# **Supplementary Online Content**

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This supplementary material has been provided by the authors to give readers additional information about their work.

### **eMethods**

### Variables included in PCE and QRISK3

We calculated mean systolic blood pressure from two measurements taken seated after two minutes rest using an appropriate cuff and an Omron HEM-7015IT digital BP monitor or manual reading. To define treated hypertension, we used information from a nurse administered questionnaire on blood pressure lowering medication. Standing height was measured using a Seca 202 device. BMI was calculated as weight (kg) measured using an electronic weighing scales (Tanita BC-418)<sup>1</sup> divided by height squared (m<sup>2</sup>). Smoking was based on self-reported information on current smokers ("most or all days" and "occasional" smokers), previous smokers and never smokers (coded as non-smokers in PCE).

Serum data were corrected for laboratory dilution effects and were excluded if they fell outside limits defined according to UK Biobank specifications<sup>2</sup>. Total serum cholesterol and HDL cholesterol were obtained from enzymatic assays (Backman Coulter AU5800) and their ratio calculated<sup>2</sup>. We corrected total and HDL cholesterol values for individuals on lipid lowering treatment by dividing the total cholesterol value by 0.73 and HDL cholesterol by 1.03<sup>3</sup>.

## <u>Definition of additional QRISK3 variables</u>

Townsend deprivation index is available for all participants corresponding to the census output area in which their residential postcode is located. smoking intensity (number of cigarettes smoked per day or amount of tobacco currently smoked). We imputed smoking intensity among current smokers where information on amount smoked ("light", "moderate", "heavy") was missing. We considered missing smoking intensity data among

self-reported current smokers as Missing Not at Random (N = 9,465) and performed imputation on 40 continuous numeric smoking-related features with predictive mean matching using the mice package in R. We then categorized the continuous feature of "number of cigarettes smoked per day" into one of three ordinal factors for current smoking intensity analogous to those used by QRISK3 (light smoker: 0-9 cigarettes/day; moderate smoker: 10-19 cigarettes/day; heavy smoker: 20 or more cigarettes/day). All current pipe smokers were considered light smokers. The features used for smoking intensity imputation included demographic/socio-economic information, age when started smoking, measures of physical activity, biometric information, alcohol consumption (translated from an ordinal feature), and all but two of the available serum markers (due to missingness > 50%). Information on prevalent diseases was obtained from the nurse administered questionnaire or from relevant International Classification of Disease (ICD) codes or Office of Population Censuses and Surveys Classification of Surgical Operations and Procedures (4th revision, OPCS-4) codes in hospital episode statistics (HES) data with a date of event preceding the date of assessment centre attendance. Erectile dysfunction was additionally defined by relevant self-reported medication. Family history of CAD in a first degree relative was constructed from self-reported history of heart disease in father, mother or sibling. Type 1 and 2 diabetes were defined as self-reported diagnoses or from ICD codes in HES data prior to date of assessment; type 2 diabetes was additionally defined by relevant self-reported medication and HbA1c measurement at baseline ≥48 mmol/mol (measured by HPLC analysis on a Bio-Rad Variant II Turbo<sup>2</sup>). The definitions of all variables according to ICD-9, ICD-10, OPCS-4 and UK Biobank field codes are shown in Supplementary Material Online, Table S1.

### Calculation of Polygenic Risk Scores (PRS)

A PRS is typically calculated as a weighted sum of an individual's risk alleles. During PRS calculation, decisions must be made on (i) the weights to use for each SNP, (ii) how many SNPs to include in the score, and (iii) how to account for correlations between the SNPs (linkage disequilibrium [LD]).

We considered two methods for PRS calculation which address these issues in different ways:

- 1. Clumping and thresholding (C+T). This approach is widely used and involves taking the estimated SNP effects (from summary statistics) from the largest available GWAS as the SNP weights. LD is accounted for using a clumping algorithm, which yields a subset of independent SNPs while preferentially selecting those that are most strongly associated with the phenotype. The number of SNPs in this subset to be included in the PRS is chosen by applying a threshold on the P-value for association from the original GWAS. Typically, many P-value thresholds are explored with the aim of maximizing the ability of the PRS to predict the target phenotype.
- 2. Lassosum. Lassosum has similar aims to an alternative method, LDpred. Both lassosum and LDpred aim to calculate a PRS that is maximally predictive of the phenotype, by optimising the weights given to each SNP whilst accounting for LD. Lassosum achieves this by using penalized regression to carry out shrinkage and selection on the SNP effects from the base GWAS, accounting for LD information from a reference panel <sup>4</sup>. Two model parameters (s and lambda) must be chosen; PRS are calculated across a grid of values of s and lambda with the aim of maximizing phenotypic prediction. LDpred uses a Bayesian model to estimate the posterior mean SNP effects based on the SNP effects from the base GWAS, a point-normal

mixture prior (which allows a proportion of SNPs to have an effect of zero) and LD information from a reference panel <sup>5</sup>. The relative performance of lassosum and LDpred will depend on whether the true distribution of SNP effects is closer to a point-normal mixture distribution or a double exponential distribution<sup>6</sup>. However, for CAD lassosum has been shown to perform similarly to or better than LDpred <sup>4,5</sup>, and lassosum is less computationally intensive<sup>5</sup>.

