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# **BMJ Open**

## The association of serum high sensitivity C-reactive protein with the mortality risk in Asian: the Health Examinees cohort

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3 4	196	The association of serum high sensitivity C-reactive protein with the mortality risk in
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6 7	197	Asian: the Health Examinees cohort
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11	200	Sang-Ah Lee <sup>1,2 *</sup> , Sung Ok Kwon <sup>1</sup> , Hyerim Park <sup>1</sup> , Xiao-Ou Shu <sup>2</sup> , Jong-Koo Lee <sup>3</sup> , Daehee Kang <sup>4</sup>
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23 24	209	
25	210	
26 27	211	ABSTRACT
28	212	<b>Objectives</b> This study aimed to examine the association of <i>hs</i> CRP with mortality risk and the attenuated effect
29 30	213	of non-communicable disease history (NCD <sub>history</sub> ) on the association.
31	214	Design Prospective cohort study.
32	215	Setting the Health Examinees (HEXA) cohort.
33 34	216	<b>Participants</b> A total of 41 070 men and 81 011 women aged $\geq$ 40 years were involved (follow-up: 6.8 years).
35	217	<b>Outcome measures</b> The data and cause of death occurring until December 31, 2015, were confirmed by death
36 37	218	statistics from the National Statistical Office. We conducted the advanced analysis after stratification by
38	219	NCD <sub>history</sub> and the sensitivity analysis after excluding death before 1 or 2 years from recruitment. Cox
39 40	220	proportional hazard and restricted cubic spline models were used to assess the association.
41	221	<b>Results</b> The association between serum <i>hs</i> CRP and the risk of all-cause mortality was observed with strong
42 43	222	linearity in both genders, which was not influenced by NCD <sub>history</sub> . Otherwise, the association of serum hsCRP
43 44	223	with cancer-mortality risk was not observed in women with NCD <sub>history</sub> , but the association with the risk of
45	224	cardiovascular disease (CVD) mortality was predominantly observed in men with NCD <sub>history</sub> .
46 47	225	<b>Conclusions</b> This study suggested the dose-response association of <i>hs</i> CRP with mortality risk, including
48	226	cancer and CVD mortality, in Korean with low serum hsCRP, although the association with cancer and CVD-
49 50	227	mortality risk could be influenced by gender and NCD <sub>history</sub> .
51	228	
52	229	
53 54	230	Strengths and limitations of this study
55 56	231	• This is the large population-based prospective study.
57 58 59	232	• We examined the effect of very high <i>hs</i> CRP concentration on mortality risk.
60	233	• The <i>hs</i> CRP level of present study was measured within 18 hours in a single institution to minimize error/bias.

1 2		2
3 4	234	• Due to due to random fluctuations of <i>hs</i> CRP, using the single measurement of <i>hs</i> CRP at baseline could reflect
5 6	235	the inaccurate status of blood <i>hs</i> CRP levels in the study participants and increase the instability of <i>hs</i> CRP.
7 8	236	This study lacked information on medication use at recruitment and during the follow-up period, and
9		
10 11	237	information on hormone-replacement therapy (HRT) among women.
12	238 239	
13 14	239	*Correspondence to: Sang-Ah Lee, Ph.D.
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## 266 INTRODUCTION

High sensitivity C-reactive protein (hsCRP) is an acute-phase response protein synthesized by the liver and the most sensitive and dynamic marker of inflammation[1]. Since hsCRP has been reported as a candidate marker for generalized atherosclerosis and cardiovascular disease (CVD)[2], many studies[3-7] have investigated the role of hsCRP levels as a predictor of mortality risk. A recent meta-analysis[8] reported the predictable role of serum hsCRP on all-cause and CVD mortality in the general population. Nevertheless, it is controversial whether the predictable role of hsCRP could be applied to the risk of mortality in Asians, whose hsCRP levels are lower than those in individuals in Western countries.

Serum *hs*CRP represents a low-grade inflammation state that is generally involved in the process of aging[9]. Several large cohorts, including Study of Women's Health Across the Nation (SWAN)[10], the Women's Health Study[11] and the Dallas Heart Study[12], reported significant differences in hsCRP levels by race and gender. In two studies of multiethnic populations residing in the USA[10, 13], the median hsCRP level in East Asians was less than half the concentration in Caucasians. Even among East Asian populations, the geometric mean of hsCRP levels varied depending on ethnic background[14]. In addition, a meta-analysis[11] reported the hsCRP levels among women of various ethnic groups living in the United States (from the Women's Health Study) on the association between hsCRP and the mortality risk; the association was observed in only men supported by the results from two cohort studies [15, 16] reported in Korea. On the other hand, the increased hsCRP may be influenced by comorbidity itself because inflammation has emerged as an important factor in the progression of non-communicable diseases (NCDs), including CVD[17], cancer[18], chronic obstructive pulmonary disease (COPD)[19], type 2 diabetes[20] and fractures[21], which contribute to increased morbidity and mortality. This study aimed to examine the association of serum hsCRP with the risk of mortality in Koreans with low

serum *hs*CRP and to evaluate the attenuated effect of non-communicable disease history (NCD<sub>*history*</sub>) on the
 association.

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#### **METHODS**

#### Study population

Details on the main objectives, rationale, study design and baseline characteristics of the Health Examinees (HEXA) study have been published elsewhere[22]. Considering the homogeneity and comparability of participants, we created a qualified dataset called HEXA-G (Health Examinees-Gem) from previously published HEXA studies[23]. In the new HEXA-G data, a total of 141 968 participants remained after the exclusion of withdrawers (n=12). In addition, 19 887 were excluded due to missing information (n=19 876) or small sample size (n=11) on any hsCRP components at the baseline survey. Ultimately, 122 081 subjects, including 41 070 men and 81 011 women, remained in the final analysis (Fig. 1). All study participants provided informed consent prior to entering the study. The Institutional Review Board of the Seoul National University Hospital, Seoul, Korea, approved it for statistical analysis (IRB No. E-1503-103-657).

#### Laboratory measurements

After at least 10 hours of overnight fasting, blood samples were obtained in the morning. Bio-specimens included fasting blood samples that were collected in a serum separator tube and two ethylenediaminetetraacetic acid (EDTA) tubes. All samples were then transported to the National Biobank of Korea and stored for future research purposes within 18 hours. hsCRP was measured using a turbidimetric immunoassay (ADVIA 1650 and ADVIA 1800; Siemens Healthineers).

#### Follow-up and ascertainment of mortality

All-cause mortality was confirmed by death statistics from the National Statistical Office, which provided the data and causes of all deaths occurring through December 31, 2015. We added the mortality data from Statistics Korea to our dataset using each participant's unique identifier. Information on death and causes of death was obtained from a record link with the national death certificate files in Korea. The main outcome of interest was all-cause mortality (defined as death from any cause), including cancers and CVD mortality. The cause of death was classified according to the International Classification of Diseases, 10th revision (ICD-10). Deaths were coded as C00-C97 for cancer and I00-I99 for CVD. 

#### 323 Baseline variables

Trained interviewers collected information on demographic, socioeconomic and lifestyle factors. Anthropometric measurements were obtained using standardized methods. Body mass index (BMI) was calculated, and all participants were defined into four classes based on the World Health Organization classification of BMI for Asian adults[24]: underweight (BMI <18.5 kg/m<sup>2</sup>), normal (18.5 BMI <23.0 kg/m<sup>2</sup>), overweight (23.0 $\leq$  BMI <25.0 kg/m<sup>2</sup>), obesity (25.0 $\leq$  BMI <29.9 kg/m<sup>2</sup>), and severe obesity (BMI  $\geq$ 30.0 kg/m<sup>2</sup>). The current study defined metabolic syndrome using the National Cholesterol Education Program Adult Treatment Panel III (NCEP-ATP III)[25], modified for the Asian guideline for waist circumference (WC ≥90 and  $\geq$ 80 cm for men and women, respectively). Nonsmokers were defined as those who had smoked less than 400 cigarettes over the course of their lifetime. Participants who had smoked were categorized into two groups: noncurrent (never/former) and current smoker. Noncurrent drinkers were defined as those who had never consumed an alcoholic drink over the course of their lifetime or those who had not consumed alcohol at recruitment, while current drinkers were defined as those who persisted in consuming alcohol. Regular exercise was classified into two groups (ves/no) as follows: "Do you currently engage in regular exercise strenuous enough to cause you to break into a sweat at least once per week?" Furthermore, considering the attenuated effect of the NCD<sub>history</sub> on the association between serum hsCRP and the risk of mortality, we performed advanced analysis after stratification by NCD<sub>history</sub>. We considered six main non-communicable diseases (hypertension, diabetes, hyperlipidemia, cancer, cardiovascular and cerebrovascular diseases, and respiratory disease) to classify healthy subjects vs. subjects with NCD<sub>history</sub>.

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## 343 Statistical analysis

For the categorical analysis, we created nine categories based on the distribution of hsCRP levels in our population: <1.00 (reference group), 1.01-1.50, 1.51-2.00, 2.01-2.50, 2.51-3.00, 3.01-4.00, 4.01-6.00, 6.01-10.0, and >10.0 mg/L. For the advanced analysis after stratification by the NCD<sub>history</sub>, the hsCRP levels were categorized as  $\leq 1.00, 1.01-2.00, 2.01-3.00, 3.01-10.0, and >10.0 mg/L$  because of the reduced sample size in each subgroup. The concentrations of hsCRP were log-transformed for analyses because of the skewed distribution. 

We calculated a follow-up time for each subject starting from the date of interview until the date of death or
 We calculated a follow-up time for each subject starting from the date of interview until the date of death or
 December 31, 2015, whichever came first. Using age as the time scale, subjects enter the risk set at the age at
 which they completed the baseline questionnaire and exit at their event/censoring age. The associations of

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4 5	353	hsCRP and all-cause mortality, as well as cancer and CVD mortality, were analyzed by Cox proportional hazard
6 7	354	models (aHR) and included adjustment for age, gender, demographic factors (education, marital status, job, BMI
8 9	355	and NCD <sub>history</sub> ), and lifestyle factors (smoking, alcohol consumption and exercise). In addition, we conducted a
10 11	356	sensitivity analysis to avoid latent period bias after excluding death before 1 year (aHR <sub>1year</sub> ) or 2 years (aHR <sub>2year</sub> )
12	357	since recruitment. We employed restricted cubic splines (RCSs) to evaluate the possibility of complex (i.e.,
13 14	358	nonlinear) hazard functions[26] using continuous values of hsCRP (aHR <sub>continuous</sub> ). We selected five hsCRP
15 16	359	concentration values as knots based on hsCRP concentration percentiles, tested the linear and nonlinear associa-
17 18	360	tions between knots using a cubic function, and presented the integrated graph smoothly. All statistical analyses
19 20	361	were performed using SAS version 9.3 (SAS Institute Inc., Cary, NC, USA) and RCS analysis was carried out
21 22	362	using the SAS LGTPHCURV9 macro. The <i>P</i> -values < 0.05 were defined as indicating statistical significance.
23 24	363	
25 26	364	Patient and public involvement
27 28	365	No patients and public were involved in the design, conducting, reporting, and dissemination plans of the present
29 30	366	study.
31 32	367	
33 34	368	
35 36	369	study. RESULTS
37 38	370	The association of demographic and lifestyle factors with the risk of all-cause mortality is presented in Table
39 40	371	1. During the follow-up period (average 6.8 years), 1 365 men and 864 women died. The median levels of
41 42	372	hsCRP were 0.77 and 0.59 mg/L for men and women, respectively. The risk of all-cause mortality was inversely
43 44	373	associated with female gender (aHR=0.38), high educated (aHR=0.65), overweight (aHR=0.81) or obesity
45 46	374	(aHR=0.83), current alcohol consumption (aHR=0.81) and regular exercise (aHR=0.83), but was positively
47 48	375	associated with single marital status (aHR=1.23), NCD <sub>history</sub> (aHR=1.57), underweight (aHR=2.05) and current
49 50	376	smoking (aHR=1.97).
51	377	
52 53	378	
54 55	0,0	
55	379	

		All	Death	All-cause m	ortality
		( <i>n</i> =122 081)	( <i>n</i> =2229)	Age,gender adjusted	adj HR <sup>a</sup>
	Age	53.1 ± 8.3	59.7 ± 8.8		
	Female	66.4	38.8	0.40 (0.36-0.43)	0.38 (0.33-0.44)
	Education (≥10 year, %)	68.2	55.4	0.67 (0.60-0.75)	0.65 (0.56-0.75)
	Blue-colored worker <sup>b</sup> (%)	32.3	33.8	1.46 (1.26-1.68)	1.16 (0.99-1.35)
	Marital status (single, %)	11.0	13.3	1.35 (1.19-1.54)	1.23 (1.07-1.40)
	NCD <sub>history</sub> (yes, %)	32.4	53.6	1.51 (1.39-1.65)	1.57 (1.42-1.72)
	Hypertension	18.9	31.5	1.18 (1.08-1.30)	1.22 (1.11-1.35
	Diabete	6.5	17.1	1.81 (1.62-2.03)	1.77 (1.57-2.00)
	Hyperlipidemia	9.2	7.6	0.73 (0.62-0.86)	0.78 (0.66-0.92
	Cancer	3.2	8.8	2.69 (2.31-3.12)	2.66 (2.27-3.11)
	Cerebral & cardiovascular disease	3.7	10.2	1.50 (1.30-1.73)	1.43 (1.23-1.66
	Respiratory disease	2.4	4.3	1.37 (1.12-1.68)	1.32 (1.06-1.64
	Body mass index (%)			× ,	× .
	<18.5	1.8	3.7	2.14 (1.69-2.69)	2.05 (1.61-2.62
	18.5-22.9	38.1	34.9	1.00 (ref.)	1.00 (ref.)
	23.0-24.9	27.8	26.0	0.82 (0.73-0.91)	0.81 (0.72-0.91
	25.0-29.9	29.5	32.5	0.90 (0.81-1.00)	0.83 (0.74-0.93
	$\geq$ 30.0	2.8	2.9	1.08 (0.83-1.39)	0.81 (0.61-1.08
	P-trend			0.0118	<.0001
	Metabolic syndrome (yes, %)	22.0	28.4	1.13 (1.03-1.24)	1.07 (0.96-1.19
	Current smoker (%)	11.7	22.7	2.04 (1.79-2.33)	1.97 (1.71-2.27
	Current drinker (%)	44.0	43.8	0.86 (0.77-0.95)	0.81 (0.73-0.91
	Regular exercise (yes, %)	53.4	49.1	0.76 (0.70-0.83)	0.83 (0.76-0.91
383 384 385 386	Regular exercise (yes, %) NCD <sub>history:</sub> Non-communicable diseas <sup>a</sup> Adjusted for age, gender, education, <sup>b</sup> Compared to white-colored worker	e history			<b>`</b>
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**Table 1** Baseline characteristics of participants by all-cause mortality

The risk of all-cause mortality was inclined with a dose-dependent pattern as increased serum hsCRP level ( $P_{trend} < 0.001$ , Supplement 1), regardless of gender ( $P_{trend} < 0.001$  in both genders), even in the sensitivity analysis  $(P_{trend} < 0.001 \text{ for aHR}_{1\text{ver}} \text{ in both genders})$ . The increased risk of female mortality with increased hsCRP levels was observed in both premenopausal ( $P_{trend}=0.020$ ) and postmenopausal women ( $P_{trend} < 0.001$ ), although the statistical significance in premenopausal women disappeared after sensitivity analysis (Ptrend=0.150 for aHR2year, Supplement 1). The integrated graph, based on the restricted cubic spline analyses, indicated a strong and linear association of serum hsCRP level with all-cause mortality in both genders (aHR<sub>continuous</sub>=1.019 and 1.013 in men and women, respectively, Fig. 2 (a)).

The dose-response association between hsCRP level and the risk of all-cause mortality was not influenced by NCD<sub>history</sub> (Supplement 2). After stratification by gender, however, the attenuated effect by NCD<sub>history</sub> on the association was observed only in women; the linearity of the relationship was observed in healthy women  $(P_{trend}=0.001 \text{ for aHR}_{2\text{vear}})$  but disappeared in women with NCD<sub>history</sub>, particularly after sensitivity analysis with the exclusion of a 2-year follow-up time ( $P_{trend}=0.084$  for aHR<sub>2vear</sub>). Based on the restricted cubic spline analyses, otherwise, the pattern of increase in the association was different depending on the NCD<sub>history</sub> (Fig. 2 (b)(c)). In the healthy subjects, the risk of all-cause mortality was increased with a gradual slope (strength) until 3.0 mg/L hsCRP, with a very steep slope until 4.5 mg/L and finally with a reduced and flattened slope after 4.5 mg/L (Fig. 2 (b)). On the other hand, the slope of the association fluctuated as the hsCRP level increased in the subjects with NCD<sub>history</sub>; the slope increased up to 3.0 mg/L hsCRP but decreased until 4.5 mg/L and rapidly increased after 4.5 mg/L (Fig. 2 (c)).

The association of serum hsCRP with the risk of cancer-mortality was not influenced by NCD<sub>history</sub>  $(P_{trend} < 0.001 \text{ regardless of NCD}_{history})$  (Table 2). Otherwise, after stratification by gender, the association was not observed in women with NCD<sub>history</sub> ( $P_{trend}$  =0.856); however, the association was not influenced by NCD<sub>history</sub> in men (Ptrend<0.001 and 0.002 for aHR in both healthy and NCDhistory) (Table 2). Although the risk of CVD mortality was linearly associated with increasing hsCRP levels, the association was dominant in men (Ptrend=0.002) and in subjects with NCDhistory (Ptrend=0.001, Table 3) after stratified by gender and NCDhistory, respectively. After stratification by gender and NCD<sub>history</sub>, otherwise, the association only appeared in individuals of both genders with NCD<sub>history</sub> (P<sub>trend</sub>=0.015 and 0.035 in men and women with NCD<sub>history</sub>, respectively); no association between hsCRP level and CVD mortality risk was found in either healthy men or women.

	Cancer-mortality							bjects at	recruitme	nt	Subjects with NCD <sub>history</sub> at recruitment				
	E	MR	aHR	HR <sub>1year</sub>	HR <sub>2year</sub>	E	MR	aHR	HR <sub>1year</sub>	HR <sub>2year</sub>	 Е	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>
Total															
≤1.00	590	10.9	Ref	Ref	Ref	270	7.9	Ref	Ref	Ref	320	16.3	Ref	Ref	Ret
1.01-2.00	232	17.1	1.25	1.23	1.17	85	13.4	1.43	1.40	1.31	147	20.3	1.19	1.13	1.09
2.01-3.00	86	20.4	1.32	1.24	1.19	29	16.0	1.38	1.34	1.35	57	23.7	1.35	1.18	1.10
3.01-10.0	149	29.4	1.83	1.76	1.72	54	24.8	2.22	2.07	2.01	95	33.0	1.75	1.59	1.55
>10.0	66	48.9	2.69	2.28	1.96	20	30.6	1.85	1.59	1.57	46	65.9	3.25	2.64	2.16
P-trend			<.001	<.001	<.001			<.001	<.001	<.001			<.001	<.001	<.001
Men						2									
≤1.00	302	18.5	Ref	Ref	Ref	169	23.6	Ref	Ref	Ref	133	14.5	Ref	Ref	Ret
1.01-2.00	144	26.6	1.36	1.36	1.32	95	32.6	1.40	1.38	1.34	49	19.7	1.31	1.34	1.31
2.01-3.00	59	34.7	1.45	1.31	1.19	40	40.4	1.54	1.37	1.16	19	26.7	1.29	1.22	1.26
3.01-10.0	111	52.7	2.17	2.10	2.00	77	64.5	2.26	2.24	2.12	34	37.3	1.98	1.80	1.70
>10.0	50	82.9	3.13	2.66	2.34	38	114.1	4.07	3.42	2.79	13	46.1	1.58	1.40	1.56
P-trend			<.001	<.001	<.001			<.001	<.001	<.001			0.002	0.009	0.015
Women										O,					
≤1.00	288	7.7	Ref	Ref	Ref	137	5.5	Ref	Ref	Ref	151	12.1	Ref	Ref	Re
1.01-2.00	88	10.8	1.13	1.08	0.99	36	9.4	1.60	1.48	1.31	52	12.1	0.86	0.86	0.81
2.01-3.00	27	10.7	1.16	1.17	1.2	10	9.1	1.48	1.50	1.47	17	12.0	0.96	0.98	1.03
3.01-10.0	38	12.9	1.31	1.24	1.29	20	15.8	2.58	2.48	2.57	18	10.7	0.75	0.71	0.74
>10.0	15	20.4	1.89	1.61	1.28	7	18.9	2.16	1.75	1.42	8	21.9	1.66	1.47	1.17
P-trend			0.019	0.074	0.161			<.001	0.001	0.002			0.856	0.635	0.538

E: Number of death, MR: Mortality rate (10 000 person year), Ref: Reference

aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise

HR<sub>1year</sub>: aHR after exclude subjects who died within 1 yr f/u time

 HR<sub>2year</sub>: aHR after exclude subjects who died within 2 yr f/u time

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	Cardiovascular disease mortality					H	Healthy subjects at recruitment					Subjects with NCD <sub>history</sub> at recruitment					
	E	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>	E	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>	E	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>		
Total																	
≤1.00	167	3.1	Ref	Ref	Ref	58	1.7	Ref	Ref	Ref	109	5.5	Ref	Ref	Ref		
1.01-2.00	79	5.8	1.35	1.37	1.23	18	2.8	1.19	1.15	0.94	64	8.4	1.42	1.46	1.36		
2.01-3.00	42	10.0	2.06	2.05	2.02	6	3.3	1.47	1.54	1.46	36	15.0	2.28	2.25	2.26		
3.01-10.0	39	7.7	1.45	1.38	1.44	8	3.7	1.44	1.50	1.70	31	1.08	1.48	1.37	1.40		
>10.0	13	9.6	1.81	1.76	1.59	3	4.6	2.02	2.10	1.58	10	14.3	1.85	1.74	1.68		
P-trend			0.001	0.002	0.004			0.130	0.100	0.162			0.001	0.006	0.009		
Men						20											
≤1.00	89	5.5	Ref	Ref	Ref	25	2.7	Ref	Ref	Ref	64	8.9	Ref	Ref	Ref		
1.01-2.00	45	8.3	1.33	1.32	1.25	12	4.8	1.30	1.22	1.22	33	11.3	1.31	1.33	1.33		
2.01-3.00	30	17.6	2.70	2.67	2.53	3	4.2	1.31	1.37	1.37	27	27.3	3.05	2.99	2.99		
3.01-10.0	24	11.4	1.43	1.36	1.46	6	6.6	1.70	1.79	1.79	18	15.1	1.42	1.21	1.21		
>10.0	8	13.0	1.90	2.02	1.70	3	10.6	3.42	3.61	3.61	5	15.0	1.59	1.62	1.62		
P-trend			0.002	0.003	0.009			0.053	0.038	0.062			0.015	0.027	0.047		
Women																	
≤1.00	78	2.1	Ref	Ref	Ref	33	1.3	Ref	Ref	Ref	45	6.3	Ref	Ref	Ref		
1.01-2.00	34	4.2	1.41	1.46	1.25	6	1.6	1.09	1.13	0.62	28	9.6	1.60	1.66	1.58		
2.01-3.00	12	4.8	1.26	1.30	1.44	3	2.7	1.65	1.70	1.86	9	9.1	1.17	1.20	1.39		
3.01-10.0	15	5.1	1.51	1.45	1.44	2	1.6	1.06	1.07	1.14	13	10.9	1.75	1.64	1.65		
>10.0	5	6.8	1.72	1.35	1.45	0	-	-	-	-	5	15.0	2.51	1.91	2.07		
P-trend			0.092	0.177	0.168			0.940	0.998	0.922			0.035	0.092	0.078		

E: Number of death, MR: Mortality rate (10 000 person year), Ref: Reference

aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise

 $HR_{1year}$ : aHR after exclude subjects who died within 1 yr f/u time

HR<sub>2year</sub>: aHR after exclude subjects who died within 2 yr f/u time

### 

# **DISCUSSION**

This study suggests that the risk of all-cause mortality was associated with elevated hsCRP levels with a dose-response manner in both gender among Asian who have reported low hsCRP levels compared to other races, and was not influenced by NCD<sub>*history*</sub>. Otherwise, the association was influenced by gender and NCD<sub>*history*</sub> although a dose-response association of hsCRP with the risk of cancer- and CVD-mortality was also observed in this population. The level of hsCRP was not associated with the risk of cancer- mortality among women with NCD<sub>*history*</sub>. The risk effect of high hsCRP level on CVD mortality was predominantly observed in men with NCD<sub>*history*</sub>.

