Association of gamma-glutamyl transferase with subclinical coronary atherosclerosis and cardiac outcomes in non-alcoholics

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In an asymptomatic population, we determined the relationship between serum gamma-glutamyl transferase (GGT) and subclinical atherosclerosis, using coronary computed tomography angiography (CCTA). This was a retrospective observational cohort study which analyzed 5120 consecutive asymptomatic individuals with no prior history of coronary artery disease or significant alcohol intake who voluntarily underwent CCTA as part of a general health examination. All subjects were stratified into tertiles based on GGT levels. Degree and extent of subclinical coronary atherosclerosis were evaluated using CCTA. Cardiac events were a composite of all-cause death, myocardial infarction, unstable angina, and coronary revascularization. After adjustment for cardiovascular risk factors, there were no significant differences among GGT tertiles in terms of adjusted odds ratios for non-calcified and mixed plaques. The risk of any atherosclerotic and calcified plaques, significant stenosis, multi-vessel disease, and significant stenosis in the left main or proximal left anterior descending artery was higher in the third GGT tertile than in the first tertile (all \( p < 0.05 \)). Over a median 5.4-year follow-up, the third GGT tertile had significant adjusted hazards ratios for cardiac events than did the first GGT tertile, even after stepwise adjustment for cardiovascular risk factors (all \( p < 0.01 \)). In asymptomatic individuals, elevated GGT was independently associated with high-risk feature atherosclerosis and poorer cardiac outcomes.

Gamma-glutamyl transferase (GGT) is a hepatobiliary enzyme synthesized in epithelial cells of the intrahepatic duct. Elevated GGT is a specific indicator of obstructive biliary diseases and excessive alcohol intake. An association between serum GGT and nonalcoholic fatty liver disease (NAFLD) has been established, and some authors consider serum GGT as a surrogate marker of NAFLD. Several studies have reported that serum GGT levels are related to cardiovascular diseases (CVD). Based on studies to date, it may be assumed that GGT represents a link between NAFLD and atherosclerosis, with insulin resistance as a common pathophysiology. Recently, coronary computed tomography angiography (CCTA) has proven to be effective in providing a comprehensive evaluation of coronary atherosclerosis. An association of elevated serum GGT and subclinical coronary atherosclerosis detected by CCTA has also been reported. However, there are limited data on clinical outcomes including future cardiac events. Therefore, the present study sought to evaluate the relationship between serum GGT and subclinical coronary atherosclerosis as well as cardiac events in a large cohort of asymptomatic Korean individuals who voluntarily underwent CCTA for early detection of coronary artery disease (CAD).

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Association between GGT levels and subclinical atherosclerosis. The association between GGT levels and subclinical atherosclerosis is described in Table 3. Univariable analyses revealed that increasing tertiles of GGT were significantly associated with subclinical coronary atherosclerosis detected by CCTA. After adjustment for cardiovascular risk factors (age, sex, obesity, diabetes mellitus, hypertension, hyperlipidemia, current smoking, family history of CAD, and hs-CRP), no statistically significant differences were observed in the adjusted odds ratios (OR) for non-calcified and mixed plaques between GGT tertiles. The risk of atherosclerotic plaque involvement, segment stenosis score, and modified Duke prognostic score increased with GGT tertile (all p < 0.001). Furthermore, the prevalence of significant coronary artery stenosis, multi-vessel disease, and significant stenosis in the left main (LM) or proximal left anterior descending (LAD) artery also increased according to GGT tertile (all p < 0.001).