### **Reclassification metrics**

We calculated the net reclassification index (NRI) at the current recommended threshold for treatment in the US (7.5%) and UK (10%), the associated integrated discrimination improvement (IDI), and the category free NRI. The NRI is calculated for nested models (PCE and PCE+PRS in our analysis) via a 2x2 classification table using defined thresholds. The 'upward' movement in categories for subjects with the outcome highlights better classification, and any 'downward movement' highlights worse reclassification. The improvement in reclassification is quantified as the sum of differences in proportions of individuals moving up minus the proportion moving down for those with the outcome, and the proportion of individuals moving down minus the proportion moving up for those without the outcome (NRI) <sup>6</sup>. Studies have shown that NRI is sensitive to the number and choice of thresholds selected 7. To account for this, the continuous (category-free) NRI has been proposed. This is the relative increase in the predicted probabilities for people who experience events and the decrease for those who do not<sup>8</sup>. Finally, IDI is proposed as a measure that integrates the NRI over all possible cut-offs for the probability of the outcome; the IDI is equivalent to the difference in discrimination slopes of 2 models <sup>6</sup>.

#### **eReferences**

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eTable 1. Definition of QRISK3 Variables in UK Biobank

	HES statistics ICD-10 and ICD9 codes (with date of event < date of visit)	Self-reported Data-Field 20002	Treatment and medication code Data-Field 20003	Other data fields
Atrial Fibrillation	ICD-10 I48; ICD-9 4273 and 4270; OPCS-4 K622 and K623	1471; 1483		
Migraine	ICD-10 G43, G440, N943; ICD-9 346	1265		
Type 1 diabetes	ICD-10 E10, O230; ICD9 25001, 25011, 25021, 25031, 25041, 25051, 25061, 25071, 25081, 25091, 25003, 25013, 25023, 25033, 25043, 25053, 25063, 25073, 25083, 25093	1222		
Type 2 diabetes	ICD-10 E11, O231; ICD-9 25000, 25010, 25020, 25030, 25040, 25050, 25060, 25070, 25080, 25090, 25002, 25012, 25022, 25032, 25042, 25052, 25062, 25072, 25082, 25092	1223, 1220	1140868902, 1140874646, 1140874674, 1140874718, 1140874744, 1140883066, 1140884600, 1141152590, 1141157284, 1141168660, 1141171646, 1141173882, 1141189090	Biobank field 2443 = 1; HbA1c measurement ≥ 48 mmol/mol
Chronic kidney disease	ICD-10 N183, N184, N185; ICD-10 5853, 5954, 5855	1192, 1193		
Rheumatoid arthritis	ICD-10 M05, M06; ICD-9 714	1464		
Systemic lupus erythematosus	ICD-10 M32; ICD-9 7100	1381		
Severe mental illness	ICD-10 F03, F068, F09, F20, F22, F23, F259, F28, F29, F31, F39, F53, F333; ICD9 295, 298, 296	1289, 1291		
Erectile disfunction	ICD-10 N484, N52; ICD-9 60784	1518	1141168936, 1141168948, 1141168944, 1141168946, 1140869100, 1140883010	
CAD in a 1st degree relative				Biobank fields 20107, 20110, 20111 = 1
Atypical antipsychotic medication			1140867420, 1140867444, 1140927956, 1140928916, 1141152848, 1141153490, 1141169714, 1141195974	
Steroid medications			1140874790, 1140874816, 1140874896.00, 1140874930, 1140874976, 1141145782, 1141173346	
Lipid lowering medications			1140861954, 1140861958, 1140888594, 1140888648, 1141146234, 1141192410, 1141192736	
Blood pressure lowering medications			1140860192, 1140860292, 1140860696, 1140860728, 1140860750, 1140860806, 1140860882, 1140860904, 1140861088, 1140861190, 1140861276, 1140866072, 1140866078, 1140866090, 1140866102, 1140866108, 1140866122, 1140866138, 1140866156, 1140866162, 1140866724, 1140866738, 114086818, 1140872568, 1140874706, 1140874744, 1140875808, 1140879758, 1140879760, 1140879818, 1140879822, 1140879806, 1140879830, 1140879818, 1140879842, 1140879866, 1140884298, 114088552, 1140888556, 1140888666, 1140988646, 1140923712, 1140910442, 1140910614, 1140916356, 114092372, 1140923336, 1140923404, 1140923712, 1140926778, 1140928226, 1141145660, 1141165470, 114116006, 1141169516, 1141171336, 1141180592, 1141180772, 1141180778, 1141184722, 1141193282, 1141194794, 1141194810	Biobank fields 6177, 6153 = 2

