

# Association of C-reactive protein with the metabolic risk factors among young and middle-aged Koreans

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## Abstract

We investigated the relationship between the clustering of risk factors for metabolic syndrome and the plasma C-reactive protein (CRP) concentration as measured by high-sensitive CRP assay. Body mass index, waist circumference, triglycerides (TGs), high-density lipoprotein cholesterol, fasting glucose, systolic and diastolic blood pressures, insulin, and CRP were measured in 1046 Korean adults (560 males; age, 18–64 years) in 2003 to 2004. There were statistically significant positive correlations for log CRP with body mass index, waist circumference, log TG, log insulin, and log homeostasis model assessment in both sexes after adjusting for age and smoking status. High-density lipoprotein cholesterol showed a significant negative correlation with log CRP in both sexes. For both sexes, the mean level of log CRP increased with increasing number of risk factors of metabolic syndrome ( $P$  for trend  $<.01$  for males and  $<.001$  for females). Stepwise multivariate linear regression analysis showed that waist circumference contributed the largest portion of the variance in CRP levels in both sexes. Log homeostasis model assessment and log TG were independently associated with log CRP levels only in females. These results indicate that CRP, a marker of inflammation that underlies atherosclerosis, is associated with the clustering of each metabolic syndrome risk factor and, furthermore, that abdominal obesity is the strongest predictor of CRP level in the Korean adult population.

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## 1. Introduction

Inflammation has been suggested as a risk factor for the development of atherosclerosis [1,2]. C-reactive protein (CRP), an acute-phase reactant and sensitive marker of subclinical inflammation, has proven to be a strong and independent predictor of both diabetes [3–7] and cardiovascular disease [8–12].

Metabolic syndrome, which is characterized by the clustering of abdominal obesity, hypertension, impaired glucose tolerance, and atherogenic dyslipidemia, is associated with increased risk of cardiovascular disease [13]. The more risk factors cluster, the greater risk of cardiovascular disease [14–16]. Recent studies have shown that elevated CRP is strongly associated with various characteristics of the metabolic syndrome [17–22].

Previous studies on the relationship of CRP and the metabolic syndrome, however, focused on the individual risk factors of metabolic syndrome, not the clustering of various risk factors [23–27]. In addition, little is known about this relationship in Asian people who exhibit more metabolic complications associated with obesity even at lower body mass index (BMI) than whites do [28–32]. Therefore, we investigated the association between plasma CRP concentration and the risk factors of metabolic syndrome in the Korean adult population.

## 2. Subjects and methods

The study population consisted of 1224 Korean men and women who visited Dankook University Hospital for health examination from December 2003 to January 2004. They responded to a self-reported questionnaire and underwent a physical examination, including height and weight, and blood pressure measured sitting after 5 minutes of rest, and

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provided a collection of fasting blood samples. We excluded 25 subjects with CRP level of 10 mg/L or higher, indicating clinically relevant inflammatory conditions; 52 subjects with diagnosed diabetes, fasting glucose of 126 mg/dL or higher, and diabetes medication; 44 subjects who were older than 65 years; and a further 57 subjects with a missing value for smoking status, which is an important confounding factor for CRP concentration. Consequently, the current analysis was confined to 560 men and 486 women with complete data on all components of the metabolic syndrome and serum CRP concentration.

### 2.1. Definition

Characteristics of the metabolic syndrome were based on the Third Report of the National Cholesterol Education Program Expert Panel on Detection, Evaluation, and Treatment of High Blood Cholesterol in Adults (Adult Treatment Panel [ATP] III) using the following cutoff limits: (1) blood pressure of 135/85 mm Hg or higher; (2) triglyceride (TG) of 1.7 mmol/L or higher (150 mg/dL); (3) low high-density lipoprotein cholesterol (HDL-C) of less than 1.0 mmol/L (40 mg/dL) for men and less than 1.3 mmol/L (50 mg/dL) for women; and (4) fasting glucose of 6.1 mmol/L or higher (110 mg/dL) [13]. Abdominal obesity was defined by Asia-Pacific cutoff limits: waist circumference of 90 cm or more for men and 80 cm or more for women because the importance of ethnic-specific cutoff points for waist circumference as a measure of central adiposity was recognized. Subjects with 3 or more risk