Table 1. Baseline characteristics of the study population according to the tertiles of gamma-glutamyl transferase. Values are presented as median (interquartile range) or number (%). Pearson's chi-square test or Fisher's exact test were used for categorical variables and one-way analysis of variance or Kruskal–Wallis test were used for numerical variables. Coronary artery disease in a first-degree relative of any age.

| Characteristics                        | Overall (n = 5,120) | Gamma-glutamyl transferase | Tertile 1 ≤ 15 IU/L (n = 1618) | Tertile 2 16–27 IU/L (n = 1791) | Tertile 3 ≥ 28 IU/L (n = 1711) | p value |
|----------------------------------------|---------------------|-----------------------------|-------------------------------|-------------------------------|-------------------------------|---------|
| Age, years                             | 53 (49–58)          | 53 (49–59)                  | 54 (50–59)                    | 53 (49–57)                    | <0.001                        |
| Male sex, no. (%)                      | 3486 (68.1)         | 618 (38.2)                  | 1364 (76.2)                   | 1504 (87.9)                   | <0.001                        |
| Body mass index, kg/m²                 | 24.3 (22.5–26.2)    | 23.0 (21.2–24.7)            | 24.5 (22.8–26.2)              | 25.5 (23.8–27.2)              | <0.001                        |
| Waist circumference, cm                | 85 (80–90)          | 80 (75–86)                  | 86 (80–90)                    | 89 (84–93)                    | <0.001                        |
| Systolic blood pressure, mmHg          | 119 (110–128)       | 115 (107–124)               | 119 (110–128)                 | 122 (113–130)                 | <0.001                        |
| Diastolic blood pressure, mmHg         | 76 (69–83)          | 72 (65–79)                  | 76 (69–83)                    | 79 (73–86)                    | <0.001                        |
| Diabetes mellitus, no. (%)             | 772 (15.1)          | 147 (9.1)                   | 280 (15.6)                    | 345 (20.2)                    | <0.001                        |
| Hypertension, no. (%)                  | 1778 (34.7)         | 406 (25.1)                  | 634 (35.4)                    | 738 (43.1)                    | <0.001                        |
| Hyperlipidemia, no. (%)                | 1559 (30.4)         | 358 (22.1)                  | 555 (31.0)                    | 646 (37.8)                    | <0.001                        |
| Current smoker, no. (%)                | 1024 (20.0)         | 136 (8.4)                   | 371 (20.7)                    | 517 (30.2)                    | <0.001                        |
| Obesity, no. (%)                       | 2114 (41.3)         | 362 (22.4)                  | 756 (42.3)                    | 996 (58.2)                    | <0.001                        |
| Family history of coronary artery disease, no. (%) | 788 (15.4) | 252 (15.6)                  | 282 (15.7)                    | 254 (14.8)                    | 0.739                          |
| Fasting blood glucose, mg/dL           | 99 (93–108)         | 96 (91–102)                 | 100 (94–109)                  | 103 (96–114)                  | <0.001                        |
| Total cholesterol, mg/dL               | 193 (172–217)       | 190 (171–214)               | 192 (171–214)                 | 198 (176–221)                 | <0.001                        |
| Low-density lipoprotein cholesterol, mg/dL | 120 (101–120)   | 117 (99–136)                | 121 (102–140)                 | 123 (104–143)                 | <0.001                        |
| High-density lipoprotein cholesterol, mg/dL | 51 (43–61)       | 57 (48–68)                  | 49 (43–60)                    | 48 (41–56)                    | <0.001                        |
| Triglycerides, mg/dL                   | 108 (78–156)        | 84 (64–113)                 | 109 (81–151)                  | 138 (100–201)                 | <0.001                        |
| Creatinine, mg/dL                      | 0.9 (0.8–1.0)       | 0.8 (0.7–0.9)               | 0.9 (0.8–1.0)                 | 0.9 (0.9–1.0)                 | <0.001                        |
| Uric acid, mg/dL                       | 5.5 (4.5–6.4)       | 4.7 (3.9–5.6)               | 5.6 (4.8–6.4)                 | 6.0 (5.2–7.0)                 | <0.001                        |
| Aspartate aminotransferase, IU/L       | 25 (21–31)          | 22 (19–26)                  | 24 (21–29)                    | 28 (24–36)                    | <0.001                        |
| Alanine aminotransferase, IU/L         | 22 (16–31)          | 17 (13–21)                  | 22 (17–29)                    | 30 (22–43)                    | <0.001                        |
| High-sensitivity C-reactive protein ≥ 2 mg/L | 47 (9.0)          | 7 (0.4)                     | 21 (1.2)                      | 19 (1.1)                      | 0.046                          |

