

## Postprandial Triglyceride Is Associated with Fasting Triglyceride and HOMA-IR in Korean Subjects with Type 2 Diabetes

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**Background:** Recent studies indicate postprandial triglyceride (TG) had a better association with cardiovascular events and metabolic syndrome than fasting TG. The authors of the present study investigated the metabolic and clinical relevance of postprandial TG.

**Methods:** In a cross-sectional retrospective study, the authors of the present study compared fasting and postprandial TG and analyzed the relationship between postprandial TG and various demographic and metabolic parameters in 639 Korean subjects with type 2 diabetes (T2D, group I,  $n=539$ ) and impaired fasting glucose (IFG, group II,  $n=100$ ) after ingestion of a standardized liquid meal (total 500 kcal, 17.5 g fat, 68.5 g carbohydrate, and 17.5 g protein).

**Results:** Fasting and postprandial TG were significantly correlated ( $r=0.973$ ,  $r=0.937$ ,  $P<0.001$ ) in group I and II, respectively. Of the variables, total cholesterol, waist circumference and body mass index were significantly correlated with fasting and postprandial TG in both groups. Only postprandial TG showed a significant correlation with glucose metabolic parameters (e.g., postprandial glucose, homeostatic model assessment of insulin resistance [HOMA-IR], and fasting C-peptide) in subjects with T2D. Multiple regression analysis showed fasting TG and HOMA-IR could be predictable variables for postprandial TG in subjects with T2D.

**Conclusion:** Postprandial TG was very strongly correlated with fasting TG. The authors of the present study suggest insulin resistance may be more associated with postprandial TG than fasting TG in Korean T2D patients on a low-fat diet.

**Keywords:** Diabetes; Diet, fat-restricted; Non-fasting triglyceride; Triglycerides

### INTRODUCTION

Well-known characteristics of diabetic dyslipidemia include elevated levels of fasting triglyceride (TG), decreased levels of high density lipoprotein cholesterol (HDL-C), and increased portion of small dense low density lipoprotein cholesterol (LDL-C) particles in LDL cholesterol [1]. Similar to the abrupt surge in postprandial glucose after an oral glucose tolerance test in type 2 (T2) diabetics, a rise in circulating TG is a com-

mon occurrence after ingestion of a high-fat meal. Although elevated postprandial TG is expected after consumption of a fat-containing meal, not all subjects present hypertriglyceridemia requiring TG-lowering therapy. The levels of postprandial or non-fasting TG are mainly influenced by not only the diet contents [2] but also the time of sampling for non-fasting TG [3]. Postprandial TG level differs according to the fat amount in a meal. Several studies reported a very low (5 g) or low (<15 g) amount of dietary fat did not significantly increase post-

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prandial TG, a moderate (30 to 50 g) amount of fat increased postprandial TG dose dependently, and a very high (>80 g) amount of fat exaggeratedly increased postprandial TG without dose-dependence [2,4].

Postprandial TG has been suggested as an independent risk factor for cardiovascular disease (CVD) in healthy subjects [5-7]. Several studies support the hypothesis postprandial TG is correlated with carotid IMT and waist circumference (WC), and furthermore superior to fasting TG in association with metabolic syndrome [8-10]. However, the potential effects of postprandial hypertriglyceridemia on the development of atherosclerosis are still controversial in subjects with type 2 diabetes (T2D) [11]. In addition, the metabolic associations of postprandial TG in Korean patients with T2D are not well elucidated, and the clinical and laboratory parameters affecting the levels of non-fasting TG also remain unclear. Thus, the authors of the present study investigated the metabolic relation and clinical implications of postprandial TG after a particular low-fat diet in Korean T2 diabetic patients who used to have a balanced low-fat diet.

## METHODS

### Patients and research design

In a clinic-based, retrospective cross-sectional study, patients who satisfied specific criteria based on their medical records were analyzed. Inclusion criteria were patients enrolled in the diabetes registry of Severance Diabetes Center between July 2009 and August 2010. The subjects were registered first-time visitors to the center, patients who performed a standardized mixed-meal stimulation test (Ensure; total 500 kcal, 17.5 g fat, 68.5 g carbohydrate, and 17.5 g protein; Meiji Dairies Co., Tokyo, Japan) or patients who had diabetes complication work-up performed. In the registry protocol, blood samples were collected at 0 and 90 minutes (basal and stimulated level, respectively) for glucose, TG, insulin, and C-peptide analyses after injection of a standardized mixed-meal stimulation test. Exclusion criteria included insulin users, severe liver or kidney disease, thyroid disorders, pregnancy, steroid therapy, heavy alcoholics, and hematologic as well as malignant diseases. The study protocol was approved by the Ethics Committee of the Yonsei University College of Medicine. The subjects were classified into 2 groups based on the American Diabetes Association (ADA) 2011 guidelines: group I, subjects with T2D; group II, subjects with impaired fasting glucose subjects (IFG; in-

creased risk for diabetes).

