Kunutsor, S., Mäkikallio, T. H., Kauhanen, J., Voutilainen, A., \& Laukkanen, J. A. (2019). Lipoprotein(a) is not associated with venous thromboembolism risk. Scandinavian Cardiovascular Journal, 53(3), 125-132. https://doi.org/10.1080/14017431.2019.1612087

Peer reviewed version

Link to published version (if available):
10.1080/14017431.2019.1612087

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# Lipoprotein(a) is not associated with venous thromboembolism risk 

Setor K. Kunutsor, ${ }^{\text {a,b,*, }}$, Timo H. Mäkikallio ${ }^{\text {c }}$, Jussi Kauhanen ${ }^{\text {d }}$, Ari Voutilainen ${ }^{\text {d }}$, Jari A. Laukkanen ${ }^{\text {de,f }}$
${ }^{a}$ National Institute for Health Research Bristol Biomedical Research Centre, University Hospitals Bristol NHS Foundation Trust and University of Bristol, Bristol, UK ${ }^{\mathrm{b}}$ Musculoskeletal Research Unit, Translational Health Sciences, Bristol Medical School, University of Bristol, Learning \& Research Building (Level 1), Southmead Hospital, Bristol, BS10 5NB, UK
${ }^{\text {c }}$ Division of Cardiology, Department of Internal Medicine, Oulu University Hospital, Oulu, Finland
${ }^{d}$ Institute of Public Health and Clinical Nutrition, University of Eastern Finland, Kuopio, Finland
${ }^{e}$ Faculty of Sport and Health Sciences, University of Jyväskylä, Jyväskylä, Finland
${ }^{\mathrm{f}}$ Central Finland Health Care District Hospital District, Jyväskylä, Finland

## *Corresponding author:

Setor K. Kunutsor, Musculoskeletal Research Unit, Translational Health Sciences, Bristol Medical School, University of Bristol, Learning \& Research Building (Level 1), Southmead Hospital, Bristol, BS10 5NB, UK; Phone: +44-7539589186; Fax: +44-1174147924; Email address: skk31@cantab.net

Word count [3013]


#### Abstract

Objectives. Evidence from case-control studies as well as meta-analyses of these study designs suggest elevated lipoprotein(a) $[\mathrm{Lp}(\mathrm{a})]$ to be associated with an increased risk of venous thromboembolism (VTE). Prospective evidence on the association is limited, uncertain, and could be attributed to regression dilution bias. We aimed to assess the prospective association of $\operatorname{Lp}(a)$ with risk of VTE and correct for regression dilution. Design. We related plasma $\operatorname{Lp}$ (a) concentrations to the incidence of VTE in 2,180 men of the Kuopio Ischemic Heart Disease cohort study. Hazard ratios (HRs) (95\% confidence intervals [CI]) were assessed and repeat measurements of $\operatorname{Lp}(a)$ at 4 and 11 years from baseline, were used to correct for within-person variability. Results. After a median follow-up of 24.9 years, 110 validated VTE cases were recorded. The regression dilution ratio of $\log _{e} \mathrm{Lp}(\mathrm{a})$ adjusted for age was 0.85 ( $95 \%$ CI: $0.82-0.89$ ). In analyses adjusted for several established risk factors and potential confounders, the $\mathrm{HR}(95 \% \mathrm{CI})$ for VTE per 1 SD (equivalent to 3.56-fold) higher baseline $\log _{\mathrm{e}} \mathrm{Lp}(\mathrm{a})$ was 1.06 (0.87-1.30). In pooled analysis of five population-based cohort studies (including the current study) comprising 66,583 participants and 1,314 VTE cases, the fully-adjusted corresponding HR for VTE was 1.00 ( $95 \%$ CI: $0.94-1.07$ ), with no evidence of heterogeneity between studies. Conclusions. Primary analysis as well as pooled evidence from previous studies suggest circulating $\mathrm{Lp}(\mathrm{a})$ is not prospectively associated with future VTE risk, indicating that evidence of associations demonstrated in case-control designs may be driven by biases such as selection bias.


## KEYWORDS

lipoprotein(a); venous thromboembolism; cohort study; risk factor; regression dilution

## Introduction

Lipoprotein (a) [Lp(a)], composed of a dual structure and has both proatherosclerotic and prothrombotic functions [1,2], is an enigmatic lipoprotein that has been the subject of research over the past two decades. The relationship existing between $\mathrm{Lp}(\mathrm{a})$ and cardiovascular disease (CVD) has been well established. Consistently, several well-designed large-scale epidemiological studies have shown $\operatorname{Lp}(a)$ to be independently associated with cardiovascular outcomes [3-6] with some suggestions of causal relationships reported [4-6]. Though Lp(a) pathophysiology in vascular disease is controversial and still not fully understood, evidence suggests that $\operatorname{Lp}(\mathrm{a})$ contributes to the aetiology of vascular diseases via proatherosclerotic and proinflammatory mechanisms [7]. Venous thromboembolism (VTE) (comprising deep vein thrombosis (DVT) and pulmonary embolism (PE)), which is an important cause of increased morbidity and premature mortality $[8,9]$, is closely linked with CVD [10-12] and both conditions share common antecedent risk factors [13]. Given the prothrombotic properties of $\operatorname{Lp}(a)$, it has been suggested that $\operatorname{Lp}(a)$ may play a role in the pathophysiology of VTE. Indeed, emerging data supports an association between elevated Lp(a) and VTE risk. Several case-control studies have shown increased VTE risk with elevated Lp(a) concentrations [1417]. Two meta-analyses of these study designs have also confirmed these associations $[18,19]$. It appears the data showing a relationship between $\operatorname{Lp}(a)$ and VTE have largely been based on case-control designs, which are characterised by selection bias and do not show a temporal relationship between $\operatorname{Lp}(a)$ and VTE risk. A number of prospective cohort studies based in the general population have consistently reported no evidence of an association between $\mathrm{Lp}(\mathrm{a})$ and future VTE risk [20,21].

Based on the emerging data, it appears $\operatorname{Lp}($ a) might not be prospectively linked to VTE risk, however more research is needed given that incident VTE rates in these previous studies were relatively small. Furthermore, there is a possibility that the inability of previous longterm follow-up cohort studies to demonstrate an association between $\operatorname{Lp}(a)$ and VTE risk could be partly attributed to regression dilution bias [22]. This is a phenomenon which potentially results in the underestimation of the true association between an exposure ( $\operatorname{Lp}(a)$
and outcome (VTE), particularly for cohorts with long-term follow-up. Regression dilution bias can be addressed by correcting the risk estimates using the regression dilution ratio (RDR) [23].