Results

Baseline characteristics. All subjects were classified into tertiles based on GGT levels (Tertile 1 ≤ 15 IU/L, n = 1618; Tertile 2, 16–27 IU/L, n = 1791; Tertile 3, ≥ 28 IU/L, n = 1711). The baseline characteristics of 5120 study participants according to the tertiles of GGT levels are summarized in Table 1. The mean age of study participants was 53.8 ± 7.6 years, and 3486 (68.1%) were males. The prevalence of males, diabetes mellitus, hypertension, hyperlipidemia, current smoking, and obesity was significantly higher in tertiles of increasing GGT levels. In addition, BMI, waist circumference, systolic and diastolic blood pressure, as well as levels of fasting blood glucose, total cholesterol, LDL cholesterol, triglyceride, uric acid, AST, and ALT increased with GGT category. In contrast, levels of HDL cholesterol were lower in tertiles of decreasing GGT levels.

CCTA findings. CCTA findings according to GGT tertiles are showed in Table 2. Mean CACS was 38.2 ± 136.4. Among the study participants, 405 (7.9%) had significant stenosis of coronary artery (≥ 50% stenosis diameter) in at least one coronary artery on CCTA. Mean CACS increased with GGT category (p < 0.001). A significant difference was presented in the prevalence of calcified, non-calcified, mixed, or any atherosclerotic plaques according to GGT level (all p < 0.001). Scores which reflect plaque burden on CCTA such as segment involvement score, segment stenosis score, and modified Duke prognostic score increased with GGT tertile (all p < 0.001). Furthermore, the prevalence of significant coronary artery stenosis, multi-vessel disease, and significant stenosis in the left main (LM) or proximal left anterior descending (LAD) artery also increased according to GGT tertile (all p < 0.001).
Table 2. Comparison of coronary computed tomography angiographic findings according to the tertiles of gamma-glutamyl transferase. Values are presented as mean ± standard deviation or number (%). Pearson’s chi-square test or Fisher’s exact test were used for categorical variables and one-way analysis of variance or Kruskal–Wallis test were used for numerical variables.