Anthropometric measurements of individuals wearing light clothing and no shoes were conducted. WC was measured with the tape measure placed horizontally at the level of the umbilicus while the participant gently exhaled. The body mass index (BMI) was calculated as weight in kilograms divided by a square of height in meters.

### Laboratory measurements

Plasma TG, total cholesterol, HDL-C, blood urea nitrogen, creatinine, AST, and ALT, were assayed by routine Hitachi 7600 autoanalyzer (Hitachi Instruments Service, Tokyo, Japan). LDL-C was calculated using the Friedewald equation. Plasma glucose was measured using the glucose oxidase method, and A1C was measured by high-performance liquid chromatography (HPLC) using Variant II Turbo (Bio-Rad Laboratories, Hercules, CA, USA). Serum insulin and C-peptide levels were measured in duplicate by using an immunoradiometric assay (IRMA) method (Beckman Coulter, Fullerton, CA, USA). Samples for laboratory measurements were simultaneously collected during a standardized mixed-meal stimulation test.

### Assessment of the $\beta$ -cell functional status

Pancreatic  $\beta$ -cell function and insulin sensitivity were obtained by homeostasis model assessment (HOMA) using HOMA of pancreatic  $\beta$ -cell function (HOMA- $\beta$ ) calculated as basal insulin ( $\mu$ IU/mL) $\times$ 20/basal glucose (mmol/L)-3.5, and HOMA of insulin resistance (HOMA-IR) calculated as basal insulin ( $\mu$ IU/mL) $\times$ glucose (mmol/L)/22.5. Pancreatic  $\beta$ -cell secretory function or insulin reserve were also determined from a mixed-meal stimulation test calculated by dividing the increase in insulin level by the increase in glucose level over the same period (insulinogenic index; IGI=[insulin (90')-insulin (0')]/[glucose (90')-glucose (0')]).

### Statistical analyses

Data are given as means  $\pm$  standard deviation (SD) for normally distributed variables, and otherwise as medians (25th and 75th percentile values). Mean values and median value were compared between diabetes and IFG patients using unpaired Student's *t*-test and Mann-Whitney *U*-test as appropriate. Pearson's correlation coefficient was used to determine the correlation between continuous parameters. Multiple linear regression analyses were performed using postprandial TG as a dependent factor. Several clinical and laboratory factors (e.g.,

age, sex, WC, glucose AC, HDL-C, and HOMA-IR) were entered as independent factors. An alpha level of 0.05 was accepted as significant for all statistical procedures. All statistical analyses were conducted using SPSS for Windows software version 18.0 (SPSS Inc., Chicago, IL, USA).

## RESULTS

### Baseline characteristics of the patients

A total of 639 patients satisfied the criteria, and were classified into 2 groups depending on their glucose status: group I, T2D patients ( $n=539$ ); group II, IFG patients ( $n=100$ ). The patients' demographic and laboratory data are summarized in Table 1 and Supplemental Table 1. The mean age was 57.5 years, and gender (364 men and 275 women) was relatively evenly distributed. The mean age was 57.5 years, and gender (364 men and 275 women) was relatively evenly distributed.