Due to the wide uncertainty in the evidence, we sought to evaluate in detail the prospective nature of the association between $\operatorname{Lp}(a)$ and future VTE risk using a population-based cohort of 2,180 men from eastern Finland followed up for over of 20 years. Secondly, repeat measurements of $\operatorname{Lp}(a)$ performed several years apart in a random sample of participants enabled quantification of within-person variability in $\mathrm{Lp}(\mathrm{a})$ levels. We also performed pooled analysis of available published prospective evidence on the association, thereby offering the opportunity to re-evaluate the nature and magnitude of the association in a larger representative sample of participants and VTE cases.

## Methods

## Study design and population

This study was conducted in accordance with STROBE (STrengthening the Reporting of OBservational studies in Epidemiology) guidelines for reporting observational studies in epidemiology (Appendix A) [24]. The study population is based on the Kuopio Ischemic Heart Disease (KIHD) risk factor study, a general population-based prospective cohort study designed to investigate risk factors for CVD and other chronic diseases. The design and recruitment methods of the KIHD study have been described in previous reports [25-30]. Participants consisted of a representative sample of men aged 42-61 years who were inhabitants of the city of Kuopio and its surrounding rural communities in eastern Finland. The actual baseline cohort consisted of 2,682 participants had baseline measurements performed between March 1984 and December 1989. In the current analysis, complete information on plasma Lp(a), relevant covariates, and VTE outcomes was available for 2,180 men. The research protocol was approved by the institutional review board of the University of Eastern Finland. All study procedures were conducted according to the Declaration of Helsinki. Written informed consent was obtained from all participants

## Assessment of $\operatorname{Lp}(a)$ and other risk markers

Assessment of data on demographics, lifestyle characteristics, physical measurements, collection of blood samples and measurement of serum lipids, lipoproteins and biochemical analytes have been described in previous reports $[27,29]$. Blood samples were taken between 8 and 10 a.m. after an overnight fast. The cholesterol content of lipoprotein fractions were measured from fresh samples after combined ultracentrifugation and precipitation, and were assessed enzymatically (Boehringer Mannheim, Mannheim, Germany) [31]. Lp(a) measurements were made from frozen plasma samples stored at $-20^{\circ} \mathrm{C}$ for 2-6 years, using a radioimmunoassay (Mercodia Apo(a) RIA, Mercodia AB, Uppsala, Sweden), with repeat measurements performed in a random subset of participants at 4 years and 11 years after the baseline measurements. Fasting plasma glucose (FPG) was measured by the glucose dehydrogenase method (Merck, Darmstadt, Germany). Serum high sensitivity C-reactive protein (hsCRP) measurements were made with an immunometric assay (Immulite High Sensitivity C-Reactive Protein Assay; DPC, Los Angeles, CA, USA). Plasma fibrinogen concentrations were determined in fresh plasma samples with excess thrombin using the Coagulometer KC4 device (Heinrich Amelung GmbH, Lemgo, Germany). For the assessments of age, lifestyle factors such as smoking and alcohol consumption, medical conditions, and medication history; participants completed self-administered questionnaires [32]. The energy expenditure of physical activity was assessed using the validated KIHD 12month leisure-time physical activity questionnaire [33,34]. Body mass index (BMI) was estimated as weight in kilograms divided by the square of height in meters.

## Ascertainment of incident VTE

We included all first lifetime VTE events that occurred from study enrollment through to 2013. These were identified by computer linkage to the National Hospital Discharge Registry data and a comprehensive review of available hospital records, wards of health centres, health practitioner questionnaires, death certificate and autopsy registers, and medico-legal reports. The diagnosis of DVT or PE required positive imaging tests. Documents were cross-checked
in detail and VTE events were validated by two physicians. No losses to follow-up were recorded as all participants in the KIHD study (using Finnish personal identification codes) are under continuous surveillance for the development of new outcomes including VTE cases.

## Statistical analysis

Prospective cohort analyses Skewed variables (hsCRP, triglycerides, and fibrinogen) were log transformed to achieve approximately symmetrical distributions. Descriptive analyses were conducted to summarize the baseline characteristics of the participants, with means (standard deviation, SD) or medians (interquartile range, IQR) reported for continuous variables and n (percentages) for categorical variables. The partial correlation coefficients were calculated using linear regression models adjusted for age, to assess the cross-sectional associations of $\operatorname{Lp}(a)$ with various risk markers. The SD of baseline loge $\mathrm{Lp}(\mathrm{a})$ concentration was 1.27 , corresponding to approximately four-fold higher circulating $\operatorname{Lp}(a)$ (ie, $\mathrm{e}^{1.27}=3.56$ ). Hazard ratios (HRs) with 95\% confidence intervals (CIs) were calculated using Cox proportional hazard models, after confirming no major departure from the assumptions of proportionality of hazards using Schoenfeld residuals.[35] Lp(a) was modelled continuously (per1 SD (ie, 3.56 fold) higher Lp (a) levels) and by quartiles defined according to the baseline distribution of plasma Lp(a) levels. Hazard ratios were calculated with adjustment for confounders in two models: i) age and ii) established risk factors and other potential confounders [BMI, systolic blood pressure (SBP), history of hypertension, prevalent coronary heart disease (CHD), smoking status, history of diabetes, total cholesterol, lipid lowering medication, estimated glomerular filtration rate (eGFR) as calculated using the Chronic Kidney Disease Epidemiology Collaboration formula [36], physical activity, alcohol consumption, prevalent cancer, fibrinogen and hsCRP. We employed formal tests of interaction to assess statistical evidence of effect modification on the association by categories of pre-specified clinically relevant individual level characteristics. To quantify and correct for within-person variability in $\operatorname{Lp}(a)$ levels, which is, the extent to which an individual's $\operatorname{Lp}(\mathrm{a})$ measurements vary around the long-term average exposure levels ("usual
levels") [37], adjusted regression dilution ratios (RDRs) were calculated by regressing available repeat measurements of $\operatorname{Lp}(a)$ on baseline values [23]. The RDR assumes that the "usual levels" of $\operatorname{Lp}(\mathrm{a})$ represents the true long-term exposure of $\mathrm{Lp}(\mathrm{a})$ levels on VTE risk.

