



The Association between Social Network Betweenness and Coronary Calcium: A Baseline Study of Patients with a High Risk of Cardiovascular Disease

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Aim: The association of social networks with cardiovascular disease (CVD) has been demonstrated through various studies. This study aimed to examine the association between *social network betweenness* – a network position of mediating *between* diverse social groups – and coronary artery calcium.

Methods: The data of 1,384 participants from the Cardiovascular and Metabolic Disease Etiology Research Center – High Risk Cohort, a prospective cohort study enrolling patients with a high risk of developing CVD (clinicaltrials.gov: NCT02003781), were analyzed. The deficiency in social network betweenness was measured in two ways: *only-family networks*, in which a respondent had networks with only family members, and *no-cutpoint networks*, in which the respondent does not function as a point of bridging between two or more social groups that are not directly connected.

Results: Participants who had higher coronary artery calcium scores (CACs) were likely to have a smaller network size ($p < 0.001$), only-family networks ($p < 0.001$), and no-cutpoint networks ($p < 0.001$). Multiple logistic regression analyses revealed no significant association between network size and CACS. Only no-cutpoint networks had a significant relationship with CACS > 400 (odds ratio, 1.72; 95% confidence interval, 1.07–2.77; $p = 0.026$). The association was stronger among older (age > 60 years) and female respondents.

Conclusion: Deficiency in social network betweenness is closely related to coronary calcium in participants with a high risk of CVD. To generalize these results to a general population, further study should be performed.

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Key words: Coronary calcium score, Social networks, Cardiovascular disease, Risk factor

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Introduction

Social networks have been considered as an important social factor associated with cardiovascular disease (CVD) in terms of social contacts, social support, or social isolation¹. A recent meta-analysis showed that the function of social networks (content

provided by social networks, such as emotional or instrumental support) strongly affected the onset and development of CVD; however, there is little evidence on the effects of the structure of social networks (objective features of social networks, such as frequency of contacts or network size)²⁾. Despite the varied results concerning network structure, investigation of the detailed features of network structure has been consistently demanded because these determine the potential role and function of social networks³⁾.

Social network betweenness is a network structure in which a member can indirectly bridge *between* two or more social actors or groups that have no direct social connections to each other⁴⁾. For example, a person who maintains intimate relationships with the two social groups of neighbors and university alumni can better take advantage of social networks than those connected with only neighbors or only alumni, because this person can (i) draw more diverse resources and information from people of two different backgrounds, and (ii) feel a sense of freedom from crossing over various social groups⁵⁾. Whereas a few studies have examined the social determinants of coronary artery calcium, such as psychosocial stress⁶⁾ or marital status⁷⁾, there were no research that explicitly examined the relationship to structural features of social networks. By adopting an elaborate social network module that reveals thorough information about a person's important social network members, we investigated the association between social network structure and coronary artery calcium, with a focus on the limitations in social network betweenness.

Methods

Study Population

This study analyzed the data from Cardiovascular and Metabolic Disease Etiology Research Center–High Risk Cohort (CMERC-HI). Briefly, CMERC-HI is a prospective cohort study aimed at developing more specific preventive strategies for patients with a high risk of CVD (clinicaltrials.gov NCT02003781). The inclusion criteria of the cohort were as follows: high-risk patients with hypertension [estimated glomerular filtration rate (eGFR) >60 mL/min/1.73 m² with target organ damage, or eGFR ≤60 mL/min/1.73 m²]; patients with diabetes mellitus with albuminuria; anuric patients with end-stage renal disease (ESRD) who were undergoing dialysis; relatives of patients with acute myocardial infarction (men <55 years old; women <65 years old); patients with asymptomatic atherosclerotic CVD (abdominal aorta diameter ≥3 cm or ankle-brachial index <0.9, or carotid plaque or carotid intima-media thickness ≥0.9 mm, or asymp-

tomatic old cerebrovascular accident, or >30% stenosis in at least one major coronary artery); patients with rheumatoid arthritis aged >40 years and taking methotrexate and steroid; patients with atrial fibrillation with CHA2DS2-VASc score ≥1; and kidney transplant recipients at >3 months after transplantation. People aged >20 years who met at least one of the inclusion criteria were enrolled. The exclusion criteria were as follows: a history of acute coronary syndrome; symptomatic coronary artery disease or symptomatic peripheral; heart failure; with a life expectancy of <6 months; pregnant women; a history of contrast allergy and related adverse effects. A total of 1,958 participants were enrolled from November 2013 to May 2016 at the Severance Hospital in Seoul, Republic of Korea. The Institutional Review Board of the Yonsei University Health System Clinical Trial Center approved the study protocol (4-2013-0581), and written informed consent was obtained from all participants. Among the participants, 312 did not undergo a coronary artery calcium scanning, and 16 did not participate in the social network survey or provided incomplete information about their networks. After excluding 23 participants with missing values in socioeconomic status, 20 in health behaviors or medical history, and 213 in laboratory analyses (203 with missing lipid profile data), we examined 1,384 participants in the final analyses.