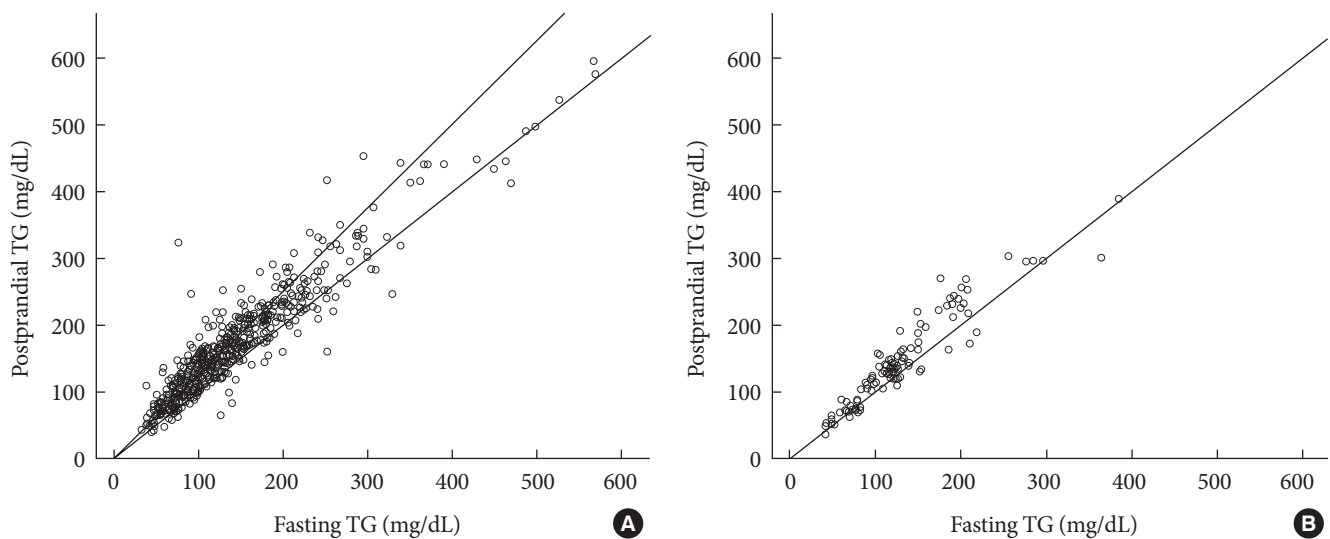
There were no significant differences in age, sex and BMI in both groups. The level of fasting and postprandial glucose, and A1c were significantly higher in group I ( $141.21 \pm 47.2$  mg/dL,  $255.30 \pm 78.14$  mg/dL,  $7.87 \pm 1.51$  mg/dL, respectively) than in group II ( $104.22 \pm 14.90$  mg/dL,  $146.46 \pm 32.24$  mg/dL,  $6.25 \pm 0.98$  mg/dL) ( $P < 0.001$  for all). Postprandial insulin, C-peptide and HOMA- $\beta$  were statistically higher in group II ( $67.30$  [10.12 to 259.96]  $\mu$ IU/mL,  $7.72 \pm 3.30$   $\mu$ g/L,  $62.28$  [9.47 to 654.00], respectively) than group I ( $50.68$  [2.77 to 350.00]  $\mu$ IU/mL,  $6.83 \pm 4.93$   $\mu$ g/L,  $46.35$  (-1,522 to 2,433), respectively) ( $P=0.015$ ,  $P=0.026$ ,  $P < 0.001$ ). WC, waist/hip ratio (WHR) and HOMA-IR were significantly higher in group I ( $88.11 \pm 9.38$  cm,  $0.91 \pm 0.058$ ,  $2.95$  [0.38 to 44.61], respectively) than in group II ( $85.75 \pm 10.08$  cm,  $0.89 \pm 0.057$ ,  $1.88$  (0.25 to 9.37), respectively) ( $P=0.030$ ,  $P=0.001$ ,  $P < 0.001$ ). However, fasting TG ( $126$  [33 to 2,010] vs.  $125.5$  [41 to 384]; group I vs. group II,  $P=0.432$ ) and postprandial TG ( $156$  [40 to 2,220] vs.  $142$  [37 to 388]; group I