Table 1  
Clinical and biochemical characteristics of subjects

Variable	Male (n = 560)	Female (n = 486)	P
Age, mean $\pm$ SD (y)	42.7 $\pm$ 9.0	42.1 $\pm$ 10.2	NS
BMI, mean $\pm$ SD (kg/m <sup>2</sup> ) <sup>a</sup>	24.0 $\pm$ 2.9	22.9 $\pm$ 2.8	<.001
Waist circumference, mean $\pm$ SD (cm) <sup>a</sup>	84.8 $\pm$ 8.0	78.1 $\pm$ 8.0	<.001
HDL-C, mean $\pm$ SD (mg/dL) <sup>a</sup>	47.1 $\pm$ 10.7	53.5 $\pm$ 12.1	<.001
Fasting glucose, mean $\pm$ SD (mg/dL) <sup>a</sup>	90.7 $\pm$ 10.0	88.0 $\pm$ 8.3	<.001
TG, median (IQR) (mg/dL) <sup>b</sup>	130 (93–185)	96 (69–140)	<.001
Systolic blood pressure, mean $\pm$ SD (mm Hg) <sup>a</sup>	121.8 $\pm$ 12.8	118.0 $\pm$ 13.5	<.001
Diastolic blood pressure, mean $\pm$ SD (mm Hg) <sup>a</sup>	83.3 $\pm$ 8.8	79.2 $\pm$ 8.9	<.001
HOMA, median (IQR) <sup>b</sup>	0.5 (0.2–1.2)	0.2 (0.2–1.0)	<.01
Insulin, median (IQR) ( $\mu$ IU/mL) <sup>b</sup>	2.4 (0.9–5.3)	0.9 (0.9–4.8)	<.01
CRP, median (IQR) (mg/L) <sup>b</sup>	0.6 (0.3–1.3)	0.4 (0.2–0.9)	<.001
Smoking status (%) <sup>c</sup>			<.001
Never	23.8	49.8	
Former	25.4	43.2	
Current	50.8	7.0	
Medication (%) <sup>b</sup>			NS
Hypertension	4.9	6.2	

<sup>a</sup> *t* Test.

<sup>b</sup> Mann-Whitney *U* test.

<sup>c</sup>  $\chi^2$  Test.

Table 2

Partial Spearman correlation coefficients between log CRP and variables of metabolic syndrome adjusted for age and smoking status

Variable	Men	Women
BMI	0.20**	0.34**
Waist circumference	0.22**	0.28**
HDL-C	−0.15**	−0.16**
Log TG	0.12*	0.21**
Fasting glucose	0.03	0.16**
Systolic blood pressure	0.05	0.09
Diastolic blood pressure	0.01	0.09
Log HOMA	0.13*	0.21**
Log insulin	0.13*	0.20**

\* *P* < .01.

\*\* *P* < .001.

factors of the above criteria were diagnosed as having the metabolic syndrome.

If the participants were using antihypertensive medication, they were considered as participants with high blood pressure. Obesity was defined as BMI of 25 kg/m<sup>2</sup> or higher according to the Asia-Pacific criteria [33]. Smoking was classified into “nonsmoker,” “former smoker,” and “current smoker.” Insulin resistance was assessed from fasting glucose and insulin values using the following homeostasis model assessment (HOMA) calculation: insulin resistance = [fasting insulin ( $\mu$ IU/mL)  $\times$  fasting glucose (mmol/L)]/22.5. This formula moderately agreed with the hyperinsulinemic euglycemic clamp [34] in Korean population [35].

