[ORIGINAL ARTICLE]

Increased Serum Dihomo- γ -linolenic Acid Levels Are Associated with Obesity, Body Fat Accumulation, and Insulin Resistance in Japanese Patients with Type 2 Diabetes

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Abstract:

Objective To clarify the associations between serum omega-6 (n-6) and omega-3 (n-3) polyunsaturated fatty acid (PUFA) levels and obesity-related metabolic abnormalities in patients with type 2 diabetes.

Methods and Materials Data from 225 Japanese patients with type 2 diabetes were cross-sectionally analyzed. The serum levels of n-6 PUFAs [dihomo- γ -linolenic acid (DGLA) and arachidonic acid (AA)] and n-3 PUFAs (eicosapentaenoic acid and docosahexaenoic acid) were measured, and the estimated Δ -5 desaturase (D5D) activity was calculated based on the AA to DGLA ratio. The associations between the composition of PUFAs and obesity-related parameters, including the body mass index (BMI), waist circumference, alanine amino transferase (ALT) level, homeostatic model assessment of insulin resistance (HOMA-IR), and body fat percentage, as measured by a bioelectrical impedance analysis, were analyzed.

Results Among the PUFAs, the DGLA level had the strongest correlations with BMI ($p < 0.001$), waist circumference ($p < 0.001$), ALT level ($p < 0.001$), HOMA-IR ($p < 0.001$), and body fat percentage ($p < 0.01$). AA was positively correlated and D5D was negatively correlated with several obesity-related parameters, while n-3 PUFAs did not have a constant correlation. A multivariate regression analysis revealed that the DGLA level was an independent determinant for HOMA-IR ($\beta = 0.195$, $p = 0.0066$) after adjusting for sex, age, BMI, and the ALT, triglyceride, and HbA1c levels.

Conclusion A high serum DGLA level was associated with obesity, body fat accumulation, a high ALT level, and insulin resistance in patients with type 2 diabetes. The measurement of the serum PUFA levels may be useful for evaluating metabolic abnormalities and estimating the dietary habits of patients.

Key words: polyunsaturated fatty acid, dihomogamma-linolenic acid, obesity, type 2 diabetes, insulin resistance

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Introduction

In the past half-century, the total and saturated fat intakes have fallen in industrial countries (1, 2). However, the ratio of omega-6 (n-6) to omega-3 (n-3) polyunsaturated fatty acids (PUFAs) has increased in parallel with the increase in the prevalence of obesity (3).

n-3 and n-6 PUFAs are essential fatty acids that must be supplied from the diet due to the lack of endogenous en-

zymes for desaturation in humans. n-3 PUFAs are mainly supplied from fish oils, whereas n-6 PUFAs are broadly available from sources ranging from vegetable oils to meat products. n-3 and n-6 PUFAs compete for common metabolic enzymes (4). Basic studies have shown that n-3 and n-6 PUFAs have opposing effects on adipogenesis (5), the browning of adipose tissue (6), the brain-gut-adipose tissue axis (7), and systemic inflammation (8). Many epidemiological or clinical studies have shown that high n-3/n-6 PUFA ratios were associated with a decreased incidence of

ischemic heart disease (9, 10).

In addition, the association between PUFA profiles and obesity or its-related disorders have been attracting attention. A previous report on 534 participants of the Woman's Health Study showed that baseline red blood cell membrane phospholipid cis n-6 fatty acid levels were positively associated and cis n-3 FAs were inversely associated with longitudinal weight gain (11). Recent reports on the EPIC-InterAct Case-Cohort Study (12) and Hitachi Health Study (13) showed that among the n-3 and n-6 PUFAs, the dihomo- γ -linolenic acid (DGLA) level had the strongest positive association with the incidence of type 2 diabetes. In addition, a cross-sectional report showed that a high DGLA level and low Δ -5 desaturase (D5D) activity, a rate-limiting enzyme involved in the formation of arachidonic acid (AA) from DGLA, were associated with hepatic steatosis (14). These previous findings suggest that PUFA metabolism-particularly in n-6 PUFA-is closely associated with obesity and its-related disorders. However, most previous studies have included participants without diabetes at baseline, and there have been few studies limited to patients with type 2 diabetes. In particular, the clinical data that have been used to analyze the association between PUFA profiles and multiple metabolic abnormalities in patients with type 2 diabetes are limited. In the present study, we aimed to clarify the associations between the composition of PUFAs and multiple parameters related to obesity-related metabolic abnormalities.

