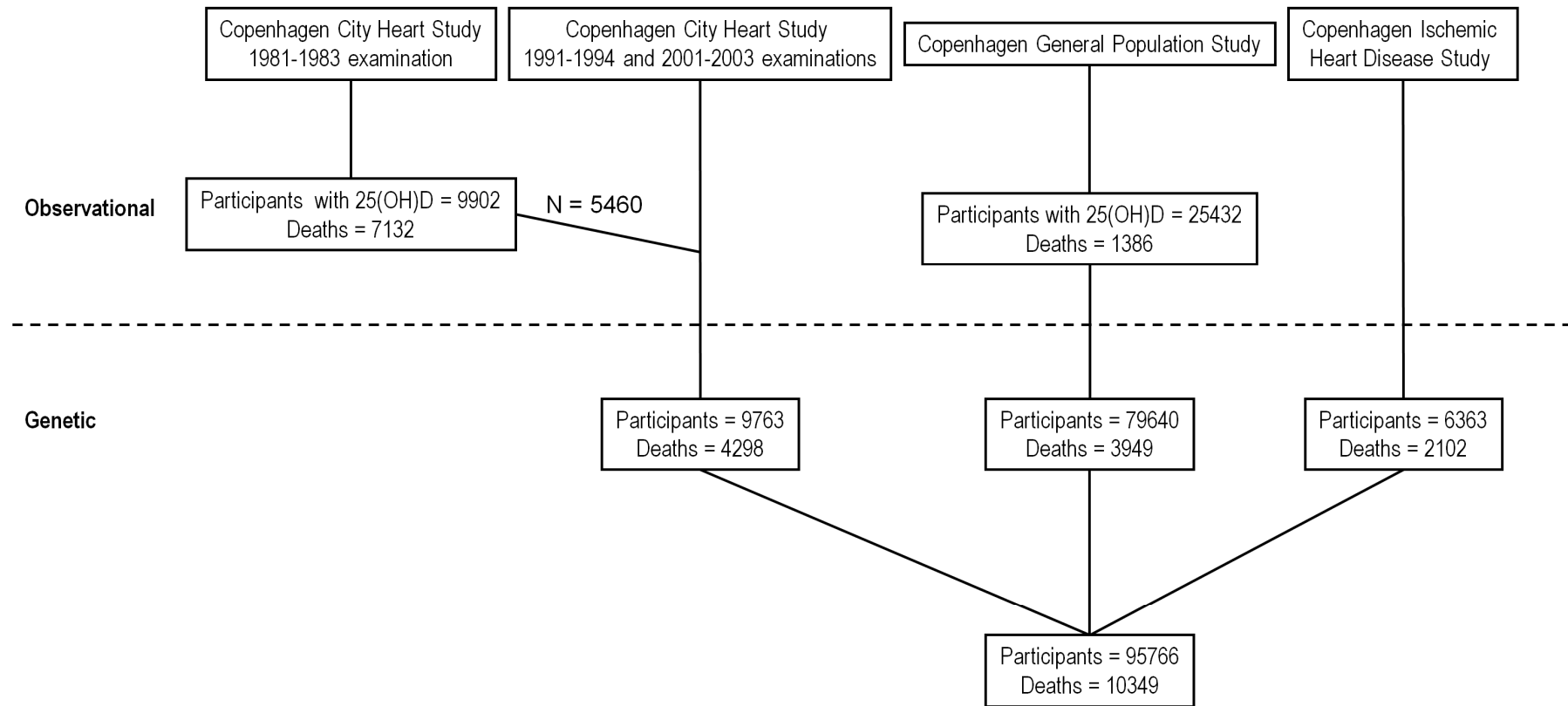


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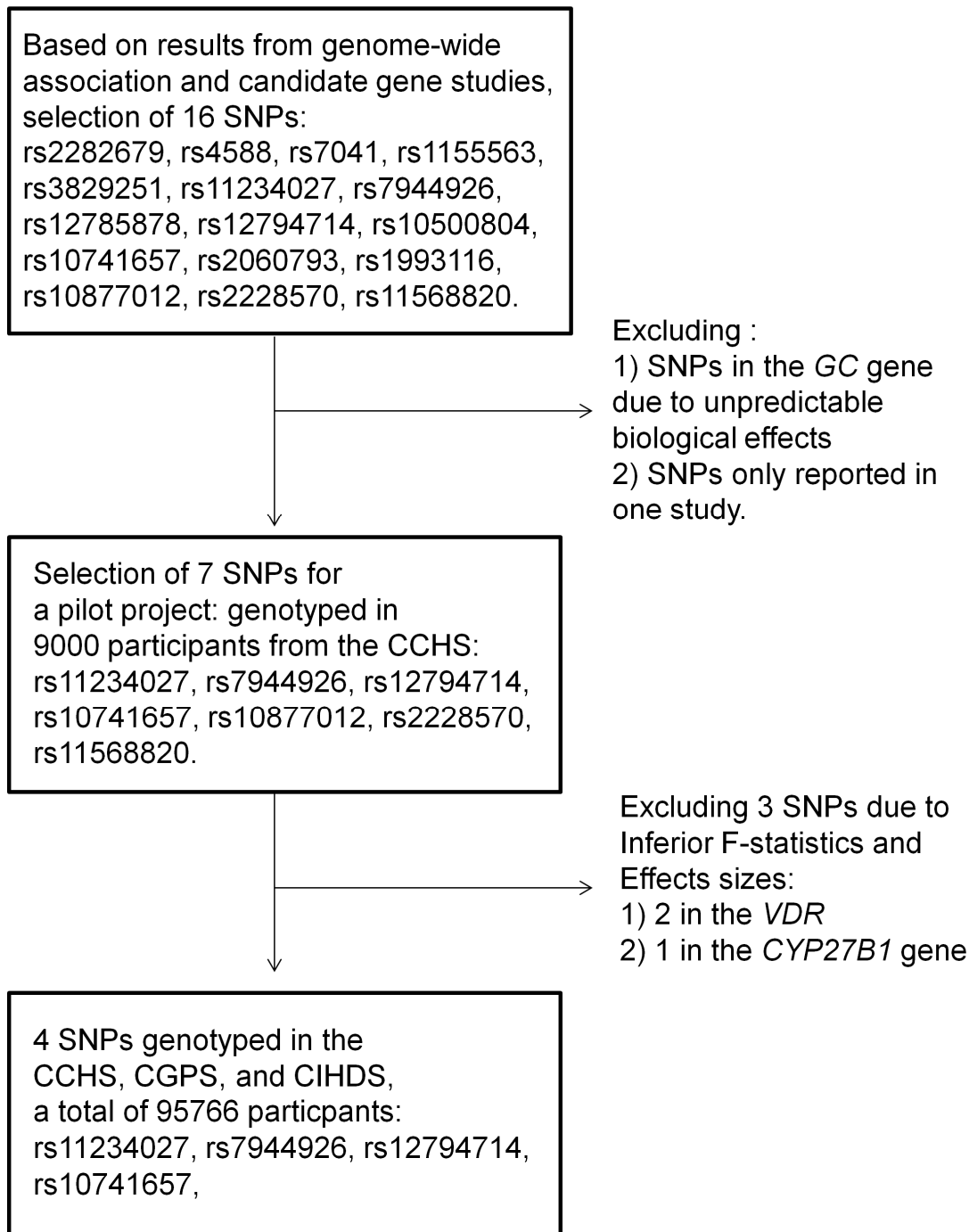
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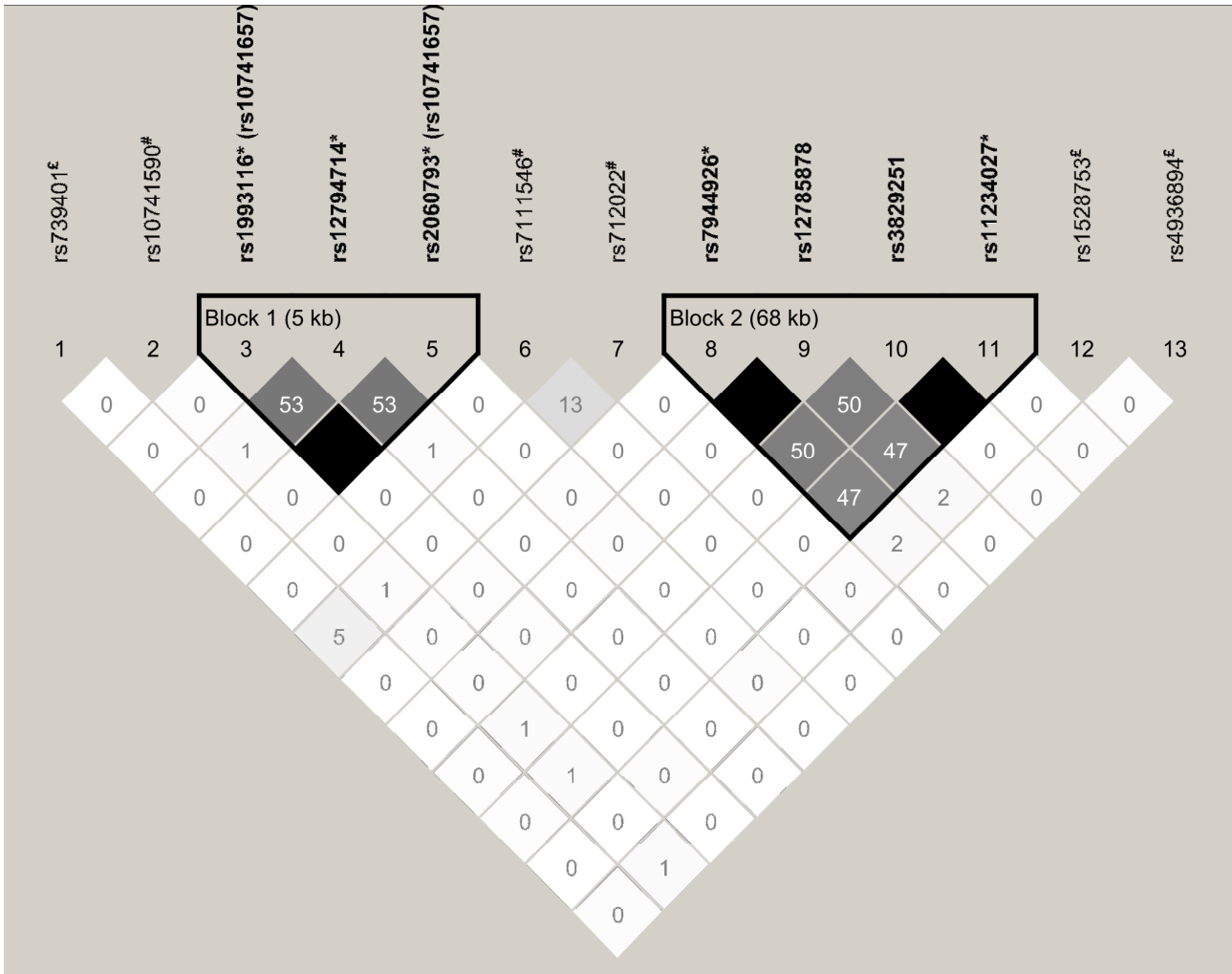
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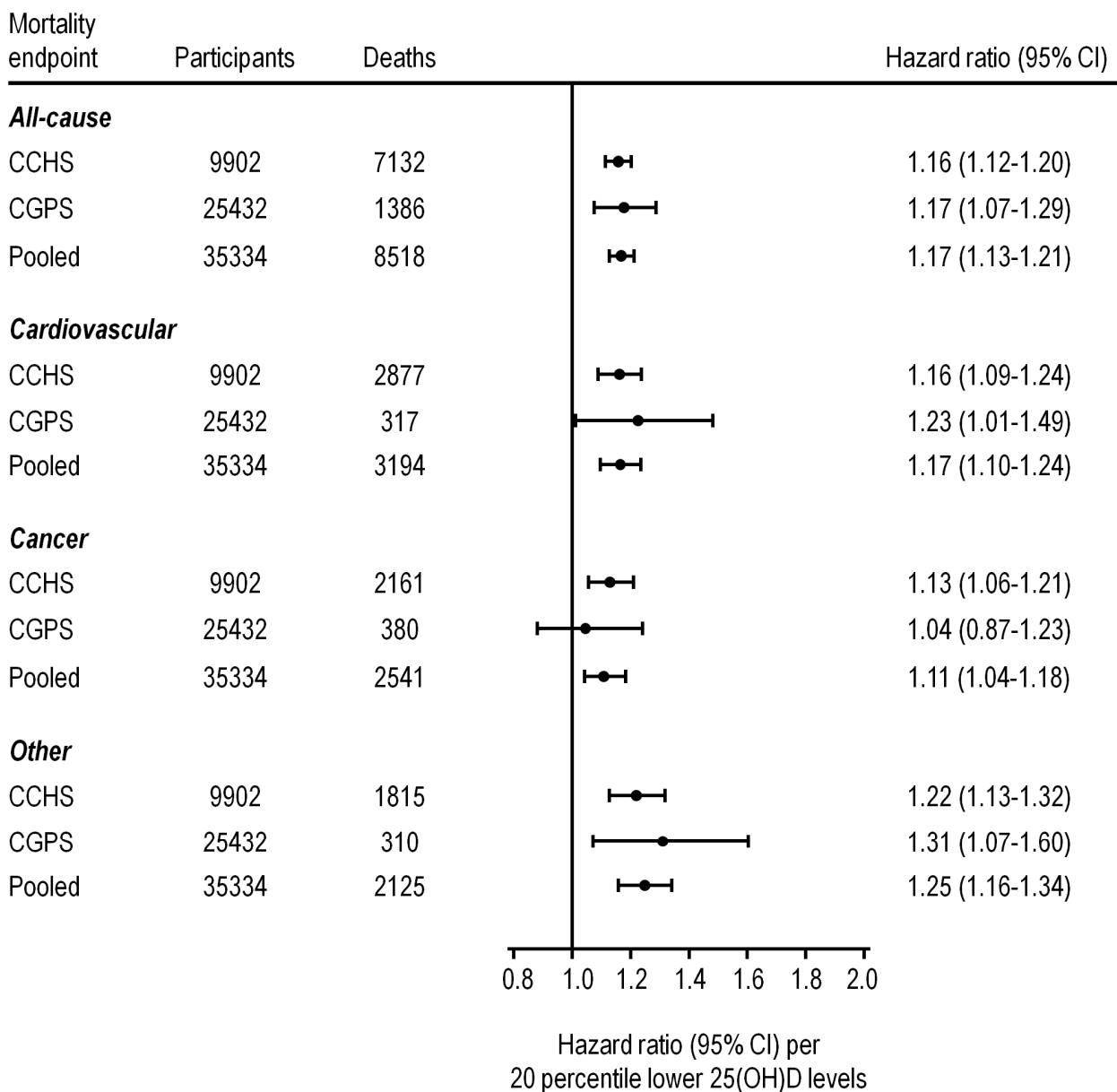
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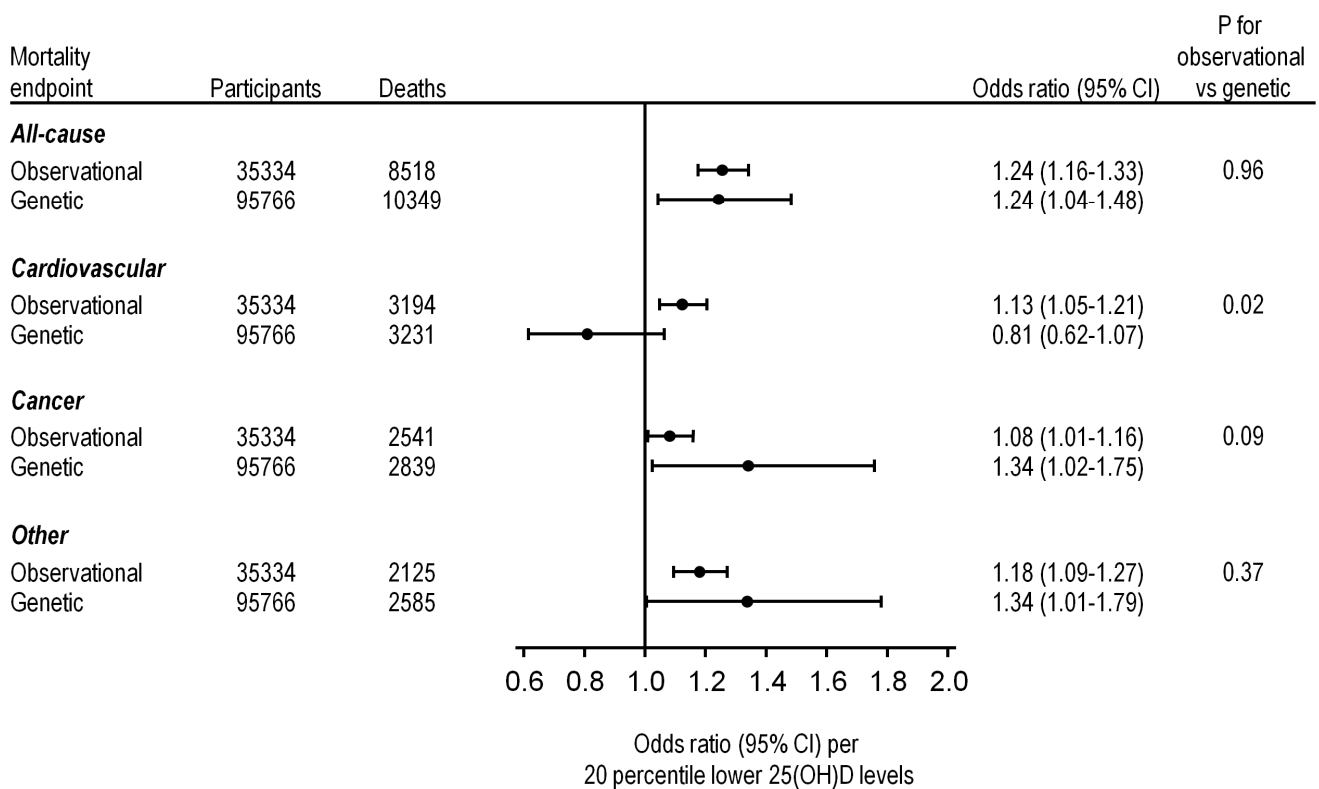
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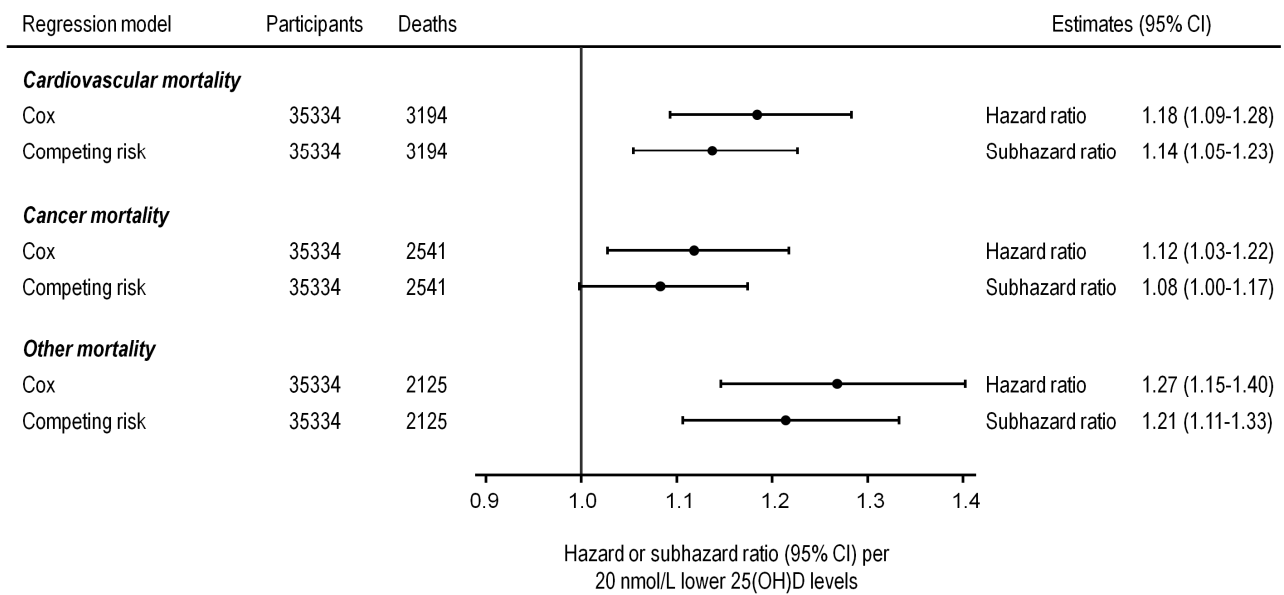
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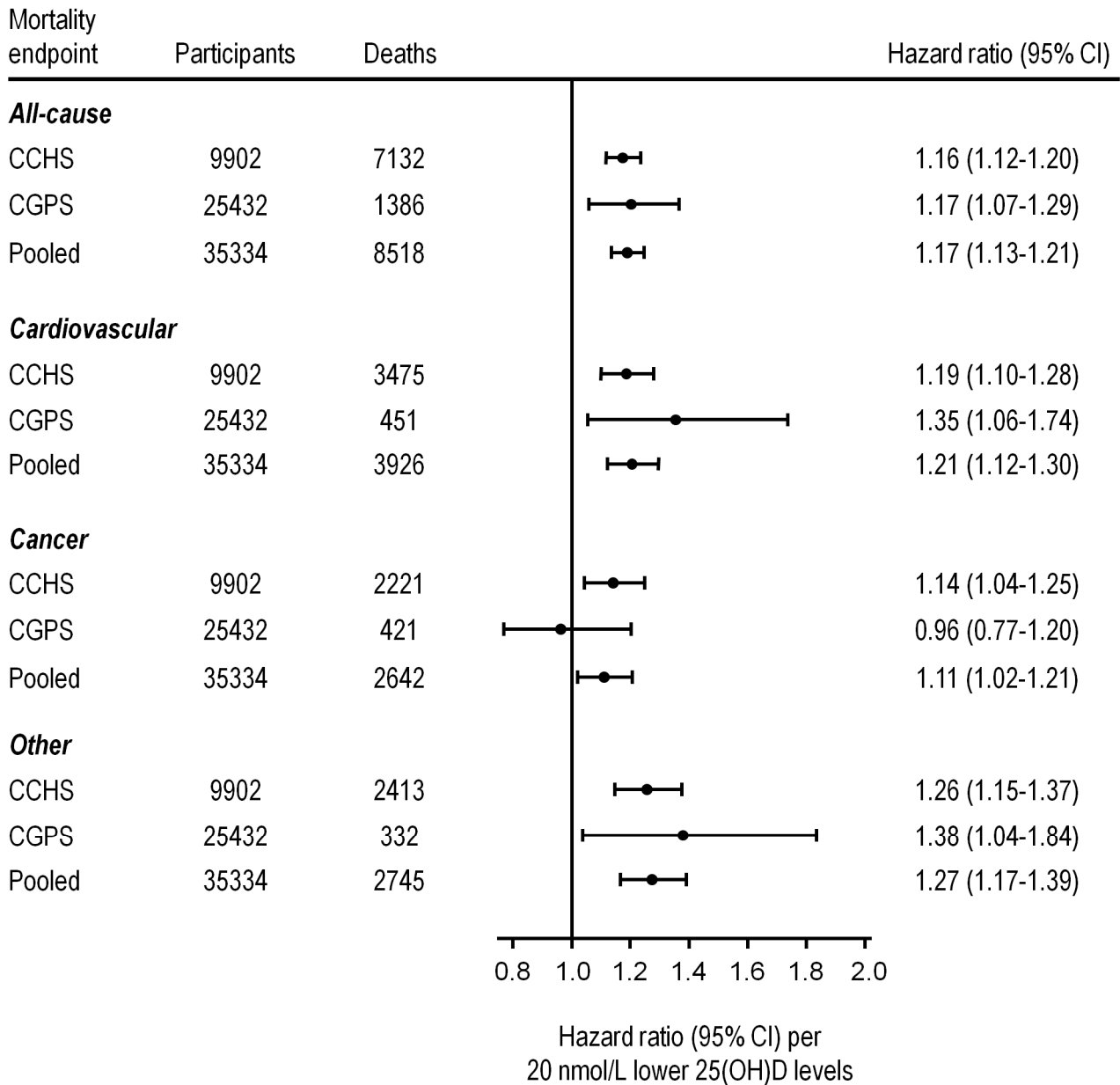
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Supplementary figure E. Instrumental variable analysis using deseasonalised plasma 25-hydroxyvitamin D. Observational and genetic risk estimates for all-cause and cause-specific mortality for 20 percentile lower 25-hydroxyvitamin D levels. Observational estimates were by logistic regression and genetic estimates by instrumental variable analyses. Observational analyses were adjusted for age, sex, smoking status, cumulative tobacco consumption, alcohol consumption, leisure time physical activity, systolic blood pressure, body mass index, income, diabetes, plasma cholesterol, and study. Genetic analyses were adjusted for age, year of birth, sex, and study. Observational estimates were based on participants from the Copenhagen City Heart Study and Copenhagen General Population Study combined, while genetic estimates were based on participants from the Copenhagen City Heart Study, Copenhagen General Population Study, and Copenhagen Ischemic Heart Disease Study combined. 25(OH)D = 25-hydroxyvitamin D. CI = confidence interval.