eTable 2. Definition of Coronary Artery Disease (CAD) and Cardiovascular Disease (CVD)

ICD-10	ICD-9	OPCS-4	Non-cancer illness code (Biobank field: 20002)	Operation code (Biobank field: 20004)	Vascular/heart problems diagnosed by doctor (Biobank field: 6150)
			CA	AD.	
121	410	K40.1-4	1075: Heart attack/myocardial infarction	1070: Coronary angioplasty	1: Heart attack
122	411	K41.1-4		1095: Coronary artery bypass grafts	
123	412	K45.1-5			
124.1	429.79	K49.1-2			
125.2		K49.8-9			
		K50.2			
		K75.1-4			
		K75.8-9			
			C\	/D	
G45	410	K40	1074: Angina	1070: Coronary angioplasty	1: Heart attack
120	411	K41	1075: Heart attack/myocardial infarction	1071: Other arterial surgery/revascularisation procedures	2: Angina
121	412	K42	1082: Transient ischaemic attack	1095: Coronary artery bypass grafts	3: Stroke
122	413	K43	1583: Ischaemic stroke	1105: Carotid artery surgery/endarterectomy	
123	414	K44		1109: Carotid artery angioplasty +/- stent	
124	434	K45		1514: Coronary angiogram	
125	436	K46			
163		K47.1			
164		K49			
		K50			
		K75			

eTable 3A. Descriptive Characteristics of Tuning (Case-Control) Set (N=80,103)

	Women	Men	Number of missing
N	25,444	54,659	12740
Age (years) (mean (SD)) *	60.0 (7.1)	59.7 (7.1)	
Mean SBP (mmHg) (mean (SD)) *	138.0 (19.2)	141.1 (18.0)	95
SD of SBP (mmHg) (mean (SD))	5.9 (4.8)	5.5 (4.5)	178
BMI (kg/m²) (mean (SD))	28.1 (5.5)	28.3 (4.4)	440
Townsend deprivation score (mean (SD))	-1.0 (3.2)	-1.2 (3.2)	103
Reported Ethnicity (%) *	, ,	, ,	88
African	140 (0.6)	282 (0.5)	
Caribbean	278 (1.1)	292 (0.5)	
		52055	
White/Not stated	24122 (94.8)	(95.2)	
Bangladeshi	12 (0.0)	38 (0.1)	
Chinese	59 (0.2)	105 (0.2)	
Indian	303 (1.2)	758 (1.4)	
Other	380 (1.5)	607 (1.1)	
Other Asian	75 (0.3)	238 (0.4)	
Pakistani	75 (0.3)	284 (0.5)	
Smoking category (%) *			2479
		23137	
Non-smoker	13852 (54.8)	(42.6)	
		24997	
Ex-smoker	9058 (35.8)	(46.0)	
Light Smoker (0-9/day)	524 (2.1)	1635 (3.0)	
Moderate Smoker (10-19/day)	1044 (4.1)	2110 (3.9)	
Heavy Smoker (20+/day)	805 (3.2)	2456 (4.5)	
Type 1 diabetes (%)	82 (0.3)	205 (0.4)	
Type 2 diabetes (%)	2213 (8.7)	6512 (11.9)	
Erectile disfunction (%)		610 (1.1)	
Atrial fibrillation (%)	480 (1.9)	1782 (3.3)	
Rheumatoid arthritis (%)	595 (2.3)	579 (1.1)	
Systemic lupus erythematosus (%)	94 (0.4)	24 (0.0)	
Migraine (%)	1063 (4.2)	774 (1.4)	
Severe mental health (%)	141 (0.6)	269 (0.5)	
Chronic kindey disease(%)	38 (0.1)	124 (0.2)	
		25411	
Blood pressure lowering medication (%) *	10237 (40.2)	(46.5)	
Systemic steroid medications (%)	398 (1.6)	661 (1.2)	
Atypical antipsychotic medications (%)	74 (0.3)	167 (0.3)	
	40	25819	
First degree relative with CHD (%)	13711 (53.9)	(47.2)	2000
Cholesterol (mean (SD)) *	242.4 (46.6)	227.7 (44.6)	3802
HDL cholesterol (mean (SD)) *	58.9 (14.9)	47.6 (12.1)	9996
Diabetes medication (%) *	1461 (5.7)	4428 (8.1)	
Obstantant made at the 1000 #	0500 (07.6)	24949	
Cholesterol medication (%) *	9569 (37.6)	(45.6)	