### 2.2. Laboratory measurements

Venous blood samples were collected from each subject after a 12-hour fast and used for assay of glucose, total cholesterol and HDL-C, and TG. Total cholesterol and TG were determined enzymatically. High-density lipoprotein cholesterol was measured with phosphotungstate method. Glucose was enzymatically determined with hexokinase method. Insulin was measured with immunoradiometric assay. C-reactive protein was measured with latex-enhanced immunonephelometry, on a Behring BNII Nephelometer (Dade Behring, Newark, DE). This assay used a calibrator traceable to World Health Organization Reference Material (Certified Reference Material 470). The lower detection limit of the assay was 0.16 mg/L. Undetectable CRP values were recorded as 0.16 mg/L. The intra- and interassay coefficients of variation for CRP were 4.0% and 2.6%, respectively.

### 2.3. Data analysis

The clinical and biochemical features of the population were presented as mean  $\pm$  SD, median (interquartile range [IQR]), or percentages. Because the distributions of CRP, insulin, TG, and HOMA were highly skewed, these variables were natural log-transformed for all other analyses. All analyses were done separately by sex because mean CRP levels differed by sex in the present population. Age and smoking status were the determinants of CRP levels in

Table 3

Geometric means of CRP (mg/L) adjusted for age and smoking status and 95% confidence interval (CI) for the variables of metabolic syndrome in men

Variable	n	Geometric means of CRP (95% CI)	P
<i>Obesity (BMI ≥ 25 kg/m<sup>2</sup>)</i>			
Yes	200	0.9 (0.7-1.0)	<.001
No	360	0.6 (0.5-0.7)	
<i>Abdominal obesity (waist circumference &gt;90 cm)</i>			
Yes	407	0.7 (0.7-0.8)	<.001
No	153	0.6 (0.5-0.6)	
<i>High blood pressure (≥ 130/85 mm Hg or antihypertensive medication)</i>			
Yes	281	0.7 (0.7-0.8)	.086
No	279	0.6 (0.6-0.7)	
<i>Low HDL-C (&lt;40 mg/dL)</i>			
Yes	137	0.8 (0.7-1.0)	<.05
No	423	0.6 (0.6-0.7)	
<i>Hyperglycemia (fasting glucose ≥ 110 mg/dL)</i>			
Yes	29	1.0 (0.7-1.4)	<.05
No	531	0.7 (0.6-0.7)	
<i>High TG (≥ 150 mg/dL)</i>			
Yes	225	0.8 (0.7-0.9)	<.05
No	335	0.6 (0.6-0.7)	

Data were calculated using analysis of covariance.

previous reports [25,26,36], and CRP levels also differed by age and smoking status in the present population; therefore, multivariate models were tailored to account for these possible confounders. Partial Spearman correlation coefficients adjusted for age and smoking status were calculated for CRP with variables of metabolic syndrome components, log insulin, and log HOMA. The adjusted mean values of log CRP according to the number of metabolic disorders were calculated by analysis of covariance. Using a general linear model, we tested the linear trends for increasing log CRP levels across the number of metabolic risk factors.

Next, stepwise multivariate linear regression analysis was fit for log CRP as a dependent variable to demonstrate the relative contribution of each of the metabolic syndrome risk factors to CRP. After age and smoking status were forced into the model as confounders, the following variables were included into the model: components of metabolic syndrome from ATP III (waist circumference, systolic blood pressure, HDL-C, log TG, and fasting glucose) and log HOMA. All statistical analyses were performed using the SPSS statistical software (version 11.5, SPSS, Chicago, IL).

### 3. Results

Table 1 presents the demographic and clinical characteristics of the 1046 subjects. Mean age of male and female subjects was 42.7 years (SD, 9.0 years) and 42.1 years (SD, 10.2 years), respectively (range, 18–64 years). The plasma concentration of CRP in males was significantly higher than

in females (median, 0.6 mg/L [IQR, 0.3–1.3 mg/L] vs median, 0.4 mg/L [IQR, 0.2–0.9 mg/L];  $P < .001$ ). The geometric mean plasma concentrations of CRP according to the smoking status were highest among current smokers and lowest among nonsmokers in both male and female subjects (data not shown).