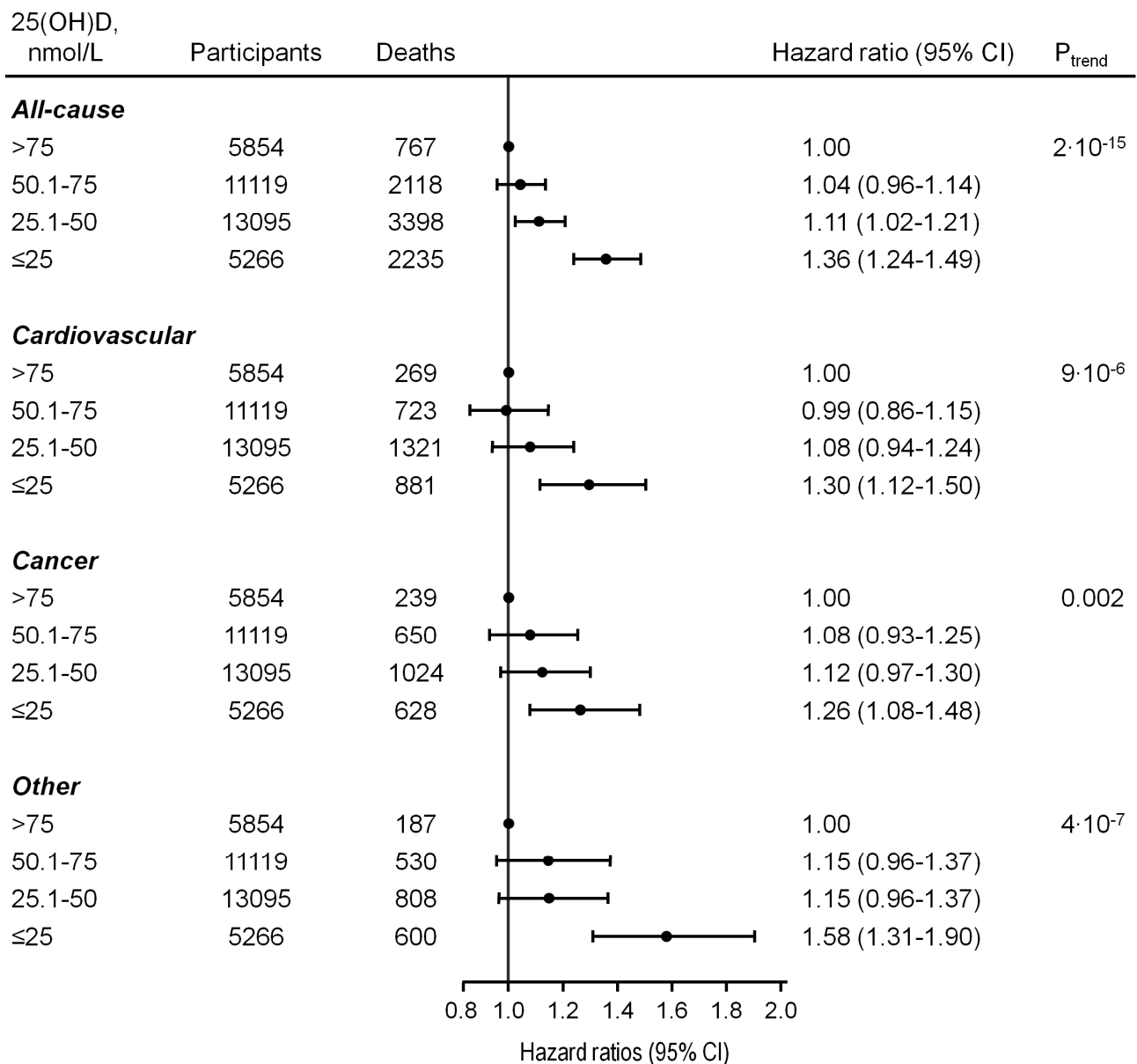


Supplementary figure F. The association of plasma 25-hydroxyvitamin D with cause-specific mortality in the general population using Cox and competing risks regression. The analyses were carried out using Cox regression or Fine and Gray's competing risk proportional subhazard models adjusted for age, sex, smoking status, cumulative tobacco consumption, alcohol consumption, leisure time physical activity, systolic blood pressure, body mass index, income, diabetes, plasma cholesterol, season (month and year of blood sample), and study. 25(OH)D = 25-hydroxyvitamin D. CI = confidence interval.