Several large cohorts [10-12, 14] have suggested that serum hsCRP levels may differ according to ethnic background, with the highest concentrations seen in African Americans, followed by Hispanic, White, Chinese and Japanese individuals. Although the reason for this ethnic difference is not clearly resolved, genetic diversity[27], the relatively low BMI in Asian populations and ethnic differences in diet and lifestyle[28] have been suggested. Although the extent to which these findings adopt to Asian populations has been unclear, several recent studies [11, 16] conducted in Asia reported a positive association of *hs*CRP with mortality risk. In this population, the hsCRP level was associated with the risk of all-cause mortality in a dose-dependent manner, even though the level of hsCRP was lower than that in the western population. A meta-analysis[29] and large cohort studies[3-6] supported the robustness of the association regardless of adjusted confounders, the cut-off point of CRP level and exclusion deaths within the first 2 years of follow-up. The reason for the discrepancy in *hs*CRP levels with respect to gender is not clearly resolved, although several studies suggested different lifestyle and metabolic risk factors between men and women[30] and genetic diversity[27]. A high level of serum hsCRP in our population was positively related to the increased risk of all-cause mortality in both genders, supported by several previous studies[8, 16, 31]. Nevertheless, several studies reported no association of hsCRP levels with all-cause mortality was observed in women[7, 16]. In particular, the association was shown in postmenopausal women only, which might suggest the protective effect of endogenous female hormones on the low level of hsCRP[32]; the average hsCRP level was 0.48 and 0.68 mg/L for premenopausal and postmenopausal women in this study. The protective effect could be supported by the

4 450 proposition that estrogen or progesterone might to some extent repress the detrimental effects of chronic

<sup>5</sup> 451 inflammation on tissue damage[33].

452 Inflammation has emerged as an important factor in the processes of NCD, including CVD[17], cancer[18],

453 type 2 diabetes[20], COPD[19, 34] and fracture[21]. In addition, medications that had taken to treat any specific

NCD, such as rennin-angiotensin system inhibitors[35] and statins and thiazolidinedione[36], could influence the level of hsCRP. The association between hsCRP and the mortality risk was not attenuated by NCD<sub>history</sub> in either gender in this study, but the statistical significance of the association disappeared in women after sensitivity analysis (aHR<sub>2vear</sub>). A dose-response relationship between hsCRP level and all-cause mortality risk was pronounced in both genders. On the other hand, the positive association of hsCRP with the risk of all-cause mortality risk was significantly observed in only men with NCD<sub>history</sub> but not in women with NCD<sub>history</sub>. The attenuated effect of NCD<sub>history</sub> on the association between hsCRP and the risk of cancer-mortality was not observed in men, consistent with results from several studies which reported the associations among healthy men[3] or cancer patients[37, 38] only. Most studies[3, 4, 6, 7, 15, 16, 31, 39] supported that CVD mortality increased with elevated hsCRP levels, predominantly in men[4, 7, 15, 16]. Although hsCRP levels are lower in our population than in other races, the level of hsCRP was positively associated with CVD mortality in men but not in women, similar to previous studies[7, 15, 16, 31, 39]. After stratification by gender and NCD<sub>history</sub>, the association between hsCRP and the risk of CVD mortality was dominant in subjects with NCD<sub>history</sub> in this study.

This study has several strengths because of the large population-based prospective study; it makes possible 1) to adjust for confounders; 2) to examine sensitivity analysis after excluding death before 1 or 2 years from recruitment; 3) to assess an advanced analysis after stratification by gender and NCD<sub>history</sub>; 4) to examine the association using various cut-off points of hsCRP considering low serum hsCRP levels in Asian populations; and 5) to evaluate the complex (i.e., nonlinear) hazard functions using restricted cubic splines on the association between continuous hsCRP levels and the risk of mortality. In particular, most previous studies excluded subjects with more than 10 mg/L hsCRP because of their relatively low sample size or reflecting acute phase reactions of severe inflammation, but we examined the effect of very high hsCRP concentration on the risk of mortality because it is possible to be more concerning for these subjects in the future. The hsCRP level of this study, in addition, was measured within 18 hours in a single institution to minimize measurement error/bias from institutional variation to avoid bias from measurement or long-term storage before analysis. Despite of those strengths, it is also has several limitations. First, the use of a single measurement of hsCRP at baseline could reflect the inaccurate status of blood hsCRP levels in the study participants and increase the

481 instability of *hs*CRP due to random fluctuations over time. Nevertheless, a report [40] on the long-term *hs*CRP

- 482 variability suggested that the *hs*CRP variability within individual is relatively small and that the variability
- 60 483 could not account for the association. Second, our study lacked information on medication use at recruitment

3		
4 5	484	and during the follow-up period. Several medications related to NCDs, including statins, angiotensin-converting
6 7	485	enzyme inhibitors, fibrates, niacin, thiazolidinedione and estrogen/progestogen hormone, could influence the
8	486	hsCRP level[37]; however, we tried to overcome this limitation through advanced analysis after stratification by
9 10 11	487	NCD <sub>history</sub> . Third, because there is no available information on hormone-replacement therapy (HRT) among
11 12	488	women, which could not examine the influence of HRT on the association of hsCRP with the risk of hormone-
13 14	489	related cancer or CVD mortality among women, we could not suggest the effect of female hormones on the
15 16	490	association.
17 18	491	In conclusion, the association of hsCRP level is dose-responsively increased with the risk of all-cause
19 20	492	mortality in men and women (particularly postmenopausal women), which was not influenced by the association
21 22	493	was not observed in women with NCD <sub>history</sub> . Otherwise, the association of hsCRP level with the risk of cancer-
23 24	494	and CVD-mortality could be attenuated by gender or NCD <sub>history</sub> .
25 26	495	
27 28	496	
29 30	497	Contributors
31	498	SAL, XS and DK: designed and conducted the research, SAL and SOK: analyzed the data and performed the
32 33	499	statistical analyses; HP and JKL: managed data mining and collection; SAL: wrote the manuscript and had primary
34 35	500	responsibility for the final content of the manuscript; and all authors: read and approved the final manuscript.
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42		
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45 46	505	
47		
48 49	506	Patient consent for publication Not required.
50 51	507	
52 53	508	Ethics approval The Institutional Review Board of the Seoul National University Hospital, Seoul, Korea,
54 55	509	approved it for statistical analysis (IRB No. E-1503-103-657).
56 57	510	
58 59	511	Drevenence and near review. Not commissioned, automatic near reviews 4
60	JTT	<b>Provenance and peer review</b> Not commissioned; externally peer reviewed.

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3 4	510		
5	512	Data availability statement	
6 7	513	No additional data available.	
8	514		
9 10 11	515		
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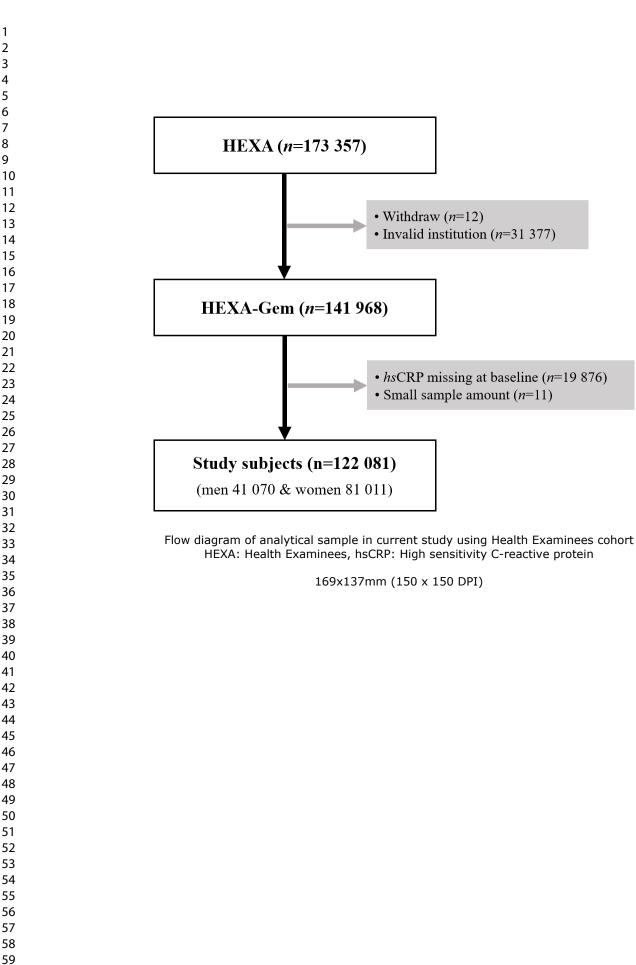
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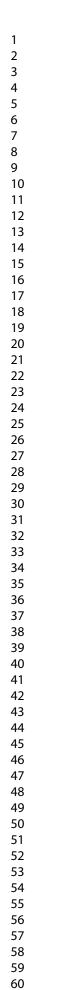
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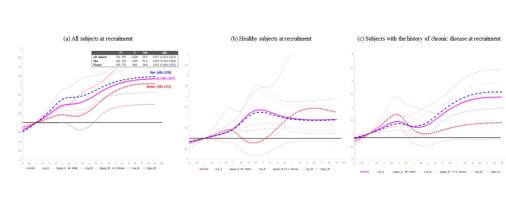
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A dose-response association between serum hsCRP level and risk of all-cause mortality by subject. PY: Person-year, E: Number of death, MR: Mortality rate (10 000 person year) aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise Low\_A and Upper\_A: 95%CI for all subjects

Low\_M and Upper\_M: 95%CI for men Low\_W and Upper\_W: 95%CI for women

330x97mm (96 x 96 DPI)

	PY	Е	MR	aHR	HR <sub>1year</sub>	HR <sub>2year</sub>
All subjects					- ,	
Continuous	781 035	2229	28.5	1.017 (1.012-1.021)	1.015 (1.010-1.020)	1.014 (1.009-1.019
1.00	539 271	1153	21.4	Reference	Reference	Reference
1.01-1.50	90 911	308	33.9	1.26 (1.10-1.45)	1.27 (1.10-1.45)	1.21 (1.04-1.40)
1.51-2.00	44 615	163	36.5	1.28 (1.08-1.53)	1.28 (1.07-1.53)	1.26 (1.04-1.52)
2.01-2.50	25 139	117	46.5	1.53 (1.25-1.89)	1.51 (1.22-1.87)	1.49 (1.19-1.87)
2.51-3.00	16 996	72	42.4	1.39 (1.08-1.80)	1.31 (1.00-1.72)	1.23 (0.92-1.65)
3.01-4.00	19 667	103	52.4	1.61 (1.29-2.01)	1.62 (1.29-2.03)	1.64 (1.30-2.08)
4.01-6.00	17 933	102	56.9	1.84 (1.48-2.28)	1.77 (1.41-2.21)	1.70 (1.34-2.16)
6.01-10.00	13 019	88	67.6	2.02 (1.59-2.56)	1.96 (1.54-2.50)	1.93 (1.49-2.51)
>10.0	13 484	123	91.2	2.59 (2.12-3.16)	2.41 (1.95-2.97)	2.26 (1.80-2.84)
P-trend				<.001	<.001	<.001
Men						
Continuous	261 321	1365	52.2	1.019 (1.014-1.025)	1.017 (1.011-1.023)	1.017 (1.010-1.023
1.00	163 068	638	39.1	Reference	Reference	Reference
1.01-1.50	36 094	190	52.6	1.27 (1.07-1.51)	1.28 (1.07-1.53)	1.22 (1.01-1.47)
1.51-2.00	17 946	103	57.4	1.34 (1.07-1.67)	1.34 (1.07-1.68)	1.35 (1.06-1.72)
2.01-2.50	10 059	77	76.5	1.56 (1.20-2.03)	1.53 (1.16-2.00)	1.47 (1.10-1.96)
2.51-3.00	6959	54	77.6	1.71 (1.27-2.29)	1.57 (1.15-2.15)	1.46 (1.04-2.05)
3.01-4.00	8177	77	94.2	1.88 (1.45-2.43)	1.94 (1.50-2.52)	1.92 (1.46-2.54)
4.01-6.00	7425	75	101.0	2.05 (1.59-2.63)	1.95 (1.49-2.53)	1.91 (1.44-2.52)
6.01-10.00	5456	59	108.1	2.03 (1.52-2.73)	1.96 (1.44-2.66)	1.85 (1.33-2.58)
>10.0	6137	92	149.9	2.84 (2.25-3.58)	2.66 (2.08-3.39)	2.58 (1.99-3.35)
<i>P</i> -trend	0157	12	149.9	<.001	<.001	<.001
Women						
Continuous	519 714	864	16.6	1.013 (1.004-1.021)	1.011(1.002-1.021)	1.010 (0.999-1.02)
1.00	376 203	515	13.7	Reference	Reference	Reference
1.01-1.50	54 817	118	21.5	1.28 (1.03-1.59)	1.27 (1.02-1.58)	1.23 (0.97-1.56)
1.51-2.00	26 669	60	22.5	1.23 (0.92-1.64)	1.21 (0.90-1.63)	1.14 (0.83-1.56)
2.01-2.50	15 080	40	26.5	1.52 (1.09-2.14)	1.52 (1.08-2.15)	1.56 (1.09-2.24)
2.51-3.00	10 037	18	17.9	0.84 (0.49-1.44)	0.87 (0.51-1.48)	0.83 (0.46-1.47)
3.01-4.00	11 490	26	22.6	1.16 (0.75-1.81)	1.09 (0.68-1.72)	1.21 (0.76-1.93)
4.01-6.00	10 508	20	25.7	1.48 (0.99-2.22)	1.47 (0.97-2.22)	1.36 (0.86-2.14)
6.01-10.00	7563	29	38.3	2.00 (1.34-2.98)	1.98 (1.32-2.98)	2.10 (1.39-3.19)
>10.0	7347	31	42.2	2.02 (1.36-3.02)	1.84 (1.21-2.81)	1.51 (0.93-2.47)
<i>P</i> -trend	/ 54 /	JI	72.2	<.001	<.001	0.001
Premenopause				<.001	<.001	0.001
1.00	141 286	96	6.8			
1.01-2.00	20 500	90 20	0.8 9.8	1.52 (0.92-2.52)	1.49 (0.89-2.50)	1.57 (0.90-2.73)
2.01-3.00			9.8 10.3			
3.01-10.0	5835	6		1.76 (0.77-4.06)	1.83 (0.79-4.22)	1.42 (0.52-3.93)
3.01-10.0 >10.0	6886 1750	6	8.7 22.7	1.51 (0.66-3.50)	1.31 (0.53-3.25) 2.63 (0.83-8.37)	1.21 (0.44-3.36)
	1759	4	22.1	2.57 (0.81-8.14)		2.09 (0.51-8.58)
<i>P</i> -trend				0.020	0.036	0.150
Postmenopause	100 164	200	10.0			
1.00	192 164	366	19.0	1.06 (1.02, 1.55)	1.05 (1.00.1.54)	1 10 (0 07 1 40)
1.01-2.00	52 897	145	27.4	1.26 (1.03-1.55)	1.25 (1.02-1.54)	1.18 (0.95-1.48)
2.01-3.00	16 943	44	26.0	1.11 (0.80-1.56)	1.12 (0.80-1.57)	1.19 (0.83-1.68)
3.01-10.0	19 687	67	34.0	1.49 (1.13-1.97)	1.47 (1.10-1.95)	1.52 (1.13-2.05)
>10.0	4828	27	55.9	2.09 (1.37-3.21)	1.88 (1.19-2.96)	1.56 (0.92-2.63)
P-trend				< 0.001	0.001 ar)	0.003

Supplement 1. The association of serum hsCRP level with the risk of all-cause mortality

aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise

HR<sub>1year</sub>: aHR after exclude subjects who died within 1 yr f/u time

HR<sub>2year</sub>: aHR after exclude subjects who died within 2 yr f/u time

<text>

	Healthy subjects at recruitment				ent	Subjects with NCD <sub>history</sub> at recruitment				
	E	MR	aHR	HR <sub>1year</sub>	HR <sub>2year</sub>	Е	MR	aHR	HR <sub>1year</sub>	HR <sub>2yea</sub>
All										
1.00	517	15.1	Ref	Ref	Ref	636	32.3	Ref	Ref	Re
1.01-2.00	145	22.9	1.20	1.19	1.16	326	45.1	1.20	1.19	1.10
2.01-3.00	53	29.3	1.38	1.37	1.32	136	56.6	1.51	1.46	1.41
3.01-10.0	102	46.8	2.22	2.15	2.15	191	66.3	1.62	1.60	1.58
>10.0	40	61.3	2.38	2.23	2.27	83	118.9	2.74	2.54	2.29
P-trend			<.001	<.001	<.001			<.001	<.001	<.00
Men										
1.00	270	29.5	Ref	Ref	Ref	368	51.4	Ref	Ref	Re
1.01-2.00	89	35.8	1.11	1.11	1.13	204	70.0	1.40	1.41	1.33
2.01-3.00	33	46.3	1.22	1.17	1.15	98	99.0	1.82	1.73	1.61
3.01-10.0	70	76.8	2.14	2.08	2.03	141	118.1	1.92	1.90	1.83
>10.0	31	110.0	2.60	2.49	2.73	61	183.1	3.05	2.83	2.58
P-trend			<.001	<.001	<.001			<.001	<.001	<.001
Women										
1.00	247	9.8	Ref	Ref	Ref	268	21.4	Ref	Ref	Re
1.01-2.00	56	14.6	1.35	1.32	1.20	122	28.3	1.19	1.20	1.19
2.01-3.00	20	18.2	1.61	1.66	1.60	38	26.9	1.06	1.06	1.1
3.01-10.0	32	25.2	2.31	2.23	2.37	50	29.7	1.16	1.14	1.17
>10.0	9	24.3	1.69	1.49	1.12	22	60.3	2.15	1.99	1.68
P-trend			<.001	<.001	0.001			0.018	0.043	0.08

**Supplement 2.** The association between serum hsCRP level and all-cause mortality by gender and noncommunicable disease history (NCD<sub>*history*</sub>) at recruitment

E: Number of death, MR: Mortality rate (10 000 person year)

aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise

 $HR_{1year}$ : aHR after exclude subjects who died within 1 yr f/u time

HR<sub>2year</sub>: aHR after exclude subjects who died within 2 yr f/u time

Section/Topic	Item #	Recommendation	Reported on page #
Title and abstract	1	(a) Indicate the study's design with a commonly used term in the title or the abstract	1
		(b) Provide in the abstract an informative and balanced summary of what was done and what was found	1-2
Introduction			
Background/rationale	2	Explain the scientific background and rationale for the investigation being reported	3
Objectives	3	State specific objectives, including any pre-specified hypotheses	3
Methods			
Study design	4	Present key elements of study design early in the paper	4
Setting	5	Describe the setting, locations, and relevant dates, including periods of recruitment, exposure, follow-up, and data collection	4-5
Participants	6	<ul> <li>(a) Cohort study—Give the eligibility criteria, and the sources and methods of selection of participants. Describe methods of follow-up</li> <li>Case-control study—Give the eligibility criteria, and the sources and methods of case ascertainment and control selection. Give the rationale for the choice of cases and controls</li> <li>Cross-sectional study—Give the eligibility criteria, and the sources and methods of selection of participants</li> </ul>	4
		(b) Cohort study—For matched studies, give matching criteria and number of exposed and unexposed Case-control study—For matched studies, give matching criteria and the number of controls per case	-
Variables 7		Clearly define all outcomes, exposures, predictors, potential confounders, and effect modifiers. Give diagnostic criteria, if applicable	5
Data sources/ measurement 8		For each variable of interest, give sources of data and details of methods of assessment (measurement). Describe comparability of assessment methods if there is more than one group	5
Bias	9	Describe any efforts to address potential sources of bias	6
Study size	10	.0 Explain how the study size was arrived at	
Quantitative variables 11		Explain how quantitative variables were handled in the analyses. If applicable, describe which groupings were chosen and why	5-6
Statistical methods	12	(a) Describe all statistical methods, including those used to control for confounding	5-6
		(b) Describe any methods used to examine subgroups and interactions	5-6
		(c) Explain how missing data were addressed	4
		(d) Cohort study—If applicable, explain how loss to follow-up was addressed Case-control study—If applicable, explain how matching of cases and controls was addressed	4

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		Cross-sectional study—If applicable, describe analytical methods taking account of sampling strategy	
		(e) Describe any sensitivity analyses	6
Results	·		
Participants	13*	(a) Report numbers of individuals at each stage of study—eg numbers potentially eligible, examined for eligibility, confirmed eligible, included in the study, completing follow-up, and analysed	4
		(b) Give reasons for non-participation at each stage	4
		(c) Consider use of a flow diagram	4
Descriptive data	14*	(a) Give characteristics of study participants (eg demographic, clinical, social) and information on exposures and potential confounders	6-7
		(b) Indicate number of participants with missing data for each variable of interest	4
		(c) Cohort study—Summarise follow-up time (eg, average and total amount)	6
Outcome data	15*	Cohort study—Report numbers of outcome events or summary measures over time	6-10
		Case-control study—Report numbers in each exposure category, or summary measures of exposure	
		Cross-sectional study—Report numbers of outcome events or summary measures	
Main results	16	(a) Give unadjusted estimates and, if applicable, confounder-adjusted estimates and their precision (eg, 95% confidence interval). Make clear which confounders were adjusted for and why they were included	6-10
		(b) Report category boundaries when continuous variables were categorized	5
		(c) If relevant, consider translating estimates of relative risk into absolute risk for a meaningful time period	-
Other analyses	17	Report other analyses done—eg analyses of subgroups and interactions, and sensitivity analyses	8
Discussion	·	•	
Key results	18	Summarise key results with reference to study objectives	11
Limitations	19	Discuss limitations of the study, taking into account sources of potential bias or imprecision. Discuss both direction and magnitude of any potential bias	
Interpretation	20	Give a cautious overall interpretation of results considering objectives, limitations, multiplicity of analyses, results from similar studies, and other relevant evidence	11-12
Generalisability	21	Discuss the generalisability (external validity) of the study results	11-12
Other information	·	·	
Funding	22	Give the source of funding and the role of the funders for the present study and, if applicable, for the original study on which the present article is based	13

\*Give information separately for cases and controls in case-control studies and, if applicable, for exposed and unexposed groups in cohort and cross-sectional studies. **Note:** An Explanation and Elaboration article discusses each checklist item and gives methodological background and published examples of transparent reporting. The STROBE checklist is best used in conjunction with this article (freely available on the Web sites of PLoS Medicine at http://www.plosmedicine.org/, Annals of Internal Medicine at http://www.annals.org/, and Epidemiology at http://www.epidem.com/). Information on the STROBE Initiative is available at www.strobe-statement.org.