Materials and Methods

Data sources and participants

We performed a retrospective, cross-sectional study using clinical records from 225 patients (male, n=165; female, n=60) who were diagnosed with type 2 diabetes according to the Japan Diabetes Society guidelines (15). The participants were admitted to the Endocrinology and Diabetes Center at Yokohama Rosai Hospital for glycemic control and diabetes education from January 2011 to December 2016. The exclusion criteria were treatment with eicosapentaenoic acid (EPA), docosahexaenoic (DHA), ezetimibe or insulin; renal insufficiency, with an estimated glomerular filtration rate (eGFR) of <30 mL/min/1.73 m²; decreased insulin secretion with a fasting serum C-peptide immunoreactivity (CPR) level of <0.17 nmol/L (=0.5 ng/dL); a history of diabetes-related ketoacidosis and diabetes-related coma within 6 months; a history of severe infection or trauma requiring hospital care within 6 months; malignant neoplasm (treated or untreated); or a history of surgery for weight loss. This study was carried out in accordance with the ethical principles of the Declaration of Helsinki and was approved by the research ethics committee of Yokohama Rosai Hospital (Approved ID: 28-58). The need for informed consent was waived because of the retrospective nature of the study and because the study utilized data provided in the course of

normal patient care.

Measurements and the calculation of indices

The history of diabetes was self-reported. At admission, the participant's weight and height were measured and were used to calculate body mass index (BMI). The waist circumference measurement was performed by the investigator at the level of the umbilicus, directly on the skin while the participant was standing erect. On the second morning of hospitalization, blood samples were collected after over 12 hours of fasting. The blood samples were collected in serum tubes containing separating gel and were centrifuged at 1785 ×g for 7 min. The supernatants were then collected and immediately transported to SRL (Tokyo, Japan) at 4°C. The serum PUFA (DGLA, AA, EPA, and DHA) levels were measured using gas chromatography-flame ionization detection. The estimated D5D activity was calculated as the ratio of AA to DGLA (13, 14). The glycated hemoglobin (HbA1c) level was measured by high-performance liquid chromatography (TOSOH, Tokyo, Japan). The immunoreactive insulin (IRI) level was measured by a chemiluminescent enzyme immunoassay. The homeostatic model assessment of insulin resistance (HOMA-IR) value was calculated as [fasting plasma glucose (mg/dL)×fasting IRI (μU/mL)/405]. The body composition was measured in 77 patients by bioelectrical impedance analysis using an InBody 720 system (16) (Biospace, Tokyo, Japan), which is approved for use as a medical instrument by Pharmaceuticals and Medical Devices Agency, Japan. The body fat percentage and lean mass weight measured by the InBody 720 were strongly correlated with the values measured by dual-energy X-ray absorptiometry in obese patients (r=0.736 and r=0.899) (17). The skeletal mass index was calculated as the appendicular skeletal muscle mass/height² (18). The parameters of atherosclerosis were assessed based on brachial-ankle pulse wave velocity (baPWV) (19) and the maximum carotid intima-media thickness (max IMT) (20).

Statistical analysis

The data are given as medians (interquartile range) because most of the continuous variables were non-parametrically distributed. Spearman's rank correlation coefficients were calculated to assess the relationships between continuous variables. The Mann-Whitney U test was used for intergroup comparisons. Multiple regression analyses were performed using a least-squares analysis of variance to clarify the independent factors associated with BMI or HOMA-IR. The independent variables included in the multiple regression analysis for BMI were participant sex, age, current smoking, habitual drinking, triglyceride (TGL) level, HbA1c level, and the PUFA levels that were significantly correlated with the BMI in a simple correlation analysis. The independent variables included in the multiple regression analysis for HOMA-IR were participant sex, age, BMI, eGFR, ALT level, TGL level, HbA1c level, and the PUFA levels that were significantly correlated with the indices of

insulin resistance in the simple correlation analysis. Missing values were excluded from the analysis. In all of the statistical analyses, *p* values of <0.05 were considered to indicate statistical significance. The JMP® 12 software program (SAS Institute, Cary, USA) was used for all of the analyses.

Table 1. The Clinical Characteristics of the Patients.