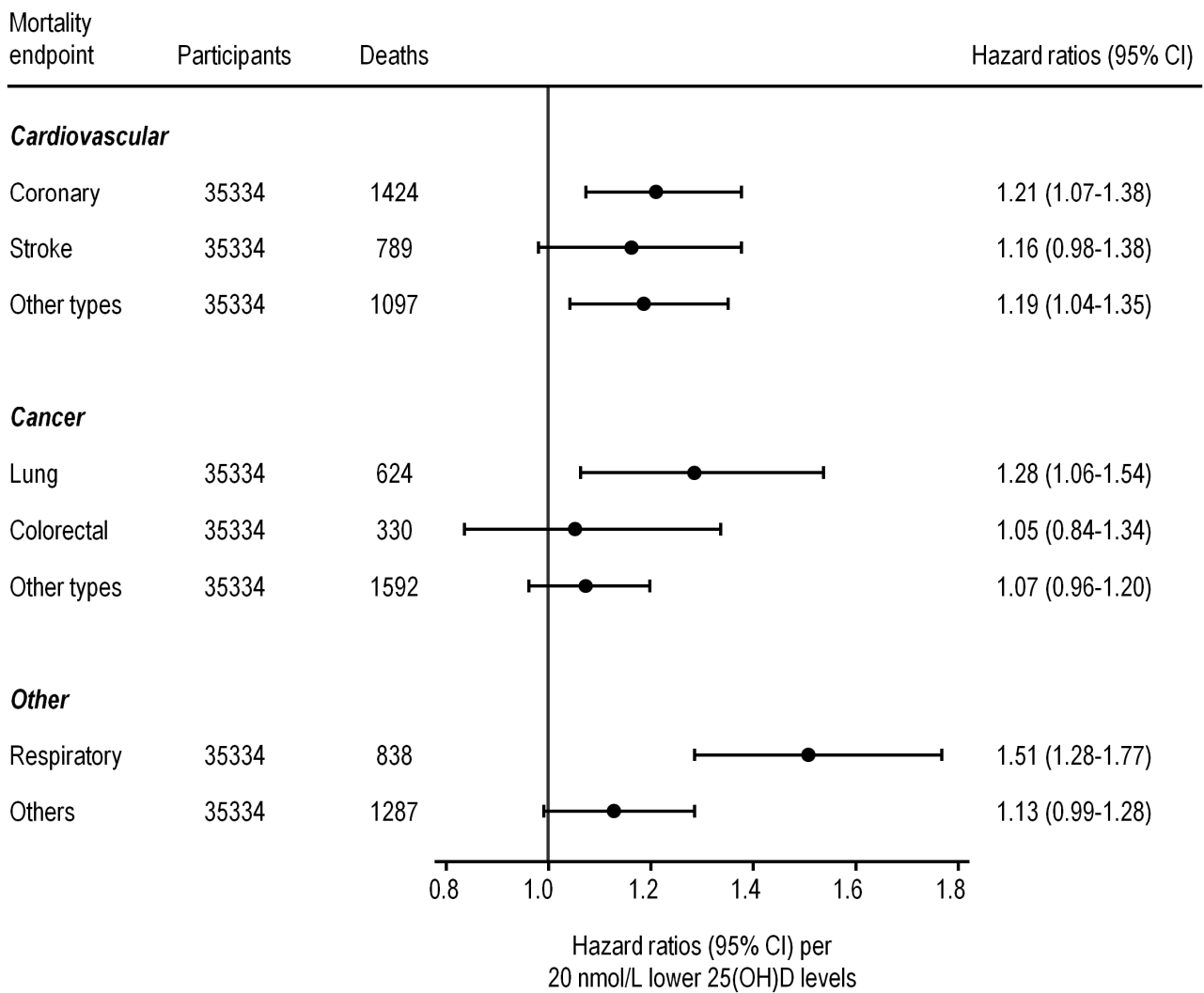


Supplementary figure G. The association of plasma 25-hydroxyvitamin D with cause-specific mortality in the general population using an alternative classification of cause-specific mortality.

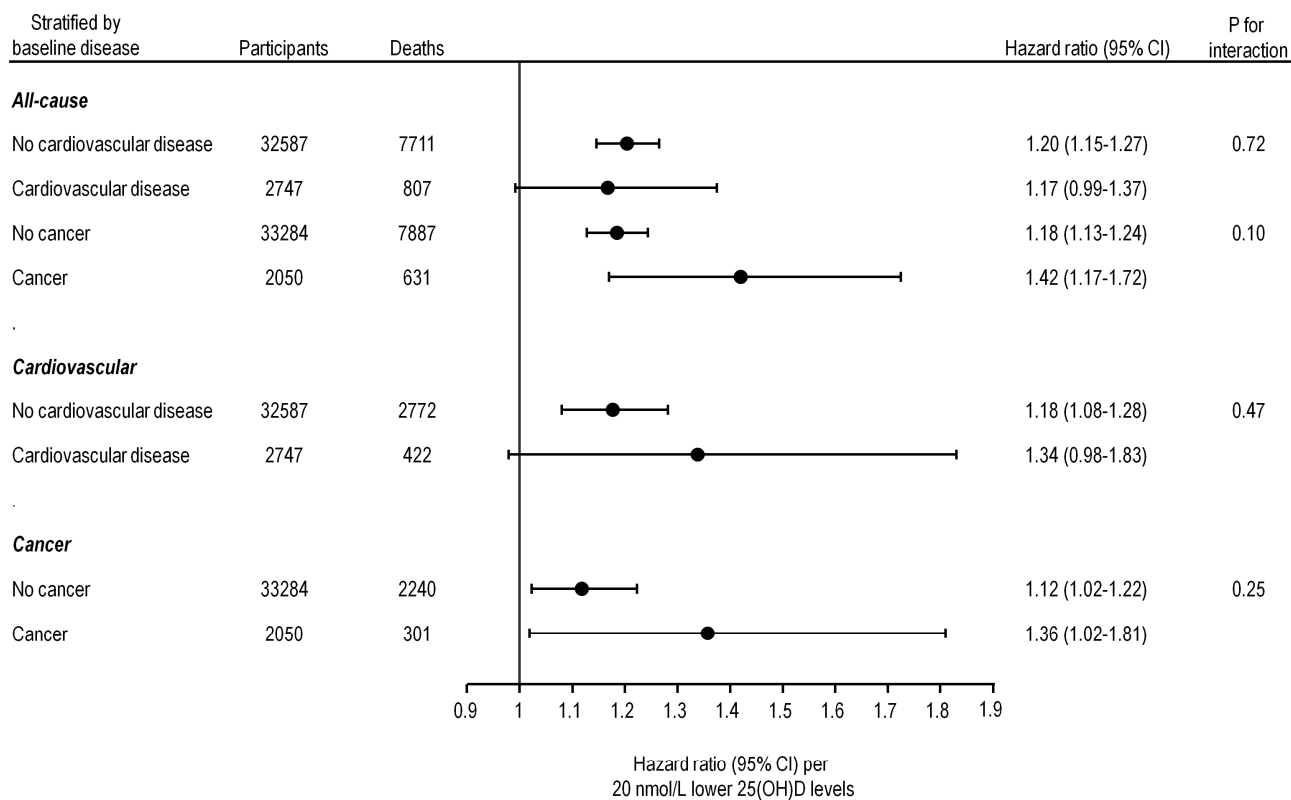
The regression model was identical to the main analyses; however, endpoints were classified in an overlapping manner, e.g., if there was both a cardiovascular and a cancer diagnosis on a death certificate that person would be included in analyses of both endpoints. Furthermore, all diagnoses were used and not only the three ranked diagnoses per person. There were up to 32 years of follow-up in the Copenhagen City Heart Study whereas it was up to 9.4 years in the Copenhagen General Population Study. 25(OH)D = 25-hydroxyvitamin D. CI = confidence interval. CCHS = Copenhagen City Heart Study. CGPS = Copenhagen General Population Study.