Partial Spearman rank correlation coefficients of log CRP with variables of the metabolic syndrome adjusted for age and smoking status are shown in Table 2. There were statistically significant positive correlations in both sexes for CRP with BMI, waist circumference, log TG, log insulin, and log HOMA. However, there was a significant negative correlation of log CRP with HDL-C in both sexes ( $P < .001$ ). The strongest correlation was observed between log CRP and BMI in females ( $r = 0.34$ ,  $P < .001$ ) and between log CRP and waist circumference in males ( $r = 0.22$ ,  $P < .001$ ).

The geometric mean CRP levels in subjects with each risk factor of metabolic syndrome were significantly higher in both sexes than in subjects without risk factors after adjusting for age and smoking status (Tables 3 and 4). Nevertheless, the geometric mean CRP levels of male subjects with high blood pressure defined by ATP III criteria were not significantly different from those of males with normal blood pressure (Table 3). The geometric mean CRP levels of female subjects with high fasting glucose were not significantly different from those of females with normal blood glucose (Table 4).

The mean levels of log CRP (SD) for those with 0, 1, 2, 3, and more than 4 components of the metabolic syndrome

Table 4

Geometric means of CRP (mg/L) adjusted for age and smoking status and 95% CI for the variables of metabolic syndrome in women

Variable	n	Geometric means of CRP (95% CI)	<i>P</i>
<i>Obesity (BMI ≥ 25 kg/ m<sup>2</sup>)</i>			
Yes	107	0.9 (0.7-1.1)	<.001
No	379	0.4 (0.4-0.4)	
<i>Abdominal obesity (waist circumference &gt; 80 cm)</i>			
Yes	185	0.7 (0.6-0.8)	<.001
No	301	0.4 (0.4-0.5)	
<i>High blood pressure (≥ 130/85 mm Hg or antihypertensive medication)</i>			
Yes	163	0.6 (0.5-0.7)	.001
No	323	0.4 (0.4-0.5)	
<i>Low HDL-C (&lt;50 mg/dL)</i>			
Yes	198	0.6 (0.6-0.7)	<.001
No	288	0.4 (0.4-0.4)	
<i>Hyperglycemia (fasting glucose ≥ 110 mg/dL)</i>			
Yes	7	0.8 (0.4-1.6)	.206
No	479	0.5 (0.4-0.5)	
<i>High TG (≥ 150 mg/dL)</i>			
Yes	104	0.7 (0.6-0.8)	<.001
No	382	0.4 (0.4-0.5)	

Data were calculated using analysis of covariance.

were  $-0.66$  (1.01),  $-0.59$  (1.00),  $-0.30$  (0.96),  $0.27$  (0.94), and  $0.02$  (0.84) mg/L in males, and  $-1.18$  (0.76),  $-0.92$  (0.93),  $-0.46$  (1.01),  $-0.07$  (1.02), and  $-0.04$  (0.98) mg/L in females, respectively. There was a linear increase in log CRP with the increasing number of metabolic syndrome risk factors in both sexes ( $P$  for trend  $<.001$  for males and  $<.001$  for females) (Fig. 1A and B).