Sex (male/female)	165/60
Age (years)	63 (49-72)
Duration of diabetes (years)	6 (0-13)
BMI (kg/m ²)	25.7 (22.8-28.8)
Waist circumference (cm)	92.0 (87.0-100.0)
HbA1c (%)	9.8 (8.4-11.2)
ALT (IU/L)	25.0 (18.0-39.0)
Total-cholesterol (mg/dL)	163 (145-189.5)
HDL-cholesterol (mg/dL)	43 (37-50.5)
TGL (mg/dL)	121 (96-156)
Non-HDL-cholesterol (mg/dL)	121 (100.5-145.5)
eGFR (mL/min/1.73m ²)	77.8 (67.3-89.3)
HOMA-IR	2.10 (1.24-3.44)
DGLA (μg/mL)	36.3 (28.2-47.4)
AA (μg/mL)	190.5 (157.0-235.3)
EPA (μg/mL)	55.0 (34.5-75.7)
DHA (μg/mL)	133.7 (106.5-171.5)
EPA/AA	0.29 (0.17-0.43)
Estimated D5D	5.0 (4.2-6.7)
Body fat percentage (%)	31.7 (22.3-37.9)
Skeletal mass index	7.2 (6.7-7.9)
Current smoking, n (%)	69 (30.7%)
Habitual drinking, n (%)	122 (47.8%)
History of cerebrovascular disease, n (%)	30 (13.3%)
Number of glucose-lowering agents used	1 (1-3)
Taking lipid-lowering agents, n (%)	89 (39.5%)
Taking antihypertensive agents, n (%)	108 (46.7%)

Data are given as medians (interquartile range). BMI: body mass index, HbA1c: glycated hemoglobin, ALT: alanine amino transferase, HDL: high-density lipoprotein, TGL: triglycerides, eGFR: estimated glomerular filtration rate, HOMA-IR: homeostatic model assessment of insulin resistance, DGLA: dihomo- γ -linolenic acid, AA: arachidonic acid, EPA: eicosapentaenoic acid, DHA: docosahexaenoic acid, D5D: Δ -5 desaturase

Results

The clinical characteristics of the patients

The characteristics of the 225 cases in this study are shown in Table 1. The median age was 63 (49-72) years, and 165 (73.3%) patients were male. Nearly half of the patients (45.3%) were obese (BMI $\geq$ 25 kg/m²). The glycemic control was poor, with a mean HbA1c level of 9.8%. Eighty-nine patients (39.5%) were taking lipid-lowering agents (statins or fibrates). The median lipid levels were almost within the normal range. The correlations of each of the serum PUFAs are shown in Table 2. No inverse correlation was observed between the levels of n-3 and n-6 PUFAs.

Correlations between the PUFA composition and clinical parameters

To clarify whether n-6 or n-3 PUFAs were associated with obesity-related parameters, we analyzed the correlations between the PUFA composition and various clinical parameters. As shown in Table 3, among the PUFAs, the DGLA level had the strongest correlations with BMI (ρ =0.316, p <0.001), waist circumference (ρ =0.227, p <0.001), ALT level (ρ =0.270, p <0.001), HOMA-IR (ρ =0.367, p <0.001), and body fat percentage (ρ =0.315, p <0.01). AA was significantly correlated with BMI (ρ =0.217, p =0.0010), waist circumference (ρ =0.165, p =0.020), ALT level (ρ =0.150, p =0.025), and HOMA-IR (ρ =0.169, p =0.022); however, the correlations were weaker than those of DGLA. The estimated D5D activity had negative correlations with BMI (ρ =-0.166, p =0.013), ALT level (ρ =-0.178, p =0.0076), and HOMA-IR (ρ =-0.255, p <0.001). EPA (ρ =-0.159, p =0.017), EPA/AA ratio (ρ =-0.246, p <0.001), and estimated D5D activity (ρ =-0.166, p =0.013) were negatively associated with BMI. Age had a negative correlation with n-6 PUFAs and positive correlations with n-3 PUFAs and the EPA/AA ratio. Age also had a negative correlation with BMI (ρ =-0.429, p <0.001). A multivariate regression analyses adjusted for sex, age, current smoking, habitual drinking, TGL level, and HbA1c level, revealed that the DGLA and AA levels were the only independent determinants of BMI (Table 4). After adjusting for sex, age, BMI, and eGFR, we observed no significant asso-

Table 2. The Correlations of Serum PUFAs.