Social Network Betweenness

The CMERC adopted a social network module called *name generator* from a nationwide study in the United States⁸⁾. The Korean version of name generator has been used in nationwide social surveys in South Korea⁹⁾, and proved to capture important network properties for health in a comparative study of the USA and South Korea¹⁰⁾. The module started as follows: “From time to time, most people discuss things that are important to them with others. For example, these may include good or bad things that happen to you, problems you are having, or important concerns you may have. Looking back over the last 12 months, who are the people with whom you most often discussed things that were important to you?” The participants listed the real names of up to five people, and reported the type of relationship with each person (e.g., spouse, child, neighbor). The participants also reported the emotional closeness between network members within each pair (“How emotionally close are NAME1 and NAME2 to each other?”) on a five-level scale (*do not know each other, not close, somewhat close, very close, or very much close*).

Fig. 1 illustrates two possible networks, where circles represent participants, squares represent social

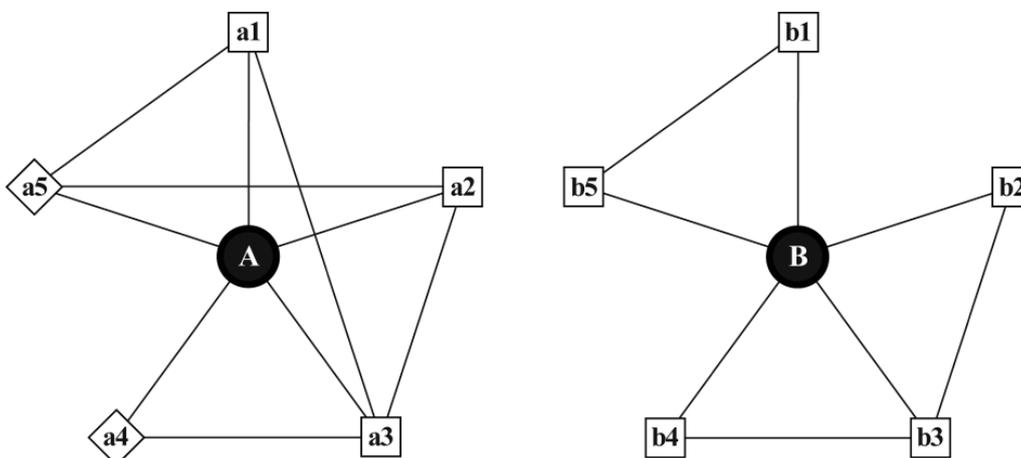


Fig. 1. Two possible social networks. Circles represent participants, squares represent social network members from a family (e.g., parent, child, relative), diamonds represent non-family network members (e.g., friend, neighbor), and lines show the social connections. Network size was defined as the total number of social network members. Only-family networks (OF) do not include any non-family member. Compared to A's networks, B's networks are limited in the family. A cutpoint is a person whose deletion breaks up the remaining group into two or more disconnected pieces. If a respondent did not function as a cutpoint of his or her networks, he or she was considered to have no-cutpoint (NC) networks. A's networks show an example of NC networks: A's network members are all connected to at least one other network member even when respondent A is missing, whereas B's networks are divided into the two groups of [b1, b5] and [b2, b3, b4] when respondent B is deleted.

network members from a family (e.g., parent, child, relative), diamonds represent non-family network members (e.g., friend, neighbor), and lines show the social connections. *Network size* was defined as the total number of social network members. A and B have five network members each; thus, their network size is 5. The deficiency in *social network betweenness* was measured by using two binary indicators. First, if a participant did not enumerate any non-family members in his or her network list, he or she was considered to have *only-family (OF) networks*. Compared with A's networks of three family members [a1, a2, a3] and two non-family members [a4, a5], B's networks are limited in the family. Participants who listed no social network member were assumed to have OF networks in this study. Second, if a respondent did not function as a cutpoint of his or her networks, he or she was considered to have *no-cutpoint (NC) networks*. In network theory, a *cutpoint* is a person whose deletion breaks up the remaining group into two or more disconnected pieces¹¹. A's networks show an example of NC networks: A's network members are all connected to at least one other network member even when respondent A is missing, whereas B's networks are divided into the two groups of [b1, b5] and [b2, b3, b4] when respondent B is deleted. Participants who listed no or only one social network member can-

not act as a cutpoint so were categorized into a NC network group. When counting the number of groups in networks, we only considered *very close* or *very much close* social connections among network members. Those who have OF or NC networks are expected to have little diversity in their social interactions, and little potential to draw resources from different social pools.