The relative contribution of each component of metabolic syndrome to CRP level was examined by stepwise linear regression analysis. After age and smoking status were forced into the model, the following variables were assessed: waist circumference, log TG, fasting blood glucose, systolic blood pressure, HDL-C, and log HOMA. Only those variables with a  $P$  value of less than .05 were included in the final fitted model. Stepwise multivariate linear regression analysis showed that waist circumference was independently associated with log CRP levels in males, whereas waist circumference, log HOMA, and log TG were associated in females. The model  $R^2$  was 8.2% for male subjects and 19.5% for female subjects. Waist

Table 5

Stepwise multiple linear regression analysis with log CRP as dependent variable

Variable	$\beta$	SE ( $\beta$ )	$P$	Partial $R^2$ (%)
<i>Male</i>				
Waist circumference	.027	.005	.000	5.0
<i>Female</i>				
Waist circumference	.028	.006	.000	9.5
Log HOMA	.167	.042	.000	3.3
Log TG	.217	.097	.026	1.4

A final predictive model was derived by stepwise linear regression analysis. After age and smoking status were forced into the model, the following candidate independent variables were assessed: waist circumference, log TG, fasting blood glucose, systolic blood pressure, HDL-C, and log HOMA. Only those variables with a  $P$  value of less than .05 were included in the final fitted model. The model  $R^2$  for male subjects was 8.2% and for female subjects was 19.5%.  $\beta$  indicates linear regression coefficient.

circumference had the largest partial  $R^2$  in both men and women (Table 5).

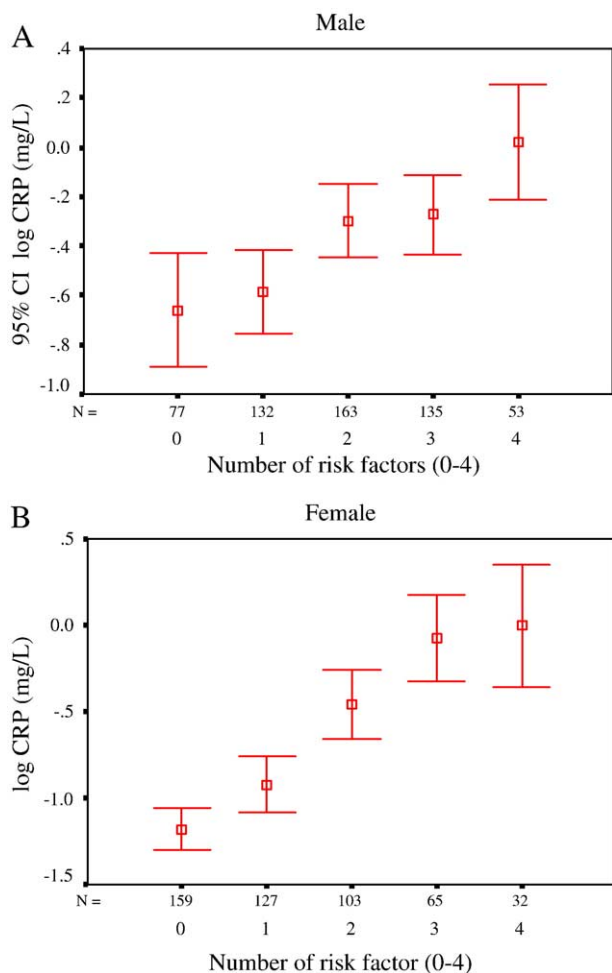


Fig. 1. Mean levels of log CRP and 95% CI adjusted for age and smoking status according to the number of components of the metabolic syndrome. A, Mean log CRP levels adjusted for age and smoking status in males ( $P_{\text{trend}} < .001$  by GLM). B, Mean log CRP levels adjusted for age and smoking status in females ( $P_{\text{trend}} < .001$  by General Linear Model).

#### 4. Discussion

The present study has demonstrated that CRP, a sensitive marker of inflammation, was associated with metabolic syndrome in the Korean adult population. After adjustment for age and smoking status, the geometric mean CRP levels in both sexes were significantly higher in those subjects who had each characteristic of the metabolic syndrome than in those who did not, except for males with high blood pressure and except for females with high fasting glucose. Previous studies [17–22] have shown that CRP levels are strongly associated with various characteristics of the metabolic syndrome, as has our present study. Only in male subjects with high blood pressure and in female subjects with high fasting glucose were the geometric mean CRP levels not significantly higher than in those with normal blood pressure and normal fasting glucose. These differences may have been caused by the single blood pressure measurement and small number of female subjects with high fasting glucose.