	DGLA	AA	EPA	DHA	EPA/AA	Estimated D5D
DGLA	-	0.446***	-0.101	0.099	-0.300***	-0.681***
AA		-	0.164*	0.305***	-0.262***	0.293***
EPA			-	0.810***	0.892***	0.250***
DHA				-	0.645***	0.143*
EPA/AA					-	0.128
Estimated D5D						-

Spearman correlation coefficient (ρ) were used to evaluate correlations between variables. * p <0.05, ** p <0.01, *** p <0.001. PUFAs: polyunsaturated fatty acids, DGLA: dihomo- γ -linolenic acid, AA: arachidonic acid, EPA: eicosapentaenoic acid, DHA: docosahexaenoic acid, D5D: Δ -5 desaturase

Table 3. Correlations between PUFA Levels and Clinical Parameters.

	DGLA	AA	EPA	DHA	EPA/AA	Estimated D5D
BMI	0.316***	0.217**	-0.159*	-0.074	-0.246***	-0.166*
Age	-0.260***	-0.328***	0.324***	0.253***	0.454***	0.033
Waist circumference	0.227***	0.165*	-0.088	-0.003	-0.152*	-0.109
ALT level	0.270***	0.150*	-0.078	0.008	-0.147*	-0.178**
TGL	0.362***	0.270***	-0.044	0.234***	-0.169*	-0.151*
HbA1c level	-0.114	0.047	-0.136*	-0.135	-0.157*	0.125
HOMA-IR	0.367***	0.169*	-0.012	0.064	-0.097	-0.255***
Body fat percentage	0.315**	0.208	-0.101	0.013	-0.188	-0.182
Skeletal mass index	0.031	0.100	-0.229*	-0.272*	-0.274*	0.030

Spearman correlation coefficient (ρ) were used to evaluate correlations between variables. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$. PUFA: polyunsaturated fatty acid, BMI: body mass index, ALT: alanine amino transferase, TGL: triglyceride, HbA1c: glycated hemoglobin, HOMA-IR: homeostatic model assessment of insulin resistance, DGLA: dihomo- γ -linolenic acid, AA: arachidonic acid, EPA: eicosapentaenoic acid, DHA: docosahexaenoic acid, D5D: Δ -5 desaturase

Table 4. Multivariate Associations of Clinical Variables with Body Mass Index.

Measure	Body mass index									
	Multivariate model									
	Model 1		Model 2		Model 3		Model 4		Model 5	
	β	p	β	p	β	p	β	p	β	p
Sex (female)	-0.006	0.92	-0.004	0.95	0.020	0.75	0.009	0.88	0.013	0.83
Age	-0.407	<0.0001	-0.397	<0.0001	-0.433	<0.0001	-0.416	<0.0001	-0.443	<0.0001
Current smoking (yes)	-0.078	0.20	-0.060	0.32	-0.069	0.26	-0.071	0.25	-0.074	0.23
Habitual drinking (yes)	-0.088	0.14	-0.104	0.062	-0.101	0.10	-0.101	0.10	-0.092	0.13
HbA1c	-0.131	0.028	-0.153	0.0094	-0.161	0.0074	-0.164	0.0062	-0.146	0.015
TGL	0.114	0.076	0.123	0.050	0.167	0.0063	0.162	0.0080	0.158	0.010
DGLA	0.153	0.022								
AA			0.148	0.025						
EPA					-0.033	0.60				
EPA/AA ratio							-0.066	0.31		
Estimated D5D									-0.062	0.31
R ²	0.275		0.273		0.258		0.260		0.260	
p	<0.0001		<0.0001		<0.0001		<0.0001		<0.0001	

ρ : Spearman correlation coefficient, β : standard partial regression coefficient. HbA1c: glycated hemoglobin, TGL: triglyceride, DGLA: dihomo- γ -linolenic acid, AA: arachidonic acid, EPA: eicosapentaenoic acid, D5D: Δ -5 desaturase

ciation between the PUFA composition and the parameters of atherosclerosis, including the baPWV and max IMT values (data now shown). In addition, there was no significant association between the use of glucose-lowering agents and the PUFA levels (data not shown).