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## The association of serum high sensitivity C-reactive protein with the risk of mortality in Asian: the Health Examinees cohort

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5	196	The association of serum high sensitivity C-reactive protein with the risk of mortality in
6 7	197	Asian: the Health Examinees cohort
8	198	
9 10	199	
10 11	200	Sang-Ah Lee <sup>1,2 *</sup> , Sung Ok Kwon <sup>1</sup> , Hyerim Park <sup>1</sup> , Xiao-Ou Shu <sup>2</sup> , Jong-Koo Lee <sup>3</sup> , Daehee Kang <sup>4</sup>
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13 14	202	
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22	208	<sup>4</sup> Department of Preventive Medicine, College of Medicine, Seoul National University, Seoul, Republic of Korea.
23 24	209	
25	210	
26 27	211	ABSTRACT
27	212	<b>Objectives</b> This study aimed to examine the association of <i>hs</i> CRP with mortality risk and the attenuated effect
29	213	of non-communicable disease history (NCD <sub>history</sub> ) on the association.
30 31	214	Design Prospective cohort study.
32	215	Setting the Health Examinees (HEXA) cohort.
33 34	216	Participants A total of 41 070 men and 81 011 women aged ≥40 years were involved (follow-up: 6.8 years).
35	217	<b>Outcome measures</b> The data and cause of death occurring until December 31, 2015, were confirmed by death
36 37	218	statistics from the National Statistical Office. We conducted the advanced analysis after stratification by
38	219	NCD <sub>history</sub> and the sensitivity analysis after excluding death before 1 or 2 years from recruitment. Cox
39 40	220	proportional hazard and restricted cubic spline models were used to assess the association.
40 41	221	<b>Results</b> The association between serum <i>hs</i> CRP and the risk of all-cause mortality was observed with strong
42	222	linearity in both genders, which was not influenced by NCD <sub>history</sub> . Otherwise, the association of serum hsCRP
43 44	223	with cancer-mortality risk was not observed in women with NCD <sub>history</sub> , but the association with the risk of
45	224	cardiovascular disease (CVD) mortality was predominantly observed in men with NCD <sub>history</sub> .
46 47	225	<b>Conclusions</b> This study suggested the dose-response association of <i>hs</i> CRP with mortality risk, including
48	226	cancer and CVD mortality, in Korean with low serum hsCRP, although the association with cancer and CVD-
49 50	227	mortality risk could be influenced by gender and NCD <sub>history</sub> .
51	228	
52	229	
53 54	230	Strengths and limitations of this study
55 56	231	• This is the large population-based prospective study.
57 58 59	232	• We examined the effect of very high <i>hs</i> CRP concentration on mortality risk.
60	233	• The <i>hs</i> CRP level of present study was measured within 18 hours in a single institution to minimize error/bias.

2 3		
4 5	234	• Due to random fluctuations of $hs$ CRP, using the single measurement of $hs$ CRP at baseline could reflect the
6 7	235	inaccurate status of blood hsCRP levels in the study participants and increase the instability of hsCRP.
, 8 9	236	· This study lacked information on medication use at recruitment and during the follow-up period, and
10	237	information on hormone-replacement therapy (HRT) among women.
11 12	238	
13	239	
14 15	240	*Correspondence to: Sang-Ah Lee, Ph.D.
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## 266 INTRODUCTION

High sensitivity C-reactive protein (hsCRP) is an acute-phase response protein synthesized by the liver and the most sensitive and dynamic marker of inflammation[1]. Since hsCRP has been reported as a candidate marker for generalized atherosclerosis and cardiovascular disease (CVD)[2], many studies[3-7] have investigated the role of hsCRP levels as a predictor of mortality risk. A recent meta-analysis[8] reported the predictable role of serum hsCRP on all-cause and CVD mortality in the general population. Nevertheless, it is controversial whether the predictable role of hsCRP could be applied to the risk of mortality in Asians, whose hsCRP levels are lower than those in individuals in Western countries.

Serum *hs*CRP represents a low-grade inflammation state that is generally involved in the process of aging[9]. Several large cohorts, including Study of Women's Health Across the Nation (SWAN)[10], the Women's Health Study[11] and the Dallas Heart Study[12], reported significant differences in hsCRP levels by race and gender. In two studies of multiethnic populations residing in the USA[10, 13], the median hsCRP level in East Asians was less than half the concentration in Caucasians. Even among East Asian populations, the geometric mean of hsCRP levels varied depending on ethnic background[14]. In addition, a meta-analysis[11] reported the hsCRP levels among women of various ethnic groups living in the United States (from the Women's Health Study) on the association between hsCRP and the mortality risk; the association was observed in only men supported by the results from two cohort studies [15, 16] reported in Korea. On the other hand, the increased hsCRP may be influenced by comorbidity itself because inflammation has emerged as an important factor in the progression of non-communicable diseases (NCDs), including CVD[17], cancer[18], chronic obstructive pulmonary disease (COPD)[19], type 2 diabetes[20] and fractures[21], which contribute to increased morbidity and mortality.

This study aimed to examine the association of serum hsCRP with the risk of mortality in Koreans with low serum hsCRP and to evaluate the attenuated effect of non-communicable disease history (NCD<sub>history</sub>) on the association.

#### **METHODS**

#### Study population

Details on the main objectives, rationale, study design and baseline characteristics of the Health Examinees (HEXA) study have been published elsewhere[22]. Considering the homogeneity and comparability of participants, we created a qualified dataset called HEXA-G (Health Examinees-Gem) from previously published HEXA studies[23]. In the new HEXA-G data, a total of 141 968 participants remained after the exclusion of withdrawers (n=12). In addition, 19 887 were excluded due to missing information (n=19 876) or small sample size (n=11) on any hsCRP components at the baseline survey. Ultimately, 122 081 subjects, including 41 070 men and 81 011 women, remained in the final analysis (Fig. 1). All study participants provided informed consent prior to entering the study. The Institutional Review Board of the Seoul National University Hospital, Seoul, Korea, approved it for statistical analysis (IRB No. E-1503-103-657).

#### Laboratory measurements

After at least 10 hours of overnight fasting, blood samples were obtained in the morning. Bio-specimens included fasting blood samples that were collected in a serum separator tube and two ethylenediaminetetraacetic acid (EDTA) tubes. All samples were then transported to the National Biobank of Korea and stored for future research purposes within 18 hours. hsCRP was measured using a turbidimetric immunoassay (ADVIA 1650 and ADVIA 1800; Siemens Healthineers).

#### Follow-up and ascertainment of mortality

All-cause mortality was confirmed by death statistics from the National Statistical Office, which provided the data and causes of all deaths occurring through December 31, 2015. We added the mortality data from Statistics Korea to our dataset using each participant's unique identifier. Information on death and causes of death was obtained from a record link with the national death certificate files in Korea. The main outcome of interest was all-cause mortality (defined as death from any cause), including cancers and CVD mortality. The cause of death was classified according to the International Classification of Diseases, 10th revision (ICD-10). Deaths were coded as C00-C97 for cancer and I00-I99 for CVD. 

## 323 Baseline variables

Trained interviewers collected information on demographic, socioeconomic and lifestyle factors. Anthropometric measurements were obtained using standardized methods. Body mass index (BMI) was calculated, and all participants were defined into four classes based on the World Health Organization classification of BMI for Asian adults[24]: underweight (BMI <18.5 kg/m<sup>2</sup>), normal (18.5 BMI <23.0 kg/m<sup>2</sup>), overweight (23.0 $\leq$  BMI <25.0 kg/m<sup>2</sup>), obesity (25.0 $\leq$  BMI <29.9 kg/m<sup>2</sup>), and severe obesity (BMI  $\geq$ 30.0 kg/m<sup>2</sup>). The current study defined metabolic syndrome using the National Cholesterol Education Program Adult Treatment Panel III (NCEP-ATP III)[25], modified for the Asian guideline for waist circumference (WC ≥90 and  $\geq$ 80 cm for men and women, respectively). Nonsmokers were defined as those who had smoked less than 400 cigarettes over the course of their lifetime. Participants who had smoked were categorized into two groups: noncurrent (never/former) and current smoker. Noncurrent drinkers were defined as those who had never consumed an alcoholic drink over the course of their lifetime or those who had not consumed alcohol at recruitment, while current drinkers were defined as those who persisted in consuming alcohol. Regular exercise was classified into two groups (ves/no) as follows: "Do you currently engage in regular exercise strenuous enough to cause you to break into a sweat at least once per week?" Furthermore, considering the attenuated effect of the NCD<sub>history</sub> on the association between serum hsCRP and the risk of mortality, we performed advanced analysis after stratification by NCD<sub>history</sub>. We considered six main non-communicable diseases (hypertension, diabetes, hyperlipidemia, cancer, cardiovascular and cerebrovascular diseases, and respiratory disease) to classify healthy subjects vs. subjects with NCD<sub>history</sub>.

<sup>10</sup> 342

## 343 Statistical analysis

For the categorical analysis, we created nine categories based on the distribution of *hs*CRP levels in our population:  $\leq 1.00$  (reference group), 1.01-1.50, 1.51-2.00, 2.01-2.50, 2.51-3.00, 3.01-4.00, 4.01-6.00, 6.01-10.0, and  $\geq 10.0$  mg/L. For the advanced analysis after stratification by the NCD<sub>*history*</sub>, the *hs*CRP levels were categorized as  $\leq 1.00$ , 1.01-2.00, 2.01-3.00, 3.01-10.0, and  $\geq 10.0$  mg/L because of the reduced sample size in each subgroup. The concentrations of *hs*CRP were log-transformed for analyses because of the skewed distribution.

We calculated a follow-up time for each subject starting from the date of interview until the date of death or
 We calculated a follow-up time for each subject starting from the date of interview until the date of death or
 December 31, 2015, whichever came first. Using age as the time scale, subjects enter the risk set at the age at
 which they completed the baseline questionnaire and exit at their event/censoring age. The associations of

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5 4 5	353	hsCRP and all-cause mortality, as well as cancer and CVD mortality, were analyzed by Cox proportional hazard
6 7	354	models (aHR) and included adjustment for age, gender, demographic factors (education, marital status, job, BMI
, 8 9	355	and NCD <sub>history</sub> ), and lifestyle factors (smoking, alcohol consumption and exercise). We used Wald tests to test
10	356	for heterogeneity of risk between serum hsCRP level groups. The proportional hazards assumption was assessed
11 12	357	on the basis of Schoenfeld residuals, and was not violated for the variables of interest in the adjusted model for
13 14	358	either cancer-mortality or cardiovascular disease mortality (P>0.05 for all categories). In addition, we conducted
15 16	359	a sensitivity analysis to avoid latent period bias after excluding death before 1 year (aHR <sub>1year</sub> ) or 2 years
17 18	360	(aHR <sub>2year</sub> ) since recruitment. Based on the Cox proportional hazard models, we made Kaplan-Meier curves and
19 20	361	log-rank analysis. We employed restricted cubic splines (RCSs) to evaluate the possibility of complex (i.e.,
21 22	362	nonlinear) hazard functions[26] using continuous values of hsCRP (aHR <sub>continuous</sub> ). We selected five hsCRP
23 24	363	concentration values as knots based on hsCRP concentration percentiles, tested the linear and nonlinear associa-
25 26	364	tions between knots using a cubic function, and presented the integrated graph smoothly. All statistical analyses
27 28 29 30 31 32	365	were performed using SAS version 9.3 (SAS Institute Inc., Cary, NC, USA) and RCS analysis was carried out
	366	using the SAS LGTPHCURV9 macro. Two-sided <i>p</i> -values < 0.05 were defined as indicating statistical
	367	significance.
33 34	368	significance. Patient and public involvement
35 36	369	Patient and public involvement
37	370	No patients and public were involved in the design, conducting, reporting, and dissemination plans of the present
38 39	371	study.
40 41	372	
42 43	373	PESIII TS
44 45	374	RESULTS
46 47 48	375	The association of demographic and lifestyle factors with the risk of all-cause mortality is presented in Table
49	376	1. During the follow-up period (average 6.8 years), 1 365 men and 864 women died. The median levels of
50 51	377	hsCRP were 0.77 and 0.59 mg/L for men and women, respectively. The risk of all-cause mortality was inversely
52 53	378	associated with female gender (aHR=0.38), high educated (aHR=0.65), overweight (aHR=0.81) or obesity
54 55	379	(aHR=0.83), current alcohol consumption (aHR=0.81) and regular exercise (aHR=0.83), but was positively
56 57	380	associated with single marital status (aHR=1.23), NCD <sub>history</sub> (aHR=1.57), underweight (aHR=2.05) and current
58 59	381	smoking (aHR=1.97).

		All	Death	All-cause m	ortality
		( <i>n</i> =122 081)	( <i>n</i> =2229)	Age,gender adjusted	adj HR <sup>a</sup>
	Age	53.1 ± 8.3	59.7 ± 8.8		
	Female	66.4	38.8	0.40 (0.36-0.43)	0.38 (0.33-0.44)
	Education (≥10 year, %)	68.2	55.4	0.67 (0.60-0.75)	0.65 (0.56-0.75)
	Blue-colored worker <sup>b</sup> (%)	32.3	33.8	1.46 (1.26-1.68)	1.16 (0.99-1.35)
	Marital status (single, %)	11.0	13.3	1.35 (1.19-1.54)	1.23 (1.07-1.40)
	NCD <sub>history</sub> (yes, %)	32.4	53.6	1.51 (1.39-1.65)	1.57 (1.42-1.72)
	Hypertension	18.9	31.5	1.18 (1.08-1.30)	1.22 (1.11-1.35
	Diabete	6.5	17.1	1.81 (1.62-2.03)	1.77 (1.57-2.00)
	Hyperlipidemia	9.2	7.6	0.73 (0.62-0.86)	0.78 (0.66-0.92
	Cancer	3.2	8.8	2.69 (2.31-3.12)	2.66 (2.27-3.11)
	Cerebral & cardiovascular disease	3.7	10.2	1.50 (1.30-1.73)	1.43 (1.23-1.66
	Respiratory disease	2.4	4.3	1.37 (1.12-1.68)	1.32 (1.06-1.64
	Body mass index (%)			× ,	× .
	<18.5	1.8	3.7	2.14 (1.69-2.69)	2.05 (1.61-2.62
	18.5-22.9	38.1	34.9	1.00 (ref.)	1.00 (ref.)
	23.0-24.9	27.8	26.0	0.82 (0.73-0.91)	0.81 (0.72-0.91
	25.0-29.9	29.5	32.5	0.90 (0.81-1.00)	0.83 (0.74-0.93
	$\geq$ 30.0	2.8	2.9	1.08 (0.83-1.39)	0.81 (0.61-1.08
	P-trend			0.0118	<.0001
	Metabolic syndrome (yes, %)	22.0	28.4	1.13 (1.03-1.24)	1.07 (0.96-1.19
	Current smoker (%)	11.7	22.7	2.04 (1.79-2.33)	1.97 (1.71-2.27
	Current drinker (%)	44.0	43.8	0.86 (0.77-0.95)	0.81 (0.73-0.91
383 384 385 386	Regular exercise (yes, %)	53.4	49.1	0.76 (0.70-0.83)	0.83 (0.76-0.91
	Regular exercise (yes, %) NCD <sub>history:</sub> Non-communicable diseas <sup>a</sup> Adjusted for age, gender, education, <sup>b</sup> Compared to white-colored worker	e history			<b>`</b>
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**Table 1** Baseline characteristics of participants by all-cause mortality

The risk of all-cause mortality was inclined with a dose-dependent pattern as increased serum hsCRP level (Ptrend<0.001, Supplement 1), regardless of gender (Ptrend<0.001 in both genders), even in the sensitivity analysis  $(P_{trend} < 0.001 \text{ for aHR}_{1\text{ver}} \text{ in both genders})$ . The increased risk of female mortality with increased hsCRP levels was observed in both premenopausal ( $P_{trend}=0.020$ ) and postmenopausal women ( $P_{trend} < 0.001$ ), although the statistical significance in premenopausal women disappeared after sensitivity analysis ( $P_{trend}$ =0.150 for aHR<sub>2year</sub>, Supplement 1). The integrated graph, based on the restricted cubic spline analyses, indicated a strong and linear association of serum hsCRP level with all-cause mortality in both genders (aHR<sub>continuous</sub>=1.019 and 1.013 in men and women, respectively, Fig. 2).

The dose-response association between hsCRP level and the risk of all-cause mortality was not influenced by NCD<sub>history</sub> (Supplement 2). After stratification by gender, however, the attenuated effect by NCD<sub>history</sub> on the association was observed only in women; the linearity of the relationship was observed in healthy women  $(P_{trend}=0.001 \text{ for aHR}_{2\text{vear}})$  but disappeared in women with NCD<sub>history</sub>, particularly after sensitivity analysis with the exclusion of a 2-year follow-up time ( $P_{trend}=0.084$  for aHR<sub>2vear</sub>). Based on the restricted cubic spline analyses, otherwise, the pattern of increase in the association was different depending on the NCD<sub>history</sub> (Fig. 3.4). In the healthy subjects, the risk of all-cause mortality was increased with a gradual slope (strength) until 3.0 mg/L hsCRP, with a very steep slope until 4.5 mg/L and finally with a reduced and flattened slope after 4.5 mg/L (Fig. 3). On the other hand, the slope of the association fluctuated as the hsCRP level increased in the subjects with NCD<sub>history</sub>; the slope increased up to 3.0 mg/L hsCRP but decreased until 4.5 mg/L and rapidly increased after 4.5 mg/L (Fig. 4).

The association of serum hsCRP with the risk of cancer-mortality was not influenced by NCD<sub>history</sub> (Ptrend<0.001 regardless of NCDhistory) (Table 2 and Fig. 5-9). Otherwise, after stratification by gender, the association was not observed in women with NCD<sub>history</sub> ( $P_{trend} = 0.856$ ); however, the association was not influenced by NCD<sub>history</sub> in men (P<sub>trend</sub><0.001 and 0.002 for aHR in both healthy and NCD<sub>history</sub>) (Table 2). Although the risk of CVD mortality was linearly associated with increasing hsCRP levels, the association was dominant in men (Ptrend=0.002) and in subjects with NCDhistory (Ptrend=0.001, Table 3) after stratified by gender and NCD<sub>history</sub>, respectively (Fig. 10-14). After stratification by gender and NCD<sub>history</sub>, otherwise, the association only appeared in individuals of both genders with NCD<sub>history</sub> (P<sub>trend</sub>=0.015 and 0.035 in men and women with NCD<sub>history</sub>, respectively); no association between hsCRP level and CVD mortality risk was found in either healthy men or women.