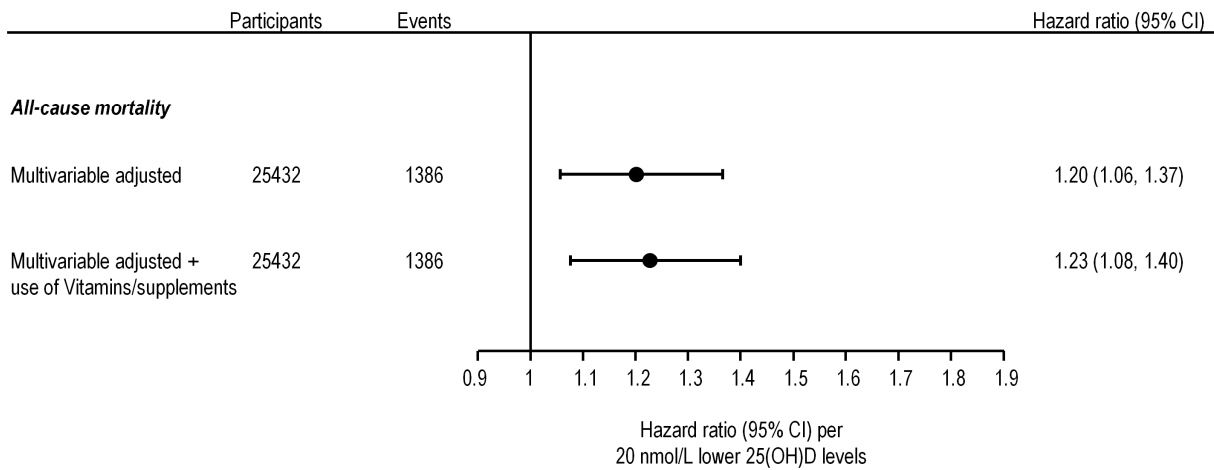
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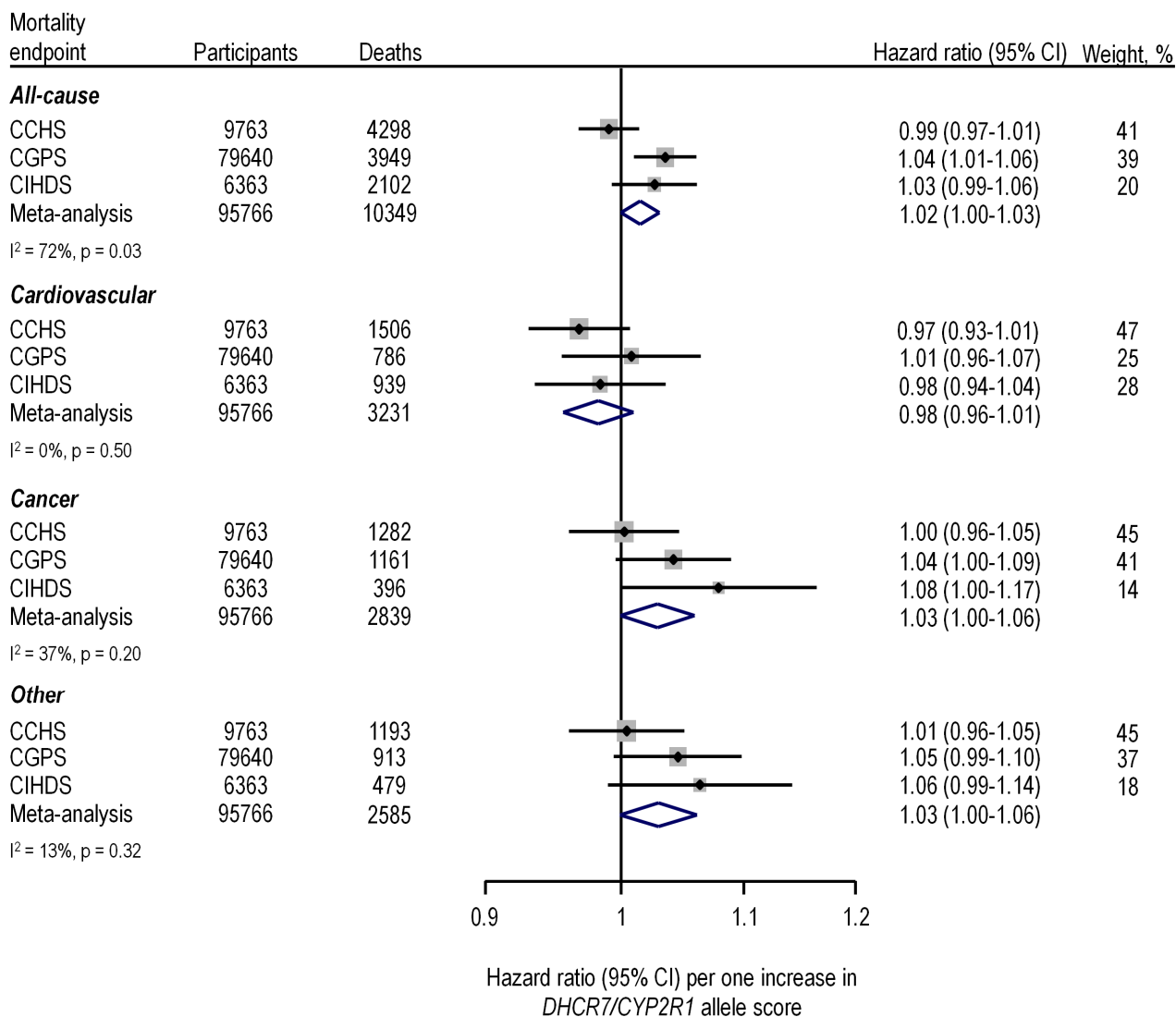
Supplementary figure I. The association of plasma 25-hydroxyvitamin D with cause-specific mortality using further subdivision of mortality endpoints. The analyses were by Cox regression adjusted for age, sex, smoking status, cumulative tobacco consumption, alcohol consumption, leisure time physical activity, systolic blood pressure, body mass index, income, diabetes, plasma cholesterol, season (month and year of blood sample), and study. 25(OH)D = 25-hydroxyvitamin D. CI = confidence interval. **Cardiovascular (ICD-8: 390-458, ICD-10: I00-I99).** Coronary (ICD-8: 410-414, ICD-10: I20-I25). Stroke (ICD-8: 430-438, ICD-10: I60-I69). **Cancer (ICD-8: 140-209, ICD-10: C00-C97).** Lung (ICD-8: 162, ICD-10: C33-C34). Colorectal (ICD-8: 153-154, ICD-10: C18-C21). **Other (remaining ICD diagnoses).** Respiratory (ICD-8: 460-519, ICD-10: J00-J99).



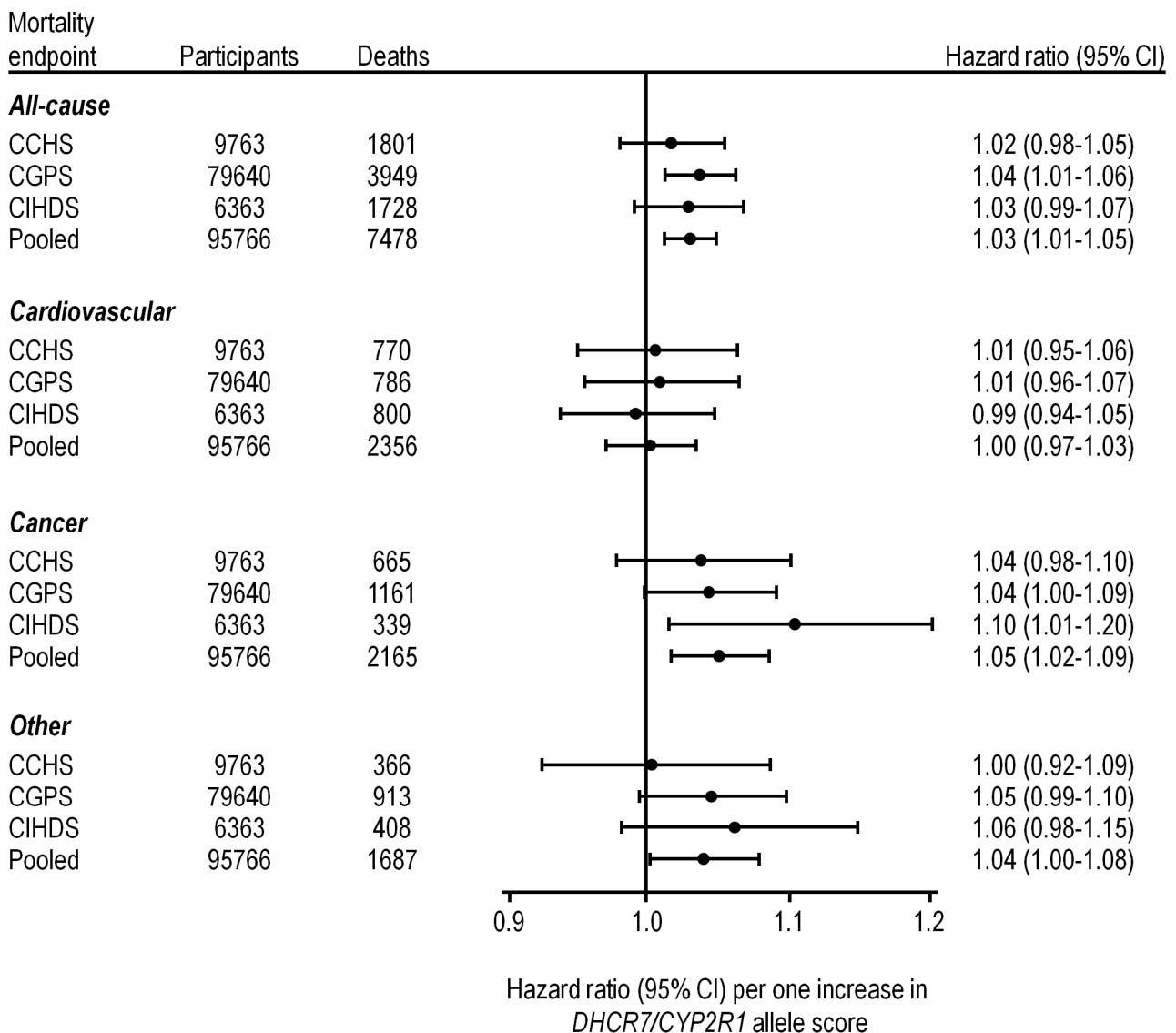
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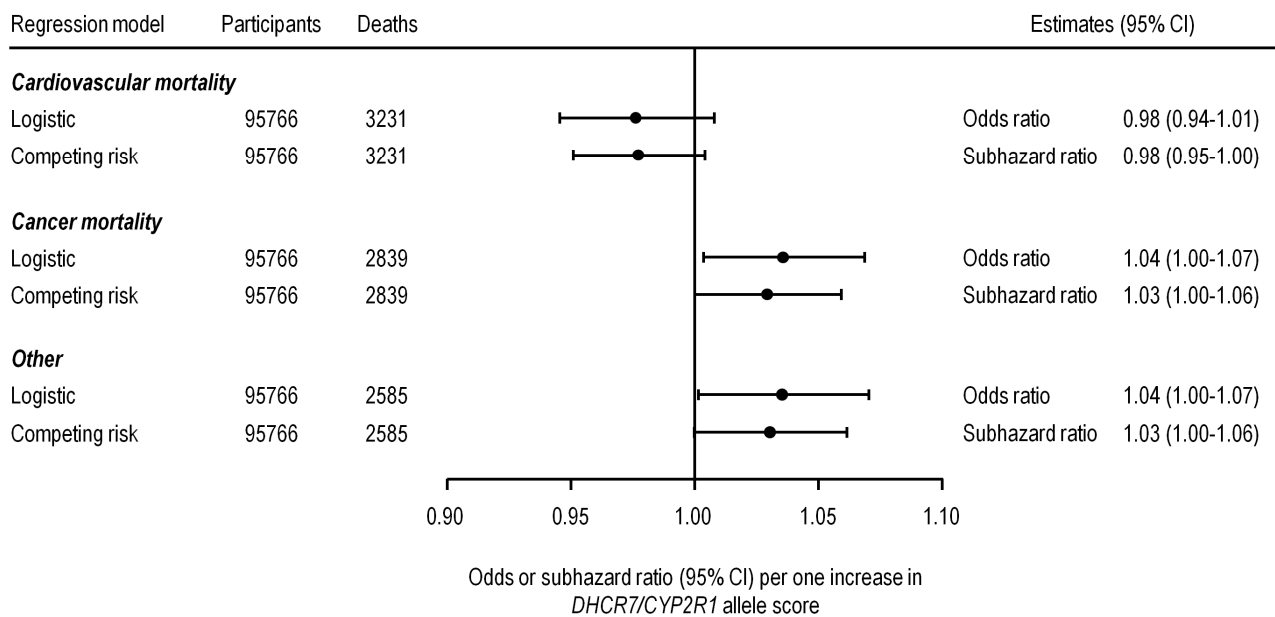
Supplementary figure K. The association of plasma 25-hydroxyvitamin D with all-cause mortality in analyses with and without adjustment for vitamin or nutritional supplement intake in CGPS. The analyses were by Cox regression adjusted for age, sex, smoking status, cumulative tobacco consumption, alcohol consumption, leisure time physical activity, systolic blood pressure, body mass index, income, diabetes, plasma cholesterol, season (month and year of blood sample), and study. 25(OH)D = 25-hydroxyvitamin D. CI = confidence interval.