| Variables                                             | Overall                        | Gamma-glutamyl transferase | p value |
|-------------------------------------------------------|--------------------------------|----------------------------|---------|
| Mean coronary artery calcium score                    | 38.2 ± 136.4                   | Tertile 1: 20.8 ± 74.9     | < 0.001 |
|                                                      |                                | Tertile 2: 42.6 ± 152.2    |         |
|                                                      |                                | Tertile 3: 50.0 ± 160.3    |         |
| Any atherosclerotic plaque, no. (%)                   | 2083 (40.7)                    | Tertile 1: 471 (29.1)      |         |
|                                                      |                                | Tertile 2: 778 (43.4)      |         |
|                                                      |                                | Tertile 3: 834 (48.7)      |         |
| Plaque characteristics, no. (%)                       |                                |                            |         |
| Calcified plaque                                      | 1371 (26.8)                    | Tertile 1: 300 (18.5)      | < 0.001 |
|                                                      |                                | Tertile 2: 521 (29.1)      |         |
|                                                      |                                | Tertile 3: 550 (32.1)      |         |
| Non-calcified plaque                                  | 940 (18.4)                     | Tertile 1: 214 (13.2)      | < 0.001 |
|                                                      |                                | Tertile 2: 346 (19.3)      |         |
|                                                      |                                | Tertile 3: 380 (22.2)      |         |
| Mixed plaque                                          | 451 (8.8)                      | Tertile 1: 91 (5.6)        | < 0.001 |
|                                                      |                                | Tertile 2: 164 (9.2)       |         |
|                                                      |                                | Tertile 3: 196 (11.5)      |         |
| Segment involvement score                             | 1.0 ± 1.7                      | Tertile 1: 0.7 ± 1.4       | < 0.001 |
|                                                      |                                | Tertile 2: 1.1 ± 1.8       |         |
|                                                      |                                | Tertile 3: 1.3 ± 1.9       |         |
| Segment stenosis score                                | 0.6 ± 1.9                      | Tertile 1: 0.3 ± 1.2       | < 0.001 |
|                                                      |                                | Tertile 2: 0.7 ± 2.0       |         |
|                                                      |                                | Tertile 3: 0.8 ± 2.3       |         |
| Modified Duke prognostic score                       | 1.2 ± 0.6                      | Tertile 1: 1.1 ± 0.4       | < 0.001 |
|                                                      |                                | Tertile 2: 1.2 ± 0.7       |         |
|                                                      |                                | Tertile 3: 1.2 ± 0.7       |         |
| Number of stenosed coronary arteries, no. (%)         |                                |                            |         |
| Significant stenosis                                  | 405 (7.9)                      | Tertile 1: 78 (4.8)        | < 0.001 |
|                                                      |                                | Tertile 2: 146 (8.2)       |         |
|                                                      |                                | Tertile 3: 181 (10.6)      |         |
| One-vessel disease                                    | 298 (5.8)                      | Tertile 1: 67 (4.1)        | < 0.001 |
|                                                      |                                | Tertile 2: 102 (5.7)       |         |
|                                                      |                                | Tertile 3: 129 (7.5)       |         |
| Multi-vessel disease                                  | 107 (2.1)                      | Tertile 1: 11 (0.7)        | < 0.001 |
|                                                      |                                | Tertile 2: 44 (2.5)        |         |
|                                                      |                                | Tertile 3: 52 (3.0)        | < 0.001 |
| Left main or proximal left anterior descending artery | 136 (2.7)                      | Tertile 1: 21 (1.3)        | < 0.001 |
|                                                      |                                | Tertile 2: 54 (3.0)        |         |
|                                                      |                                | Tertile 3: 61 (3.6)        |         |

Table 3. Association between gamma-glutamyl transferase levels and coronary computed tomography angiographic findings. CI: confidence interval. Covariates in the multivariable model include age, sex, obesity, diabetes mellitus, hypertension, hyperlipidemia, creatinine, uric acid, current smoking, family history of coronary artery disease, and high-sensitivity C-reactive protein ≥ 2 mg/L.
1.22 − 4.98, \( p = 0.012 \), and significant stenosis in the LM or proximal LAD artery (OR 1.94, 95% CI: 1.11 − 3.38, \( p = 0.020 \)) was higher in the third GGT tertile than in the first GGT tertile.

Clinical outcomes. During the follow-up period (median 5.4 years [interquartile range, 4.4–6.3 years]), a total of 165 cardiac events occurred in 154 patients: 52 all-cause deaths, three myocardial infarctions, 10 unstable angina requiring hospitalization, and 100 coronary revascularizations (Table 4). After adjustment for stepwise cardiovascular risk factors, the third GGT tertile had a significant adjusted hazard ratios for cardiac event, which was defined as a composite of all-cause death, myocardial infarction, unstable angina requiring hospitalization, or coronary revascularization, relative to that of the first GGT tertile (all \( p < 0.05 \), Table 5).

Discussion. The main findings of the present study were as follows: (1) in asymptomatic individuals, elevated serum GGT levels were significantly associated with atherosclerotic plaques, especially calcified plaques, even after adjustment for cardiovascular risk factors; (2) high serum GGT levels were an independent predictor of significant coronary atherosclerosis such as significant stenosis in at least one coronary artery, multi-vessel disease, and significant stenosis in the LM or proximal LAD; (3) during a follow-up of median 5.4 years, individuals with high serum GGT levels experienced more cardiac events.