Acquisition of Computed Tomographic Images for Coronary Artery Calcium and Image Analysis

The patients were scanned by using a 320-detector row computed tomography (CT) scanner (Aquilion ONE; Toshiba Medical Systems Corporation, Tokyo, Japan). A non-enhanced prospective electrocardiogram-gated CT scan was performed to measure the coronary artery calcium score (CACS) with the following parameters: rotation time 275 ms, slice collimation 0.5 mm, slice thickness 3.0 mm, tube voltage 100 kV, and automatic tube current modulation (SURE Exposure 3D standard; Toshiba Medical Systems Corporation, Otawara, Japan). Images were analyzed in a core workstation by using dedicated software (TeraRecon V. 4.4.11.82.3430.Beta, Foster City, CA, USA). CACS was calculated by using the Agatston method^{12, 13}.

Statistical Analysis

The participants were divided into four groups according to CACS, as follows: (i) 0, (ii) 1–100, (iii) 101–400, and (iv) >400. Bivariate relationships were considered among CACS and various characteristics. Continuous variables were expressed as mean \pm SD, and compared by using one-way analysis of variance (ANOVA). Categorical variables were expressed as number and percentage, and compared with the chi-square test. In multiple logistic regression analyses, the odds of CACS >400 were computed according to network size and deficiency in network betweenness with covariates of age, sex, smoking, history of hypertension, history of diabetes, ESRD, exercise assessed by using the International Physical Activity Questionnaire¹⁴), depression assessed with Beck Depression Inventory¹⁵), statin use, body mass index (BMI), systolic blood pressure (SBP), low-density lipoprotein (LDL) cholesterol, triglyceride, fasting glucose, eGFR, level of education, household income, working status, marital status, and number of co-habitants. Five models were examined with different sets of social network variables: model 1 with network size; model 2 with network size and OF networks; model 3 with network size and NC networks; model 4 with network size, OF, and NC networks; and model 5 with three dummy variables for those who had OF networks alone, NC networks alone, and both OF and NC networks. Based on the results from model 5, we also did the Wald test ($H_0: \beta_{OF \text{ alone}} + \beta_{NC \text{ alone}} = \beta_{OF+NC}$) to examine if there was a multiplicative association of OF and NC networks. If the sum of coefficients for OF alone and NC alone is significantly larger than that for both OF and NC, the association can be considered multiplicative. Using subsamples of age ≤ 60 years, age > 60 years, male, or female, we re-examined models 2 and 3 and reported the ORs for OF networks and NC networks as figures. Additionally, we considered several models to address the robustness of the results. First, as problems in the coronary artery could reversely affect social networks, we considered only long-lasting social networks and re-examined the associations between CACS and the deficiency in social network betweenness in model As and Bs. Based on the information about how many years the participants had known their social network members, we eliminated recently-formed social networks and re-calculated network size and betweenness using only long-lasting network members based on two different thresholds (≥ 5 years or ≥ 10 years), and tested if the associations of network betweenness were sustained. By removing those young networks, we aimed to rule out the associations due to the network changes caused by the present health condition from our prediction. Second, for a more accurate

assessment of causal effects, we adopted the coarsened exact matching (CEM) method in Model Cs¹⁶). Unlike other matching methods based on propensity scores, CEM matches observations with those with “exactly” the same values of covariates, and performs regression analyses using a subsample of matched observations where the treated and untreated groups are weighted to have the same sample size. Since it is hard to find observations with exactly the same covariates in a given sample, CEM “coarsens” covariates to a small number of categories before matching. Under the assumption that this set of variables determine the present social networks, we can consider the associations between CACS and networks from CEM as causal. As it was hard to retain enough sample size for statistical inference when treating with multivalued treatment, we did not test a CEM model with three dummy variables for network betweenness. In this analysis, continuous variables were coarsened into three groups, and categorical variables were adopted as original categories. As covariates for matching, we examined those that had significant bivariate relationships with network betweenness (age, diabetes, depression, SBP, LDL-cholesterol, glucose, eGFR, education, income, working, and marriage). When doing regression analyses after finding a subsample of matched observations, we used the same set of covariates as other multiple logistic regression analyses except that the categorical variables that were not coarsened during matching were excluded. Third, we restricted our sample to those with social network size > 1 and performed the same logistic regression analyses in Model Ds. Since social network betweenness can arise from at least two social network members, some sociologists emphasized a qualitative difference between those who have so small networks (≤ 1) to cultivate network betweenness and the others with big networks¹⁷). By restricting our sample to those with social network size > 1 , we could focus on the disadvantage of deficiency in network betweenness among participants whose networks were big enough to have network betweenness. The analyses were performed by using STATA 14 (StataCorp LP, College Station, TX, USA).