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	Cancer-mortality					Healthy subjects at recruitment				Subjects with NCD <sub>history</sub> at recruitment					
	E	MR	aHR	HR <sub>1year</sub>	HR <sub>2year</sub>	E	MR	aHR	HR <sub>1year</sub>	HR <sub>2year</sub>	 Е	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>
Total															
≤1.00	590	10.9	Ref	Ref	Ref	270	7.9	Ref	Ref	Ref	320	16.3	Ref	Ref	Ret
1.01-2.00	232	17.1	1.25	1.23	1.17	85	13.4	1.43	1.40	1.31	147	20.3	1.19	1.13	1.09
2.01-3.00	86	20.4	1.32	1.24	1.19	29	16.0	1.38	1.34	1.35	57	23.7	1.35	1.18	1.10
3.01-10.0	149	29.4	1.83	1.76	1.72	54	24.8	2.22	2.07	2.01	95	33.0	1.75	1.59	1.55
>10.0	66	48.9	2.69	2.28	1.96	20	30.6	1.85	1.59	1.57	46	65.9	3.25	2.64	2.16
P-trend			<.001	<.001	<.001			<.001	<.001	<.001			<.001	<.001	<.001
Men						2									
≤1.00	302	18.5	Ref	Ref	Ref	169	23.6	Ref	Ref	Ref	133	14.5	Ref	Ref	Ret
1.01-2.00	144	26.6	1.36	1.36	1.32	95	32.6	1.40	1.38	1.34	49	19.7	1.31	1.34	1.31
2.01-3.00	59	34.7	1.45	1.31	1.19	40	40.4	1.54	1.37	1.16	19	26.7	1.29	1.22	1.26
3.01-10.0	111	52.7	2.17	2.10	2.00	77	64.5	2.26	2.24	2.12	34	37.3	1.98	1.80	1.70
>10.0	50	82.9	3.13	2.66	2.34	38	114.1	4.07	3.42	2.79	13	46.1	1.58	1.40	1.56
P-trend			<.001	<.001	<.001			<.001	<.001	<.001			0.002	0.009	0.015
Women										O,					
≤1.00	288	7.7	Ref	Ref	Ref	137	5.5	Ref	Ref	Ref	151	12.1	Ref	Ref	Re
1.01-2.00	88	10.8	1.13	1.08	0.99	36	9.4	1.60	1.48	1.31	52	12.1	0.86	0.86	0.81
2.01-3.00	27	10.7	1.16	1.17	1.2	10	9.1	1.48	1.50	1.47	17	12.0	0.96	0.98	1.03
3.01-10.0	38	12.9	1.31	1.24	1.29	20	15.8	2.58	2.48	2.57	18	10.7	0.75	0.71	0.74
>10.0	15	20.4	1.89	1.61	1.28	7	18.9	2.16	1.75	1.42	8	21.9	1.66	1.47	1.17
P-trend			0.019	0.074	0.161			<.001	0.001	0.002			0.856	0.635	0.538

E: Number of death, MR: Mortality rate (10 000 person year), Ref: Reference

aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise

HR<sub>1year</sub>: aHR after exclude subjects who died within 1 yr f/u time

 HR<sub>2year</sub>: aHR after exclude subjects who died within 2 yr f/u time

Total

≤1.00

1.01-2.00

2.01-3.00

3.01-10.0

>10.0

Men

≤1.00

1.01-2.00

2.01-3.00

3.01-10.0

>10.0

*P*-trend

Women

1.01-2.00

2.01-3.00

3.01-10.0

>10.0

*P*-trend

≤1.00

*P*-trend

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Cardiovascular disease mortality					Healthy subjects at recruitment					Subjects with NCD <sub>history</sub> at recruitment				
Е	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>	E	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>	E	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>
67	3.1	Ref	Ref	Ref	58	1.7	Ref	Ref	Ref	109	5.5	Ref	Ref	Ref
79	5.8	1.35	1.37	1.23	18	2.8	1.19	1.15	0.94	64	8.4	1.42	1.46	1.36
42	10.0	2.06	2.05	2.02	6	3.3	1.47	1.54	1.46	36	15.0	2.28	2.25	2.26
39	7.7	1.45	1.38	1.44	8	3.7	1.44	1.50	1.70	31	1.08	1.48	1.37	1.40
13	9.6	1.81	1.76	1.59	3	4.6	2.02	2.10	1.58	10	14.3	1.85	1.74	1.68
		0.001	0.002	0.004			0.130	0.100	0.162			0.001	0.006	0.009
					20									
89	5.5	Ref	Ref	Ref	25	2.7	Ref	Ref	Ref	64	8.9	Ref	Ref	Ref
45	8.3	1.33	1.32	1.25	12	4.8	1.30	1.22	1.22	33	11.3	1.31	1.33	1.33
30	17.6	2.70	2.67	2.53	3	4.2	1.31	1.37	1.37	27	27.3	3.05	2.99	2.99
24	11.4	1.43	1.36	1.46	6	6.6	1.70	1.79	1.79	18	15.1	1.42	1.21	1.21
8	13.0	1.90	2.02	1.70	3	10.6	3.42	3.61	3.61	5	15.0	1.59	1.62	1.62
		0.002	0.003	0.009			0.053	0.038	0.062			0.015	0.027	0.047
78	2.1	Ref	Ref	Ref	33	1.3	Ref	Ref	Ref	45	6.3	Ref	Ref	Ref
34	4.2	1.41	1.46	1.25	6	1.6	1.09	1.13	0.62	28	9.6	1.60	1.66	1.58

Table 3. The association between serum hsCRP level and cardiovascular disease mortality by gender and non-communicable disease history (NCD <sub>history</sub> ) at recruit	itment
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E: Number of death, MR: Mortality rate (10 000 person year), Ref: Reference

1.26

1.51

1.72

0.092

1.30

1.45

1.35

0.177

1.44

1.44

1.45

0.168

aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise

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0

2.7

1.6

1.65

1.06

0.940

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1.70

1.07

0.998

1.86

1.14

0.922

9

13

5

9.1

10.9

15.0

1.17

1.75

2.51

0.035

1.20

1.64

1.91

0.092

1.39

1.65

2.07

0.078

HR<sub>1vear</sub>: aHR after exclude subjects who died within 1 yr f/u time

167

79

42

39

13

89

45

30

24

78

34

12

15

5

4.8

5.1

6.8

HR<sub>2vear</sub>: aHR after exclude subjects who died within 2 yr f/u time

## 

# **DISCUSSION**

This study suggests that the risk of all-cause mortality was associated with elevated hsCRP levels with a dose-response manner in both gender among Asian who have reported low hsCRP levels compared to other races, and was not influenced by NCD<sub>*history*</sub>. Otherwise, the association was influenced by gender and NCD<sub>*history*</sub> although a dose-response association of hsCRP with the risk of cancer- and CVD-mortality was also observed in this population. The level of hsCRP was not associated with the risk of cancer- mortality among women with NCD<sub>*history*</sub>. The risk effect of high hsCRP level on CVD mortality was predominantly observed in men with NCD<sub>*history*</sub>.

Several large cohorts [10-12, 14] have suggested that serum hsCRP levels may differ according to ethnic background, with the highest concentrations seen in African Americans, followed by Hispanic, White, Chinese and Japanese individuals. Although the reason for this ethnic difference is not clearly resolved, genetic diversity[27], the relatively low BMI in Asian populations and ethnic differences in diet and lifestyle[28] have been suggested. Although the extent to which these findings adopt to Asian populations has been unclear, several recent studies [11, 16] conducted in Asia reported a positive association of *hs*CRP with mortality risk. In this population, the hsCRP level was associated with the risk of all-cause mortality in a dose-dependent manner, even though the level of hsCRP was lower than that in the western population. A meta-analysis[29] and large cohort studies[3-6] supported the robustness of the association regardless of adjusted confounders, the cut-off point of CRP level and exclusion deaths within the first 2 years of follow-up. The reason for the discrepancy in *hs*CRP levels with respect to gender is not clearly resolved, although several studies suggested different lifestyle and metabolic risk factors between men and women[30] and genetic diversity[27]. A high level of serum hsCRP in our population was positively related to the increased risk of all-cause mortality in both genders, supported by several previous studies[8, 16, 31]. Nevertheless, several studies reported no association of hsCRP levels with all-cause mortality was observed in women[7, 16]. In particular, the association was shown in postmenopausal women only, which might suggest the protective effect of endogenous female hormones on the low level of hsCRP[32]; the average hsCRP level was 0.48 and 0.68 mg/L

2 449 for premenopausal and postmenopausal women in this study. The protective effect could be supported by the

4 450 proposition that estrogen or progesterone might to some extent repress the detrimental effects of chronic

<sup>5</sup> 451 inflammation on tissue damage[33].

452 Inflammation has emerged as an important factor in the processes of NCD, including CVD[17], cancer[18],

453 type 2 diabetes[20], COPD[19, 34] and fracture[21]. In addition, medications that had taken to treat any specific

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NCD, such as rennin-angiotensin system inhibitors[35] and statins and thiazolidinedione[36], could influence the level of hsCRP. The association between hsCRP and the mortality risk was not attenuated by NCD<sub>history</sub> in either gender in this study, but the statistical significance of the association disappeared in women after sensitivity analysis (aHR<sub>2vear</sub>). A dose-response relationship between hsCRP level and all-cause mortality risk was pronounced in both genders. On the other hand, the positive association of hsCRP with the risk of all-cause mortality risk was significantly observed in only men with NCD<sub>history</sub> but not in women with NCD<sub>history</sub>. The attenuated effect of NCD<sub>history</sub> on the association between hsCRP and the risk of cancer-mortality was not observed in men, consistent with results from several studies which reported the associations among healthy men[3] or cancer patients[37, 38] only. Most studies[3, 4, 6, 7, 15, 16, 31, 39] supported that CVD mortality increased with elevated hsCRP levels, predominantly in men[4, 7, 15, 16]. Although hsCRP levels are lower in our population than in other races, the level of hsCRP was positively associated with CVD mortality in men but not in women, similar to previous studies[7, 15, 16, 31, 39]. After stratification by gender and NCD<sub>history</sub>, the association between hsCRP and the risk of CVD mortality was dominant in subjects with NCD<sub>history</sub> in this study. Although many interventional studies have been conducted recently on anti-inflammatory drugs for the prevention of cardiovascular disease, the results are controversial. According to the results of our study, elevated inflammatory markers in people with chronic disease were associated with an increased risk of CVD mortality. This suggests that CVD-mortality in people with chronic diseases might be reduced by use of anti-inflammatory medication. This study has several strengths because of the large population-based prospective study; it makes possible 1) to adjust for confounders; 2) to examine sensitivity analysis after excluding death before 1 or 2 years from recruitment; 3) to assess an advanced analysis after stratification by gender and NCD<sub>history</sub>; 4) to examine the association using various cut-off points of hsCRP considering low serum hsCRP levels in Asian populations; and 5) to evaluate the complex (i.e., nonlinear) hazard functions using restricted cubic splines on the association between continuous hsCRP levels and the risk of mortality. In particular, most previous studies excluded subjects with more than 10 mg/L hsCRP because of their relatively low sample size or reflecting acute phase reactions of severe inflammation, but we examined the effect of very high hsCRP concentration on the risk of mortality because it is possible to be more concerning for these subjects in the future. The hsCRP level of this study, in addition, was measured within 18 hours in a single institution to minimize measurement error/bias from institutional variation to avoid bias from measurement or long-term storage before analysis. 

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3		
4 5	483	Despite of those strengths, it is also several limitations. First, the use of a single measurement of <i>hs</i> CRP at
6 7	484	baseline could reflect the inaccurate status of blood hsCRP levels in the study participants and increase the
8 9	485	instability of hsCRP due to random fluctuations over time. Nevertheless, a report [40] on the long-term hsCRP
10	486	variability suggested that the hsCRP variability within individual is relatively small and that the variability
11 12	487	could not account for the association. Second, our study lacked information on medication use at recruitment
13 14	488	and during the follow-up period. Several medications related to NCDs, including statins, angiotensin-converting
15 16	489	enzyme inhibitors, fibrates, niacin, thiazolidinedione and estrogen/progestogen hormone, could influence the
17 18	490	hsCRP level[37]; however, we tried to overcome this limitation through advanced analysis after stratification by
19 20	491	NCD <sub>history</sub> . Third, because there is no available information on hormone-replacement therapy (HRT) among
21 22	492	women, which could not examine the influence of HRT on the association of hsCRP with the risk of hormone-
23 24	493	related cancer or CVD mortality among women, we could not suggest the effect of female hormones on the
25 26	494	association. In addition, further studies are needed on the effects of obesity although the inverse relationship
27 28	495	between all-cause mortality with obesity in our population was consistent to Wei's report in Asian[41]. On the
29 30	496	other hand, the inverse association of alcohol drinking with all-cause mortality couldn't interpret directly
31 32	497	because our report wasn't separated the distinguish between mild drinkers and abuse alcohol drinker, which
33 34	498	requires additional research for our population in the future.
35	499	In conclusion, the association of hsCRP level is dose-responsively increased with the risk of all-cause
36 37	500	mortality in men and women (particularly postmenopausal women), which was not influenced by the association
38 39	501	was not observed in women with NCD <sub>history</sub> . Otherwise, the association of hsCRP level with the risk of cancer-
40 41	502	and CVD-mortality could be attenuated by gender or NCD <sub>history</sub> .
42 43	503	
44 45	504	
46 47	505	Figure 1 Flow diagram of analytical sample in current study using Health Examinees cohort.
48 49	506	Figure 2 A dose-response association between serum <i>hs</i> CRP level and risk of all-cause mortality in all
50 51	507	subjects at recruitment.
52 53	508	Figure 3 A dose-response association between serum <i>hs</i> CRP level and risk of all-cause mortality in healthy
54 55	509	subjects at recruitment.
56 57	510	Figure 4 A dose-response association between serum <i>hs</i> CRP level and risk of all-cause mortality in subjects
58 59 60	511	with non-communicable disease history (NCD <sub>history</sub> ) at recruitment.

1 2		14
3 4	512	<b>Figure 5</b> Kaplan-Meier crude survival curves for cancer-mortality according to serum <i>hs</i> CRP level in all
5 6	513	subjects at recruitment.
7 8	514	Figure 6 Kaplan-Meier crude survival curves for cancer-mortality according to serum <i>hs</i> CRP level in men at
9 10	515	recruitment.
11 12	516	Figure 7 Kaplan-Meier crude survival curves for cancer-mortality according to serum <i>hs</i> CRP level in women
13 14	517	at recruitment.
15 16	518	Figure 8 Kaplan-Meier crude survival curves for cancer-mortality according to serum <i>hs</i> CRP level in healthy
17 18	519	subjects at recruitment.
19 20	520	Figure 9 Kaplan-Meier crude survival curves for cancer-mortality according to serum <i>hs</i> CRP level in subjects
21 22	521	with non-communicable disease history (NCD <sub>history</sub> ) at recruitment.
23 24	522	Figure 10 Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum hsCRP
25 26	523	level in all subjects at recruitment.
27 28	524	Figure 11 Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum <i>hs</i> CRP
29 30	525	level in men at recruitment.
31 32	526	Figure 12 Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum <i>hs</i> CRP
33 34	527	level in women at recruitment.
34 35 36	528	Figure 13 Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum <i>hs</i> CRP
37	529	level in healthy subjects at recruitment.
38 39	530	Figure 14 Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum <i>hs</i> CRP
40 41	531	level in subjects with non-communicable disease history (NCD <sub>history</sub> ) at recruitment.
42 43	532	
44 45	533	
46 47	534	Contributors
48 49	535	SAL, XS and DK: designed and conducted the research, SAL and SOK: analyzed the data and performed the
50 51	536	statistical analyses; HP and JKL: managed data mining and collection; SAL: wrote the manuscript and had primary
52	537	responsibility for the final content of the manuscript; and all authors: read and approved the final manuscript.
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58 59	540	
60	541	Competing interests None declared.

1 2		15
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12 13	546	<b>Provenance and peer review</b> Not commissioned; externally peer reviewed.
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16	548	No additional data available.
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19	549	
20 21	550	
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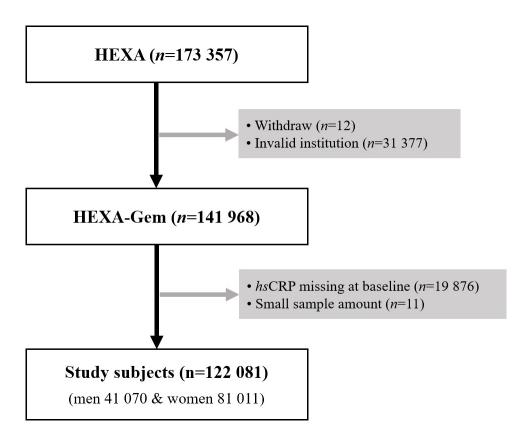
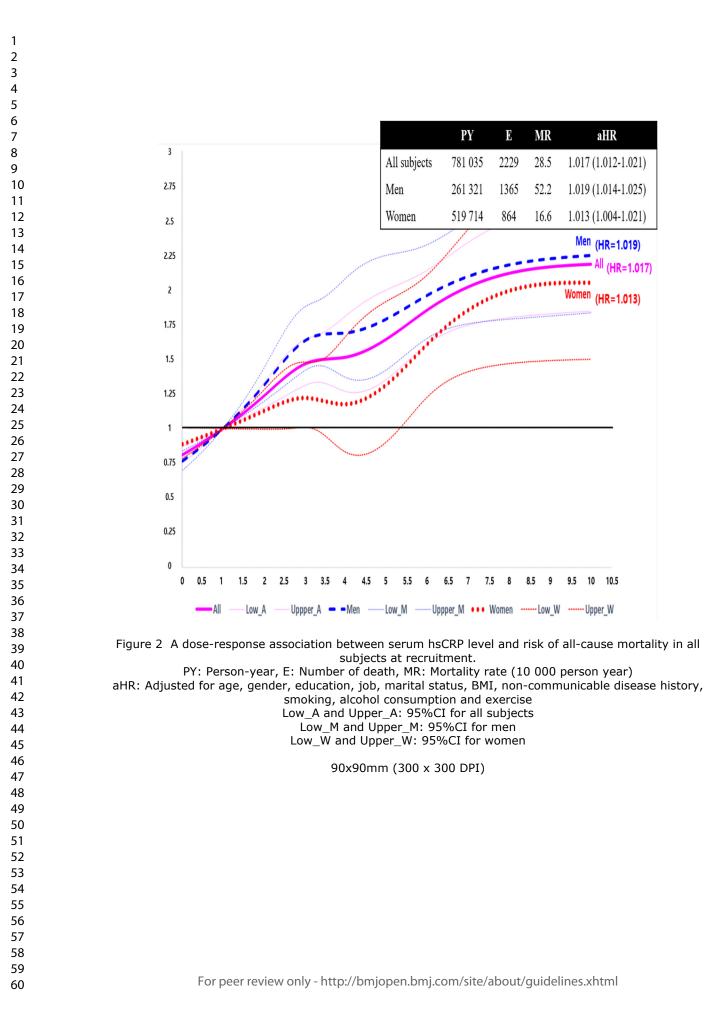
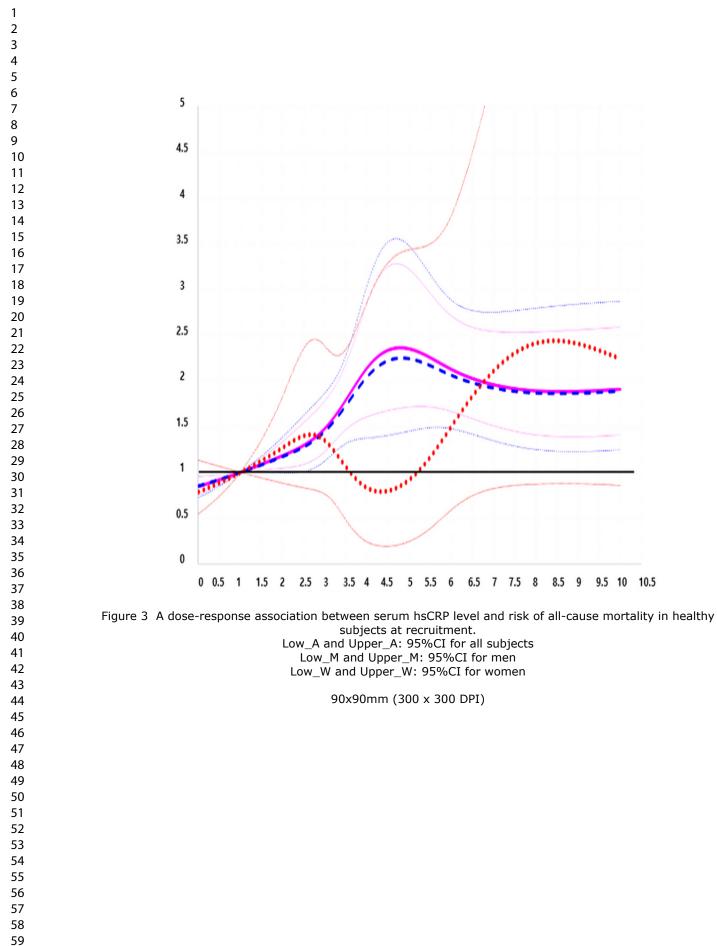


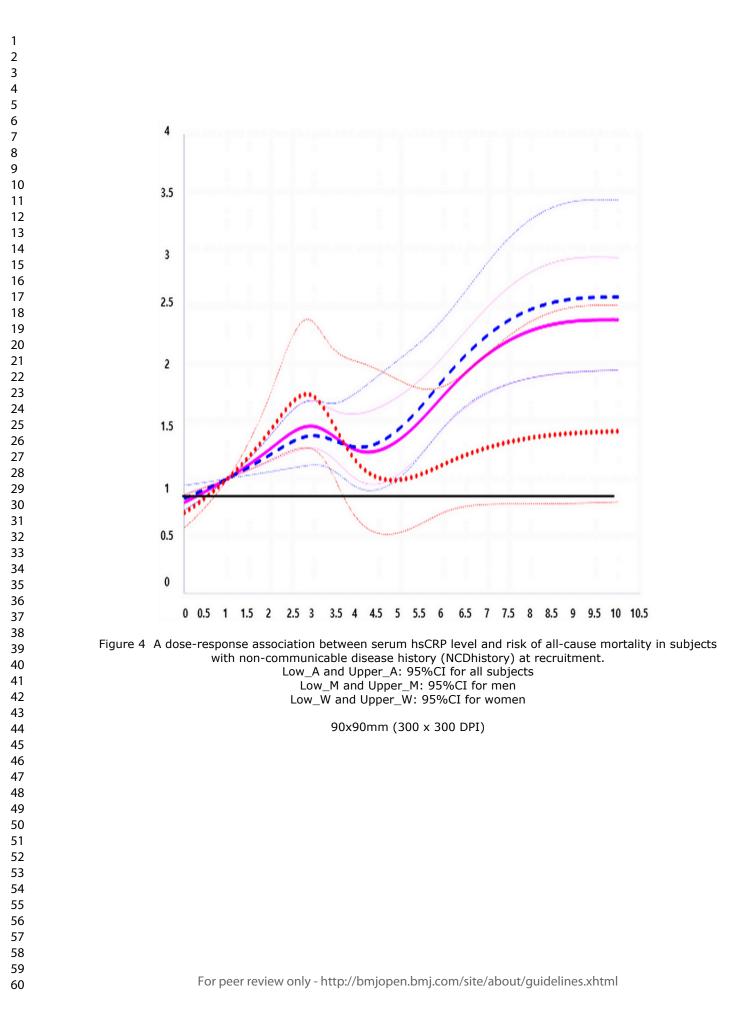
Figure 1 Flow diagram of analytical sample in current study using Health Examinees cohort. HEXA: Health Examinees, hsCRP: High sensitivity C-reactive protein

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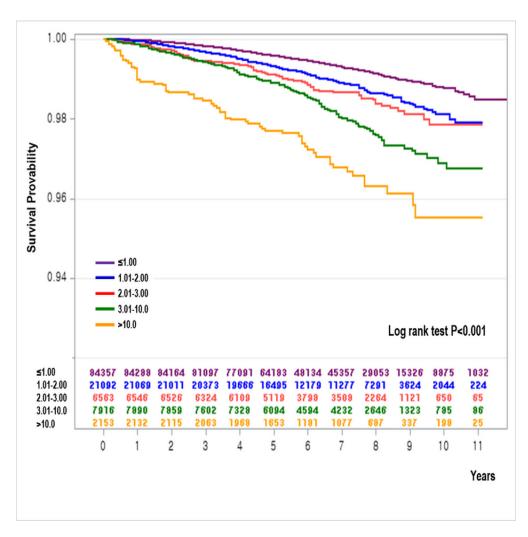
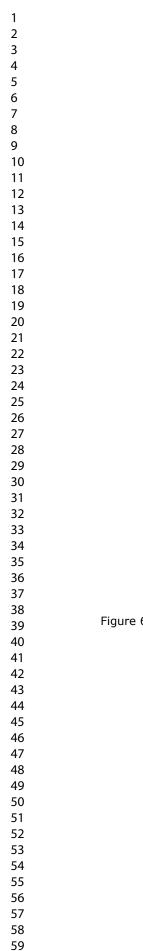


Figure 5 Kaplan-Meier crude survival curves for cancer-mortality according to serum hsCRP level in all subjects at recruitment.