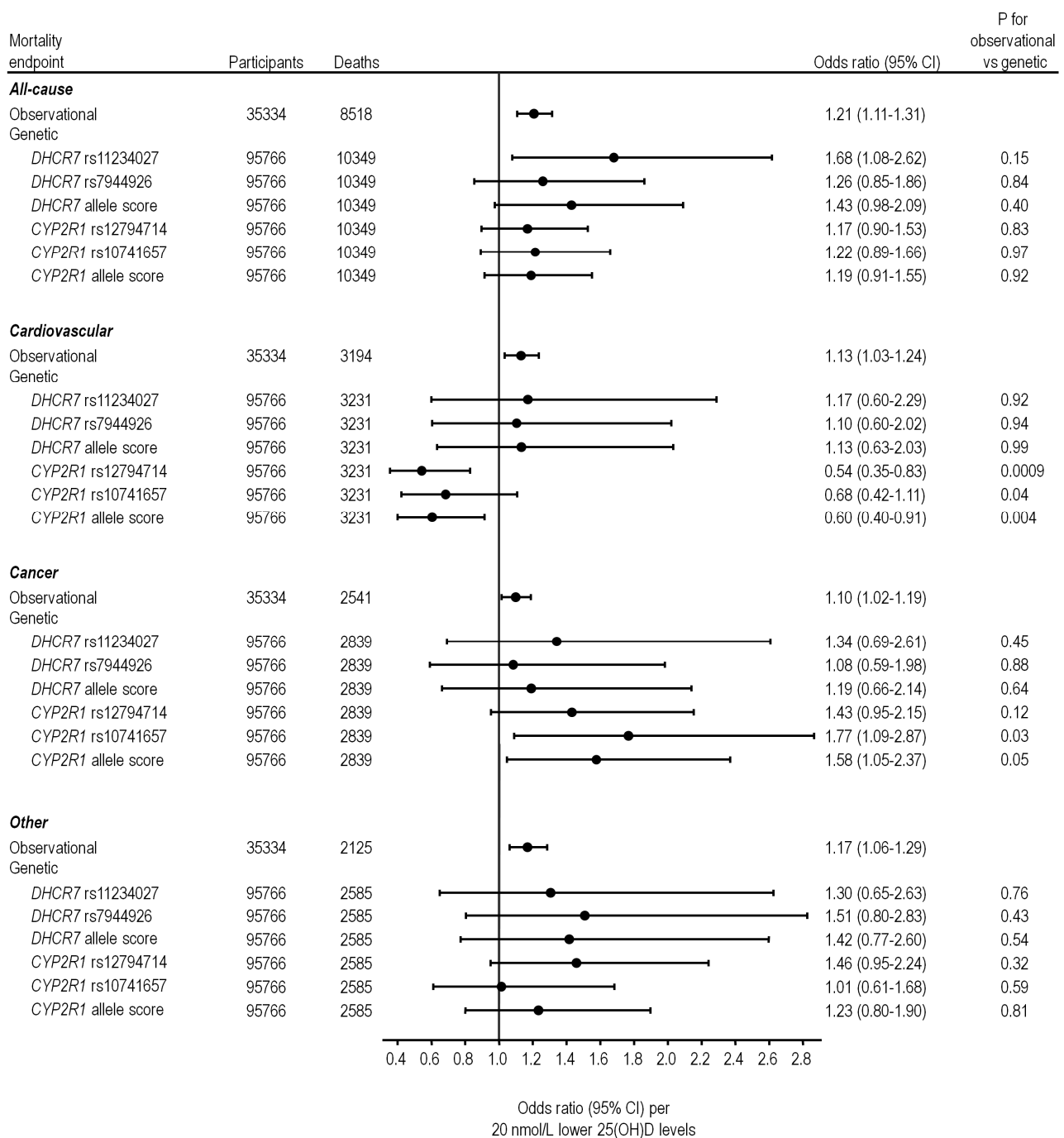
Supplementary figure L. All-cause and cause-specific mortality according to the *DHCR7/CYP2R1* allele score using meta-analysis with fixed effect estimates instead of pooling of the studies. The analyses were carried out using Cox regression adjusted for age, year of birth, and sex. 25(OH)D = 25-hydroxyvitamin D. CI = confidence interval. CCHS = Copenhagen City Heart Study. CGPS = Copenhagen General Population Study. CIHDS = Copenhagen Ischemic Heart Disease Study.



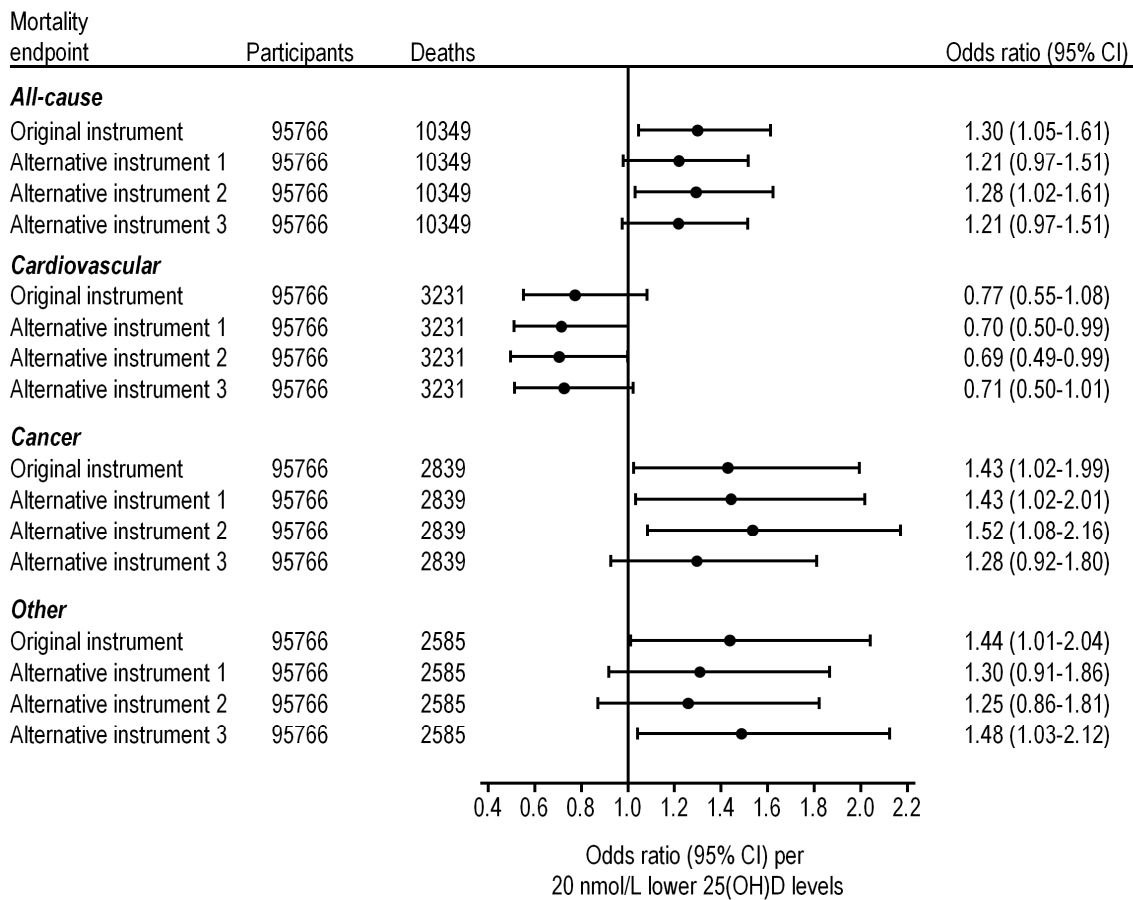
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Supplementary figure N. All-cause and cause-specific mortality according to the *DHCR7/CYP2R1* allele score using logistic or competing risks regression. The analyses were carried out using logistic regression or Fine and Gray's competing risk proportional subhazard models adjusted for age, year of birth, sex, and study; the latter only in the pooled analyses. 25(OH)D = 25-hydroxyvitamin D. CI = confidence interval.