The association between elevated GGT levels and risk of CAD or CVD has been suggested in large epidemiologic studies. Subsequent studies have reported that higher GGT levels were significantly associated with endothelial dysfunction, carotid artery plaques, and arterial stiffness. Moreover, recent studies have indicated that GGT levels were significantly related to high CACS and its progression, which reflects total atherosclerotic burden and is a surrogate marker of future CAD events. However, the absence of coronary calcification does not exclude the presence of clinically significant and potentially vulnerable atherosclerotic plaques. Although CCTA provides more comprehensive information regarding coronary atherosclerosis, there are limited data on the association between GGT and subclinical coronary atherosclerosis evaluated by CCTA. An observational cohort study with CCTA investigated the association between GGT levels and coronary atherosclerotic plaques. However, it analyzed limited data without clinical outcomes. Therefore, the present study aimed to evaluate the influence of GGT levels on the risk of subclinical coronary atherosclerosis on CCTA and cardiac outcomes.

In the present study, individuals with higher GGT levels had a higher prevalence, extent, and severity of coronary atherosclerosis detected by CCTA. Even after adjustments for clinical and laboratory variables, higher GGT levels were an independent predictor of coronary atherosclerotic plaques, especially calcified plaques.
Thus, our results were consistent with previous studies evaluating CACS\(^1\). Notably, higher GGT levels were an independent risk factor for high-risk feature CAD such as multi-vessel disease and significant stenosis in the LM or proximal LAD, which are known to be associated with poorer prognosis\(^8\). As a result, in the current study, individuals with higher GGT levels experienced more cardiac events. Therefore, our findings suggest that subjects with high serum GGT levels may benefit from preemptive cardiovascular risk evaluation and potentially, regular cardiovascular risk surveillance to guard against future cardiovascular events.

Although the exact mechanisms underpinning the association between serum GGT and CVD remain unclear, one potential pathophysiology is that they may be linked via NAFLD. Clinically, GGT is a hepatic enzyme that reflects the degree of hepatic inflammation or fibrosis\(^20\) and has been suggested as a surrogate marker for NAFLD\(^22\). NAFLD is regarded as the hepatic manifestation of metabolic syndrome and insulin resistance, which may increase the risk of atherosclerosis\(^7,8,21\). A chronic activation of systemic and hepatic inflammation may also act as a common pathway for NAFLD and CVD\(^22\). NAFLD is progressive across simple fatty livers to steatohepatitis, fibrosis, and finally liver cirrhosis. In particular, steatohepatitis with the potential of hepatic inflammation or fibrosis has been reported to increase the risk of cardiovascular mortality and morbidity\(^23\). To diagnose NAFLD, a liver biopsy is required which has several limitations including invasiveness, complications, sampling variability, and cost\(^26,27\). Ultrasonography typically usually employed as a substitute for liver biopsy, but there is a limit to the distinction between simple fatty liver and steatohepatitis\(^28\). Instead, noninvasive methods to determine the presence of hepatic inflammation or fibrosis have been developed. Of these, NAFLD fibrosis score (NFS) and Fibrosis-4 (FIB-4) are representative scoring systems that have been well-validated\(^30\). NFS include factors representing metabolic syndrome such as BMI or impaired glucose tolerance/diabetes mellitus, while FIB-4 includes age and serum markers. Although several studies have presented a significant association between NFS or FIB-4 and the prevalence or future risk of CVD, the results were not fully adjusted for other cardiovascular risk factors\(^32\). This study sought to evaluate the relationship between GGT levels as a noninvasive marker for hepatic inflammation or fibrosis and characteristics of atherosclerosis with their prognosis in a large cohort of asymptomatic Korean individuals who voluntarily underwent CCTA, which has proven to be effective in providing a comprehensive evaluation of coronary atherosclerosis\(^10\).

Our study has several limitations. First, the present study was performed at a single center, and all study participants voluntarily visited the hospital for a general health examination. Therefore, there was a potential of selection bias. Second, our study participants were exclusively Korean, limiting the applicability of our findings to other ethnic groups. Third, as the current study was a retrospective cohort study, these data may not fully reflect patient outcomes. Additionally, we did not specify the cause of death. Fourth, calcified plaques and higher CACS may lead to overestimation of significant coronary artery stenosis. Finally, CCTA has potential drawbacks, including radiation hazards, use of contrast, and higher cost. Therefore, although this study enrolled only volunteers, the use of CCTA in asymptomatic individuals cannot be justified.