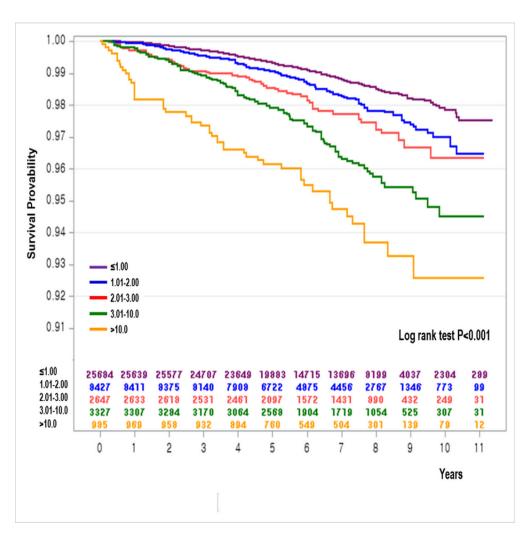
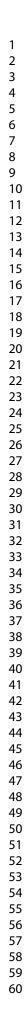


Figure 6 Kaplan-Meier crude survival curves for cancer-mortality according to serum hsCRP level in men at recruitment.



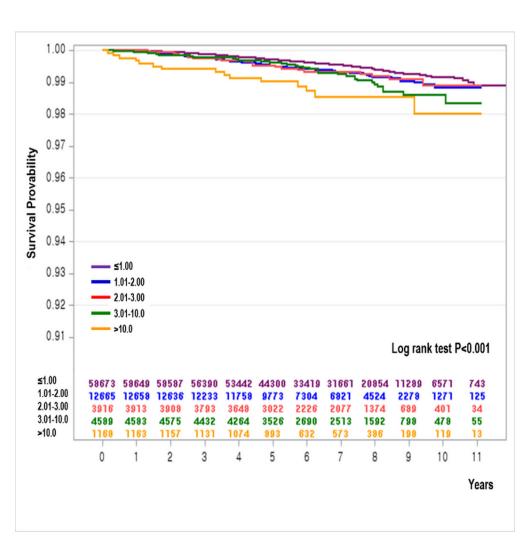


Figure 7 Kaplan-Meier crude survival curves for cancer-mortality according to serum hsCRP level in women at recruitment.

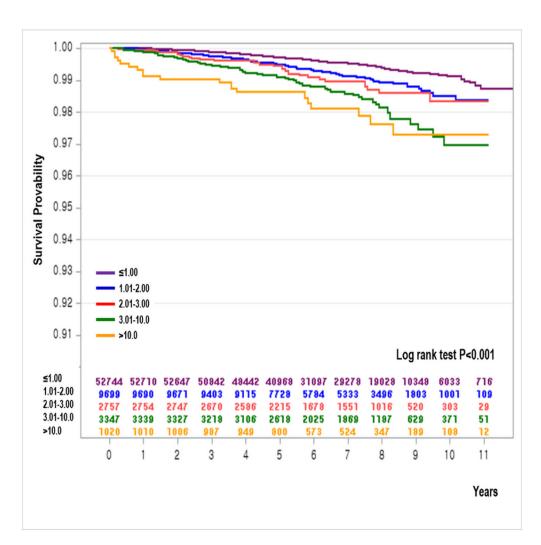
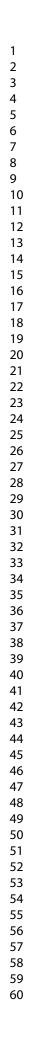


Figure 8 Kaplan-Meier crude survival curves for cancer-mortality according to serum hsCRP level in healthy subjects at recruitment.



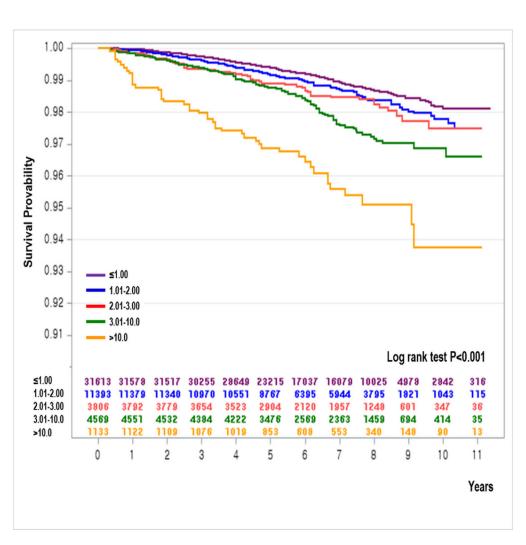


Figure 9 Kaplan-Meier crude survival curves for cancer-mortality according to serum hsCRP level in subjects with non-communicable disease history (NCDhistory) at recruitment.

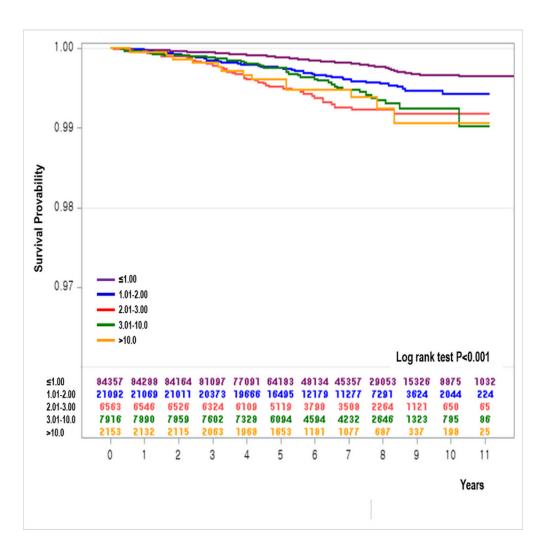


Figure 10 Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum hsCRP level in all subjects at recruitment.

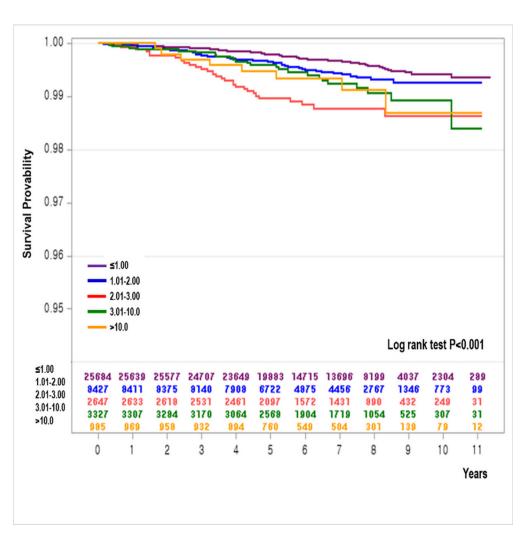


Figure 11 Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum hsCRP level in men at recruitment.

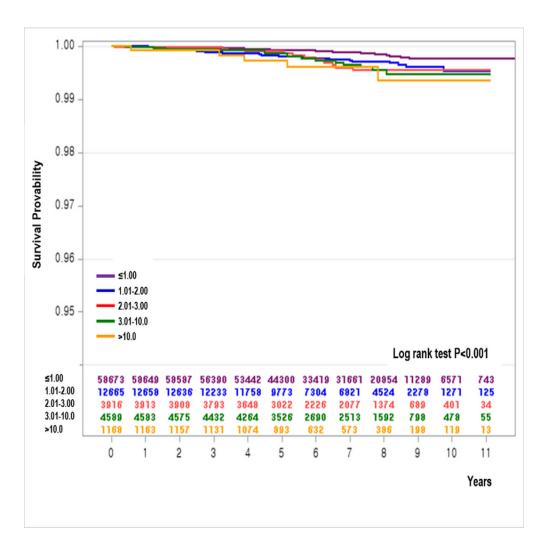
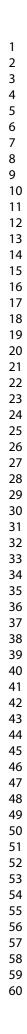


Figure 12 Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum hsCRP level in women at recruitment.



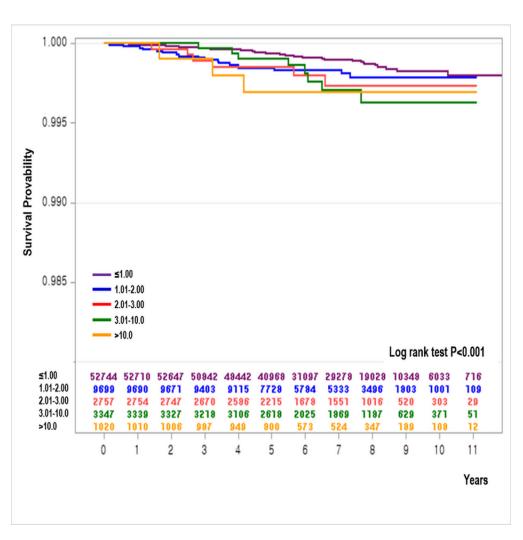


Figure 13 Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum hsCRP level in healthy subjects at recruitment.

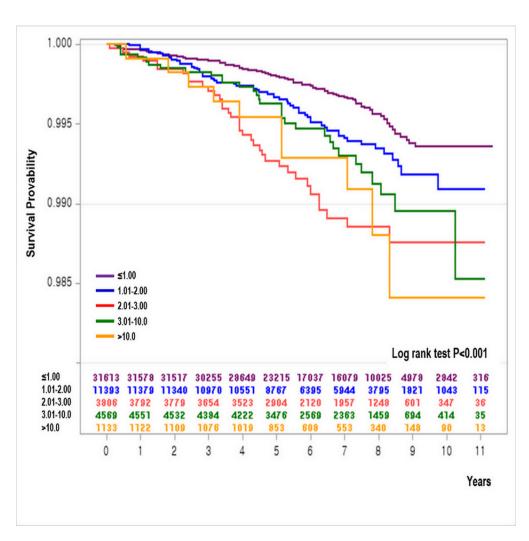


Figure 14 Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum hsCRP level in subjects with non-communicable disease history (NCDhistory) at recruitment.

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	PY	Е	MR	aHR	HR <sub>1year</sub>	HR <sub>2year</sub>
All subjects					•	-
Continuous	781 035	2229	28.5	1.017 (1.012-1.021)	1.015 (1.010-1.020)	1.014 (1.009-1.019
≤1.00	539 271	1153	21.4	Reference	Reference	Reference
1.01-1.50	90 911	308	33.9	1.26 (1.10-1.45)	1.27 (1.10-1.45)	1.21 (1.04-1.40)
1.51-2.00	44 615	163	36.5	1.28 (1.08-1.53)	1.28 (1.07-1.53)	1.26 (1.04-1.52)
2.01-2.50	25 139	117	46.5	1.53 (1.25-1.89)	1.51 (1.22-1.87)	1.49 (1.19-1.87)
2.51-3.00	16 996	72	42.4	1.39 (1.08-1.80)	1.31 (1.00-1.72)	1.23 (0.92-1.65)
3.01-4.00	19 667	103	52.4	1.61 (1.29-2.01)	1.62 (1.29-2.03)	1.64 (1.30-2.08)
4.01-6.00	17 933	102	56.9	1.84 (1.48-2.28)	1.77 (1.41-2.21)	1.70 (1.34-2.16)
6.01-10.00	13 019	88	67.6	2.02 (1.59-2.56)	1.96 (1.54-2.50)	1.93 (1.49-2.51)
>10.0	13 484	123	91.2	2.59 (2.12-3.16)	2.41 (1.95-2.97)	2.26 (1.80-2.84)
P-trend				<.001	<.001	<.001
Men						
Continuous	261 321	1365	52.2	1.019 (1.014-1.025)	1.017 (1.011-1.023)	1.017 (1.010-1.023
≤1.00	163 068	638	39.1	Reference	Reference	Reference
1.01-1.50	36 094	190	52.6	1.27 (1.07-1.51)	1.28 (1.07-1.53)	1.22 (1.01-1.47)
1.51-2.00	17 946	103	57.4	1.34 (1.07-1.67)	1.34 (1.07-1.68)	1.35 (1.06-1.72)
2.01-2.50	10 059	77	76.5	1.56 (1.20-2.03)	1.53 (1.16-2.00)	1.47 (1.10-1.96)
2.51-3.00	6959	54	77.6	1.71 (1.27-2.29)	1.57 (1.15-2.15)	1.46 (1.04-2.05)
3.01-4.00	8177	77	94.2	1.88 (1.45-2.43)	1.94 (1.50-2.52)	1.92 (1.46-2.54)
4.01-6.00	7425	75	101.0	2.05 (1.59-2.63)	1.95 (1.49-2.53)	1.91 (1.44-2.52)
6.01-10.00	5456	59	108.1	2.03 (1.52-2.73)	1.96 (1.44-2.66)	1.85 (1.33-2.58)
>10.0	6137	92	149.9	2.84 (2.25-3.58)	2.66 (2.08-3.39)	2.58 (1.99-3.35)
P-trend				<.001	<.001	<.001
Women						
Continuous	519 714	864	16.6	1.013 (1.004-1.021)	1.011(1.002-1.021)	1.010 (0.999-1.02)
≤1.00	376 203	515	13.7	Reference	Reference	Reference
1.01-1.50	54 817	118	21.5	1.28 (1.03-1.59)	1.27 (1.02-1.58)	1.23 (0.97-1.56)
1.51-2.00	26 669	60	22.5	1.23 (0.92-1.64)	1.21 (0.90-1.63)	1.14 (0.83-1.56)
2.01-2.50	15 080	40	26.5	1.52 (1.09-2.14)	1.52 (1.08-2.15)	1.56 (1.09-2.24)
2.51-3.00	10 037	18	17.9	0.84 (0.49-1.44)	0.87 (0.51-1.48)	0.83 (0.46-1.47)
3.01-4.00	11 490	26	22.6	1.16 (0.75-1.81)	1.09 (0.68-1.72)	1.21 (0.76-1.93)
4.01-6.00	10 508	27	25.7	1.48 (0.99-2.22)	1.47 (0.97-2.22)	1.36 (0.86-2.14)
6.01-10.00	7563	29	38.3	2.00 (1.34-2.98)	1.98 (1.32-2.98)	2.10 (1.39-3.19)
>10.0	7347	31	42.2	2.02 (1.36-3.02)	1.84 (1.21-2.81)	1.51 (0.93-2.47)
P-trend				<.001	<.001	0.001
Premenopause						
≤1.00	141 286	96	6.8			
1.01-2.00	20 500	20	9.8	1.52 (0.92-2.52)	1.49 (0.89-2.50)	1.57 (0.90-2.73)
2.01-3.00	5835	6	10.3	1.76 (0.77-4.06)	1.83 (0.79-4.22)	1.42 (0.52-3.93)
3.01-10.0	6886	6	8.7	1.51 (0.66-3.50)	1.31 (0.53-3.25)	1.21 (0.44-3.36)
>10.0	1759	4	22.7	2.57 (0.81-8.14)	2.63 (0.83-8.37)	2.09 (0.51-8.58)
<i>P</i> -trend		·		0.020	0.036	0.150
Postmenopause						
≤1.00	192 164	366	19.0			
1.01-2.00	52 897	145	27.4	1.26 (1.03-1.55)	1.25 (1.02-1.54)	1.18 (0.95-1.48)
2.01-3.00	16 943	44	26.0	1.11 (0.80-1.56)	1.12 (0.80-1.57)	1.19 (0.83-1.68)
3.01-10.0	19 687	67	34.0	1.49 (1.13-1.97)	1.47 (1.10-1.95)	1.52 (1.13-2.05)
>10.0	4828	27	55.9	2.09 (1.37-3.21)	1.88 (1.19-2.96)	1.56 (0.92-2.63)
<i>P</i> -trend	-020	21	55.7	<0.001	0.001	0.003

Supplement 1 The association of ser m hsCRP level with the risk of all-cause mortality

PY: Person-year, E: Number of death, MR: Mortality rate (10,000 person year)

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- aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise
  - $HR_{1year}$ : aHR after exclude subjects who died within 1 yr f/u time
  - HR<sub>2year</sub>: aHR after exclude subjects who died within 2 yr f/u time

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	Healthy subjects at recruitment				Sub	Subjects with NCD <sub>history</sub> at recruitment				
	E	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>	E	MR	aHR	$HR_{1year}$	HR <sub>2yea</sub>
All										
$\leq 1.00$	517	15.1	Ref	Ref	Ref	636	32.3	Ref	Ref	Re
1.01-2.00	145	22.9	1.20	1.19	1.16	326	45.1	1.20	1.19	1.1
2.01-3.00	53	29.3	1.38	1.37	1.32	136	56.6	1.51	1.46	1.4
3.01-10.0	102	46.8	2.22	2.15	2.15	191	66.3	1.62	1.60	1.5
>10.0	40	61.3	2.38	2.23	2.27	83	118.9	2.74	2.54	2.2
P-trend			<.001	<.001	<.001			<.001	<.001	<.00
Men										
$\leq 1.00$	270	29.5	Ref	Ref	Ref	368	51.4	Ref	Ref	Re
1.01-2.00	89	35.8	1.11	1.11	1.13	204	70.0	1.40	1.41	1.3
2.01-3.00	33	46.3	1.22	1.17	1.15	98	99.0	1.82	1.73	1.6
3.01-10.0	70	76.8	2.14	2.08	2.03	141	118.1	1.92	1.90	1.8
>10.0	31	110.0	2.60	2.49	2.73	61	183.1	3.05	2.83	2.5
P-trend			<.001	<.001	<.001			<.001	<.001	<.00
Women										
$\leq 1.00$	247	9.8	Ref	Ref	Ref	268	21.4	Ref	Ref	Re
1.01-2.00	56	14.6	1.35	1.32	1.20	122	28.3	1.19	1.20	1.1
2.01-3.00	20	18.2	1.61	1.66	1.60	38	26.9	1.06	1.06	1.1
3.01-10.0	32	25.2	2.31	2.23	2.37	50	29.7	1.16	1.14	1.1
>10.0	9	24.3	1.69	1.49	1.12	22	60.3	2.15	1.99	1.6
P-trend			<.001	<.001	0.001			0.018	0.043	0.08

**Supplement 2.** The association between serum *hs*CRP level and all-cause mortality by gender and noncommunicable disease history (NCD<sub>*history*</sub>) at recruitment

E: Number of death, MR: Mortality rate (10 000 person year)

aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise

 $HR_{1year}\!\!:aHR$  after exclude subjects who died within 1 yr f/u time

HR<sub>2year</sub>: aHR after exclude subjects who died within 2 yr f/u time

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Section/Topic	Item #	Recommendation	Reported on page
Title and abstract	1	(a) Indicate the study's design with a commonly used term in the title or the abstract	1
		(b) Provide in the abstract an informative and balanced summary of what was done and what was found	1-2
Introduction			
Background/rationale	2	Explain the scientific background and rationale for the investigation being reported	3
Objectives	3	State specific objectives, including any pre-specified hypotheses	3
Methods			
Study design	4	Present key elements of study design early in the paper	4
Setting	5	Describe the setting, locations, and relevant dates, including periods of recruitment, exposure, follow-up, and data collection	4-5
Participants	6	<ul> <li>(a) Cohort study—Give the eligibility criteria, and the sources and methods of selection of participants. Describe methods of follow-up</li> <li>Case-control study—Give the eligibility criteria, and the sources and methods of case ascertainment and control selection. Give the rationale for the choice of cases and controls</li> <li>Cross-sectional study—Give the eligibility criteria, and the sources and methods of selection of participants</li> </ul>	4
		(b) Cohort study—For matched studies, give matching criteria and number of exposed and unexposed Case-control study—For matched studies, give matching criteria and the number of controls per case	Not Applicable
Variables	7	Clearly define all outcomes, exposures, predictors, potential confounders, and effect modifiers. Give diagnostic criteria, if applicable	5
Data sources/ measurement	8*	For each variable of interest, give sources of data and details of methods of assessment (measurement). Describe comparability of assessment methods if there is more than one group	5
Bias	9	Describe any efforts to address potential sources of bias	6
Study size	10	Explain how the study size was arrived at	4
Quantitative variables	11	Explain how quantitative variables were handled in the analyses. If applicable, describe which groupings were chosen and why	5-6
Statistical methods	12	(a) Describe all statistical methods, including those used to control for confounding	5-6
		(b) Describe any methods used to examine subgroups and interactions	5-6
		(c) Explain how missing data were addressed	4
		(d) Cohort study—If applicable, explain how loss to follow-up was addressed Case-control study—If applicable, explain how matching of cases and controls was addressed	4

		Cross-sectional study—If applicable, describe analytical methods taking account of sampling strategy	
		(e) Describe any sensitivity analyses	6
Results	•		
Participants	13*	3* (a) Report numbers of individuals at each stage of study—eg numbers potentially eligible, examined for eligibility, confirmed eligible, included in the study, completing follow-up, and analysed	
		(b) Give reasons for non-participation at each stage	4
		(c) Consider use of a flow diagram	4
Descriptive data	14*	(a) Give characteristics of study participants (eg demographic, clinical, social) and information on exposures and potential confounders	6-7
		(b) Indicate number of participants with missing data for each variable of interest	4
		(c) Cohort study—Summarise follow-up time (eg, average and total amount)	6
Outcome data	15*	Cohort study—Report numbers of outcome events or summary measures over time	6-10
		Case-control study—Report numbers in each exposure category, or summary measures of exposure	Not Applicable
		Cross-sectional study—Report numbers of outcome events or summary measures	Not Applicable
Main results	16	( <i>a</i> ) Give unadjusted estimates and, if applicable, confounder-adjusted estimates and their precision (eg, 95% confidence interval). Make clear which confounders were adjusted for and why they were included	6-10
		(b) Report category boundaries when continuous variables were categorized	5
		(c) If relevant, consider translating estimates of relative risk into absolute risk for a meaningful time period	Not Applicable
Other analyses	17	Report other analyses done—eg analyses of subgroups and interactions, and sensitivity analyses	8
Discussion		·	
Key results	18	Summarise key results with reference to study objectives	11
Limitations	19	Discuss limitations of the study, taking into account sources of potential bias or imprecision. Discuss both direction and magnitude of any potential bias	12-13
Interpretation	20	Give a cautious overall interpretation of results considering objectives, limitations, multiplicity of analyses, results from similar studies, and other relevant evidence	11-13
Generalisability	21	Discuss the generalisability (external validity) of the study results	11-12
Other information		•	
Funding	22	Give the source of funding and the role of the funders for the present study and, if applicable, for the original study on which the present article is based	14

\*Give information separately for cases and controls in case-control studies and, if applicable, for exposed and unexposed groups in cohort and cross-sectional studies. **Note:** An Explanation and Elaboration article discusses each checklist item and gives methodological background and published examples of transparent reporting. The STROBE checklist is best used in conjunction with this article (freely available on the Web sites of PLoS Medicine at http://www.plosmedicine.org/, Annals of Internal Medicine at http://www.annals.org/, and Epidemiology at http://www.epidem.com/). Information on the STROBE Initiative is available at www.strobe-statement.org.