**Table 1.** Baseline characteristics of the study subjects

Characteristic	All subjects ( $n=639$ )	Group 1 (T2D) ( $n=539$ )	Group 2 (IFG) ( $n=100$ )	P value
Age, yr	$57.57 \pm 11.15$	$57.7 \pm 11.2$	$56.59 \pm 10.77$	NS
Sex (male), %	57	59	52	NS
BMI, kg/m <sup>2</sup>	$25.52 \pm 3.66$	$25.22 \pm 3.75$	$25.58 \pm 3.64$	NS
WC, cm	$87.75 \pm 9.52$	$88.11 \pm 9.38$	$85.75 \pm 10.08$	0.030
WHR	$0.91 \pm 0.058$	$0.91 \pm 0.058$	$0.89 \pm 0.057$	0.001
Glucose at 0 min, mg/dL	$135.42 \pm 45.81$	$141.21 \pm 47.2$	$104.22 \pm 14.90$	<0.001
Glucose at 90 min, mg/dL	$238.39 \pm 82.91$	$255.30 \pm 78.14$	$146.46 \pm 32.24$	<0.001
A1c, %	$7.62 \pm 1.56$	$7.87 \pm 1.51$	$6.25 \pm 0.98$	<0.001
C-peptide at 0 min, $\mu$ g/L	$2.55 \pm 3.55$	$2.61 \pm 3.81$	$2.22 \pm 1.48$	NS
C-peptide at 90 min, $\mu$ g/L	$6.97 \pm 4.72$	$6.83 \pm 4.93$	$7.72 \pm 3.30$	0.026
Insulin at 0 min, $\mu$ IU/mL <sup>a</sup>	11.10 (1.00-81.76)	8.99 (1.27-81.76)	7.03 (1.00-43.60)	0.040
Insulin at 90 min, $\mu$ IU/mL <sup>a</sup>	65.52 (2.77-350.00)	50.68 (2.77-350.00)	67.30 (10.12-259.96)	0.015
Cholesterol, mg/dL	$177.59 \pm 63.44$	$175.74 \pm 65.85$	$188.60 \pm 45.38$	0.029
TG at 0 min, mg/dL <sup>a</sup>	148.4 (33-2,010)	126.0 (33-2,010)	125.5 (41-384)	NS
TG at 90 min, mg/dL <sup>a</sup>	175.3 (37-2,220)	156.0 (40-2,220)	142.0 (37-388)	NS
HDL-C	$63.60 \pm 34.14$	$64.26 \pm 34.7$	$59.70 \pm 30.57$	NS
LDL-C	$82.48 \pm 39.95$	$79.23 \pm 38.10$	$101.82 \pm 45.2$	<0.001
HOMA-IR <sup>a</sup>	3.71 (0.00-44.61)	2.95 (0.38-44.61)	1.88 (0.25-9.37)	<0.001
HOMA- $\beta$ <sup>a</sup>	75.28 (-1,522-2,433)	46.35 (-1,522-2,433)	62.28 (9.47-654.00)	<0.001
IGI <sup>a</sup>	15.78 (-640-1,251)	6.67 (-247-1,251)	22.98 (-640.80-475.65)	<0.001

T2D, type 2 diabetes; IFG, impaired fasting glucose; NS, not significant; BMI, body mass index; WC, waist circumference; WHR, waist/hip ratio; TG, triglyceride; HDL-C, high density lipoprotein cholesterol; LDL-C, low density lipoprotein cholesterol; HOMA-IR, homeostatic model assessment of insulin resistance; HOMA- $\beta$ , homeostatic model assessment of pancreatic  $\beta$ -cell function; IGI, insulinogenic index.

<sup>a</sup>Differences between the 2 groups were analyzed by Mann-Whitney *U*-test. Data were presented as the median and 25th and 75th percentile values.



**Fig. 1.** Correlation between plasma fasting triglyceride (TG) levels and postprandial TG levels in subjects with type 2 diabetes (A) and impaired fasting glucose (B).

vs. group II,  $P=0.052$ ) were not significantly different between the 2 groups.

#### Independent correlation between plasma TG levels (fasting and postprandial) and clinical and metabolic parameters

A very strong (at least 0.9) linear relationship was observed between fasting and postprandial TG in T2D and IFG groups ( $r=0.973$ ,  $r=0.937$ , respectively,  $P<0.001$  for both; Fig. 1). In group I, fasting TG and postprandial TG were statistically correlated with total cholesterol ( $r=0.309$ ,  $r=0.305$ , respectively), fasting glucose ( $r=0.151$ ,  $r=0.162$ ), A1c ( $r=0.143$ ,  $r=0.162$ ), WC ( $r=0.138$ ,  $r=0.123$ ), and BMI ( $r=0.118$ ,  $r=0.108$ ). Only postprandial TG was correlated with postprandial glucose ( $r=0.114$ ,  $P=0.009$ ), HOMA-IR ( $r=0.091$ ,  $P=0.037$ ), and fasting C-peptide ( $r=0.089$ ,  $P=0.042$ ).