In conclusion, this large retrospective cohort study in asymptomatic individuals undergoing CCTA demonstrated that elevated serum GGT levels were independently associated with high-risk subclinical atherosclerosis, resulting in poorer cardiac outcomes. Therefore, cardiovascular risk evaluation and surveillance should be considered in subjects with high GGT levels to prevent future cardiovascular disease. These findings should be further investigated and validated in future studies.

**Materials and methods**

**Study population.** This was a retrospective observational cohort study which analyzed 9269 consecutive South Korean individuals aged ≥ 20 years who had undergone self-referred CCTA evaluation as part of a general health examination at the Health Screening and Promotion Center in the Asan Medical Center between January 2007 and December 2011. Among these, 7129 (76.9%) agreed to participate in the present study. Possible risks associated with CCTA were explained, and written informed consent was obtained from each participant. Exclusion criteria included subjects with (1) a previous history of significant alcohol intake with ≥ 210 g/week in males and 140 g/week in females\(^34\), (2) a previous history of angina or myocardial infarction; (3) abnormal resting electrocardiography results, i.e., pathological Q waves, ischemic ST segments or T-wave changes, or left bundle-branch blocks; (4) insufficient medical records; (5) structural heart disease; (6) unmeasured GGT; (7) a previous cardiac procedure; (8) a previous history of open-heart surgery or percutaneous coronary intervention; or (9) renal insufficiency (creatinine > 1.5 mg/dL). A total final of 5120 subjects were enrolled (Fig. 1). The study was approved by the local Institutional Review Board of the Asan Medical Center, Seoul, Korea (2016–1068). Written informed consent under the ‘Ethics, consent and permissions’ was obtained from each participant. This study was carried out in accordance with Good Clinical Practice (GCP) guidelines and the Declaration of Helsinki.

**Clinical and laboratory measurements.** Basic demographic data of study participants were acquired from a database maintained by the Health Screening and Promotion Center at the Asan Medical Center. Medical history including angina, myocardial infarction, stroke, structural heart disease, open heart surgery, percutaneous coronary intervention, previous cardiac procedures, diabetes mellitus, hypertension, hyperlipidemia, or smoking status, was obtained from the responses in the systemized self-report questionnaire issued prior to the general health examination\(^17\).

Height and weight were obtained with subjects wearing light clothing and no shoes. Body mass index (BMI) was calculated as weight in kilograms divided by the square of height in meters (kg/m\(^2\)). Waist circumference (cm) was measured mid-way between the lower costal margin and iliac crest at the end of a normal expiration of breath by a well-trained nurse. Blood pressure was measured on the right arm after a ≥ 5 min rest using an automatic manometer and an appropriate cuff size. After overnight fasting, early morning blood samples were drawn from the antecubital vein into vacuum tubes and subsequently analyzed in the central certified laboratory.
of the Asan Medical Center. The concentrations of GGT, aspartate aminotransferase (AST), alanine aminotransferase (ALT), fasting plasma glucose, creatinine, uric acid, fasting total cholesterol, triglycerides, high-density lipoprotein (HDL) cholesterol, low-density lipoprotein (LDL) cholesterol, and high-sensitivity C-reactive protein (hs-CRP) were measured.

Obesity was defined as a BMI ≥ 25 kg/m² on the basis of an Asian-specific cutoff point as recommended in the World Health Organization. Diabetes mellitus was defined as subjects with a self-reported history of diabetes and/or treatment with dietary modification, use of anti-diabetic medication as indicated on the systemized questionnaire, or those with a fasting plasma glucose ≥ 126 mg/dL. Hypertension was defined as blood pressure ≥ 140/90 mmHg or a self-reported history of hypertension and/or use of anti-hypertensive medication. Hyperlipidemia was defined as total cholesterol ≥ 240 mg/dL or use of an anti-hyperlipidemic medication. A family history of CAD was defined as having a first-degree relative of any age with CAD based on the self-report questionnaire.