The multivariate analysis of the association between PUFA levels and HOMA-IR

We performed a multivariate regression analysis to clarify whether the DGLA level and estimated the D5D activity were independent determinants of HOMA-IR. After adjusting for sex, age, BMI, eGFR level, ALT level, TGL level, and HbA1c level, the DGLA level and the estimated D5D activity were found to be independently associated with HOMA-IR ($\beta=0.195$, $p=0.0066$ and $\beta=-0.147$, $p=0.024$). The AA level was not independently associated with HOMA-IR ($\beta=0.065$, $p=0.38$) (Table 5). We confirmed similar results in

a multivariate regression analysis using log-transformed HOMA-IR as a dependent variable (data not shown).

Discussion

The results of the present study showed that a high serum DGLA level were associated with obesity, body fat accumulation, a high ALT level, and insulin resistance in Japanese patients with type 2 diabetes. The measurement of the serum PUFA levels, especially the n-6 PUFA levels, may be useful in evaluating obesity-related metabolic abnormalities.

We found that the DGLA level had significant positive correlations with the body mass index, waist circumference, ALT level, HOMA-IR, and body fat percentage. In addition, the AA level was positively correlated with several obesity-related parameters. These results suggest that an excessive intake of n-6 PUFAs is associated with obesity and its-

Table 5. Univariate and Multivariate Associations of Clinical Variables with HOMA-IR.

Measure	HOMA-IR							
	Univariate model		Multivariate model					
			Model 1		Model 2		Model 3	
	β	p	β	p	β	p	β	p
Sex (female)			0.087	0.1935	0.100	0.14	0.107	0.11
Age	-0.092	0.270	0.163	0.0615	0.155	0.09	0.136	0.12
BMI	0.349	<0.0001	0.217	0.0033	0.223	0.0032	0.234	0.0016
eGFR	0.079	0.290	0.034	0.67	0.045	0.58	0.049	0.535
ALT	0.448	<0.0001	0.334	<0.0001	0.352	<0.0001	0.342	<0.0001
TGL	0.235	0.0013	0.112	0.10	0.154	0.03	0.158	0.02
HbA1c	-0.099	0.18	-0.001	0.98	-0.030	0.66	-0.001	0.99
DGLA	0.349	<0.0001	0.195	0.0066				
AA	0.221	0.0026			0.065	0.38		
Estimated D5D	-0.216	0.0032					-0.147	0.024
R ²	-		0.3219		0.296		0.313	
p	-		<0.0001		<0.0001		<0.0001	

ρ : Spearman correlation coefficient, β : standard partial regression coefficient. HOMA-IR: homeostatic model assessment of insulin resistance, BMI: body mass index, eGFR: estimated glomerular filtration rate, ALT: alanine amino transferase, TGL: triglyceride, HbA1c: glycated hemoglobin, DGLA: dihomo- γ -linolenic acid, AA: arachidonic acid, D5D: Δ -5 desaturase

related metabolic abnormalities. The intake of n-6 PUFAs is mainly derived from linoleic acid (LA), which is plentiful in vegetable oils. LA is metabolized to γ -linoleic acid (GLA) by Δ -6 desaturase, and then progressively elongated to DGLA. GLA and DGLA are hardly supplied from a normal diet. Thus, high serum DGLA is usually associated with the excessive intake of LA. DGLA itself is thought to have an anti-inflammatory effect (21, 22). However, DGLA is further converted to AA, which promotes body fat gain through the mechanisms of adipogenesis (5, 23), lipid homeostasis (24-26), and the brain-gut-adipose tissue axis (7, 27-29). In addition, AA-derived prostaglandin E2 and leukotriene B4 have pro-inflammatory properties (21). Thus, the excessive intake of LA contributes to the development of obesity and a chronic inflammatory state via AA (30). In addition, AA is supplied directly from the diet, including grain-fed animals, dairy, and eggs. Patients with type 2 diabetes should avoid the high intake of n-6 PUFAs, including LA and AA, for the control of body weight and the prevention of cardiovascular disease. By checking the serum levels of PUFAs, physicians can estimate the dietary habits of patients and provide appropriate dietary advice.