In group II, fasting TG and postprandial TG were statistically correlated with postprandial insulin ( $r=0.328$ ,  $r=0.330$ , respectively), postprandial C-peptide ( $r=0.312$ ,  $r=0.320$ ), WC ( $r=0.303$ ,  $r=0.246$ ), BMI ( $r=0.275$ ,  $r=0.246$ ), total cholesterol ( $r=0.238$ ,  $r=0.246$ ), and HOMA-IR ( $r=0.226$ ,  $r=0.228$ , Table 2).

#### Multiple linear regression analysis (postprandial TG as a dependent variable)

Of the several clinical and laboratory variables, only age, sex, HOMA-IR, WC, fasting and postprandial TG, HDL, and A1c showed a multicollinearity referring to a situation in which 2

or more explanatory variables were highly linearly related during a multiple regression model. In multiple regression analysis, the postprandial TG was entered as the dependent variable. Independent variables were age, sex, fasting TG, HOMA-IR, HDL-C, glucose AC and WC. As shown in Table 3 the variables independently associated with postprandial TG were fasting TG in both groups ( $\beta=0.957$ ,  $P<0.001$  in the T2D group and  $\beta=0.906$ ,  $P<0.001$  in the IFG group) and HOMA-IR in the T2D group ( $\beta=0.827$ ,  $P=0.024$ ). In this regard, fasting TG and HOMA-IR could be a predictable variable for postprandial TG in subjects with T2D.

## DISCUSSION

The gold standard measurement for hypertriglyceridemia is the fasting TG. Although controversial, fasting TG is known to be an independent risk factor for CVD [5,12]. However, 15 to 18 hours a day are spent in the postprandial state and postprandial TG significantly contributes to overall TG exposure. In this regard, the use of fasting TG as a predictor for CVD does not provide information on diurnal TG profiles and individual metabolism [9]. Accordingly, postprandial or non-fasting TG levels are increasingly used as a predictor for CVD [13].

While traditional Korean meals are usually not high in fat, the Korean diet has become more westernized. The increase of postprandial TG is known as mainly influenced by diet composition [2,4,14]. Additionally, patient's characteristics such as

**Table 2.** Pearson's correlation coefficients between fasting and postprandial TG and metabolic parameters

Variable	T2D (n=539)		IFG (n=100)	
	Fasting TG	Postprandial TG	Fasting TG	Postprandial TG
Age, yr	- 0.124 (0.004)	- 0.125 (0.004)	NS	NS
BMI, kg/m <sup>2</sup>	0.118 (0.006)	0.108 (0.013)	0.275 (0.007)	0.246 (0.016)
WC, cm	0.138 (0.002)	0.123 (0.005)	0.303 (0.004)	0.266 (0.011)
Fasting TG, mg/dL		0.973 (<0.001)		0.937 (<0.001)
Postprandial TG, mg/dL	0.973 (<0.001)		0.937 (<0.001)	
Glucose at 0 min, mg/dL	0.151 (<0.001)	0.162 (<0.001)	NS	NS
Glucose at 90 min, mg/dL	NS	0.114 (0.009)	NS	NS
A1c, %	0.143 (0.001)	0.162 (<0.001)	NS	NS
C-peptide at 0 min, µg/L	NS	0.089 (0.042)	0.219 (0.029)	0.198 (0.051)
C-peptide at 90 min, µg/L	NS	NS	0.312 (0.002)	0.320 (0.001)
Insulin at 90 min, µIU/mL	NS	NS	0.328 (0.001)	0.330 (0.001)
Total cholesterol, mg/dL	0.309 (<0.001)	0.305 (<0.001)	0.238 (0.031)	0.246 (0.027)
HOMA-IR	NS	0.091 (0.037)	0.226 (0.024)	0.228 (0.023)

The numbers in parentheses refer to *P* values.

TG, triglyceride; T2D, type 2 diabetes; IFG, impaired fasting glucose; NS, not significant; BMI, body mass index; WC, waist circumference; HOMA-IR, homeostatic model assessment of insulin resistance.

