance C18:1n9t (OR = 12.94) correlated positively with all of the components of the MS, particularly with WC ($r = 0.53$, $P < 0.01$); and C24:1 (OR = 14.17) showed a fair association with TG ($r = 0.39$, $P < 0.01$). The same behavior was observed with the protective FAs, we can appreciate that the FAs with the lowest ORs have the strongest correlations; for example C18:1n7t (OR = 0.14) correlated inversely with all of the components of the MS, predominantly with FG, TG, and WC (0.47, 0.48, 0.49, respectively, $P < 0.01$); and C14 (OR = 0.14) showed a moderate correlation with most of the Syndrome's components.

Since there is no information available about what kind of FA profile to expect in a healthy child, in order to provide ORs we needed to suggest a cut-off point for each FA's concentration. We can assume that selecting a cut-off point obtained from a ROC curve to analyze OR is a valid approach, this is supported by the high sensitivity, specificity, and area under the curve that we found and thus suggests that selected FAs could be useful as MS biomarkers.

As we previously mentioned, elaidic (C18:1n9t) and linoelaidic acids (C18:2n7t) are *trans* fats, formed in the

food industry during the partial hydrogenation of vegetable oils; when analyzing the quality and quantity of the fat present in the Mexican diet, Villalpando *et al* found a high content of this two TFAs in a variety of fast foods, margarines, crackers, fries, and doughnuts widely commercialized in Mexico (>2% g/100g) [32]; as we now know, the consumption of industrial TFAs is associated with weight gain and visceral fat accumulation in male adult subjects [33] and it may also increase LDL-C and TG and reduce HDL-C [17]. Although no differences were found in linoelaidic acid (C18:2n6t) concentration between study groups, MS Group had a higher concentration of elaidic acid (C18:1n9t), and this tFA also showed an important association with MS (OR = 12.94).

On the other hand, vaccenic and conjugated linolenic acids are TFAs from ruminants that are present in smaller amounts in dairy products and meat; current evidence suggests that TFAs from ruminants have limited implications against health [20]. Our results are consistent with previously reported data; and we did find a negative association between vaccenic acid (C18:1n7t) and all of the components of MS. Interestingly, the sum of TFAs (C18:1n7t, C18:1n9t, C18:2n6t, and C18:2n7t) had an OR of 4.71 (CI 95%: 1.73 - 12.8) but no association was found with MS or its components; so it seems that analyzing each TFA separately instead of considering the total sum is a better approach.

When we analyzed the data obtained, we realized that nervonic acid (C24:1) had the highest association with MS (OR = 14.17), however, to our knowledge, there aren't any previously reported studies that involve this FA with MS. Straczkowski *et al* found that skeletal muscle ceramides have an important proportion of C24:1 (5.8%) and that this proportion correlates inversely with insulin sensitivity ($r = -0.39$, $P = 0.047$ vs clamp) [34]. Ceramide is the main second messenger derived from the hydrolysis of membrane sphingomyelin and is directly involved in cell differentiation, inhibition of cell proliferation, induction of apoptosis, and more recently with

glucose uptake stimulated by insulin action [35]; probably, the higher amount of C24:1 found in the erythrocytes' membranes of our study subjects with MS, is closely related with an insulin resistance state.

Although MS Group has a detrimental metabolic condition it is noteworthy that no differences were found in linolenic acid (C18:3n3), eicosapentaenoic acid (C20:5n3), docosahexaenoic acid (C22:6n3), sum of PUFAs, or n-6/n-3 ratio when compared to the Control Group. Since the beneficial role of n-3 family in lipid metabolism is well known, we expected to find a lower proportion of these FAs in the MS Group. Despite there is evidence that supports the fact that the Mexican diet is insufficient in n-3 and in n-6 in nearly the 52% of Mexican adolescents [29]; when we analyzed the proportion of C18:3n3, C20:4n6, C20:5n3, C22:6n3, Σ PUFAs, and n-6/n-3 ratio, in both groups (75 subjects) we found that Mexican children and adolescents have a similar FA profile to the one previously reported in comparable populations of Hungarian and Australian children [36,37]. These results suggest that there is an increased n-3 synthesis in Mexican Adolescents that successfully compensates the insufficient dietary intake.

Description of the FA profile in erythrocytes' membranes of the Control Group is an important contribution of the present study, since there isn't enough data about the FA profile in healthy young populations; we dare to suggest that the described profile in lean adolescents may be used as reference data for further studies.

In summary, the proportion of specific FAs in erythrocytes' membranes differs between adolescents with MS and healthy controls; these FA not only showed a strong association with MS, but also correlated with most of its individual components. Additionally TFAs displayed an antagonistic behavior; these results suggest that analyzing each TFA separately will constitute a more accurate approach to determine the role of TFAs in the pathogenesis of MS or other related metabolic disorders.

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