Results

Table 1 shows the clinical characteristics of participants according to CACS. Participants who were included in a higher CACS group were more likely to be older, male, currently smoking, having diabetes, a sedentary lifestyle, ESRD or a history of receiving kidney transplantation, higher SBP, lower DBP, and higher level of fasting plasma glucose. These participants had lower serum cholesterol, possibly because more statin

Table 1. Characteristics of participants by coronary artery calcium score ($N=1384$)

Variable	Category	Coronary calcium score (AU)				<i>P</i>
		0 ($n=545$)	1-100 ($n=364$)	101-400 ($n=266$)	>400 ($n=209$)	
Age, year		54.9 ± 412.0	61.1 ± 10	63.9 ± 10.2	63.3 ± 10.6	<0.001
Sex, n (%)	Male	259 (48%)	212 (58%)	144 (54%)	145 (69%)	<0.001
	Female	286 (52%)	152 (42%)	122 (46%)	64 (31%)	
Current smoker, n (%)		213 (39%)	172 (47%)	130 (49%)	119 (57%)	<0.001
Hypertension, n (%)		456 (84%)	307 (84%)	229 (86%)	184 (88%)	0.451
Diabetes, n (%)		153 (28%)	150 (41%)	139 (52%)	119 (57%)	<0.001
End-stage renal disease, n (%)		84 (15%)	47 (13%)	34 (13%)	58 (28%)	<0.001
Kidney transplantation, n (%)		36 (7%)	19 (5%)	19 (7%)	30 (14%)	0.001
Exercise, MET minute/week		1928.6 ± 2493.9	2224.2 ± 3211.8	1887.0 ± 2747.6	1535.5 ± 2155.1	0.032
Depression, n (%)		34 (6%)	27 (7%)	24 (9%)	24 (11%)	0.096
Statin use, n (%)		235 (43%)	188 (52%)	154 (58%)	119 (57%)	<0.001
Body mass index, kg/m ²		25.0 ± 3.8	25.5 ± 3.6	25.3 ± 3.4	24.8 ± 3.9	0.079
Systolic blood pressure, mmHg		123.7 ± 14.8	129.1 ± 16.4	130.7 ± 18.1	132.2 ± 21.8	<0.001
Diastolic blood pressure, mmHg		76.8 ± 10.3	77.2 ± 10.4	74.3 ± 9.5	73.9 ± 10.1	<0.001
Total cholesterol, mg/dL		179.7 ± 35.3	172.9 ± 35.7	166.5 ± 35.9	162.7 ± 34.5	<0.001
HDL-cholesterol, mg/dL		50.9 ± 13.2	48.0 ± 12.5	48.5 ± 22	45.6 ± 11.9	<0.001
LDL-cholesterol, mg/dL		101.0 ± 30.5	95.1 ± 28.8	90.1 ± 27.6	86.6 ± 27.9	<0.001
Triglyceride, mg/dL		133.7 ± 71.5	140.8 ± 76.9	144.7 ± 104.7	142.3 ± 83.5	0.251
Fasting glucose, mg/dL		107.2 ± 28.5	112.6 ± 29.6	116.5 ± 31	120.7 ± 41.4	<0.001
Estimated GFR, ml/min/1.73 m ²		68.4 ± 36.1	66.2 ± 33.6	64.2 ± 30.9	53.0 ± 32.5	<0.001
Education, n (%)	≤ Middle school	132 (24%)	111 (30%)	94 (35%)	71 (34%)	0.013
	High school	192 (35%)	110 (30%)	82 (31%)	71 (34%)	
	≥ College	221 (41%)	143 (39%)	90 (34%)	67 (32%)	
Household income, n (%)	< US\$3000	64 (12%)	50 (14%)	52 (20%)	37 (18%)	0.013
	US\$3001-6000	140 (26%)	75 (21%)	66 (25%)	39 (19%)	
	≥ US\$6000	122 (22%)	94 (26%)	46 (17%)	40 (19%)	
	DK or refusal	219 (40%)	145 (40%)	102 (38%)	93 (44%)	
Current worker, n (%)		298 (55%)	180 (49%)	104 (39%)	93 (44%)	<0.001
Married, n (%)		432 (79%)	304 (84%)	220 (83%)	155 (74%)	0.034
Number of household members		1.3 ± 1.3	1.2 ± 1.2	1.0 ± 1.2	1.0 ± 1.2	<0.001
Network size		2.2 ± 1.4	2.0 ± 1.4	1.9 ± 1.4	1.7 ± 1.3	<0.001
Only-family networks, n (%)		283 (52%)	214 (59%)	180 (68%)	154 (74%)	<0.001
No-cutpoint networks, n (%)		332 (61%)	242 (66%)	198 (74%)	173 (83%)	<0.001

Data are presented as mean (SD) for continuous variables or n (%) for categorical variables. The *p*-values are from ANOVA tests for continuous variables or chi-square tests for categorical variables.

AU, Agatston unit; MET, metabolic equivalent of task; DK, do not know.