Systematic review and meta-analysis We conducted a meta-analysis of published prospective cohort studies reporting on the association between $\operatorname{Lp}(a)$ and risk of VTE, using a predefined protocol and reported in accordance with PRISMA and MOOSE guidelines $[38,39]$
(Appendix B and C). Published observational population-based prospective (cohort, case cohort, or nested case-control) studies that evaluated the associations between baseline levels of Lp(a) and risk of first VTE in the adult general population up to July 2018, were sought using computer-based databases (MEDLINE, EMBASE, and Web of Science). Case-control study designs were not part of the inclusion criteria. The computer-based searches combined free and MeSH search terms and combined key words related to the exposure (e.g., "lipoprotein(a)") and outcome (e.g., "venous thromboembolism", "deep vein thrombosis", "pulmonary embolism"). We placed no restrictions on language or the publication date. Details of the search strategy are reported in Appendix D. We assessed study quality using the nine-star Newcastle-Ottawa Scale (NOS)[40] as described previously [41]. Summary measures were presented as relative risks (RRs) with $95 \%$ confidence intervals (CIs). Following Cornfield's rare disease assumption [42], hazard ratios and odds ratios were assumed to approximate the same measure of RR. To enable a consistent approach to the meta-analysis and enhance comparison with the primary analysis, reported study-specific risk estimates were also transformed to per SD increase in $\mathrm{Lp}($ a) or as extreme quartiles of $\mathrm{Lp}(\mathrm{a})$ using standard statistical methods [43,44], which have been described in detail previously [45,46]. Summary RRs were pooled using a random effects model to minimize the effect of between-study heterogeneity [47]. Subsidiary analysis used fixed effects models. Statistical heterogeneity between studies was quantified using standard chi-square tests and the $\mathrm{I}^{2}$ statistic [48]. All statistical analyses were conducted using Stata version 15 (Stata Corp, College Station, Texas).

## Results

## Baseline characteristics of Lp(a) and correction for within-person variability

The mean baseline age of study participants was 53 (SD, 5) years and the median (IQR) of $\mathrm{Lp}($ a) at baseline was 9.66 (3.75-22.27) $\mathrm{mg} / \mathrm{dl}$ (Table 1). Plasma Lp (a) levels were weakly correlated with several risk markers. There were inverse correlations of $\operatorname{Lp}(a)$ with BMI, triglycerides, and FPG; whereas, positive correlations were observed for total cholesterol, creatinine, fibrinogen, and hsCRP. Repeat measurements of $\operatorname{Lp}(a)$ taken 4 years and 11 years after baseline were available in a random sample of 691 men, providing a total of 1,360 repeat measurements of $\operatorname{Lp}(a)$. Overall, the regression RDR of loge $\operatorname{Lp}(a)$, adjusted for age, was 0.85 ( $95 \% \mathrm{CI}: 0.82$ to 0.89 ), suggesting that the associations using baseline measurements of $\mathrm{Lp}(\mathrm{a})$ with VTE would under-estimate the association by $[(1 / 0.85)-1] * 100=18 \%$.

## Lipoprotein(a) and risk of VTE

Prospective cohort results During a median follow-up of 24.9 (interquartile range, 17.9-27.1) years, 110 VTE cases (annual rate 2.32/1,000 person-years at risk; $95 \% \mathrm{CI}$ : 1.93 to 2.80 ) were recorded. The HR per 1 SD change in baseline $\log _{\mathrm{e}} \mathrm{Lp}(\mathrm{a})$ concentration was $1.06(95 \% \mathrm{CI}$ : 0.88 to $1.29 ; p=0.530$ ) in age-adjusted analysis, which remained consistent on further adjustment for several established risk factors and potential confounders 1.06 ( $95 \%$ CI: 0.87 to $1.30 ; p=0.537$ ) (Table 2). The null associations were maintained in analyses by quartiles of the baseline distribution of $\operatorname{Lp}(a)$ levels (Table 2). The findings were also similar on correction for regression dilution (Table 2). In further analysis that compared $\operatorname{Lp}(a)$ concentrations > $30 \mathrm{mg} / \mathrm{dl}$ with that $\leq 30 \mathrm{mg} / \mathrm{dl}$, no evidence of any association was observed. Hazard ratios did not vary importantly by several relevant clinical characteristics (Figure 1).

Meta-analysis of published studies We identified four population-based prospective cohort studies reporting on the associations between circulating Lp(a) and VTE risk (Appendices E and F).[20,21,49,50] Including the current study, the pooled analysis involved five studies
comprising 66,583 participants and 1,314 VTE cases. The pooled RR for VTE per 1 SD higher baseline $\log _{e} \mathrm{Lp}(\mathrm{a})$ in fully-adjusted analyses was $1.00(95 \% \mathrm{CI}: 0.94$ to 1.07$)\left(I^{2}=0 \%\right.$, $95 \%$ CI: 0 to $79 \% ; P=0.576$ ) (Figure 2). The corresponding RR was 1.00 ( $95 \% \mathrm{CI}: 0.84$ to 1.19) when comparing the top versus bottom quartiles of $\mathrm{Lp}(\mathrm{a})$ levels. When a fixed effect model was employed, the summary RRs were identical to that of random-effects metaanalysis.

## Discussion

## Summary of main findings

In this population-based prospective study of middle-aged men without a history of VTE at study entry, our analysis showed no evidence of an association of circulating $L p(a)$ with risk of VTE. The association did also not vary importantly across several clinically relevant subgroups. Our reproducibility studies of $\operatorname{Lp}(\mathrm{a})$ yielded a high RDR which indicates that $\mathrm{Lp}(\mathrm{a})$ concentration is consistent within individuals over several years. Pooled estimates of five prospective studies (including the current study) confirmed our finding of no evidence of an association in the primary cohort analysis and there was no evidence of heterogeneity between the contributing studies.