# **BMJ Open**

# The association of serum high sensitivity C-reactive protein with the risk of mortality in an Asian population: the Health Examinees cohort

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3 4 5	196	The association of serum high sensitivity C-reactive protein with the risk of mortality in
6	197	an Asian population: the Health Examinees cohort
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10 11	200	Sang-Ah Lee <sup>1,2</sup> *, Sung Ok Kwon <sup>1</sup> , Hyerim Park <sup>1</sup> , Xiao-Ou Shu <sup>2</sup> , Jong-Koo Lee <sup>3</sup> , Daehee Kang <sup>4</sup>
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22	208	<sup>4</sup> Department of Preventive Medicine, College of Medicine, Seoul National University, Seoul, Republic of Korea.
23 24	209	
25	210	
26 27	211	ABSTRACT
28	212	<b>Objectives</b> This study aimed to examine the association of <i>hs</i> CRP with mortality risk and the attenuated effect
29	213	of non-communicable disease history (NCD <sub>history</sub> ) on the association.
30 31 32 33 34 35	214	Design Prospective cohort study.
	215	Setting the Health Examinees (HEXA) cohort.
	216	<b>Participants</b> A total of 41 070 men and 81 011 women aged $\geq$ 40 years were involved (follow-up: 6.8 years).
	217	<b>Outcome measures</b> The data and cause of death occurring until December 31, 2015, were confirmed by death
36 37	218	statistics from the National Statistical Office. We conducted the advanced analysis after stratification by
38	219	NCD <sub>history</sub> and the sensitivity analysis after excluding death before 1 or 2 years from recruitment. Cox
39 40 41 42 43 44	220	proportional hazard and restricted cubic spline models were used to assess the association.
	221	<b>Results</b> The association between serum <i>hs</i> CRP and the risk of all-cause mortality was observed with strong
	222	linearity in both genders, which was not influenced by NCD <sub>history</sub> . Otherwise, the association of serum hsCRP
	223	with cancer-mortality risk was not observed in women with NCD <sub>history</sub> , but the association with the risk of
45 46	224	cardiovascular disease (CVD) mortality was predominantly observed in men with NCD <sub>history</sub> .
46 47	225	<b>Conclusions</b> This study suggested the dose-response association of <i>hs</i> CRP with mortality risk, including
48	226	cancer and CVD mortality, in Korean with low serum hsCRP, although the association with cancer and CVD-
49 50 51 52 53 54 55 56 57 58	227	mortality risk could be influenced by gender and NCD <sub>history</sub> .
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	229	
	230	Strengths and limitations of this study
	231	This is the large population-based prospective study.
	232	• We examined the effect of very high <i>hs</i> CRP concentration on mortality risk.
59 60	233	• The $hs$ CRP level of present study was measured within 18 hours in a single institution to minimize error/bias.

2 3		
4 5	234	• Due to random fluctuations of $hs$ CRP, using the single measurement of $hs$ CRP at baseline could reflect the
6 7	235	inaccurate status of blood hsCRP levels in the study participants and increase the instability of hsCRP.
, 8 9	236	· This study lacked information on medication use at recruitment and during the follow-up period, and
10	237	information on hormone-replacement therapy (HRT) among women.
11 12	238	
13	239	
14 15	240	*Correspondence to: Sang-Ah Lee, Ph.D.
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# 266 INTRODUCTION

High sensitivity C-reactive protein (hsCRP) is an acute-phase response protein synthesized by the liver and the most sensitive and dynamic marker of inflammation[1]. Since hsCRP has been reported as a candidate marker for generalized atherosclerosis and cardiovascular disease (CVD)[2], many studies[3-7] have investigated the role of hsCRP levels as a predictor of mortality risk. A recent meta-analysis[8] reported the predictable role of serum hsCRP on all-cause and CVD mortality in the general population. Nevertheless, it is controversial whether the predictable role of hsCRP could be applied to the risk of mortality in Asians, whose hsCRP levels are lower than those in individuals in Western countries.

Serum *hs*CRP represents a low-grade inflammation state that is generally involved in the process of aging[9]. Several large cohorts, including Study of Women's Health Across the Nation (SWAN)[10], the Women's Health Study[11] and the Dallas Heart Study[12], reported significant differences in hsCRP levels by race and gender. In two studies of multiethnic populations residing in the USA[10, 13], the median hsCRP level in East Asians was less than half the concentration in Caucasians. Even among East Asian populations, the geometric mean of hsCRP levels varied depending on ethnic background[14]. In addition, a meta-analysis[11] reported the hsCRP levels among women of various ethnic groups living in the United States (from the Women's Health Study) on the association between hsCRP and the mortality risk; the association was observed in only men supported by the results from two cohort studies [15, 16] reported in Korea. On the other hand, the increased hsCRP may be influenced by comorbidity itself because inflammation has emerged as an important factor in the progression of non-communicable diseases (NCDs), including CVD[17], cancer[18], chronic obstructive pulmonary disease (COPD)[19], type 2 diabetes[20] and fractures[21], which contribute to increased morbidity and mortality.

This study aimed to examine the association of serum hsCRP with the risk of mortality in Koreans with low serum hsCRP and to evaluate the attenuated effect of non-communicable disease history (NCD<sub>history</sub>) on the association.

#### **METHODS**

#### Study population

Details on the main objectives, rationale, study design and baseline characteristics of the Health Examinees (HEXA) study have been published elsewhere[22]. Considering the homogeneity and comparability of participants, we created a qualified dataset called HEXA-G (Health Examinees-Gem) from previously published HEXA studies[23]. In the new HEXA-G data, a total of 141 968 participants remained after the exclusion of withdrawers (n=12). In addition, 19 887 were excluded due to missing information (n=19 876) or small sample size (n=11) on any hsCRP components at the baseline survey. Ultimately, 122 081 subjects, including 41 070 men and 81 011 women, remained in the final analysis (Fig. 1). All study participants provided informed consent prior to entering the study. The Institutional Review Board of the Seoul National University Hospital, Seoul, Korea, approved it for statistical analysis (IRB No. E-1503-103-657).

#### Laboratory measurements

After at least 10 hours of overnight fasting, blood samples were obtained in the morning. Bio-specimens included fasting blood samples that were collected in a serum separator tube and two ethylenediaminetetraacetic acid (EDTA) tubes. All samples were then transported to the National Biobank of Korea and stored for future research purposes within 18 hours. hsCRP was measured using a turbidimetric immunoassay (ADVIA 1650 and ADVIA 1800; Siemens Healthineers).

#### Follow-up and ascertainment of mortality

All-cause mortality was confirmed by death statistics from the National Statistical Office, which provided the data and causes of all deaths occurring through December 31, 2015. We added the mortality data from Statistics Korea to our dataset using each participant's unique identifier. Information on death and causes of death was obtained from a record link with the national death certificate files in Korea. The main outcome of interest was all-cause mortality (defined as death from any cause), including cancers and CVD mortality. The cause of death was classified according to the International Classification of Diseases, 10th revision (ICD-10). Deaths were coded as C00-C97 for cancer and I00-I99 for CVD. 

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## 323 Baseline variables

Trained interviewers collected information on demographic, socioeconomic and lifestyle factors. Anthropometric measurements were obtained using standardized methods. Body mass index (BMI) was calculated, and all participants were defined into four classes based on the World Health Organization classification of BMI for Asian adults[24]: underweight (BMI <18.5 kg/m<sup>2</sup>), normal (18.5 BMI <23.0 kg/m<sup>2</sup>), overweight (23.0 $\leq$  BMI <25.0 kg/m<sup>2</sup>), obesity (25.0 $\leq$  BMI <29.9 kg/m<sup>2</sup>), and severe obesity (BMI  $\geq$ 30.0 kg/m<sup>2</sup>). The current study defined metabolic syndrome using the National Cholesterol Education Program Adult Treatment Panel III (NCEP-ATP III)[25], modified for the Asian guideline for waist circumference (WC ≥90 and  $\geq$ 80 cm for men and women, respectively). Nonsmokers were defined as those who had smoked less than 400 cigarettes over the course of their lifetime. Participants who had smoked were categorized into two groups: noncurrent (never/former) and current smoker. Noncurrent drinkers were defined as those who had never consumed an alcoholic drink over the course of their lifetime or those who had not consumed alcohol at recruitment, while current drinkers were defined as those who persisted in consuming alcohol. Regular exercise was classified into two groups (ves/no) as follows: "Do you currently engage in regular exercise strenuous enough to cause you to break into a sweat at least once per week?" Furthermore, considering the attenuated effect of the NCD<sub>history</sub> on the association between serum hsCRP and the risk of mortality, we performed advanced analysis after stratification by NCD<sub>history</sub>. We considered six main non-communicable diseases (hypertension, diabetes, hyperlipidemia, cancer, cardiovascular and cerebrovascular diseases, and respiratory disease) to classify healthy subjects vs. subjects with NCD<sub>history</sub>.

<sup>10</sup> 342

## 343 Statistical analysis

For the categorical analysis, we created nine categories based on the distribution of *hs*CRP levels in our population:  $\leq 1.00$  (reference group), 1.01-1.50, 1.51-2.00, 2.01-2.50, 2.51-3.00, 3.01-4.00, 4.01-6.00, 6.01-10.0, and  $\geq 10.0$  mg/L. For the advanced analysis after stratification by the NCD<sub>*history*</sub>, the *hs*CRP levels were categorized as  $\leq 1.00$ , 1.01-2.00, 2.01-3.00, 3.01-10.0, and  $\geq 10.0$  mg/L because of the reduced sample size in each subgroup. The concentrations of *hs*CRP were log-transformed for analyses because of the skewed distribution.

We calculated a follow-up time for each subject starting from the date of interview until the date of death or
 We calculated a follow-up time for each subject starting from the date of interview until the date of death or
 December 31, 2015, whichever came first. Using age as the time scale, subjects enter the risk set at the age at
 which they completed the baseline questionnaire and exit at their event/censoring age. The associations of

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3 4 5	353	hsCRP and all-cause mortality, as well as cancer and CVD mortality, were analyzed by Cox proportional hazard
5 6 7	354	models (aHR) and included adjustment for age, gender, demographic factors (education, marital status, job, BMI
7 8	355	and NCD <sub>history</sub> ), and lifestyle factors (smoking, alcohol consumption and exercise). We used Wald tests to test
9 10	356	for heterogeneity of risk between serum hsCRP level groups. The proportional hazards assumption was assessed
11 12	357	on the basis of Schoenfeld residuals, and was not violated for the variables of interest in the adjusted model for
13 14	358	either cancer-mortality or cardiovascular disease mortality (P>0.05 for all categories). In addition, we conducted
15 16	359	a sensitivity analysis to avoid latent period bias after excluding death before 1 year (aHR <sub>1year</sub> ) or 2 years
17 18	360	(aHR <sub>2year</sub> ) since recruitment. Based on the Cox proportional hazard models, we made Kaplan-Meier curves and
19 20	361	log-rank analysis after adjustment for age, gender, demographic factors (education, marital status, job, BMI and
21 22	362	NCD <sub>history</sub> ), and lifestyle factors (smoking, alcohol consumption and exercise). We employed restricted cubic
23 24	363	splines (RCSs) to evaluate the possibility of complex (i.e., nonlinear) hazard functions[26] using continuous
25 26	364	values of hsCRP (aHR <sub>continuous</sub> ). We selected five hsCRP concentration values as knots based on hsCRP concen-
27 28	365	tration percentiles, tested the linear and nonlinear associations between knots using a cubic function, and
29 30	366	presented the integrated graph smoothly. All statistical analyses were performed using SAS version 9.3 (SAS
31	367	Institute Inc., Cary, NC, USA) and RCS analysis was carried out using the SAS LGTPHCURV9 macro. Two-
32 33 34 35	368	sided <i>p</i> -values <0.05 were defined as indicating statistical significance.
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36 37	370	Patient and public involvement
38 39	371	No patients and public were involved in the design, conducting, reporting, and dissemination plans of the present
40 41	372	study.
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## **RESULTS**

384	The association of demographic and lifestyle factors with the risk of all-cause mortality is presented in Table
385	1. During the follow-up period (average 6.8 years), 1 365 men and 864 women died. The median levels of
386	hsCRP were 0.77 and 0.59 mg/L for men and women, respectively. The risk of all-cause mortality was inversely
387	associated with female gender (aHR=0.38), high educated (aHR=0.65), overweight (aHR=0.81) or obesity
388	(aHR=0.83), current alcohol consumption (aHR=0.81) and regular exercise (aHR=0.83), but was positively
389	associated with single marital status (aHR=1.23), NCD <sub>history</sub> (aHR=1.57), underweight (aHR=2.05) and current
390	smoking (aHR=1.97).
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397	associated with single market status (arite=1.2.5), ite <i>D<sub>history</sub></i> (arite=1.57), under weight (arite=2.05) and eariest smoking (aHR=1.97).
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		All	Death	All-cause m	ortality
		( <i>n</i> =122 081)	( <i>n</i> =2229)	Age,gender adjusted	adj HRª
	Age	$53.1\pm8.3$	$59.7\pm8.8$		
	Female	66.4	38.8	0.40 (0.36-0.43)	0.38 (0.33-0.44)
	Education ( $\geq 10$ year, %)	68.2	55.4	0.67 (0.60-0.75)	0.65 (0.56-0.75)
	Blue-colored worker <sup>b</sup> (%)	32.3	33.8	1.46 (1.26-1.68)	1.16 (0.99-1.35)
	Marital status (single, %)	11.0	13.3	1.35 (1.19-1.54)	1.23 (1.07-1.40)
	NCD <sub>history</sub> (yes, %)	32.4	53.6	1.51 (1.39-1.65)	1.57 (1.42-1.72)
	Hypertension	18.9	31.5	1.18 (1.08-1.30)	1.22 (1.11-1.35)
	Diabete	6.5	17.1	1.81 (1.62-2.03)	1.77 (1.57-2.00)
	Hyperlipidemia	9.2	7.6	0.73 (0.62-0.86)	0.78 (0.66-0.92)
	Cancer	3.2	8.8	2.69 (2.31-3.12)	2.66 (2.27-3.11)
	Cerebral & cardiovascular disease	3.7	10.2	1.50 (1.30-1.73)	1.43 (1.23-1.66)
	Respiratory disease	2.4	4.3	1.37 (1.12-1.68)	1.32 (1.06-1.64)
	Body mass index (%)				
	<18.5	1.8	3.7	2.14 (1.69-2.69)	2.05 (1.61-2.62)
	18.5-22.9	38.1	34.9	1.00 (ref.)	1.00 (ref.)
	23.0-24.9	27.8	26.0	0.82 (0.73-0.91)	0.81 (0.72-0.91)
	25.0-29.9	29.5	32.5	0.90 (0.81-1.00)	0.83 (0.74-0.93)
	$\geq$ 30.0	2.8	2.9	1.08 (0.83-1.39)	0.81 (0.61-1.08)
	P-trend			0.0118	<.0001
	Metabolic syndrome (yes, %)	22.0	28.4	1.13 (1.03-1.24)	1.07 (0.96-1.19)
	Current smoker (%)	11.7	22.7	2.04 (1.79-2.33)	1.97 (1.71-2.27)
	Current drinker (%)	44.0	43.8	0.86 (0.77-0.95)	0.81 (0.73-0.91)
	Regular exercise (yes, %)	53.4	49.1	0.76 (0.70-0.83)	0.83 (0.76-0.91)
413 414	NCD <sub><i>history:</i></sub> Non-communicable diseas <sup>a</sup> Adjusted for age, gender, education.	-	tus BMI and n	on-communicable diseas	e history
415	<sup>b</sup> Compared to white-colored worker	, job, maritar sta	tus, Divir and I		e mstory
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 Table 1. Baseline characteristics of participants by all-cause mortality.

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The risk of all-cause mortality was inclined with a dose-dependent pattern as increased serum hsCRP level (Ptrend<0.001, Supplement 1), regardless of gender (Ptrend<0.001 in both genders), even in the sensitivity analysis  $(P_{trend} < 0.001 \text{ for aHR}_{1\text{ver}} \text{ in both genders})$ . The increased risk of female mortality with increased hsCRP levels was observed in both premenopausal ( $P_{trend}=0.020$ ) and postmenopausal women ( $P_{trend} < 0.001$ ), although the statistical significance in premenopausal women disappeared after sensitivity analysis ( $P_{trend}$ =0.150 for aHR<sub>2year</sub>, Supplement 1). The integrated graph, based on the restricted cubic spline analyses, indicated a strong and linear association of serum hsCRP level with all-cause mortality in both genders (aHR<sub>continuous</sub>=1.019 and 1.013 in men and women, respectively, Fig. 2 (a)).

The dose-response association between hsCRP level and the risk of all-cause mortality was not influenced by NCD<sub>history</sub> (Supplement 2). After stratification by gender, however, the attenuated effect by NCD<sub>history</sub> on the association was observed only in women; the linearity of the relationship was observed in healthy women  $(P_{trend}=0.001 \text{ for aHR}_{2\text{vear}})$  but disappeared in women with NCD<sub>history</sub>, particularly after sensitivity analysis with the exclusion of a 2-year follow-up time ( $P_{trend}=0.084$  for aHR<sub>2vear</sub>). Based on the restricted cubic spline analyses, otherwise, the pattern of increase in the association was different depending on the NCD<sub>history</sub> (Fig. 2 (b), (c)). In the healthy subjects, the risk of all-cause mortality was increased with a gradual slope (strength) until 3.0 mg/L hsCRP, with a very steep slope until 4.5 mg/L and finally with a reduced and flattened slope after 4.5 mg/L (Fig. 2 (b)). On the other hand, the slope of the association fluctuated as the hsCRP level increased in the subjects with NCD<sub>history</sub>; the slope increased up to 3.0 mg/L hsCRP but decreased until 4.5 mg/L and rapidly increased after 4.5 mg/L (Fig. 2 (c)).

The association of serum hsCRP with the risk of cancer-mortality was not influenced by NCD<sub>history</sub> (Ptrend<0.001 regardless of NCDhistory) (Table 2 and Fig. 3 (a-e)). Otherwise, after stratification by gender, the association was not observed in women with NCD<sub>history</sub> ( $P_{trend} = 0.856$ ); however, the association was not influenced by NCD<sub>history</sub> in men (P<sub>trend</sub><0.001 and 0.002 for aHR in both healthy and NCD<sub>history</sub>) (Table 2). Although the risk of CVD mortality was linearly associated with increasing hsCRP levels, the association was dominant in men (Ptrend=0.002) and in subjects with NCDhistory (Ptrend=0.001, Table 3) after stratified by gender and NCD<sub>history</sub>, respectively (Fig. 4 (a-e)). After stratification by gender and NCD<sub>history</sub>, otherwise, the association only appeared in individuals of both genders with NCD<sub>history</sub> (P<sub>trend</sub>=0.015 and 0.035 in men and women with NCD<sub>history</sub>, respectively); no association between hsCRP level and CVD mortality risk was found in either healthy men or women.