Table 2. Characteristics of participants by social network betweenness (N=1384)

Variable	Category	Only-family networks			No-cutpoint networks		
		No (n=553)	Yes (n=831)	P	No (n=439)	Yes (n=945)	P
Coronary calcium score (AU)		176.3 ± 527.0	270.6 ± 565.0	0.002	146.3 ± 485.7	273.2 ± 575.9	<0.001
Age, year		55.5 ± 11.8	62.2 ± 10.7	<0.001	55.1 ± 11.4	61.6 ± 11.1	<0.001
Sex, n (%)	Male	299 (54%)	461 (55%)	0.606	231 (53%)	529 (56%)	0.242
	Female	254 (46%)	370 (45%)		208 (47%)	416 (44%)	
Current smoker, n (%)		254 (46%)	380 (46%)	0.941	190 (43%)	444 (47%)	0.198
Hypertension, n (%)		459 (83%)	717 (86%)	0.094	368 (84%)	808 (86%)	0.417
Diabetes, n (%)		166 (30%)	395 (48%)	<0.001	139 (32%)	422 (45%)	<0.001
End-stage renal disease, n (%)		89 (16%)	134 (16%)	0.988	69 (16%)	154 (16%)	0.785
Kidney transplantation, n (%)		43 (8%)	61 (7%)	0.764	34 (8%)	70 (7%)	0.825
Exercise, MET minute/week		2067.7 ± 2620.1	1853.3 ± 2766.9	0.150	1887.3 ± 2451.7	1963.0 ± 2823.3	0.629
Depression, n (%)		32 (6%)	77 (9%)	0.019	24 (5%)	85 (9%)	0.023
Statin use, n (%)		285 (52%)	411 (49%)	0.449	220 (50%)	476 (50%)	0.929
Body mass index, kg/m ²		25.2 ± 3.7	25.1 ± 3.7	0.498	25.1 ± 3.7	25.2 ± 3.7	0.869
Systolic blood pressure, mmHg		125.5 ± 15.6	129.2 ± 18.4	<0.001	125.0 ± 15.3	129.0 ± 18.2	<0.001
Diastolic blood pressure, mmHg		77.1 ± 10.3	75.2 ± 10.1	0.001	76.7 ± 10.4	75.6 ± 10.2	0.059
Total cholesterol, mg/dL		177.9 ± 35.9	169.4 ± 35.6	<0.001	179.6 ± 36.9	169.7 ± 35.1	<0.001
HDL-cholesterol, mg/dL		50.7 ± 17.7	47.7 ± 12.9	<0.001	51.0 ± 18.6	47.9 ± 12.9	<0.001
LDL-cholesterol, mg/dL		98.1 ± 29.6	93.2 ± 29.5	0.003	99.1 ± 29.9	93.3 ± 29.3	0.001
Triglyceride, mg/dL		140.6 ± 88	137.9 ± 77.9	0.549	140.3 ± 86.6	138.4 ± 79.9	0.693
Fasting glucose, mg/dL		107.5 ± 24.7	115.7 ± 35.5	<0.001	106.7 ± 23	115.1 ± 34.9	<0.001
Estimated GFR, ml/min/1.73 m ²		68.0 ± 34.9	62.5 ± 33.8	0.003	67.1 ± 34.4	63.5 ± 34.3	0.069
Education, n (%)	≤ Middle school	107 (19%)	301 (36%)	<0.001	85 (19%)	323 (34%)	<0.001
	High school	174 (31%)	281 (34%)		128 (29%)	327 (35%)	
	≥ College	272 (49%)	249 (30%)		226 (51%)	295 (31%)	
Household income, n (%)	< US\$3000	62 (11%)	141 (17%)	<0.001	50 (11%)	153 (16%)	<0.001
	US\$3001-6000	148 (27%)	172 (21%)		125 (28%)	195 (21%)	
	≥ US\$6000	165 (30%)	137 (16%)		142 (32%)	160 (17%)	
	DK or refusal	178 (32%)	381 (46%)		122 (28%)	437 (46%)	
Current worker, n (%)		311 (56%)	364 (44%)	<0.001	246 (56%)	429 (45%)	<0.001
Married, n (%)		427 (77%)	684 (82%)	0.020	369 (84%)	742 (79%)	0.016
Number of household members		1.2 ± 1.1	1.2 ± 1.3	0.569	1.3 ± 1.1	1.1 ± 1.3	0.054
Network size		3.0 ± 1.2	1.3 ± 1.1	<0.001	3.2 ± 1.1	1.5 ± 1.2	<0.001

Data are presented as mean ± SD for continuous variables or n (%) for categorical variables. The p-values are from ANOVA tests for continuous variables or chi-square tests for categorical variables.

AU, Agatston unit; MET, metabolic equivalent of task; DK, do not know.

Table 3. Multiple logistic regression analyses of coronary artery calcium score >400 ($N=1384$)

Variable	Model 1		Model 2		Model 3		Model 4		Model 5	
	OR (95% CI)	<i>P</i>								
Network size	0.95 (0.83-1.08)	0.412	1.03 (0.88-1.20)	0.730	1.03 (0.89-1.19)	0.690	1.05 (0.90-1.24)	0.510	1.06 (0.90-1.24)	0.509
No deficiency + NC alone ($n=553$)			1.00 (reference)	–			1.00 (reference)	–		
OF ($n=831$)			1.50 (0.95-2.35)	0.080			1.21 (0.72-2.03)	0.467		
No deficiency + OF alone ($n=439$)					1.00 (reference)	–	1.00 (reference)	–		
NC ($n=945$)					1.72 (1.07-2.77)	0.026	1.56 (0.90-2.69)	0.114		
No deficiency ($n=383$)									1.00 (reference)	–
OF alone ($n=56$)									1.19 (0.39-3.68)	0.755
NC alone ($n=170$)									1.55 (0.83-2.88)	0.166
OF + NC ($n=775$)									1.88 (1.10-3.23)	0.022

All models were controlled for age, sex, smoking, hypertension, diabetes, ESRD, exercise, depression, statin use, BMI, SBP, LDL-cholesterol, glucose, eGFR, education, income, working, marriage, and number of co-habitants.