## Comparison with previous work

Several reports based on case-control designs have reported on the associations between circulating $\mathrm{Lp}(\mathrm{a})$ and VTE risk. Though the findings from these reports have been mixed, majority have generally shown an increased risk of VTE with elevated $\operatorname{Lp}(a)$ [14-17]. There have also been efforts to aggregate these data resulting in two published reviews on the topic. In the earlier review, Sofi and colleagues pooled the results of six case-control studies and showed a significant association between high Lp(a) levels and VTE risk [18]. In a more recent review, Dentali and colleagues pooled the results of 14 studies and also demonstrated $\operatorname{Lp}(a)$ to be associated with an increased risk of VTE [19]. Of all 14 studies included in this review, only one prospective cohort was included and this was the study conducted by

Kamstrup et al [20]. Indeed, data showing evidence of an association between circulating $\mathrm{Lp}(\mathrm{a})$ and VTE risk seems to be based on case-control study designs. Unfortunately, these study designs are characterised by selection bias and are not able to adequately address temporality. Prior to the current study, four large-scale prospective cohort studies based in the general population and with long-term follow-up for VTE events have all consistently shown that circulating $\operatorname{Lp}(a)$ is not associated with VTE risk [20,21,49,50]. Though these previous studies did not correct for regression dilution bias, our current analysis shows that risk estimates based on baseline and repeated measures corresponded well. Results from the KIHD prospective study as well as pooled analysis of available prospective evidence indicate that $\mathrm{Lp}(\mathrm{a})$ is not associated with risk of VTE.

## Possible explanations for findings

As with all observational cohort studies, exposure or risk factor levels are usually assessed at study entry and related to outcomes which occur after several years. However, due to random measurement errors, temporary fluctuations and changes in the exposure over time, the effect and value of the exposure changes with time leading to regression dilution bias [22]. This potentially results in the underestimation of the true association between an exposure and outcome, particularly for cohorts with long-term follow-up. It can be argued that the absence of an association between $\operatorname{Lp}(a)$ and VTE in previous cohorts could be potentially explained by the phenomenon of regression dilution. However, this is unlikely given that we found no evidence of an association despite correcting for regression dilution. Furthermore, reproducibility substudies of $\mathrm{Lp}(\mathrm{a})$ in the KIHD and that of other large-scale cohort studies[3] indicate that analyses using only single baseline measurements of $\mathrm{Lp}(\mathrm{a})$ does not underestimate the associations between $\operatorname{Lp}(\mathrm{a})$ and outcomes. There is established evidence that $\operatorname{Lp}(a)$ is associated with CVD outcomes and it has been suggested that the pathophysiological mechanisms underlying the associations may relate to the pro-atherogenic, prothrombotic, and pro-inflammatory properties of $\operatorname{Lp}(a)$ [7]. Given the prothrombotic and antifibrinolytic properties of $\mathrm{Lp}(\mathrm{a})$ [2], the closely linked nature of CVD and VTE [10-12],
and the emerging evidence from both epidemiological and clinical studies; there is a growing debate that $\operatorname{Lp}(a)$ may also be linked to the development of VTE. The current data which is based on prospective evidence does not support this suggestion and it is possible that $\mathrm{Lp}(\mathrm{a})$ may not be an emerging risk factor for VTE. Spence and Koschinsky also argue that the effects of $\operatorname{Lp}(a)$ on VTE risk may only be evident at the highest concentrations of $\operatorname{Lp}(a)$ [51], which we were not able to prove in the current study. However, mechanistic conclusions underlying the association between $\mathrm{Lp}(\mathrm{a})$ and VTE cannot be drawn from observational epidemiological studies and further studies on mechanisms are warranted.

## Strengths and limitations

Compared to previous prospective cohort studies, the current study had the advantage of being a well-characterised cohort of men who were nationally representative; involved a high response rate, a long-follow-up period of over 20 years with no loss to follow-up; and comprehensive analysis with adjustment for a broad panel of lifestyle and biological markers as well as stratified analyses by several clinical relevant characteristics. An important strength of the current study is that repeat measurements of $\mathrm{Lp}(\mathrm{a})$ made within a random subset of individuals over time after baseline were available, which enabled correction for the extent of within-person variability in $\operatorname{Lp}(a)$ over the long period of follow-up. Finally, we were able to conduct a pooled analysis of previous studies including the current study, to put the findings into wider context. In our pooled analysis, there was no evidence of heterogeneity between contributing studies. Our study was characterized by the following limitations: (i) we included only middle-aged men based on a predominantly white-European population from eastern Finland and given that plasma levels of $\mathrm{Lp}(\mathrm{a})$ may vary substantially between different populations [52], our findings therefore cannot be generalized to women, the young, and other ethnicities; (ii) we had data on only all VTEs which precluded the ability to conduct subgroup analyses of type of specific VTE outcomes such as idiopathic VTE or that due to cancer; and (iii) inability to adjust for other potential confounders such incident cancer, family history of VTE, and use of antithrombotic drugs.

## Conclusions

Primary cohort analysis as well as pooled evidence from previous studies suggest that circulating $\operatorname{Lp}(a)$ is not prospectively associated with future VTE risk. This comprehensive report indicates that the associations demonstrated in previous studies may be driven by features and limitations of study designs employed.

## Acknowledgements

We thank the staff of the Kuopio Research Institute of Exercise Medicine and the Research Institute of Public Health and University of Eastern Finland, Kuopio, Finland for the data collection in the study.

## Disclosure statement

No potential conflict of interest was reported by the authors.

## Funding sources

Prof. Jari Laukkanen acknowledges support from The Finnish Foundation for Cardiovascular Research, Helsinki, Finland. Dr. Kunutsor acknowledges support from the NIHR Biomedical Research Centre at University Hospitals Bristol NHS Foundation Trust and the University of Bristol. The views expressed in this publication are those of the authors and not necessarily those of the NHS, the National Institute for Health Research or the Department of Health and Social Care. These sources had no role in design and conduct of the study; collection, management, analysis, and interpretation of the data; and preparation, review, or approval of the manuscript.