**Table 3.** Multiple linear regression analysis (postprandial TG as the dependent variable)

Variable	T2D (n=539)		IFG (n=100)	
	Beta	<i>P</i> value	Beta	<i>P</i> value
Age, yr	- 0.017	0.212	-0.017	0.714
WC, cm	-0.008	0.572	-0.070	0.219
Fasting TG, mg/dL	0.957	<0.001	0.906	<0.001
Glucose at 0 min, mg/dL	0.018	0.214	0.021	0.435
HDL-C, mg/dL	0.014	0.297	0.021	0.342
HOMA-IR	0.827	0.024	0.065	0.248

TG, triglyceride; T2D, type 2 diabetes; IFG, impaired fasting glucose; WC, waist circumference; HDL-C, high density lipoprotein cholesterol; HOMA-IR, homeostatic model assessment of insulin resistance.

the use of lipid-lowering agents [15], WC [16,17], glycemic status [18,19], menopausal status and age [9] also affect the non-fasting TG profiles. Based on previous reports, the authors of the present study questioned the metabolic associations and clinical implications of postprandial TG in Korean patients with IFG and T2D. To answer these questions, the results of a standardized mixed-meal stimulation test were retrospectively analyzed in order to first identify the close correlation between fasting and postprandial TG and second, the clinical and labo-

ratory relevance of postprandial TG. In the present analysis, the mixed-meal stimulation test (Ensure; total 500 kcal, 17.5 g fat, 68.5 g carbohydrate, and 17.5 g protein) allowed for standardization of the calories, fat contents, and the times for blood sampling (basal and 90 minutes after meal loading). In addition, the influential effects of glucose status and insulin secretion or resistance on both fasting and non-fasting TG levels could be analyzed.

The present study shows 3 main results. First, fasting and postprandial TG demonstrated a very strong (at least 0.9) linear relationship in all subjects regardless of glucose status. In addition, the TG increment after a mixed-meal load was not contrasting. The likely explanation for these results is the type of low calorie and fat content diet (500 kcal, 17.5 g fat liquid loading) and a 30-minute earlier check of peak TG level. Specifically, 2 to 4 hours after a high-fat diet loading was the necessary time point for postprandial TG to reach the highest level [3,9,12-14,20]. Only 1 well-designed study that enrolled Korean young women showed postprandial TG reached the peak time at 120 minutes after a high-fat meal ingestion [3]. However, no increase in postprandial TG after eating a high carbohydrate meal was found [3]. Second, although both fasting and postprandial TG showed a similar correlation with other demographic and laboratory parameters in subjects with IFG,

only postprandial TG showed a significant correlation with glucose metabolic parameters such as postprandial glucose, HOMA-IR and fasting C-peptide in subjects with T2D. In both T2D and IFG groups, fasting and postprandial TG were statistically correlated with total cholesterol and central obesity parameters such as WC and BMI. Third, fasting TG and HOMA-IR could be predictable variables for postprandial TG in subjects with T2D. In subjects with impaired fasting glucose, only fasting TG level predicted the level of postprandial TG.

Regarding the characteristics of enrolled subjects, patients with T2D showed the typical pathophysiologic features with increased insulin resistance (WC, WHR, and HOMA-IR) and decreased insulin secretory function (HOMA- $\beta$ , insulin and C-peptide levels at 90 minutes) resulting in increased glucose and A1c values. However, levels of total cholesterol and LDL-C were significantly lower in subjects with T2D than with IFG (Table 1). The likely explanation involves the 50% and 27% usage of lipid-lowering drugs in T2D and IFG subjects, respectively, which may affect the lower levels of cholesterol.

There are several limitations of the present study. First, the study is a retrospective cross-sectional study, and the measurements were taken after a single liquid meal. Second, subjects were diabetic patients who are usually taking some type of drug which were not individually analyzed in the present study. However, the study protocol using the low-fat mixed-meal test and measuring postprandial TG at 90 minutes is appropriate to analyze the influential effects of glucose status and insulin secretion or resistance on both fasting and non-fasting TG levels in Korean subjects who traditionally do not eat a high-fat diet. This is the first, well-designed and relatively large scale study on clinical implication of postprandial TG on Korean T2 diabetics. Further studies are essential to investigate the postprandial TG in all drug-naïve pre-diabetic and diabetic patients.

In conclusion, postprandial TG was strongly correlated with fasting TG. In addition, the present study shows a low-fat meal may have largely similar dietary effects on fasting and postprandial TG in Koreans with T2D. Especially, insulin resistance may be more associated with postprandial TG than fasting TG in Korean diabetic patients on a low-fat diet.

## CONFLICTS OF INTEREST

No potential conflict of interest relevant to this article was reported.

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