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		C	ancer-mo	ortality		Н	Subjects with NCD <sub>history</sub> at recruitment								
	Е	MR	aHR	HR <sub>1year</sub>	HR <sub>2year</sub>	E	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>	Е	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>
Total															
≤1.00	590	10.9	Ref	Ref	Ref	270	7.9	Ref	Ref	Ref	320	16.3	Ref	Ref	Ref
1.01-2.00	232	17.1	1.25	1.23	1.17	85	13.4	1.43	1.40	1.31	147	20.3	1.19	1.13	1.09
2.01-3.00	86	20.4	1.32	1.24	1.19	29	16.0	1.38	1.34	1.35	57	23.7	1.35	1.18	1.10
3.01-10.0	149	29.4	1.83	1.76	1.72	54	24.8	2.22	2.07	2.01	95	33.0	1.75	1.59	1.55
>10.0	66	48.9	2.69	2.28	1.96	20	30.6	1.85	1.59	1.57	46	65.9	3.25	2.64	2.16
P-trend			<.001	<.001	<.001			<.001	<.001	<.001			<.001	<.001	<.001
Men						RL									
≤1.00	302	18.5	Ref	Ref	Ref	169	23.6	Ref	Ref	Ref	133	14.5	Ref	Ref	Ref
1.01-2.00	144	26.6	1.36	1.36	1.32	95	32.6	1.40	1.38	1.34	49	19.7	1.31	1.34	1.31
2.01-3.00	59	34.7	1.45	1.31	1.19	40	40.4	1.54	1.37	1.16	19	26.7	1.29	1.22	1.26
3.01-10.0	111	52.7	2.17	2.10	2.00	77	64.5	2.26	2.24	2.12	34	37.3	1.98	1.80	1.70
>10.0	50	82.9	3.13	2.66	2.34	38	114.1	4.07	3.42	2.79	13	46.1	1.58	1.40	1.56
P-trend			<.001	<.001	<.001			<.001	<.001	<.001			0.002	0.009	0.015
Women															
≤1.00	288	7.7	Ref	Ref	Ref	137	5.5	Ref	Ref	Ref	151	12.1	Ref	Ref	Ref
1.01-2.00	88	10.8	1.13	1.08	0.99	36	9.4	1.60	1.48	1.31	52	12.1	0.86	0.86	0.81
2.01-3.00	27	10.7	1.16	1.17	1.2	10	9.1	1.48	1.50	1.47	17	12.0	0.96	0.98	1.03
3.01-10.0	38	12.9	1.31	1.24	1.29	20	15.8	2.58	2.48	2.57	18	10.7	0.75	0.71	0.74
>10.0	15	20.4	1.89	1.61	1.28	7	18.9	2.16	1.75	1.42	8	21.9	1.66	1.47	1.17
P-trend			0.019	0.074	0.161			<.001	0.001	0.002			0.856	0.635	0.538

E: Number of death, MR: Mortality rate (10 000 person year), Ref: Reference

aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise

HR<sub>1vear</sub>: aHR after exclude subjects who died within 1 yr f/u time

 $HR_{2year}$ : aHR after exclude subjects who died within 2 yr f/u time

	Car	rdiovascu	ılar disea	se mortali	ty	Н	Healthy subjects at recruitment						Subjects with NCD <sub>history</sub> at recruitment					
	E	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>	E	MR	aHR	HR <sub>1year</sub>	HR <sub>2year</sub>	E	MR	aHR	$HR_{1year}$	HR <sub>2yea</sub>			
Total																		
≤1.00	167	3.1	Ref	Ref	Ref	58	1.7	Ref	Ref	Ref	109	5.5	Ref	Ref	Ret			
1.01-2.00	79	5.8	1.35	1.37	1.23	18	2.8	1.19	1.15	0.94	64	8.4	1.42	1.46	1.36			
2.01-3.00	42	10.0	2.06	2.05	2.02	6	3.3	1.47	1.54	1.46	36	15.0	2.28	2.25	2.26			
3.01-10.0	39	7.7	1.45	1.38	1.44	8	3.7	1.44	1.50	1.70	31	1.08	1.48	1.37	1.40			
>10.0	13	9.6	1.81	1.76	1.59	3	4.6	2.02	2.10	1.58	10	14.3	1.85	1.74	1.68			
P-trend			0.001	0.002	0.004			0.130	0.100	0.162			0.001	0.006	0.009			
Men						Co												
≤1.00	89	5.5	Ref	Ref	Ref	25	2.7	Ref	Ref	Ref	64	8.9	Ref	Ref	Ret			
1.01-2.00	45	8.3	1.33	1.32	1.25	12	4.8	1.30	1.22	1.22	33	11.3	1.31	1.33	1.33			
2.01-3.00	30	17.6	2.70	2.67	2.53	3	4.2	1.31	1.37	1.37	27	27.3	3.05	2.99	2.99			
3.01-10.0	24	11.4	1.43	1.36	1.46	6	6.6	1.70	1.79	1.79	18	15.1	1.42	1.21	1.21			
>10.0	8	13.0	1.90	2.02	1.70	3	10.6	3.42	3.61	3.61	5	15.0	1.59	1.62	1.62			
P-trend			0.002	0.003	0.009			0.053	0.038	0.062			0.015	0.027	0.047			
Women																		
≤1.00	78	2.1	Ref	Ref	Ref	33	1.3	Ref	Ref	Ref	45	6.3	Ref	Ref	Re			
1.01-2.00	34	4.2	1.41	1.46	1.25	6	1.6	1.09	1.13	0.62	28	9.6	1.60	1.66	1.58			
2.01-3.00	12	4.8	1.26	1.30	1.44	3	2.7	1.65	1.70	1.86	9	9.1	1.17	1.20	1.39			
3.01-10.0	15	5.1	1.51	1.45	1.44	2	1.6	1.06	1.07	1.14	13	10.9	1.75	1.64	1.65			
>10.0	5	6.8	1.72	1.35	1.45	0	-	-	-	-	5	15.0	2.51	1.91	2.07			
P-trend			0.092	0.177	0.168			0.940	0.998	0.922			0.035	0.092	0.078			

Table 3. The association between serum hsCRP level and cardiovascular disease mortality by gender and non-communicable disease history (NCD <sub>history</sub> ) at	recruitment.
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E: Number of death, MR: Mortality rate (10 000 person year), Ref: Reference aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise HR<sub>1year</sub>: aHR after exclude subjects who died within 1 yr f/u time HR<sub>2year</sub>: aHR after exclude subjects who died within 2 yr f/u time

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## **DISCUSSION**

This study suggests that the risk of all-cause mortality was associated with elevated hsCRP levels with a dose-response manner in both gender among Asian who have reported low hsCRP levels compared to other races, and was not influenced by NCD<sub>history</sub>. Otherwise, the association was influenced by gender and NCD<sub>history</sub> although a dose-response association of hsCRP with the risk of cancer- and CVD-mortality was also observed in this population. The level of hsCRP was not associated with the risk of cancer- mortality among women with NCD<sub>history</sub>. The risk effect of high hsCRP level on CVD mortality was predominantly observed in men with NCD<sub>history</sub>.

Several large cohorts [10-12, 14] have suggested that serum hsCRP levels may differ according to ethnic background, with the highest concentrations seen in African Americans, followed by Hispanic, White, Chinese and Japanese individuals. Although the reason for this ethnic difference is not clearly resolved, genetic diversity[27], the relatively low BMI in Asian populations and ethnic differences in diet and lifestyle[28] have been suggested. Although the extent to which these findings adopt to Asian populations has been unclear, several recent studies [11, 16] conducted in Asia reported a positive association of *hs*CRP with mortality risk. In this population, the hsCRP level was associated with the risk of all-cause mortality in a dose-dependent manner, even though the level of hsCRP was lower than that in the western population. A meta-analysis[29] and large cohort studies[3-6] supported the robustness of the association regardless of adjusted confounders, the cut-off point of CRP level and exclusion deaths within the first 2 years of follow-up. The reason for the discrepancy in *hs*CRP levels with respect to gender is not clearly resolved, although several studies suggested different lifestyle and metabolic risk factors between men and women[30] and genetic diversity[27]. A high level of serum hsCRP in our population was positively related to the increased risk of all-cause mortality in both genders, supported by several previous studies[8, 16, 31]. Nevertheless, several studies reported no association of hsCRP levels with all-cause mortality was observed in women[7, 16]. In particular, the association was shown in postmenopausal women only, which might suggest the protective effect of endogenous female hormones on the low level of hsCRP[32]; the average hsCRP level was 0.48 and 0.68 mg/L for premenopausal and postmenopausal women in this study. The protective effect could be supported by the proposition that estrogen or progesterone might to some extent repress the detrimental effects of chronic inflammation on tissue damage[33].

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Inflammation has emerged as an important factor in the processes of NCD, including CVD[17], cancer[18],

type 2 diabetes[20], COPD[19, 34] and fracture[21]. In addition, medications that had taken to treat any specific NCD, such as rennin-angiotensin system inhibitors[35] and statins and thiazolidinedione[36], could influence the level of hsCRP. The association between hsCRP and the mortality risk was not attenuated by NCD<sub>history</sub> in either gender in this study, but the statistical significance of the association disappeared in women after sensitivity analysis (aHR<sub>2vear</sub>). A dose-response relationship between hsCRP level and all-cause mortality risk was pronounced in both genders. On the other hand, the positive association of hsCRP with the risk of all-cause mortality risk was significantly observed in only men with NCD<sub>history</sub> but not in women with NCD<sub>history</sub>. The attenuated effect of NCD<sub>history</sub> on the association between hsCRP and the risk of cancer-mortality was not observed in men, consistent with results from several studies which reported the associations among healthy men[3] or cancer patients[37, 38] only. Most studies[3, 4, 6, 7, 15, 16, 31, 39] supported that CVD mortality increased with elevated hsCRP levels, predominantly in men[4, 7, 15, 16]. Although hsCRP levels are lower in our population than in other races, the level of hsCRP was positively associated with CVD mortality in men but not in women, similar to previous studies [7, 15, 16, 31, 39]. After stratification by gender and NCD<sub>history</sub>, the association between hsCRP and the risk of CVD mortality was dominant in subjects with NCD<sub>history</sub> in this study. Although many interventional studies have been conducted recently on anti-inflammatory drugs for the prevention of cardiovascular disease, the results are controversial. According to the results of our study, elevated inflammatory markers in people with chronic disease were associated with an increased risk of CVD mortality. This suggests that CVD-mortality in people with chronic diseases might be reduced by use of anti-inflammatory medication. This study has several strengths because of the large population-based prospective study; it makes possible 1) to adjust for confounders; 2) to examine sensitivity analysis after excluding death before 1 or 2 years from recruitment; 3) to assess an advanced analysis after stratification by gender and NCD<sub>bistory</sub>; 4) to examine the association using various cut-off points of hsCRP considering low serum hsCRP levels in Asian populations; and 5) to evaluate the complex (i.e., nonlinear) hazard functions using restricted cubic splines on the association between continuous hsCRP levels and the risk of mortality. In particular, most previous studies excluded subjects with more than 10 mg/L hsCRP because of their relatively low sample size or reflecting acute phase reactions of severe inflammation, but we examined the effect of very high hsCRP concentration on the risk of mortality because it is possible to be more concerning for these subjects in the future. The hsCRP level of this 

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4 5	511	study, in addition, was measured within 18 hours in a single institution to minimize measurement error/bias
6 7	512	from institutional variation to avoid bias from measurement or long-term storage before analysis.
8 9	513	Despite of those strengths, it is also several limitations. First, the use of a single measurement of hsCRP at
10 11	514	baseline could reflect the inaccurate status of blood hsCRP levels in the study participants and increase the
12	515	instability of hsCRP due to random fluctuations over time. Nevertheless, a report [40] on the long-term hsCRP
13 14	516	variability suggested that the hsCRP variability within individual is relatively small and that the variability
15 16	517	could not account for the association. Second, our study lacked information on medication use at recruitment
17 18	518	and during the follow-up period. Several medications related to NCDs, including statins, angiotensin-converting
19 20	519	enzyme inhibitors, fibrates, niacin, thiazolidinedione and estrogen/progestogen hormone, could influence the
21 22	520	hsCRP level[37]; however, we tried to overcome this limitation through advanced analysis after stratification by
23 24	521	NCD <sub>history</sub> . Third, because there is no available information on hormone-replacement therapy (HRT) among
25 26	522	women, which could not examine the influence of HRT on the association of hsCRP with the risk of hormone-
27 28	523	related cancer or CVD mortality among women, we could not suggest the effect of female hormones on the
29	524	association. In addition, further studies are needed on the effects of obesity although the inverse relationship
30 31	525	between all-cause mortality with obesity in our population was consistent to Wei's report in Asian[41]. On the
32 33	526	other hand, the inverse association of alcohol drinking with all-cause mortality couldn't interpret directly
34 35	527	because our report wasn't separated the distinguish between mild drinkers and abuse alcohol drinker, which
36 37	528	requires additional research for our population in the future.
38 39	529	In conclusion, the association of hsCRP level is dose-responsively increased with the risk of all-cause
40 41	530	mortality in men and women (particularly postmenopausal women), which was not influenced by the association
42 43	531	was not observed in women with NCD <sub>history</sub> . Otherwise, the association of hsCRP level with the risk of cancer-
44 45	532	and CVD-mortality could be attenuated by gender or NCD <sub>history</sub> .
46 47	533	
48	534	
49 50	535	Figure 1 Flow diagram of analytical sample in current study using Health Examinees cohort.
51 52	536	<b>Figure 2</b> A dose-response association between serum <i>hs</i> CRP level and risk of all-cause mortality in all (a),
53 54	537	healthy subjects at recruitment (b), and subjects with non-communicable disease history (NCD <sub>history</sub> ) at
55 56	538	recruitment (c).
57 58	539	<b>Figure 3</b> Kaplan-Meier crude survival curves for cancer-mortality according to serum <i>hs</i> CRP level in all (a)
59 60		

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3 4	540	men (b), women (c), healthy subjects at recruitment (d), and subjects with non-communicable disease history
5 6	541	(NCD <sub><i>history</i></sub> ) at recruitment (e).
7 8	542	<b>Figure 4</b> Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum <i>hs</i> CRP
9 10	543	level in all (a), men (b), women (c), healthy subjects at recruitment (d), and subjects with non-communicable
11 12	544	disease history (NCD <sub>history</sub> ) at recruitment (e).
13 14	545	
15	546	
16 17	547	Contributors
18 19		
20	548	SAL, XS and DK: designed and conducted the research, SAL and SOK: analyzed the data and performed the
21 22	549	statistical analyses; HP and JKL: managed data mining and collection; SAL: wrote the manuscript and had primary
23 24	550	responsibility for the final content of the manuscript; and all authors: read and approved the final manuscript.
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40 41	558	
42 43	559	Provenance and peer review Not commissioned; externally peer reviewed.
44 45	560	Data availability statement
46 47	561	No additional data available.
48 49	562	
50	563	
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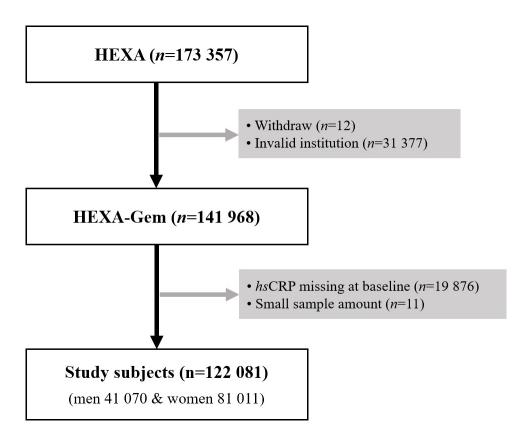


Figure 1 Flow diagram of analytical sample in current study using Health Examinees cohort. HEXA: Health Examinees, hsCRP: High sensitivity C-reactive protein

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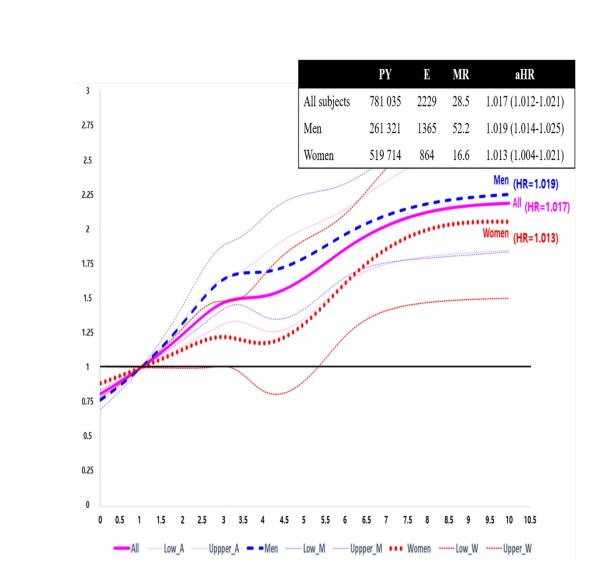


Figure 2 (a) A dose-response association between serum hsCRP level and risk of all-cause mortality in all subjects at recruitment.PY: Person-year, E: Number of death, MR: Mortality rate (10 000 person year)aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exerciseLow\_A and Upper\_A: 95%CI for all subjectsLow\_M and Upper\_M: 95%CI for menLow\_W and Upper\_W: 95%CI for women

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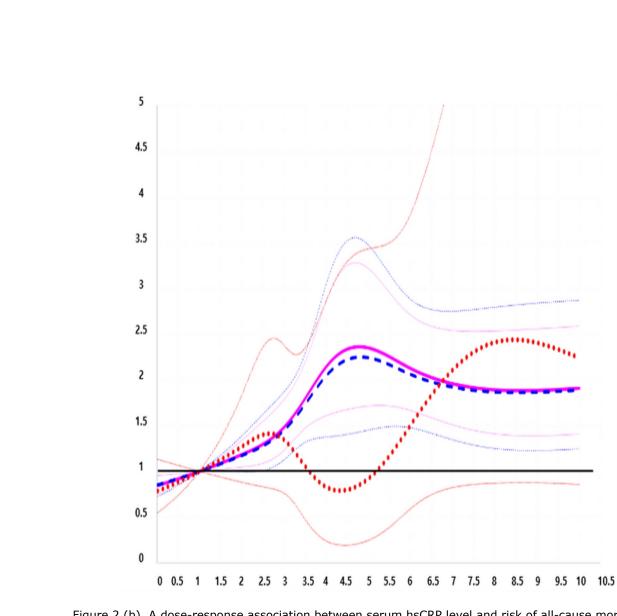


Figure 2 (b) A dose-response association between serum hsCRP level and risk of all-cause mortality in healthy subjects at recruitment.Low\_A and Upper\_A: 95%CI for all subjects Low\_M and Upper\_M: 95%CI for menLow\_W and Upper\_W: 95%CI for women

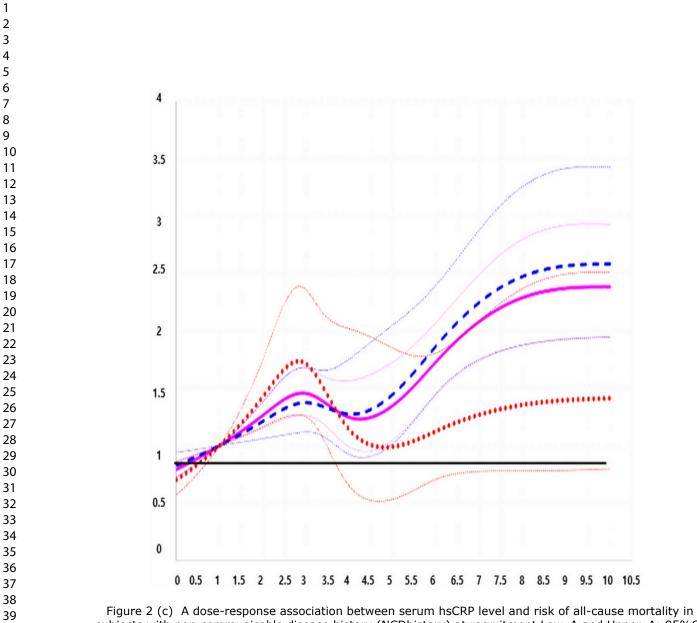


Figure 2 (c) A dose-response association between serum hsCRP level and risk of all-cause mortality in subjects with non-communicable disease history (NCDhistory) at recruitment.Low\_A and Upper\_A: 95%CI for all subjectsLow\_M and Upper\_M: 95%CI for menLow\_W and Upper\_W: 95%CI for women

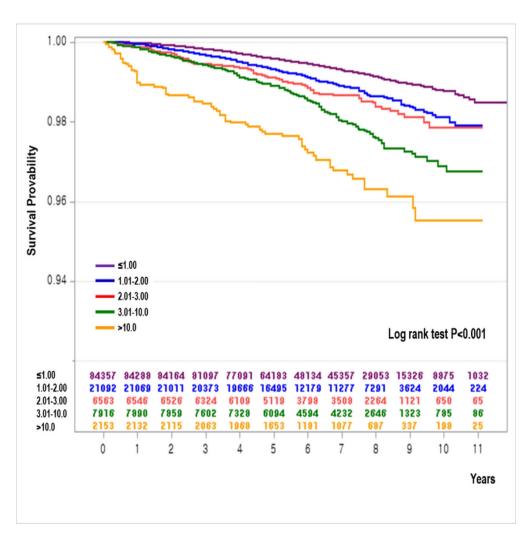
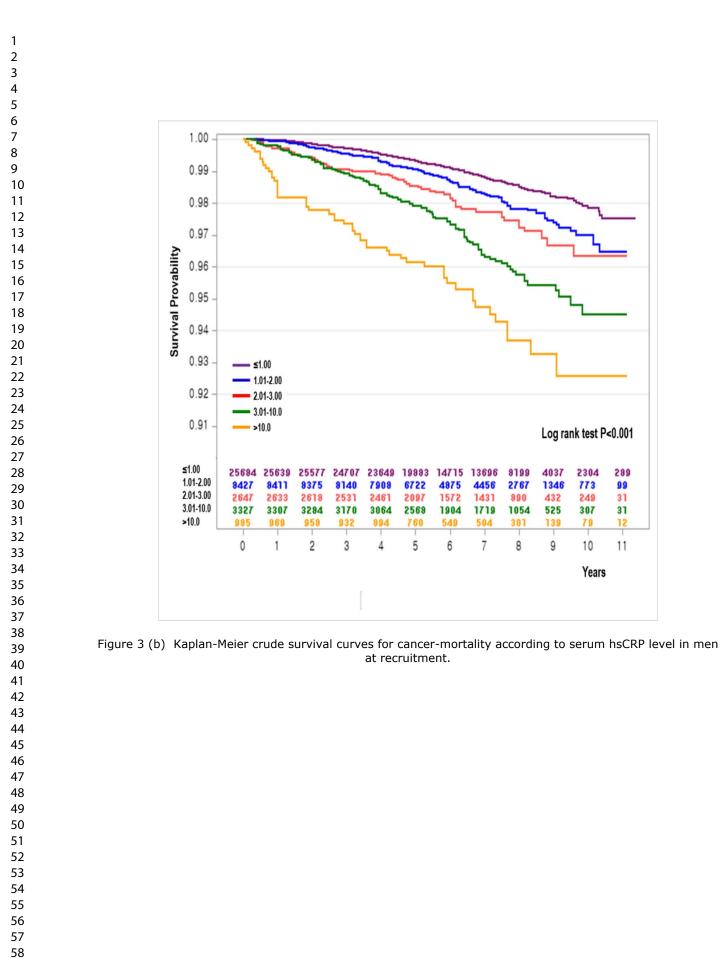


Figure 3 (a) Kaplan-Meier crude survival curves for cancer-mortality according to serum hsCRP level in all subjects at recruitment.