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Table 1. Baseline participant characteristics of the KIHD cohort overall and by quartiles of lipoprotein(a)

|  | Overall $(\mathrm{N}=\mathbf{2 , 1 8 0})$ <br> Mean (SD) or median (IQR) or n (\%) | Quartile 1 Mean (SD) or median (IQR) or n (\%) | Quartile 2 Mean (SD) or median (IQR) or n (\%) | Quartile 3 Mean (SD) or median (IQR) or n (\%) | Quartile 4 Mean (SD) or median (IQR) or n (\%) | $\begin{aligned} & \text { Pearson correlation } \\ & \quad \mathbf{r}(\mathbf{9 5 \%} \mathbf{C I}) \dagger \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Lipoprotein(a) (mg/dl) | 9.66 (3.75-22.27) | 1.61 (0.90-2.73) | 6.23 (4.88-7.84) | 15.0 (12.3-18.5) | 36.8 (28.5-51.8) | - |
| Questionnaire/Prevalent conditions |  |  |  |  |  |  |
| Age at survey (years) | 53 (5) | 53 (5) | 53 (5) | 53 (5) | 53 (5) | 0.01 (-0.03, 0.05) |
| Alcohol consumption ( $\mathrm{g} / \mathrm{week}$ ) | 31.9 (6.4-92.8) | 30.1 (6.4-96.0) | 28.4 (6.1-88.4) | 32.9 (6.1-96.9) | 35.5 (7.6-88.5) | -0.00 (-0.05, 0.04) |
| History of diabetes | 87 (4.0) | 31 (5.7) | 26 (4.8) | 20 (3.7) | 10 (1.8) | - |
| Current smokers | 669 (30.7) | 161 (29.5) | 165 (30.2) | 168 (30.9) | 175 (32.1) | - |
| History of hypertension | 653 (30.0) | 178 (32.7) | 162 (29.7) | 148 (27.2) | 165 (30.3) | - |
| History of CHD | 546 (25.1) | 157 (28.8) | 131 (24.0) | 122 (22.4) | 136 (25.0) | - |
| History of cancer | 36 (1.7) | 11 (2.0) | 11 (2.0) | 7 (1.3) | 7 (1.3) | - |
| Lipid lowering medication | 14 (0.6) | 5 (0.9) | 2 (0.4) | 1 (0.2) | 6 (1.1) | - |
| Physical measurements |  |  |  |  |  |  |
| BMI (kg/m²) | 26.9 (3.5) | 27.6 (3.9) | 26.7 (3.4) | 26.8 (3.3) | 26.6 (3.4) | -0.10 (-0.14, -0.06)*** |
| SBP (mmHg) | 134 (17) | 135 (17) | 133 (16) | 134 (17) | 133 (17) | -0.04 (-0.08, -0.00) |
| DBP (mmHg) | 89 (10) | 89 (11) | 88 (10) | 89 (11) | 88 (10) | -0.04 (-0.08, 0.00) |
| Physical activity (kj/day) | 1192 (621-1987) | 1231 (662-1991) | 1160 (669-1998) | 1104 (58.-1891) | 1275 (612-2021) | -0.03 (-0.07, 0.02) |
| Lipid markers |  |  |  |  |  |  |
| Total cholesterol ( $\mathrm{mmol} / \mathrm{l}$ ) | 5.91 (1.08) | 5.73 (1.02) | 5.88 (1.09) | 5.88 (1.12) | 6.12 (1.05) | $0.12(0.08,0.16)^{* * *}$ |
| HDL-C (mmol/l) | 1.30 (0.30) | 1.31 (0.32) | 1.30 (0.28) | 1.30 (0.30) | 1.29 (0.30) | -0.01 (-0.05, 0.03) |
| Triglycerides ( $\mathrm{mmol} / \mathrm{l}$ ) | 1.10 (0.81-1.56) | 1.15 (0.84-1.68) | 1.08 (0.78-1.58) | 1.10 (0.82-1.52) | 1.10 (0.79-1.55) | -0.06 (-0.10, -0.02)* |
| Metabolic, renal, and inflammatory markers |  |  |  |  |  |  |
| Fasting plasma glucose ( $\mathrm{mmol} / \mathrm{l}$ ) | 5.35 (1.26) | 5.50 (1.52) | 5.34 (1.20) | 5.34 (1.19) | 5.23 (1.07) | -0.07 (-0.11, -0.03)** |
| Serum creatinine ( $\mu \mathrm{mol} / 1$ ) | 89.4 (13.7) | 88.6 (12.1) | 89.1 (13.5) | 89.0 (14.9) | 91.0 (14.0) | 0.04 (0.00, 0.08)* |
| Estimated GFR (ml/min/1.73 m²) | 86.9 (17.1) | 87.4 (15.0) | 87.7 (20.5) | 87.5 (16.2) | 85.2 (16.2) | -0.03 (-0.08, 0.01) |
| Fibrinogen (g/l) | 2.96 (2.63-3.33) | 2.91 (2.58-3.28) | 2.95 (2.63-3.31) | 2.97 (2.68-3.32) | 3.00 (2.67-3.44) | 0.08 (0.03, 0.12)** |
| High sensitivity CRP (mg/l) | 1.27 (0.70-2.38) | 1.17 (0.61-2.25) | 1.20 (0.65-2.34) | 1.34 (0.76-2.37) | 1.35 (0.80-2.73) | $0.09(0.05,0.13) * * *$ |

BMI, body mass index; CHD, coronary heart disease; CRP, C-reactive protein; DBP, diastolic blood pressure; Lp(a),
lipoprotein(a)
HDL-C, high-density lipoprotein cholesterol; KIHD, Kuopio Ischemic Heart Disease; SD, standard deviation; SBP, systolic
blood pressure;
VTE, venous thromboembolism; asterisks indicate the level of statistical
significance: ${ }^{*}, \mathrm{p}<0.05 ; * *, \mathrm{p}<0.01 ; * * *, \mathrm{p}<0.001, \dagger$ Pearson correlation coefficients
between $\log _{\mathrm{e}} \mathrm{Lp}(\mathrm{a})$ and the row variables