<sup>\*</sup>Variables used in PCE. Note: PCE is considering Black vs. all other ethnicities, current smokers vs. non-current smokers and diabetes as a binary variable

eTable 3B. Descriptive Characteristics of the PCE Prospective Cohort/Testing Set (N=352,660)

	Full pop	oulation	Incide	nt CAD	Incide	nt CVD
	Women	Men	Women	Men	Women	Men
N	205,297	147,363	1779	4493	5158	8595
Age (years) (mean (SD))	56.00 (8.01)	55.79 (8.35)	60.84 (6.43)	59.60 (6.92)	60.64 (6.63)	59.87 (6.96)
	135.05 (19.16)	140.92 (17.29)	146.16	147.08	143.81	146.73
Mean SBP (mmHg) (mean (SD))	133.03 (13.10)	140.92 (17.29)	(20.99)	(18.10)	(20.54)	(18.48)
Reported Ethnicity (%)						
Black	3865 (1.9)	2567 (1.7)	15 (0.8)	30 (0.7)	82 (1.6)	86 (1.0)
White/Others	201432 (98.1)	144796 (98.3)	1764 (99.2)	4463 (99.3)	5076 (98.4)	8509 (99.0)
Smoking category (%)						
Current smokers	16680 (8.1)	17140 (11.6)	895 (50.3)	2635 (58.6)	2473 (47.9)	5025 (58.5)
Non-current smokers	13852 (91.9)	130223 (88.4)	884 (49.7)	1858 (41.4)	2685 (52.0)	3570 (41.5)
Type 1 diabetes (%)	7322 (3.6)	9540 (6.5)	159 ( 8.9)	514 (11.4)	473 ( 9.2)	1072 (12.5)
Blood pressure lowering medication (%)	35291 (17.2)	30743 (20.9)	615 (34.6)	1410 (31.4)	1784 (34.6)	2913 (33.9)
	235.52 (44.8)	227.03 (42.86)	257.24	241.85	251.52	236.30
Cholesterol (mean (SD))	233.32 (44.0)	227.03 (42.00)	(50.59)	(46.39)	(48.29)	(45.88)
	61.39 (14.7)	49.42 (11.97)	56.39	46.83 (10.76)	57.92	47.55 (11.87)
HDL cholesterol (mean (SD))	01.55 (14.7)	75.72 (11.57)	(13.71)	70.05 (10.70)	(14.48)	77.55 (11.67)
Person-years of observation (mean (SD))	8.01 (1.04)	7.89 (1.31)	4.60 (2.37)	4.33 (2.38)	4.50 (2.38)	4.36 (2.38)

**eTable 3C**. Descriptive Characteristics of the PCE Prospective Cohort and Excluded Participants.

	Prospective coh	ort (N=352,660)	E	xcluded (N=69,77	3)
	Women	Men	Women	Men	Missing counts
N	205,297	147,363	42661	27112	64670
Age (years) (mean (SD))	56.00 (8.01)	55.79 (8.35)	55.87 (8.05)	56.01 (8.36)	
Mean SBP (mmHg) (mean (SD))	135.05 (19.16)	140.92 (17.29)	134.96 (19.35)	140.54 (17.50)	1230
Reported Ethnicity (%)					
Black	3865 (1.9)	2567 (1.7)	988 (2.3)	537 (2.0)	
White/Others	201432 (98.1)	144796 (98.3)	41673 (97.7)	26575 (98.0)	
Smoking category (%)					2216
Current smokers	16680 (8.1)	17140 (11.6)	16614 (40.1)	13007 (49.8)	
Non-current smokers	13852 (91.9)	130223 (88.4)	24820 (59.9)	13116 (50.2)	
Diabetes (%)	7322 (3.6)	9540 (6.5)	1708 (4.0)	1984 (7.3)	
Blood pressure lowering medication (%)	35291 (17.2)	30743 (20.9)	7630 (17.9)	6024 (22.2)	
Cholesterol (mean (SD))	235.52 (44.8)	227.03 (42.86)	234.69 (44.73)	226.14 (43.98)	29119
HDL cholesterol (mean (SD))	61.39 (14.7)	49.42 (11.97)	59.91 (14.63)	48.51 (12.25)	62646
Person-years of observation (mean (SD))	8.01 (1.04)	7.89 (1.31)	8.06 (1.22)	7.94 (1.52)	