The present study showed that the DGLA level had a stronger association with obesity-related parameters than that of AA. It may be related to the downregulated activity of D5D, an enzyme converting DGLA to AA. The estimated D5D activity is downregulated in individuals with obesity or insulin resistance (31, 32). Actually, the DGLA levels and estimated D5D activity tended to show mirror images with regard to the obesity-related parameters in the present study. In addition, the estimated activity of D6D, an enzyme converting GLA to DGLA, was shown to be positively correlated with obesity or insulin resistance (31). Unfortunately,

the GLA level was not measured in the present study, and we could not calculate the estimated D6D activity. Overall, the DGLA level tends to increase in the state of obesity or insulin resistance. DLGA level may be a representative marker reflecting not only an excessive intake of n-6 PUFAs, but also obesity-related metabolic abnormalities.

There was no consistent correlation between the serum levels of n-3 PUFAs and obesity-related parameters. EPA had a significant negative correlation with the BMI; however, this association was diminished after adjusting for potential confounders. Previous studies showed that in contrast to n-6 PUFAs, n-3 PUFAs have protective effects against the development of obesity (33). However, clinical intervention studies to examine the relationship between n-3 supplementation and the body composition have reported conflicting results (34, 35). One possible reason for the ambiguous results of n-3 supplementation on obesity is that n-3 supplementation in most interventional studies was performed without restricting the n-6 PUFA intake. In our study, there was no inverse correlation between the serum levels of n-3 and n-6 PUFAs (Table 2), suggesting that patients with a high intake of n-3 PUFAs did not always have a reduced intake of n-6 PUFAs. Hence, n-3 PUFA supplementation may not be effective if it is performed without restricting the n-6 PUFA intake.

In the present study, we only measured the levels of four serum fatty acids. This method is associated with some limitations. First, there was no information about the levels of fatty acids other than DGLA, AA, EPA, and DHA. In addition, the total food and fat intakes were not assessed. Thus, we could not determine whether the serum DGLA levels were directly associated with obesity-related parameters or if the association was intermediated by the other serum fatty

acids. We assumed that in comparison to other fatty acids, DGLA has a stronger association with obesity-related parameters, because high DGLA levels showed the strongest positive association with the incidence of type 2 diabetes among the various types of fatty acids that were assessed in the EPIC-InterAct Case-Cohort Study (12) and the Hitachi Health Study (13). Second, serum samples were used in this study. The fatty acid composition in serum can be affected by the dietary fatty acid intake patterns of the previous few days (36). By measuring the fatty acid composition of the erythrocyte membrane or blood phospholipids, which reflect the meals of the previous several weeks (37), the associations could be more accurately assessed. Considering the significant correlation between the fatty acid levels of serum and red blood cells in healthy Japanese subjects (38), we do not believe that there was a large margin of error. Third, we used the absolute values of the fatty acid levels in our analysis; these values might have been affected by the levels of triglycerides or the amount of body fat. An analysis using the molar percentage of each fatty acid may attenuate the influence of these factors. In this study, multivariate analyses showed that the DGLA levels were a significant determinant of both the BMI and HOMA-IR values, independent from the triglyceride levels. We did not select body fat percentage as an independent variable for the multivariate analysis, because of the limited number of patients whose body composition was measured. We cannot deny that the associations between the DGLA levels and the BMI and HOMA-IR values are influenced by the amount of body fat.

The strength of the present study is that it analyzed the associations between PUFA levels and metabolic abnormalities by checking multiple clinical parameters. Few studies have investigated the association between the PUFA levels and the body composition of patients with type 2 diabetes. However, the present study is associated with several limitations in addition to the abovementioned limitations. First, this study employed a cross-sectional design. Thus, it is difficult to show a causal relationship between the PUFA composition and obesity-related metabolic abnormalities. Prospective studies are needed to confirm the results of the present study. Second, the amount of exercise was not assessed in this study. In addition, the amount of smoking and drinking was not quantitatively assessed. A self-administered semi-quantitative questionnaire could be used to estimate these parameters. Third, the participants were limited to inpatients with uncontrolled glucose levels, and a median HbA1c level of 9.8%. The data from patients with well-controlled glucose levels were limited. We selected inpatients as participants because we attempted to collect various clinical parameters. We think that hospitalization had little influence on the circulating serum PUFA levels because the blood samples were collected on the second morning of hospitalization.

In conclusion, the results of the present study showed that a high serum DGLA level was associated with obesity, body fat accumulation, a high ALT level, and insulin resistance, in

Japanese patients with type 2 diabetes. The measurement of the serum PUFA levels may be useful in evaluating for obesity-related metabolic abnormalities and estimating the dietary habits of the patients.

The authors state that they have no Conflict of Interest (COI).

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