Model 1 evaluated the association of network size. Model 2 included network size and OF. No deficiency + NC alone was reference of OF. Model 3 included network size and NC. No deficiency + OF alone was a reference to NC. Model 4 included network size, OF, and NC. No deficiency + NC alone was a reference to OF. No deficiency + OF alone was a reference to NC. Model 5 used three dummy variables for social network betweenness (OF alone, NC alone, and OF + NC) comparing with “no deficiency” as reference.

OF, only-family networks; NC, no-cutpoint networks

users were included in the higher CACS group. Concerning socio-economic status, participants with higher CACS were less educated, not married, and living with fewer household members; however, household income and working status were significantly different across CACS categories with no consistent patterns. Concerning social network characteristics, participants with higher CACS had significantly smaller ($p<0.001$) OF ($p<0.001$) and NC ($p<0.001$) networks.

Table 2 shows the association between deficiency of social network betweenness and other covariates. Participants with OF or NC networks were more likely to be older and have diabetes, depression, higher SBP, higher total cholesterol, lower HDL-cholesterol, lower LDL-cholesterol, and higher fasting plasma glucose levels. eGFR was positively associated with OF networks, whereas current smokers were more likely to have NC networks. Concerning socio-economic status, deficiency in both types of social network betweenness was related to a lower level of education, lower household income, and not working status. Married participants were

more likely to have OF networks but less likely to have NC networks. Those who had smaller networks were more likely to have both OF and NC networks.

Table 3 shows the results of multiple logistic regression analyses for identifying whether each social network component was a determinant for CACS >400. The association between CACS and network size (odds ratio [OR], 0.95; 95% confidence interval [CI], 0.83–1.08; $p=0.412$) or OF networks (OR, 1.50; 95% CI, 0.95–2.35; $p=0.080$) was not statistically significant at the 0.05 level; however, NC networks had significant association with CACS >400 when separately considered in model 3 (OR, 1.72; 95% CI, 1.07–2.77; $p=0.026$). When including two types of deficiency in network betweenness together in model 4, both had p -values >0.05. When examining three dummy variables (OF networks alone, NC networks alone, and both OF and NC networks) in model 5, participants who had both OF and NC networks had significantly greater odds of CACS >400 than those with no deficiency in network betweenness (OR,

Table 4. Robustness checks for the multiple logistic regression analyses of coronary artery calcium score >400

	Networks known for ≥5 years			Networks known for ≥10 years			Coarsened-exact-matching analysis			Network size > 1		
	n	OR (95% CI)	P	n	OR (95% CI)	P	n	OR (95% CI)	P	n	OR (95% CI)	P
	Model A-1 (n=1,384)			Model B-1 (n=1,384)			Model C-1 (n=933)			Model D-1 (n=694)		
No deficiency + NC alone	553	1.00 (reference)	-	553	1.00 (reference)	-	393	1.00 (reference)	-	553	1.00 (reference)	-
OF	831	1.65 (1.05-2.60)	0.030	831	1.67 (1.06-2.64)	0.028	540	1.60 (0.96-2.66)	0.070	831	1.41 (0.80-2.48)	0.239
	Model A-2 (n=1,384)			Model B-2 (n=1,384)			Model C-2 (n=920)			Model D-2 (n=694)		
No deficiency + OF alone	439	1.00 (reference)	-	439	1.00 (reference)	-	332	1.00 (reference)	-	439	1.00 (reference)	-
NC	945	1.98 (1.22-3.24)	0.006	945	1.98 (1.20-3.26)	0.007	588	2.78 (1.53-5.05)	0.001	945	1.74 (1.02-2.99)	0.044
	Model A-3 (n=1,384)			Model B-3 (n=1,384)						Model D-3 (n=694)		
No deficiency	383	1.00 (reference)	-	383	1.00 (reference)	-				383	1.00 (reference)	-
OF alone	56	1.22 (0.39-3.80)	0.728	56	1.23 (0.39-3.85)	0.724				56	1.64 (0.49-5.50)	0.427
NC alone	170	1.73 (0.92-3.25)	0.090	170	1.69 (0.89-3.22)	0.110				170	1.93 (0.97-3.83)	0.062
OF + NC	775	2.24 (1.28-3.90)	0.004	775	2.26 (1.28-3.98)	0.005				775	1.77 (0.92-3.38)	0.086

All models were controlled for age, sex, smoking, hypertension, diabetes, ESRD, exercise, depression, statin use, BMI, SBP, LDL-cholesterol, glucose, eGFR, education, income, working, marriage, and number of co-habitants.