Supplementary figure O. All-cause and cause-specific mortality when using each genotype as an instrument separately. Genetic estimates were by instrumental variable analyses for 20 nmol/L lower 25-hydroxyvitamin D levels. 25(OH)D = 25-hydroxyvitamin D. CI = confidence interval.



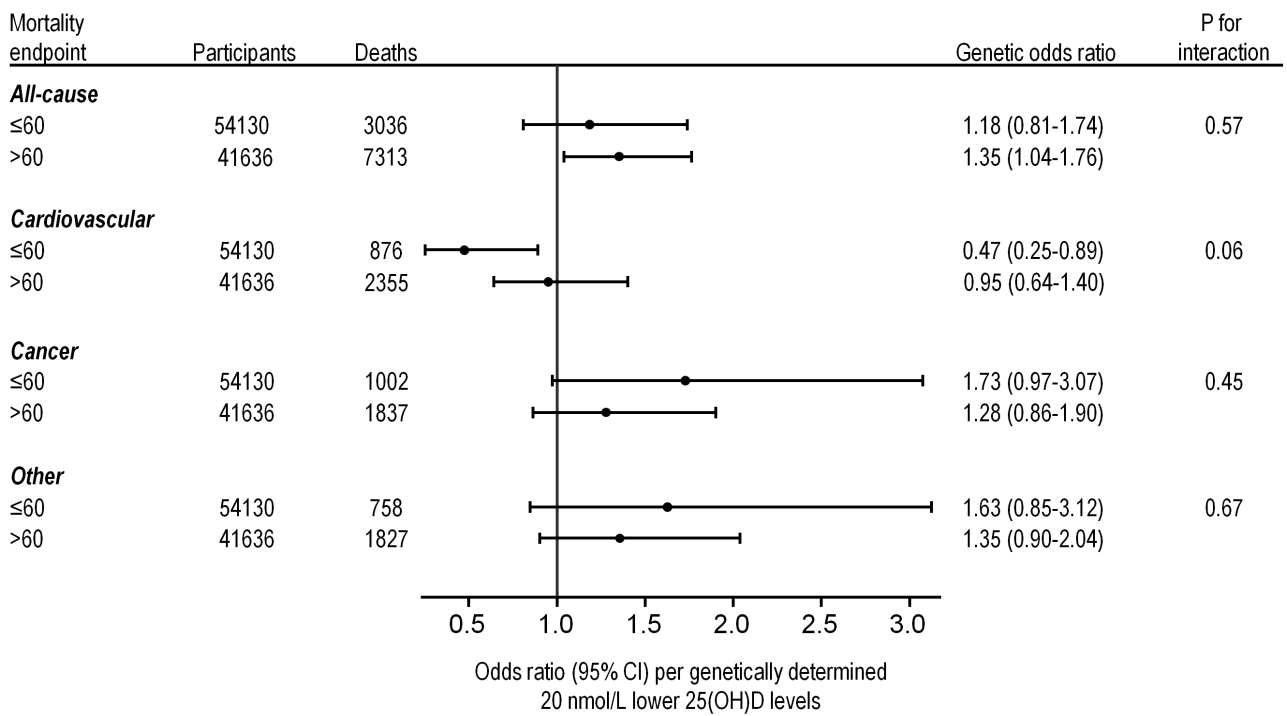
Supplementary figure P. All-cause and cause-specific mortality when using alternative definitions of the instrument. Genetic estimates were by instrumental variable analyses for 20 nmol/L lower 25-hydroxyvitamin D levels.

25(OH)D = 25-hydroxyvitamin D. CI = confidence interval.

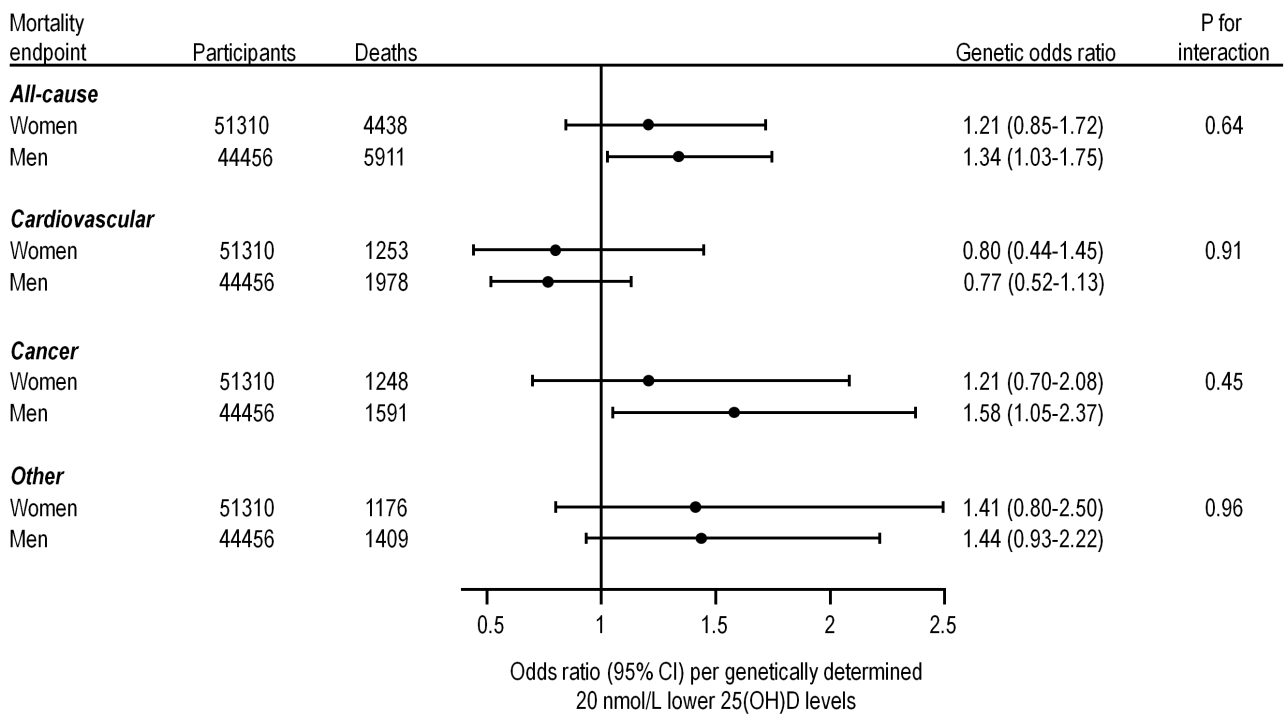
Alternative instrument 1 = allele score based on *DHCR7* rs7944926 + *CYP2R1* rs10741657 + *CYP2R1* rs12794714

Alternative instrument 2 = allele score based on *DHCR7* rs11234027 + *CYP2R1* rs10741657 + *CYP2R1* rs12794714