CCTA image acquisition and analysis. CCTA was conducted using either single-source 64-slice CT (LightSpeed VCT, GE, Milwaukee, WI, USA) or dual-source CT (Somatom Definition, Siemens, Erlangen, Germany). A standard scanning protocol was used, as previously described. All CCTA scans were analyzed using a dedicated workstation (Advantage Workstation, GE; or Volume Wizard, Siemens) by experienced cardiovascular radiologists (DHY, JWK, and THL). According to the guidelines of the Society of Cardiovascular Computed Tomography, a 16-segment coronary artery tree model was used. A coronary artery calcium score (CACS) was measured and categorized by scores of 0, 1 to 10, 11 to 100, 101 to 400, and > 400. Plaques containing calcified tissue involving more than 50% of the plaque area (density > 130 HU) were classified as calcified, plaques with < 50% calcium were classified as mixed, and plaques without calcium were classified as non-calcified lesions. The contrast-enhanced portion of the coronary lumen was semi-automatically traced at the site of maximal stenosis and compared with the mean value of the proximal and distal reference sites. Stenosis ≥ 50% was defined as significant. The overall plaque burden was determined from coronary artery plaque scores calculated from modified Duke prognostic scores, segment stenosis scores, and segment involvement scores, as described previously.

Clinical outcomes. Follow-up clinical data were obtained by a review of medical records or telephone interviews using trained personnel through to the end of June 2017. A cardiac event was defined as a composite of all-cause death, myocardial infarction, unstable angina requiring hospitalization, or coronary revascularization. The diagnosis of myocardial infarction was based on the presence of new Q waves in at least two contiguous leads, or an elevation of creatine kinase or its myocardial band isoenzyme to at least three times the upper limit.

Figure 1. Overview of the study population. CCTA coronary computed tomographic angiography; GGT gamma-glutamyl transferase; MI myocardial infarction; PCI percutaneous coronary intervention.
of the normal range at follow-up. Revascularization was performed if there was a stenosis of at least 50% of the diameter noted on invasive coronary angiography with a positive stress test result or if there was a stenosis of at least 70% observed on invasive coronary angiography\(^4\).

### Statistical analysis.

Categorical variables are expressed as frequencies with percentages. Continuous variables are expressed as the mean and standard deviation. Between-group comparisons were performed using Pearson’s chi-square test or Fisher's exact test for categorical variables and one-way analysis of variance or Kruskal–Wallis test for numerical variables, as appropriate. Univariable and multivariable analyses were performed using a logistic regression model to analyze the association between serum GGT levels and subclinical coronary atherosclerosis on CCTA. Based on previous epidemiologic studies\(^4,45\), we selected covariates in the multivariable model according to their clinical importance and statistical significance, which included age, sex, obesity, diabetes mellitus, hypertension, hyperlipidemia, creatinine, uric acid, current smoking, family history of CAD, and hs-CRP ≥ 2 mg/L. Unadjusted and adjusted odds ratios with 95% confidence intervals for the logistic regression were calculated. To investigate the associations of serum GGT levels and cardiac events, we also conducted the Cox proportional hazards regression analyses with adjustment for potential confounders. Stepwise multivariable models were determined by the backward variable selection approach. All reported \(p\) values are two-sided, and \(p < 0.05\) was considered statistically significant. Data manipulation and statistical analyses were performed using SPSS software (Version 18; SPSS Inc., Chicago, IL, USA).

### Data availability

The datasets used and/or analyzed during the current study available from the corresponding author on request.

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Y.G.K., S.B.L., and G.M.P. were responsible for the concept and design of the study, data acquisition, analysis and interpretation, and drafting of the manuscript. D.H.Y., J.W.K., and T.H.L. analyzed the coronary computed tomography angiography. H.K.K., J.C., S.W.L., and Y.H.K. critically revised the manuscript and provided important intellectual content.

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Competing interests

The authors declare no competing interests.

Additional information

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