Model As used 5 years and model Bs used 10 years as thresholds of network duration, respectively. In Model Cs using the coarsened-exact-matching method, observations were matched based on age, diabetes, depression, SBP, LDL-cholesterol, glucose, eGFR, education, income, working, and marriage. Continuous variables were coarsened into three groups, and categorical variables were adopted as original categories. Model Ds were logistic regressions which used subsamples with social network size > 1.

OF, only-family networks; NC, no-cutpoint networks

1.88; 95% CI, 1.10–3.23; $p=0.022$). Even though we did not find any evidence for the multiplicative association between OF and NC networks from the Wald test ($\chi^2=0.00$, $p=0.98$), we could observe that the ORs for those with both OF and NC networks was the largest among three types of deficiency in network betweenness.

The results of robustness checks are shown in **Table 4**. First, when considering social networks known for ≥5 years in model A-1 to A-3, OF networks (OR, 1.65; 95% CI, 1.05–2.60; $p=0.030$), NC networks (OR, 1.98; 95% CI, 1.22–3.24; $p=0.006$), and both OF and NC networks (OR, 2.24; 95% CI, 1.28–3.90; $p=0.004$) showed significant associations with CACS >400. When adopting the threshold of ≥10 years in model B-1 to B-3, we also observed signifi-

cant associations of OF networks (OR, 1.67; 95% CI, 1.06–2.64; $p=0.028$), NC networks (OR, 1.98; 95% CI, 1.20–3.26; $p=0.007$), and both OF and NC networks (OR, 2.26; 95% CI, 1.28–3.98; $p=0.005$). Second, according to the CEM analyses in model C-1 to C-2, the p -value for OF networks reached >0.05 (OR, 1.60; 95% CI, 0.96–2.66; $p=0.070$), whereas NC networks (OR, 2.78; 95% CI, 1.53–5.05; $p=0.001$) were significantly associated with CACS >400. Lastly, in model D-1 to D-3 in which those with no or only one network member were excluded, OF networks (OR, 1.41; 95% CI, 0.80–2.48; $p=0.239$) and both OF and NC networks (OR, 1.77; 95% CI, 0.92–3.38; $p=0.086$) did not show significant associations with CACS >400, whereas NC networks (OR, 1.74; 95% CI, 1.02–2.99; $p=0.044$) were significantly associated

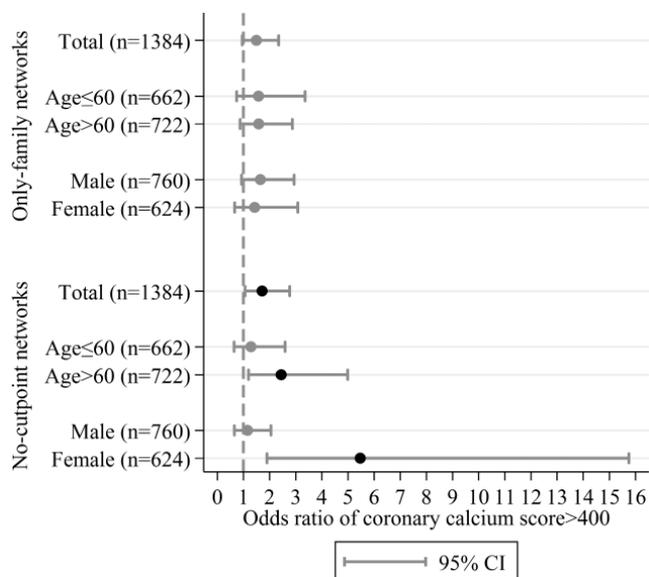


Fig. 2. ORs and 95% CIs for only-family networks and no-cutpoint network from multiple logistic regression analyses of coronary calcium score >400 in total participants and subgroups. We predicted ORs for only-family networks or no-cutpoint networks after controlling for age, sex, smoking, hypertension, diabetes, ESRD, exercise, depression, statin use, BMI, SBP, LDL-cholesterol, glucose, eGFR, education, income, working, marriage, number of co-habitants, and network size.

with CACS >400 in model D-2.

To examine the difference by age and sex, we reassessed models 2 and 3 in **Table 3** by using subsamples of age ≤60 years, age >60 years, male, or female. As seen in **Fig. 2**, there was no difference in the associations of OF networks by age and sex. However, NC networks were more disadvantageous for older (OR, 2.64; 95% CI, 1.27–5.51; $p=0.009$) and female (OR, 6.34; 95% CI, 2.02–19.96; $p=0.002$) participants.