Table 2. Association of $\operatorname{Lp}(a)$ and venous thromboembolism in the KIHD cohort

| Plasma Lp(a) (mg/dl) | Events/ Total | Person-time at risk | Model 1 | Model 2 |
| :---: | :---: | :---: | :---: | :---: |
|  |  |  | HR (95\% CI) | HR (95\% CI) |
| Baseline Lp(a) |  |  |  |  |
| Per 1 SD increase in log | 110/2,180 | 47,400 | 1.06 (0.88 to 1.29) | 1.06 (0.87 to 1.30$)$ |
| Lp(a) |  |  |  |  |
| Q1 (0.56-3.74) | $25 / 545$ | 11,810 | ref | ref |
| Q2 (3.75-9.66) | $30 / 546$ | 11,980 | 1.19 (0.70 to 2.03) | 1.24 (0.72 to 2.12) |
| Q3 (9.67-22.26) | $25 / 544$ | 11,914 | 0.99 (0.57 to 1.72) | 1.02 (0.58 to 1.79) |
| Q4 (> 22.26) | $30 / 545$ | 11,695 | 1.23 (0.72 to 2.09) | 1.25 (0.72 to 2.15) |
|  | Usual Lp(a)* |  |  |  |
| Per 1 SD increase in log | 110/2,180 | 47,400 | 1.07 (0.86 to 1.35) | 1.08 (0.85 to 1.36) |
| $\mathrm{Lp}(\mathrm{a})$ |  |  |  |  |
| Q1 (0.56-3.74) | 25 / 545 | 11,810 | ref | ref |
| Q2 (3.75-9.66) | $30 / 546$ | 11,980 | 1.23 (0.66 to 2.30) | 1.29 (0.68 to 2.42) |
| Q3 (9.67-22.26) | 25 / 544 | 11,914 | 0.98 (0.51 to 1.89) | 1.02 (0.52 to 1.98) |
| Q4 (> 22.26) | $30 / 545$ | 11,695 | 1.28 (0.68 to 2.38) | 1.29 (0.68 to 2.47) |

CI, confidence interval; HR, hazard ratio; KIHD, Kuopio Ischemic Heart Disease; Lp(a), lipoprotein(a); Q, quartile; ref, reference; SD , standard deviation;
*, indicates correction for within-person variability in values of $\operatorname{Lp}(a)$, that is, the extent to which an individual's Lp (a) measurements vary around a long-term average value ("usual Lp (a) values"); the SD of loge $\mathrm{Lp}(\mathrm{a})$ concentration is 1.27 , corresponding to approximately four-fold higher circulating $\operatorname{Lp}(a)\left(i e, e^{1.27}=3.56\right)$
Model 1: Adjusted for age
Model 2: Model 1 plus body mass index, systolic blood pressure, history of hypertension, prevalent coronary heart disease, smoking status, history of diabetes, total cholesterol, triglycerides, lipid medication, estimated glomerular filtration rate, physical activity, alcohol consumption, prevalent cancer, fibrinogen, and high sensitivity C-reactive protein

## Figure legends

Figure 1. Hazard ratios for baseline levels of lipoprotein(a) and venous thromboembolism risk by several participant level characteristics in the KIHD cohort


Hazard ratios are reported per 1 standard deviation increase in $\log _{\mathrm{e}}$ lipoprotein(a); hazard ratios were adjusted for age, body mass index, systolic blood pressure, history of hypertension, prevalent coronary heart disease, smoking status, history of diabetes, total cholesterol, triglycerides, lipid medication, estimated glomerular filtration rate, physical activity, alcohol consumption, prevalent cancer, fibrinogen, and high sensitivity C-reactive protein; CHD, coronary heart disease; CI, confidence interval; GFR, glomerular filtration rate; HDL, high-density lipoprotein; HR, hazard ratio; KIHD, Kuopio Ischemic Heart Disease; Lp(a) lipoprotein(a); SD, standard deviation; *, $P$-value for interaction

Figure 2. Prospective studies of lipoprotein(a) and risk of venous thromboembolism included in meta-analysis


The summary estimates presented were calculated using random effects models; relative risks are reported per 1 standard deviation (SD) increase in lipoprotein(a); sizes of data markers are proportional to the inverse of the variance of the relative ratio; CI, confidence interval (bars); RR, relative risk; VTE, venous thromboembolism

## SUPPLEMENTARY MATERIAL

| Appendix A | STROBE Statement |
| :--- | :--- |
| Appendix B | PRISMA checklist |
| Appendix C | MOOSE checklist |
| Appendix D | Literature search strategy |
| Appendix E | Flow of studies included in pooled analysis |
| Appendix F | Characteristics of prospective studies included in meta-analysis |

## Appendix A. STROBE Statement

| 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 | Page 1 |
|  |  | (b) Provide in the abstract an informative and balanced summary of what was done and what was found | Page 2 |
| Introduction |  |  |  |
| Background/rationale | 2 | Explain the scientific background and rationale for the investigation being reported | Page 3-4 |
| Objectives | 3 | State specific objectives, including any pre-specified hypotheses | Page 3-4 |
| Methods |  |  |  |
| Study design | 4 | Present key elements of study design early in the paper | Study design and population |
| Setting | 5 | Describe the setting, locations, and relevant dates, including periods of recruitment, exposure, follow-up, and data collection | Study design and population |
| Participants | 6 | (a) Give the eligibility criteria, and the sources and methods of selection of participants. Describe methods of follow-up | Study design and population |
|  |  | (b) For matched studies, give matching criteria and number of exposed and unexposed | Not applicable |
| Variables | 7 | Clearly define all outcomes, exposures, predictors, potential confounders, and effect modifiers. Give diagnostic criteria, if applicable | Assessment of $\operatorname{Lp}(a)$ and other risk markers |
| 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 | Assessment of $\operatorname{Lp}(\mathrm{a})$ and other risk markers |
| Bias | 9 | Describe any efforts to address potential sources of bias | Statistical analysis |
| Study size | 10 | Explain how the study size was arrived at | Statistical analysis |
| Quantitative variables | 11 | Explain how quantitative variables were handled in the analyses. If applicable, describe which groupings were chosen and why | Statistical analysis |
| Statistical methods | 12 | (a) Describe all statistical methods, including those used to control for confounding | Statistical analysis |
|  |  | (b) Describe any methods used to examine subgroups and interactions | Statistical analysis |
|  |  | (c) Explain how missing data were addressed | Not applicable |
|  |  | (d) If applicable, explain how loss to follow-up was addressed | Not applicable |
|  |  | (e) Describe any sensitivity analyses | Statistical analysis |
| 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 | Study design and population |