There was a strong linear increase in log CRP in both sexes as the number of components of metabolic disorders increased. Previous studies have shown a positive association between CRP and components of the metabolic syndrome cluster. Tamakoshi et al [21] reported that the odds ratio for elevated CRP is increased with an increasing number of metabolic syndrome factors. Aronson et al [20], Festa et al [17], and Frohlich et al [18] also showed that there is a strong linear increase in log CRP with an increasing number of metabolic syndrome factors among various populations, as we found with young and middle-aged Koreans. The mechanism through which each component of metabolic syndrome increases CRP levels directly has not yet been established. Each component of metabolic syndrome is a well-known risk factor in the atherosclerotic



process. Our results suggest that metabolic syndrome, especially the clustering of its components, is associated with a systemic low-grade inflammation, which causes the progression of atherosclerosis.

Multiple linear regression analysis, which uses log CRP levels as the dependent variable and all components of metabolic syndrome and log HOMA as the independent variable, showed that waist circumference contributes most of the variance in CRP levels in both males and females. Because many other factors than metabolic risk variables such as sex, smoking, age, ethnicity, insulin resistance, and nonspecific inflammation may affect the level of CRP, the total variance of CRP explained by them was small. Our finding suggests that the centrally distributed body fat, as evaluated by waist circumference, is a stronger predictor of CRP level than other components of the metabolic syndrome and insulin resistance, as measured by HOMA. The association between central adiposity and increased CRP has been noted in previous studies [17–19]. Experimental studies suggest that abdominal adipose tissue is a major source of cytokines, including interleukin 6, which is an important determinant of hepatic CRP synthesis [22,37]. Because waist circumference is the best surrogate of visceral adiposity across a wide age range [38], it is possible to say that CRP level is significantly associated with visceral adipose tissue, as evaluated by waist circumference, in Korean subjects. In other words, accumulation of visceral adipose tissue is a key factor of both features of the metabolic syndrome and low-grade chronic inflammation. Findings from other studies that CRP levels decreased in response to weight loss [39–44] also support our results.

Some studies have described an independent inverse association between measures of insulin resistance and CRP [4,45]. By contrast, other studies failed to identify a relationship between serum markers of inflammation and measures of insulin resistance after controlling for the effect of obesity [20,46–48]. In our study, log HOMA was independently associated with CRP only in females. Nevertheless, its contribution was also small compared with waist circumference. Future research is needed to identify the relationship between insulin resistance and CRP independent of body mass.

Our study had a few limitations. First, as the study subjects comprised only those who visited the hospital for a routine health check, the findings might not be generalized to the general Korean population. Second, CRP levels in women were low compared with other studies in western countries [49–51]. Our study population of women were relatively healthy and young, and had smaller BMI (only 23.5% of women had BMI >25) compared with those in studies from western countries [49], and ethnic differences between whites and Asians [12] including Koreans might have contributed this difference. Third, the levels of fasting insulin were extremely low compared with other studies. It may be related with the variability of insulin measurements from different laboratories [52]. Relatively lean and young

subjects in our studies than in other studies from western countries may contribute to these low insulin levels. Lastly, the HOMA, as a marker of insulin resistance, only showed moderate association in Korean population including diabetic subjects. Further studies are needed to examine the association of the HOMA and the hyperinsulinemic euglycemic clamp in nondiabetic Korean population.

In conclusion, we found that CRP level was associated with the clustering of each metabolic syndrome risk factor and that waist circumference was the strongest predictor of CRP levels in Korean adults. This finding suggests that metabolic syndrome is associated with a systemic inflammatory response, which plays an important pathogenetic role in atherothrombotic disease. We can confidently conclude that reducing waist circumference is probably the most important intervention to ameliorate the proinflammatory state associated with the metabolic syndrome.

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