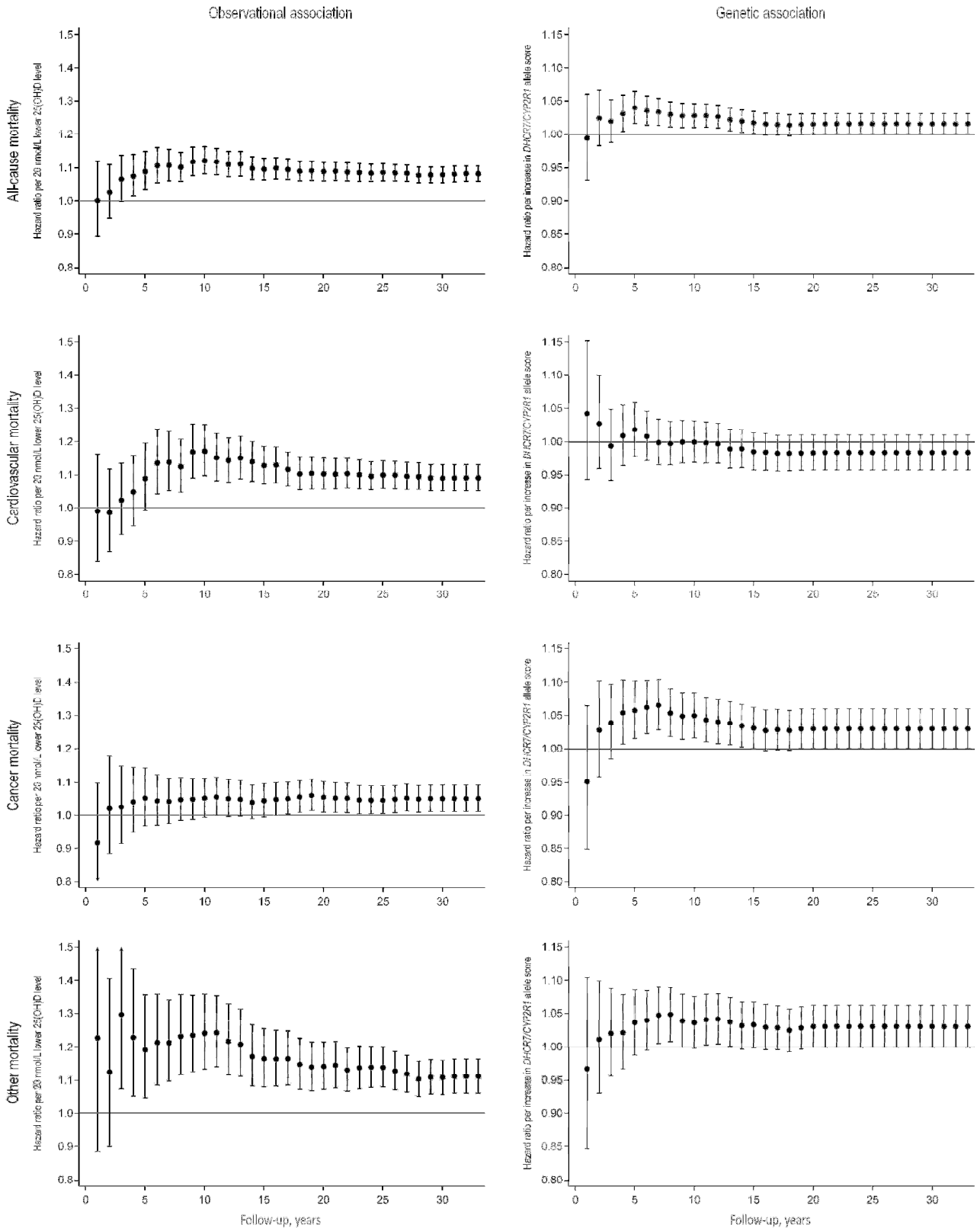
Alternative instrument 3 = allele score based on *DHCR7* rs7944926 + *CYP2R1* rs12794714



Supplementary figure Q. Genetic risk estimates for all-cause and cause-specific mortality according to age stratified analyses. Genetic estimates were by instrumental variable analyses for a 20 nmol/L lower 25-hydroxyvitamin D levels. Genetic estimates were based on participants from the Copenhagen City Heart Study, Copenhagen General Population Study, and Copenhagen Ischemic Heart Disease Study combined. 25(OH)D = 25-hydroxyvitamin D. CI = confidence interval.



Supplementary figure R. Genetic risk estimates for all-cause and cause-specific mortality according to sex stratified analyses. Genetic estimates were by instrumental variable analyses for a 20 nmol/L lower 25-hydroxyvitamin D levels. Genetic estimates were based on participants from the Copenhagen City Heart Study, Copenhagen General Population Study, and Copenhagen Ischemic Heart Disease Study combined. 25(OH)D = 25-hydroxyvitamin D. CI = confidence interval.



Supplementary figure S. Observational and genetic analyses according to restriction of follow-up from 1 to 32 years to evaluate the dependence of risk estimates on follow-up time. In observational analyses, Cox regression models were adjusted for age, sex, smoking status, cumulative tobacco consumption, alcohol consumption, leisure time physical activity, systolic blood pressure, body mass index, income,

diabetes, plasma cholesterol, season, and study. Observational estimates were based on participants from the Copenhagen City Heart Study and Copenhagen General Population Study combined (n = 35334). The genetic risk factor was one increase in *DHCR7/CYP2R1* allele score. In genetic analyses, models were adjusted for age, year of birth, sex, and study. Genetic estimates were based on participants from the Copenhagen City Heart Study, Copenhagen General Population Study, and Copenhagen Ischemic Heart Disease Study combined (n = 95766). Filled dots represent the point estimates and the whiskers represent the 95% confidence intervals. 25(OH)D = 25-hydroxyvitamin D.

Tables

Supplementary table A. Baseline characteristics in the general population and their association with age, sex, season, and study adjusted 25-hydroxyvitamin D (25(OH)D) quintiles.

	Adjusted 25(OH)D quintiles					P values ^a
	1 st (Lowest) N = 7068	2 nd N = 7066	3 rd N = 7068	4 th N = 7066	5 th (Highest) N = 7066	
Age, years	58 (49-66)	57 (48-66)	58 (49-67)	59 (49-67)	58 (49-67)	0.62
Men, %	43	46	47	45	43	0.75
Current smoker, %	40	33	29	27	26	2·10 ⁻¹⁰⁰
Cumulative tobacco consumption, pack years ^c	24 (12-38)	20 (10-34)	19 (8-34)	18 (8-32)	18 (8-32)	9·10 ⁻⁴²
Alcohol consumption, units/week ^d	6 (2-14)	7 (2-14)	7 (3-14)	7 (3-14)	7 (3-14)	0.12
Leisure-time physical activity <2 hours/week, %	16	10	9	7	6	3·10 ⁻⁸⁵
Systolic blood pressure, mmHg	139 (125-153)	136 (122-150)	136 (123-150)	135 (121-150)	133 (120-148)	8·10 ⁻³²
Body mass index, kg/m ²	26 (23-30)	26 (23-29)	25 (23-28)	25 (23-28)	24 (22-27)	1·10 ⁻²²³
Low income, % ^e	46	39	38	37	36	4·10 ⁻³⁰
Diabetes, %	6	4	4	3	3	4·10 ⁻¹³
Cholesterol, mmol/L	5.8 (5.1-6.6)	5.8 (5.0-6.4)	5.7 (5.0-6.4)	5.6 (4.9-6.4)	5.4 (4.7-6.2)	1·10 ⁻¹¹⁹
Vitamin D, nmol/L	23 (16-32)	38 (28-46)	49 (39-57)	62 (52-70)	84 (73-98)	<1·10 ⁻³⁰⁰

Continuous variables are summarised as median and interquartile range.

^aP-values were calculated using linear regression or logistic regression as appropriate.

^cIn current and former smokers only

^d1 unit ~ 12 g alcohol

^eIncome was classified differently in the 2 cohorts: 3 groups in CCHS and 4 groups in CGPS.

Supplementary table B. Baseline characteristics in the general population and their association with the DHCR7/CYP2R1 allele score.

	DHCR7/CYP2R1 allele score					P values ^a
	0-1 N = 15560 ^b	2 N = 21842	3 N = 16172	4-5 N = 27986	6-8 N = 7843	
Age, years	57 (47-66)	57 (47-66)	57 (47-66)	57 (47-66)	57 (47-66)	0.06
Men, %	45	45	45	45	45	0.61
Current smoker, %	23	23	23	23	23	0.11
Cumulative tobacco consumption, pack years ^c	24 (12-38)	20 (10-34)	19 (8-34)	18 (8-32)	18 (8-32)	0.37
Alcohol consumption, units/week ^d	8 (3-15)	8 (3-15)	8 (3-15)	7 (3-15)	8 (3-15)	0.49
Leisure-time physical activity <2 hours/week, %	7	7	7	7	7	0.80
Systolic blood pressure, mmHg	136 (123-150)	135 (122-150)	136 (122-150)	136 (122-150)	136 (122-150)	0.03 ^{NS}
Body mass index, kg/m ²	26 (23-28)	25 (23-28)	26 (23-28)	26 (23-28)	25 (23-28)	0.96
Low income, % ^e	36	37	36	37	37	0.25
Diabetes, %	4	4	4	4	4	0.99
Cholesterol, mmol/L	5.6 (4.9-6.3)	5.6 (4.9-6.3)	5.6 (4.9-6.3)	5.6 (4.9-6.3)	5.6 (4.9-6.3)	0.65
Vitamin D, nmol/L ^f	55 (16-32)	53 (28-46)	51 (39-57)	49 (52-70)	47 (73-98)	4·10 ⁻⁷²

Continuous variables are summarised as median and interquartile range.

^aP-values were calculated using linear regression or logistic regression as appropriate.

^b5460 participants attended both the 1981–1983 examination with 25(OH)D measurements and the 1991–1994 and/or 2001–2003 examinations with DNA available.

^cIn current and former smokers only

^d1 unit ~ 12 g alcohol

^eIncome was classified differently in the 2 cohorts: 3 groups in CCHS and 4 groups in CGPS.

^{NS}Not significant after correcting for 11 parallel tests (required $p = 0.05/11 = 0.0045$)

^f30792 participants only

References

- (1) Murea M, Lu L, Ma L, Hicks PJ, Divers J, McDonough CW et al. Genome-wide association scan for survival on dialysis in African-Americans with type 2 diabetes. *Am J Nephrol* 2011; 33:502-509.
- (2) Morrison AC, Felix JF, Cupples LA, Glazer NL, Loehr LR, Dehghan A et al. Genomic variation associated with mortality among adults of European and African ancestry with heart failure: the cohorts for heart and aging research in genomic epidemiology consortium. *Circ Cardiovasc Genet* 2010; 3:248-255.
- (3) Walter S, Atzmon G, Demerath EW, Garcia ME, Kaplan RC, Kumari M et al. A genome-wide association study of aging. *Neurobiol Aging* 2011; 32:2109-2128.
- (4) Yashin AI, Wu D, Arbeevev KG, Ukraintseva SV. Joint influence of small-effect genetic variants on human longevity. *Aging (Albany NY)* 2010; 2:612-620.
- (5) Lunetta KL, D'Agostino RB, Sr., Karasik D, Benjamin EJ, Guo CY, Govindaraju R et al. Genetic correlates of longevity and selected age-related phenotypes: a genome-wide association study in the Framingham Study. *BMC Med Genet* 2007; 8 Suppl 1:S13.