## Appendix B. PRISMA check-list

| Section/topic | Item No | Checklist item | Reported on page No |
| :---: | :---: | :---: | :---: |
| Title |  |  |  |
| Title | 1 | Identify the report as a systematic review, meta-analysis, or both | 1 |
| Abstract |  |  |  |
| Structured summary | 2 | Provide a structured summary including, as applicable, background, objectives, data sources, study eligibility criteria, participants, interventions, study appraisal and synthesis methods, results, limitations, conclusions and implications of key findings, systematic review registration number | 2 |
| Introduction |  |  |  |
| Rationale | 3 | Describe the rationale for the review in the context of what is already known | 4 |
| Objectives | 4 | Provide an explicit statement of questions being addressed with reference to participants, interventions, comparisons, outcomes, and study design (PICOS) | 4 |
| Methods |  |  |  |
| Protocol and registration | 5 | Indicate if a review protocol exists, if and where it can be accessed (such as web address), and, if available, provide registration information including registration number | Methods |
| Eligibility criteria | 6 | Specify study characteristics (such as PICOS, length of follow-up) and report characteristics (such as years considered, language, publication status) used as criteria for eligibility, giving rationale | Methods |
| Information sources | 7 | Describe all information sources (such as databases with dates of coverage, contact with study authors to identify additional studies) in the search and date last searched | Methods |
| Search | 8 | Present full electronic search strategy for at least one database, including any limits used, such that it could be repeated | Appendix D |
| Study selection | 9 | State the process for selecting studies (that is, screening, eligibility, included in systematic review, and, if applicable, included in the meta-analysis) | Methods |
| Data collection process | 10 | Describe method of data extraction from reports (such as piloted forms, independently, in duplicate) and any processes for obtaining and confirming data from investigators | Methods |
| Data items | 11 | List and define all variables for which data were sought (such as PICOS, funding sources) and any assumptions and simplifications made | Methods |
| Risk of bias in individual studies | 12 | Describe methods used for assessing risk of bias of individual studies (including specification of whether this was done at the study or outcome level), and how this information is to be used in any data synthesis | Methods |
| Summary measures | 13 | State the principal summary measures (such as risk ratio, difference in means). | Methods |
| Synthesis of results | 14 | Describe the methods of handling data and combining results of studies, if done, including measures of consistency (such as $\mathrm{I}^{2}$ statistic) for each meta-analysis | Methods |
| Risk of bias across studies | 15 | Specify any assessment of risk of bias that may affect the cumulative evidence (such as publication bias, selective reporting within studies) | Methods |
| Additional analyses | 16 | Describe methods of additional analyses (such as sensitivity or subgroup analyses, metaregression), if done, indicating which were pre-specified | Methods |
| Results |  |  |  |
| Study selection | 17 | Give numbers of studies screened, assessed for eligibility, and included in the review, with reasons for exclusions at each stage, ideally with a flow diagram | Appendix E |
| Study characteristics | 18 | For each study, present characteristics for which data were extracted (such as study size, PICOS, follow-up period) and provide the citations | Appendix F |
| Risk of bias | 19 |  | Appendix F |

Risk of bias within studies
Results of
individual
studies
Synthesis of
results
Risk of bias across studies
Additional analysis

## Discussion

Summary of evidence

3 Describe the rationale for the review in the context of what is already known 4
4 Provide an explicit statement of questions being addressed with reference to participants, 4 interventions, comparisons, outcomes, and study design (PICOS)

5 Indicate if a review protocol exists, if and where it can be accessed (such as web address), Methods and, if available, provide registration information including registration number characteristics (such as years considered, language, publication status) used as criteria for eligibility, giving rationale
7 Describe all information sources (such as databases with dates of coverage, contact with
Methods

Appendix D

Methods

Methods

Methods

Methods

Methods
Methods

Methods

Methods

Appendix E

Appendix F

Appendix F
Figure 2

Results and Figure 2
22 Present results of any assessment of risk of bias across studies (see item 15) Not applicable

23 Give results of additional analyses, if done (such as sensitivity or subgroup analyses, meta- Not applicable regression) (see item 16)

24 Summarise the main findings including the strength of evidence for each main outcome; Discussion consider their relevance to key groups (such as health care providers, users, and policy makers)

|  | Item |  |  |
| :--- | :---: | :--- | :--- |
| Section/topic | No | Checklist item | Reported <br> on page No |
| Limitations | 25 | Discuss limitations at study and outcome level (such as risk of bias), and at review level <br> (such as incomplete retrieval of identified research, reporting bias) | Discussion |
| Conclusions | 26 | Provide a general interpretation of the results in the context of other evidence, and <br> implications for future research | Discussion |
| Funding <br> Funding | 27 | Describe sources of funding for the systematic review and other support (such as supply of <br> data) and role of funders for the systematic review | None |

## Appendix C. MOOSE checklist

| Criteria |  | Brief description of how the criteria were handled in the review |
| :---: | :---: | :---: |
| Reporting of background |  |  |
| $\checkmark$ | Problem definition | Elevated circulating lipoprotein(a) has been suggested to the linked to the development of venous thromboembolism (VTE), but the prospective nature of the association is uncertain. |
| $\checkmark$ | Hypothesis statement | There is no prospective association between $\mathrm{Lp}(\mathrm{a})$ and VTE risk. |
| $\sqrt{ }$ | Description of study outcomes | VTE |
| $\sqrt{ }$ | Type of exposure | Blood levels of Lp(a) |
| $\checkmark$ | Type of study designs used | Prospective (cohort, case-cohort or "nested case control") populationbased studies |
| $\checkmark$ | Study population | Approximately general populations with no prevalent VTE at baseline |
| Reporting of search strategy should include |  |  |
| $\checkmark$ | Qualifications of searchers | Setor Kunutsor, MD PhD; Jari Laukkanen, MD |
| $\checkmark$ | Search strategy, including time period included in the synthesis and keywords | Time period: from inception of MEDLINE, EMBASE, Web of Science to 18 July 2018. <br> Search strategy: <br> In Appendix 4. |
| $\sqrt{ }$ | Databases and registries searched | MEDLINE, EMBASE, and Web of Science |
| $\sqrt{ }$ | Search software used, name and version, including special features | Ovid was used to search EMBASE Endnote used to manage references |
| $\sqrt{ }$ | Use of hand searching | We searched bibliographies of retrieved papers |
| $\checkmark$ | List of citations located and those excluded, including justifications | Details of the literature search process are outlined in the flow chart. The citation list for excluded studies is available upon request. |
| $\checkmark$ | Method of addressing articles published in languages other than English | We placed no restrictions on language |
| $\checkmark$ | Method of handling abstracts and unpublished studies | None found |
| $\checkmark$ | Description of any contact with authors | Not applicable |
| Reporting of methods should include |  |  |
| $\checkmark$ | Description of relevance or appropriateness of studies assembled for assessing the hypothesis to be tested | Detailed inclusion and exclusion criteria are described in the Methods section. |
| $\checkmark$ | Rationale for the selection and coding of data | Data extracted from each of the studies were relevant to the population characteristics, study design, exposure, outcome, and possible effect modifiers of the association. |
| $\checkmark$ | Assessment of confounding | We assessed confounding by ranking individual studies on the basis of different adjustment levels and performed sub-group analyses to evaluate differences in the overall estimates according to levels of adjustment. |
| $\checkmark$ | Assessment of study quality, including blinding of quality assessors; stratification or regression on possible predictors of study results | Study quality was assessed based on the nine-star Newcastle-Ottawa Scale using pre-defined criteria namely: population representativeness, comparability (adjustment of confounders), ascertainment of outcome. Sensitivity analyses by several quality indicators such as study size, duration of follow-up, and adjustment factors. |
| $\checkmark$ | Assessment of heterogeneity | Heterogeneity of the studies was explored with $\mathrm{I}^{2}$ statistic that provides the relative amount of variance of the summary effect due to the between-study heterogeneity. |
| $\checkmark$ | Description of statistical methods in sufficient detail to be replicated | Description of methods of meta-analyses are detailed in the methods. We performed random effects meta-analysis with Stata 15. |