Discussion

The key findings from this study were as follows: (i) social network betweenness was closely related with health and socio-economic status; (ii) no-cutpoint networks, in which network members form a connected group even when excluding a respondent, were significantly associated with a higher risk of CACS >400; and (iii) the disadvantage of no-cutpoint networks was larger for older and female participants. While previous studies have investigated whether biomarkers such as pulse wave velocity or inflammatory markers were associated with the presence of subclinical atherosclerosis^{18, 19}, the novelty of this study is that sociological factors may be closely related with subclinical atherosclerosis, characterized by coronary artery calcification.

In previous studies, the disadvantage of poor social networks was considered to parallel the deficiency in social support, which could directly cause the onset and progression of CVD²⁰, or have indirect effects through buffering the negative influence of acute stress²¹ and depressive symptoms²². In this study, more detailed properties of social networks were investigated by adopting the network structure of betweenness. First, network betweenness captures the accessibility to diverse origins of social resources, which is a different aspect from the content or size of networks. For example, diverse networks more effectively provide medical information and first-hand experience about certain diseases, and are known to have special importance for those with health problems²³. Considering that the sample of this study was at a high risk of CVD, the strong disadvantage for those without network betweenness is not surprising. Second, various sources represent the diversity in the activities they perform with their network members. Network betweenness allows the potential to enjoy at least two different activities and thoughts, which affords a stronger sense of freedom and accomplishment than being confined to a narrow social space. Those benefits are reported to have more importance for the subjective well-being of the older population⁵, as retirement status and little participation in organizational activities make them more dependent on selective interpersonal relationships²⁴. The strong correlation between network betweenness and depression, and the stronger associations of betweenness with the CACS older adults observed in this study are consistent with previous findings.

Among the two types of deficiency in social network, between and no-cutpoint networks had stronger and more robust associations with CACS than only-family networks. Even though these two had no significant associations with CACS at a level of 0.05 when included together in the same model due to a strong correlation between themselves ($\chi^2=599.24$, $p<0.001$), no-cutpoint networks were significantly associated with CACS in all models for robustness checks, whereas only-family networks showed little evidence of associations in the models using the CEM method and the subsample with two or more social networks. Considering that family members are the first sources of immediate support for people with health problems²⁵, the associations of only-family networks could largely originate from the previous health status, which was partly proved in the results from the matching analyses. Furthermore, the results from the subsample with network size >1 implicate that the disadvantage of only-family networks is not critical among those who have big networks, but mixed with the fundamental restriction of small network size. The

robust results from the analyses of no-cutpoint networks, on the other hand, show that they could be more strong indicators for CACS than only-family networks. Additionally, no-cutpoint networks were more accurate in detecting the bridging potential of participants, as these are based on direct information about the social network structure. Despite its importance, there are few studies on the relationship between network betweenness and CVD because of the shortage of data about the details of network structure³⁾. While a recent study showed that older women are more likely to establish their social autonomy via social network betweenness²⁶⁾, we observed a similar pattern that no-cutpoint networks were more unfavorable for the CACS of older and female participants. Cumulative findings from future studies on network structure and CVD may provide a clearer explanation about the sex difference in the associations of network betweenness.

This study has several limitations. First, owing to the cross-sectional design, we do not know the causality of social network betweenness with the progression of coronary artery calcium and/or the development of cardiovascular disease. For addressing this issue, we recalculated network indices using only long-lasting social networks and observed robust associations between CACS and both types of deficiency in social network betweenness. If we assume that the formation of long-lasting social networks precedes the progression of coronary artery calcification, the results possibly implicate the causal effects of network betweenness. In addition, we also found the significant associations of no-cutpoint networks from the CEM analyses. However, these results cannot fully support our causal inference since we do not have the accurate information about the temporal order among variables, and examined the outcome and covariates at the same time point. As CMERC-HI is a prospective cohort study, the prognostic implications of social network betweenness in this cohort can be clarified in the future. Second, as the study population comprised subjects with a relatively high risk of developing CVD, the results of this study are vulnerable to generalizability. Third, the results showed that age and socio-economic status were different according to CACS. Considering that age and income had strong associations with both social network betweenness and cardiovascular outcome, we could expect that our estimates could be biased from residual confounding. Even though we controlled for age and income in all multiple regression models, 1) there could be some omitted information such as medical access which might be limited to those with younger age and higher income, and 2) our indicators for income could be crude to sufficiently

capture variations in economic status. Future studies are required to collect more thorough data of economic and medical backgrounds for unbiased estimates of network effects on CVD. Fourth, this study assumed that CACS >400 is a surrogate marker of CVD. Although higher CACS is closely correlated with having more other cardiovascular risk factors and CACS >400 is known as a good predictor of future CVD in the western population, clinical implication of CACS >400 needs to be further validated in the Asian population²⁷⁾.

Conclusion

Participants who had higher coronary artery calcium score were likely to have a smaller network size and deficiencies in social network betweenness (only-family network, and no-cutpoint network). Especially, no-cutpoint networks had a significant and robust relationship with CACS >400, a powerful surrogate marker of future CVD.

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Conflict of Interest

The authors have no conflict of interest to disclose.

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