19883 14715 13696

Log rank test P<0.001

Years



≤1.00

1.01-2.00

2.01-3.00

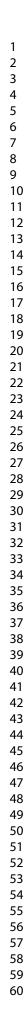
3.01-10.0

>10.0

25684 25639

25577 24707 23649

at recruitment.



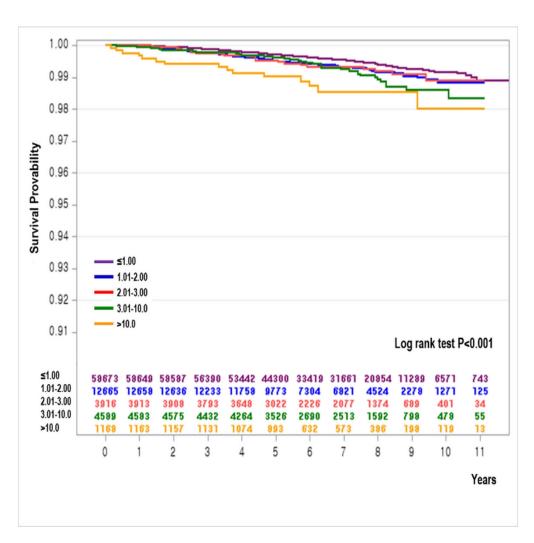


Figure 3 (c) Kaplan-Meier crude survival curves for cancer-mortality according to serum hsCRP level in women at recruitment.

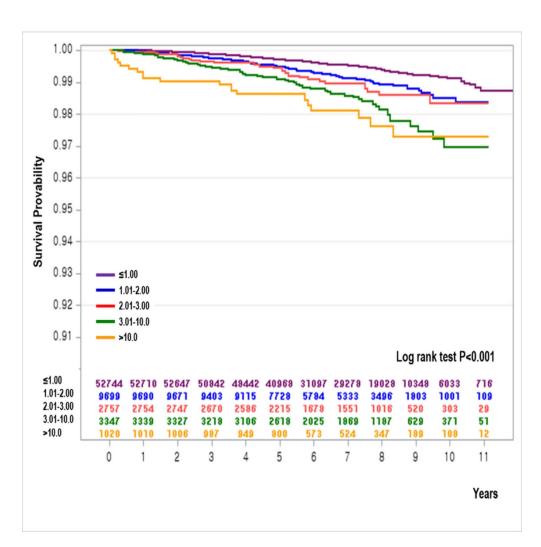
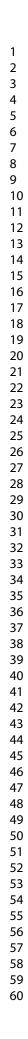


Figure 3 (d) Kaplan-Meier crude survival curves for cancer-mortality according to serum hsCRP level in healthy subjects at recruitment.



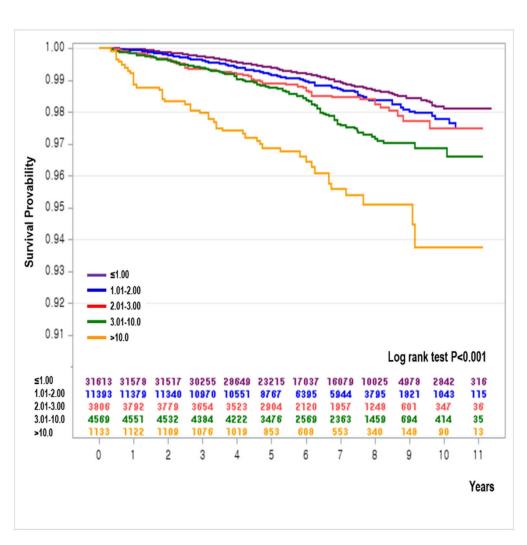


Figure 3 (e) Kaplan-Meier crude survival curves for cancer-mortality according to serum hsCRP level in subjects with non-communicable disease history (NCDhistory) at recruitment.

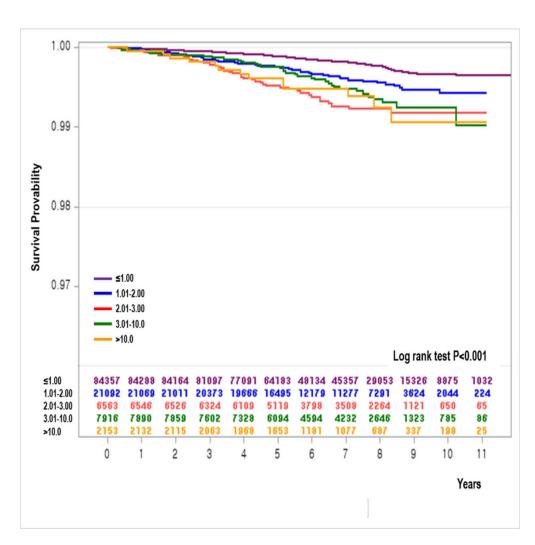


Figure 4 (a) Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum hsCRP level in all subjects at recruitment.

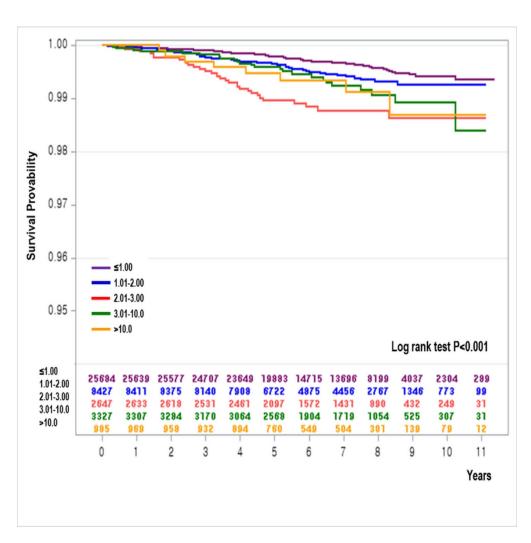


Figure 4 (b) Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum hsCRP level in men at recruitment.

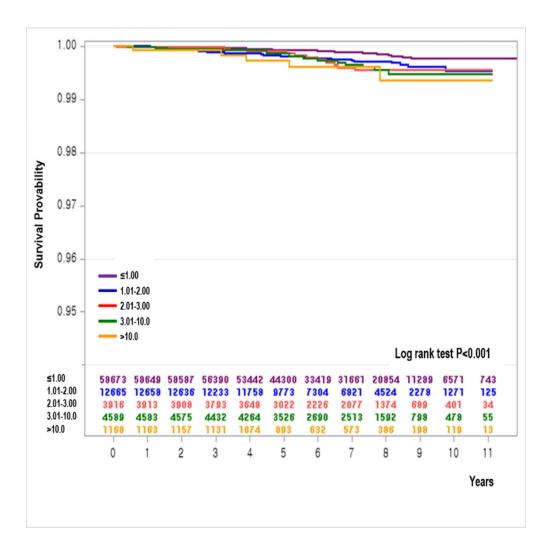
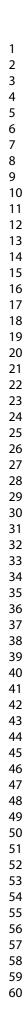


Figure 4 (c) Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum hsCRP level in women at recruitment.



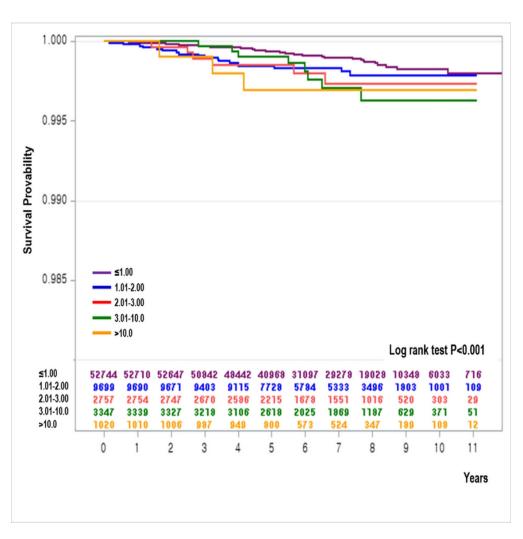


Figure 4 (d) Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum hsCRP level in healthy subjects at recruitment.

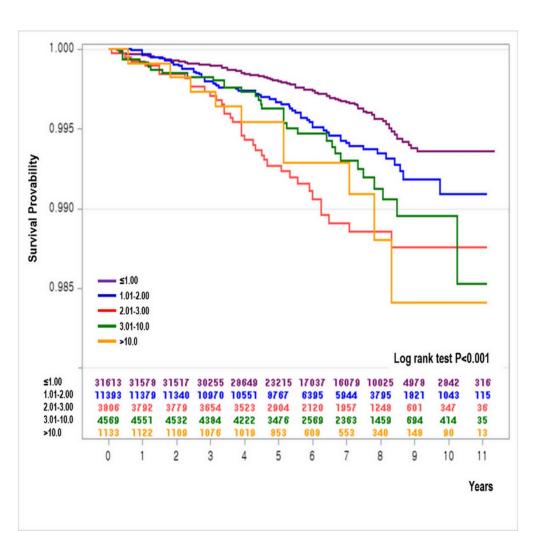


Figure 4 (e) Kaplan-Meier crude survival curves for cardiovascular disease mortality according to serum hsCRP level in subjects with non-communicable disease history (NCDhistory) at recruitment.

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	PY	Е	MR	aHR	HR <sub>1year</sub>	HR <sub>2year</sub>
All subjects					-	
Continuous	781 035	2229	28.5	1.017 (1.012-1.021)	1.015 (1.010-1.020)	1.014 (1.009-1.019
≤1.00	539 271	1153	21.4	Reference	Reference	Reference
1.01-1.50	90 911	308	33.9	1.26 (1.10-1.45)	1.27 (1.10-1.45)	1.21 (1.04-1.40)
1.51-2.00	44 615	163	36.5	1.28 (1.08-1.53)	1.28 (1.07-1.53)	1.26 (1.04-1.52)
2.01-2.50	25 139	117	46.5	1.53 (1.25-1.89)	1.51 (1.22-1.87)	1.49 (1.19-1.87)
2.51-3.00	16 996	72	42.4	1.39 (1.08-1.80)	1.31 (1.00-1.72)	1.23 (0.92-1.65)
3.01-4.00	19 667	103	52.4	1.61 (1.29-2.01)	1.62 (1.29-2.03)	1.64 (1.30-2.08)
4.01-6.00	17 933	102	56.9	1.84 (1.48-2.28)	1.77 (1.41-2.21)	1.70 (1.34-2.16)
6.01-10.00	13 019	88	67.6	2.02 (1.59-2.56)	1.96 (1.54-2.50)	1.93 (1.49-2.51)
>10.0	13 484	123	91.2	2.59 (2.12-3.16)	2.41 (1.95-2.97)	2.26 (1.80-2.84)
P-trend				<.001	<.001	<.001
Men						
Continuous	261 321	1365	52.2	1.019 (1.014-1.025)	1.017 (1.011-1.023)	1.017 (1.010-1.023
≤1.00	163 068	638	39.1	Reference	Reference	Reference
1.01-1.50	36 094	190	52.6	1.27 (1.07-1.51)	1.28 (1.07-1.53)	1.22 (1.01-1.47)
1.51-2.00	17 946	103	57.4	1.34 (1.07-1.67)	1.34 (1.07-1.68)	1.35 (1.06-1.72)
2.01-2.50	10 059	77	76.5	1.56 (1.20-2.03)	1.53 (1.16-2.00)	1.47 (1.10-1.96)
2.51-3.00	6959	54	77.6	1.71 (1.27-2.29)	1.57 (1.15-2.15)	1.46 (1.04-2.05)
3.01-4.00	8177	77	94.2	1.88 (1.45-2.43)	1.94 (1.50-2.52)	1.92 (1.46-2.54)
4.01-6.00	7425	75	101.0	2.05 (1.59-2.63)	1.95 (1.49-2.53)	1.91 (1.44-2.52)
6.01-10.00	5456	59	108.1	2.03 (1.52-2.73)	1.96 (1.44-2.66)	1.85 (1.33-2.58)
>10.0	6137	92	149.9	2.84 (2.25-3.58)	2.66 (2.08-3.39)	2.58 (1.99-3.35)
P-trend				<.001	<.001	<.001
Women						
Continuous	519 714	864	16.6	1.013 (1.004-1.021)	1.011(1.002-1.021)	1.010 (0.999-1.02)
≤1.00	376 203	515	13.7	Reference	Reference	Reference
1.01-1.50	54 817	118	21.5	1.28 (1.03-1.59)	1.27 (1.02-1.58)	1.23 (0.97-1.56)
1.51-2.00	26 669	60	22.5	1.23 (0.92-1.64)	1.21 (0.90-1.63)	1.14 (0.83-1.56)
2.01-2.50	15 080	40	26.5	1.52 (1.09-2.14)	1.52 (1.08-2.15)	1.56 (1.09-2.24)
2.51-3.00	10 037	18	17.9	0.84 (0.49-1.44)	0.87 (0.51-1.48)	0.83 (0.46-1.47)
3.01-4.00	11 490	26	22.6	1.16 (0.75-1.81)	1.09 (0.68-1.72)	1.21 (0.76-1.93)
4.01-6.00	10 508	27	25.7	1.48 (0.99-2.22)	1.47 (0.97-2.22)	1.36 (0.86-2.14)
6.01-10.00	7563	29	38.3	2.00 (1.34-2.98)	1.98 (1.32-2.98)	2.10 (1.39-3.19)
>10.0	7347	31	42.2	2.02 (1.36-3.02)	1.84 (1.21-2.81)	1.51 (0.93-2.47)
P-trend				<.001	<.001	0.001
Premenopause						
≤1.00	141 286	96	6.8			
1.01-2.00	20 500	20	9.8	1.52 (0.92-2.52)	1.49 (0.89-2.50)	1.57 (0.90-2.73)
2.01-3.00	5835	6	10.3	1.76 (0.77-4.06)	1.83 (0.79-4.22)	1.42 (0.52-3.93)
3.01-10.0	6886	6	8.7	1.51 (0.66-3.50)	1.31 (0.53-3.25)	1.21 (0.44-3.36)
>10.0	1759	4	22.7	2.57 (0.81-8.14)	2.63 (0.83-8.37)	2.09 (0.51-8.58)
<i>P</i> -trend		·		0.020	0.036	0.150
Postmenopause						
≤1.00	192 164	366	19.0			
1.01-2.00	52 897	145	27.4	1.26 (1.03-1.55)	1.25 (1.02-1.54)	1.18 (0.95-1.48)
2.01-3.00	16 943	44	26.0	1.11 (0.80-1.56)	1.12 (0.80-1.57)	1.19 (0.83-1.68)
3.01-10.0	19 687	67	34.0	1.49 (1.13-1.97)	1.47 (1.10-1.95)	1.52 (1.13-2.05)
>10.0	4828	27	55.9	2.09 (1.37-3.21)	1.88 (1.19-2.96)	1.56 (0.92-2.63)
<i>P</i> -trend	1020			<0.001	0.001	0.003

Supplement 1 The association of ser m hsCRP level with the risk of all-cause mortality

PY: Person-year, E: Number of death, MR: Mortality rate (10,000 person year)

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- aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise
  - $HR_{1year}$ : aHR after exclude subjects who died within 1 yr f/u time
  - HR<sub>2year</sub>: aHR after exclude subjects who died within 2 yr f/u time

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		Healthy subjects at recruitment					Subjects with NCD <sub>history</sub> at recruitment				
	E	MR	aHR	$HR_{1year}$	HR <sub>2year</sub>	E	MR	aHR	$HR_{1year}$	HR <sub>2yea</sub>	
All											
$\leq 1.00$	517	15.1	Ref	Ref	Ref	636	32.3	Ref	Ref	Re	
1.01-2.00	145	22.9	1.20	1.19	1.16	326	45.1	1.20	1.19	1.1	
2.01-3.00	53	29.3	1.38	1.37	1.32	136	56.6	1.51	1.46	1.4	
3.01-10.0	102	46.8	2.22	2.15	2.15	191	66.3	1.62	1.60	1.5	
>10.0	40	61.3	2.38	2.23	2.27	83	118.9	2.74	2.54	2.2	
P-trend			<.001	<.001	<.001			<.001	<.001	<.00	
Men											
$\leq 1.00$	270	29.5	Ref	Ref	Ref	368	51.4	Ref	Ref	Re	
1.01-2.00	89	35.8	1.11	1.11	1.13	204	70.0	1.40	1.41	1.3	
2.01-3.00	33	46.3	1.22	1.17	1.15	98	99.0	1.82	1.73	1.6	
3.01-10.0	70	76.8	2.14	2.08	2.03	141	118.1	1.92	1.90	1.8	
>10.0	31	110.0	2.60	2.49	2.73	61	183.1	3.05	2.83	2.5	
P-trend			<.001	<.001	<.001			<.001	<.001	<.00	
Women											
$\leq 1.00$	247	9.8	Ref	Ref	Ref	268	21.4	Ref	Ref	Re	
1.01-2.00	56	14.6	1.35	1.32	1.20	122	28.3	1.19	1.20	1.1	
2.01-3.00	20	18.2	1.61	1.66	1.60	38	26.9	1.06	1.06	1.1	
3.01-10.0	32	25.2	2.31	2.23	2.37	50	29.7	1.16	1.14	1.1	
>10.0	9	24.3	1.69	1.49	1.12	22	60.3	2.15	1.99	1.6	
P-trend			<.001	<.001	0.001			0.018	0.043	0.08	

**Supplement 2.** The association between serum *hs*CRP level and all-cause mortality by gender and noncommunicable disease history (NCD<sub>*history*</sub>) at recruitment

E: Number of death, MR: Mortality rate (10 000 person year)

aHR: Adjusted for age, gender, education, job, marital status, BMI, non-communicable disease history, smoking, alcohol consumption and exercise

 $HR_{1year}\!\!:aHR$  after exclude subjects who died within 1 yr f/u time

HR<sub>2year</sub>: aHR after exclude subjects who died within 2 yr f/u time

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Section/Topic	Item #	Recommendation					
Title and abstract	1	(a) Indicate the study's design with a commonly used term in the title or the abstract					
		(b) Provide in the abstract an informative and balanced summary of what was done and what was found	1-2				
Introduction							
Background/rationale	2	Explain the scientific background and rationale for the investigation being reported					
Objectives	3	State specific objectives, including any pre-specified hypotheses					
Methods							
Study design	4	Present key elements of study design early in the paper	4				
Setting	5	Describe the setting, locations, and relevant dates, including periods of recruitment, exposure, follow-up, and data collection					
Participants	6	<ul> <li>(a) Cohort study—Give the eligibility criteria, and the sources and methods of selection of participants. Describe methods of follow-up</li> <li>Case-control study—Give the eligibility criteria, and the sources and methods of case ascertainment and control selection. Give the rationale for the choice of cases and controls</li> <li>Cross-sectional study—Give the eligibility criteria, and the sources and methods of selection of participants</li> </ul>	4				
		(b) Cohort study—For matched studies, give matching criteria and number of exposed and unexposed Case-control study—For matched studies, give matching criteria and the number of controls per case	Not Applicable				
Variables	7	Clearly define all outcomes, exposures, predictors, potential confounders, and effect modifiers. Give diagnostic criteria, if applicable					
Data sources/ measurement	8*	For each variable of interest, give sources of data and details of methods of assessment (measurement). Describe comparability of assessment methods if there is more than one group					
Bias	9	Describe any efforts to address potential sources of bias	6				
Study size	10	Explain how the study size was arrived at	4				
Quantitative variables	11	Explain how quantitative variables were handled in the analyses. If applicable, describe which groupings were chosen and why					
Statistical methods	12	(a) Describe all statistical methods, including those used to control for confounding	5-6				
		(b) Describe any methods used to examine subgroups and interactions	5-6				
		(c) Explain how missing data were addressed	4				
		(d) Cohort study—If applicable, explain how loss to follow-up was addressed Case-control study—If applicable, explain how matching of cases and controls was addressed	4				

		Cross-sectional study—If applicable, describe analytical methods taking account of sampling strategy	
		(e) Describe any sensitivity analyses	6
Results	•		
Participants	13*	(a) Report numbers of individuals at each stage of study—eg numbers potentially eligible, examined for eligibility, confirmed eligible, included in the study, completing follow-up, and analysed	
		(b) Give reasons for non-participation at each stage	4
		(c) Consider use of a flow diagram	4
Descriptive data	14*	(a) Give characteristics of study participants (eg demographic, clinical, social) and information on exposures and potential confounders	6-7
		(b) Indicate number of participants with missing data for each variable of interest	4
		(c) Cohort study—Summarise follow-up time (eg, average and total amount)	6
Outcome data	15*	Cohort study—Report numbers of outcome events or summary measures over time	6-10
		Case-control study—Report numbers in each exposure category, or summary measures of exposure	Not Applicable
		Cross-sectional study—Report numbers of outcome events or summary measures	Not Applicable
Main results	16	( <i>a</i> ) Give unadjusted estimates and, if applicable, confounder-adjusted estimates and their precision (eg, 95% confidence interval). Make clear which confounders were adjusted for and why they were included	7-11
		(b) Report category boundaries when continuous variables were categorized	5
		(c) If relevant, consider translating estimates of relative risk into absolute risk for a meaningful time period	Not Applicable
Other analyses	17	Report other analyses done—eg analyses of subgroups and interactions, and sensitivity analyses	9
Discussion		·	
Key results	18	Summarise key results with reference to study objectives	12
Limitations	19	Discuss limitations of the study, taking into account sources of potential bias or imprecision. Discuss both direction and magnitude of any potential bias	14
Interpretation	20	Give a cautious overall interpretation of results considering objectives, limitations, multiplicity of analyses, results from similar studies, and other relevant evidence	12-14
Generalisability	21	Discuss the generalisability (external validity) of the study results	12-14
Other information		•	
Funding	22	Give the source of funding and the role of the funders for the present study and, if applicable, for the original study on which the present article is based	15

\*Give information separately for cases and controls in case-control studies and, if applicable, for exposed and unexposed groups in cohort and cross-sectional studies. **Note:** An Explanation and Elaboration article discusses each checklist item and gives methodological background and published examples of transparent reporting. The STROBE checklist is best used in conjunction with this article (freely available on the Web sites of PLoS Medicine at http://www.plosmedicine.org/, Annals of Internal Medicine at http://www.annals.org/, and Epidemiology at http://www.epidem.com/). Information on the STROBE Initiative is available at www.strobe-statement.org.