| $\sqrt{ }$ | Provision of appropriate tables and <br> graphics | Figure 2 and Appendix F |
| :--- | :--- | :--- |
| Reporting of results should include |  |  |
| $\sqrt{ }$ | Graph summarizing individual study <br> estimates and overall estimate | Figure 2 |
| $\sqrt{ }$ | Table giving descriptive information <br> for each study included | Appendix F |
| $\sqrt{ }$ | Results of sensitivity testing | Not applicable |
| $\sqrt{ }$ | Indication of statistical uncertainty of <br> findings | $95 \%$ confidence intervals were presented with all summary estimates, <br> I <br> values and results of sensitivity analyses |
| Reporting of discussion should include | There was no evidence of heterogeneity between contributing studies. |  |
| $\sqrt{ }$ | Quantitative assessment of bias | All studies were excluded based on the pre-defined inclusion criteria <br> in methods section. |
| $\sqrt{ }$ | Justification for exclusion | Brief discussion included in 'Methods' section |
| $\sqrt{ }$ | Assessment of quality of included <br> studies | Discussed in the context of the results. <br> Reporting of conclusions should include <br> $\sqrt{ }$Consideration of alternative <br> explanations for observed results |
| $\sqrt{ }$ | Generalization of the conclusions | Discussed in the context of the results. |
| $\sqrt{ }$ | Guidelines for future research | Assessment of the mechanistic pathways underlying the link between <br> Lp(a) and VTE |
| $\sqrt{ }$ | Disclosure of funding source | No separate funding was necessary for the undertaking of this <br> systematic review. |

## Appendix D. Literature search strategy

Relevant studies published before 18 July, 2018 (date last searched), were identified through electronic searches not limited to the English language using MEDLINE, EMBASE, and the Science Citation Index databases. Electronic searches were supplemented by scanning reference lists of articles identified for all relevant studies (including review articles) and by hand searching of relevant journals. The computer-based searches combined search terms related to lipoprotein(a) and venous thromboembolism without language restriction.

1 exp "Lipoprotein(a)"/ (4776)
2 exp Venous Thromboembolism/ (8367)
3 exp Venous Thrombosis/ (51541)
4 exp Pulmonary Embolism/ (36297)
$5 \quad 2$ or 3 or 4 (86213)
$6 \quad 1$ and 5 (79)
7 limit 6 to humans (79)

Parts i, ii and iii were combined using 'AND' to search MEDLINE. Each part was specifically translated for searching alternative databases.

Appendix E. Flow of studies included in pooled analysis


4 Articles included, based on 4 unique studies

Appendix F. Characteristics of prospective studies included in meta-analysis

| Lead author, publication year [Reference] | Name of study | Location of study | Year(s) of baseline survey | Baseline age (years) | $\begin{aligned} & \% \\ & \text { male } \end{aligned}$ | Mean/median duration of follow-up (years) | Total no. of participants | No. of VTE cases | Covariates adjusted for | Study quality |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Tsai, 2002 [21] | LITE | USA | 1987-1998 | 59.0 | 45.0 | 8.0 | 19,293 | 215 | Age, race, and sex | 8 |
| Kamstrup, 2012 [20] | CCHS | Denmark | 1991-1994 | 58.0 | 44.0 | 13.0 | 9,138 | 440 | Age, sex, BMI, smoking, physical activity, menopausal status, HRT, oral contraceptives | 9 |
| van Schouwenburg, 2012 [50] | PREVEND | Netherlands | 1997-1998 | 49.0 | 49.0 | 10.5 | 7,627 | 110 | Age, sex, hypertension, DM, CRP, BMI, eGFR, smoking | 7 |
| Danik, 2013 [49] | WHS | USA | 1992-1995 | 54.2 | 0.0 | 14.4 | 28,345 | 439 | Age, smoking, BMI, hormone therapy status, exercise level and randomization treatment arms | 7 |
| Current study, 2018 | KIHD | Finland | 1984-1989 | 42-61 | 100.0 | 24.9 | 2,180 | 110 | Age, BMI, SBP, history of hypertension, prevalent CHD, smoking, history of DM, total cholesterol, Triglycerides, lipid medication, eGFR, physical activity, alcohol consumption, prevalent cancer, fibrinogen, hsCRP | 8 |
| Total |  |  |  |  |  |  | 66,583 | 1,314 |  |  |

BMI, body mass index; CHD, coronary heart disease; CCHS, Copenhagen City Heart Study; DM, diabetes mellitus; eGFR, estimated glomerular filtration rate; HRT, hormone replacement therapy; hsCRP, high sensitivity C-reactive protein; KIHD, Kuopio Ischemic Heart Disease; LITE, The Longitudinal Investigation of Thromboembolism Etiology; NR, not reported; PREVEND, Prevention of Renal and Vascular End-stage Disease; SBP, systolic blood pressure; WHS, Women's Health Study