eTable 3D. Descriptive Characteristics of the QRISK3 Prospective Cohort/Testing Set (N=350,730)

aracteristics of the QMSKS Frospe		oulation		nt CAD	Incide	nt CVD
	Women	Men	Women	Men	Women	Men
N	204157	146573	1768	4471	5114	8534
Age (years) (mean (SD))	56.0 (8.0)	55.8 (8.3)	60.8 (6.4)	59.6 (6.9)	60.7 (6.6)	59.9 (7.0)
Mean SBP (mmHg) (mean (SD))	135.0 (19.1)	140.9 (17.3)	146.0 (20.8)	147.1 (18.1)	143.8 (20.4)	146.7 (18.5)
SD of SBP (mmHg) (mean (SD))	5.5 (4.5)	5.3 (4.3)	6.1 (5.1)	5.6 (4.6)	6.1 (5.0)	5.7 (4.7)
BMI (kg/m²) (mean (SD))	26.9 (5.1)	27.7 (4.2)	28.1 (5.1)	28.4 (4.1)	28.3 (5.4)	28.5 (4.3)
Townsend deprivation score (mean (SD))	-1.4 (3.0)	-1.3 (3.1)	-1.0 (3.2)	-1.3 (3.1)	-1.0 (3.2)	-1.2 (3.2)
Reported Ethnicity (%)						
African	1178 (0.6)	1154 (0.8)	3 (0.2)	8 (0.2)	21 (0.4)	35 (0.4)
Bangladeshi	48 (0.0)	97 (0.1)	1 (0.1)	4 (0.1)	3 (0.1)	6 (0.1)
Caribbean	2055 (1.0)	1110 (0.8)	12 (0.7)	12 (0.3)	51 (1.0)	35 (0.4)
Chinese	745 (0.4)	391 (0.3)	3 (0.2)	3 (0.1)	10 (0.2)	6 (0.1)
Indian	2039 (1.0)	1751 (1.2)	22 (1.2)	84 (1.9)	51 (1.0)	148 (1.7)
Other	3330 (1.6)	2066 (1.4)	20 (1.1)	46 (1.0)	72 (1.4)	80 (0.9)
Other Asian	627 (0.3)	631 (0.4)	9 (0.5)	22 (0.5)	20 (0.4)	39 (0.5)
Pakistani	500 (0.2)	682 (0.5)	5 (0.3)	36 (0.8)	22 (0.4)	56 (0.7)
White/Not stated	193635 (94.8)	138691 (94.6)	1693 (95.8)	4256 (95.2)	4864 (95.1)	8129 (95.3)
Smoking category (%)						
Non-smoker	121888 (59.7)	74007 (50.5)	881 (49.8)	1849 (41.4)	2664 (52.1)	3543 (41.5)
Ex-smoker	65692 (32.2)	55521 (37.9)	587 (33.2)	1929 (43.1)	1772 (34.6)	3689 (43.2)
Light Smoker (0-9/day)	4725 (2.3)	4762 (3.2)	49 (2.8)	152 (3.4)	119 (2.3)	266 (3.1)
Moderate Smoker (10-19/day)	7171 (3.5)	6025 (4.1)	127 (7.2)	239 (5.3)	302 (5.9)	455 (5.3)
Heavy Smoker (20+/day)	4681 (2.3)	6258 (4.3)	124 (7.0)	302 (6.8)	257 (5.0)	581 (6.8)
Type 1 diabetes (%)	283 (0.1)	262 (0.2)	13 (0.7)	18 (0.4)	35 (0.7)	40 (0.5)
Type 2 diabetes (%)	7544 (3.7)	9310 (6.4)	153 (8.7)	493 (11.0)	448 (8.8)	1021 (12.0)
Erectile disfunction (%)		971 (0.7)		52 (1.2)		98 (1.1)
Atrial fibrillation (%)	1022 (0.5)	1825 (1.2)	13 (0.7)	78 (1.7)	101 (2.0)	223 (2.6)
Rheumatoid arthritis (%)	2914 (1.4)	1027 (0.7)	58 (3.3)	55 (1.2)	125 (2.4)	108 (1.3)
Systemic lupus erythematosus (%)	406 (0.2)	43 (0.0)	9 (0.5)	4 (0.1)	23 (0.4)	7 (0.1)
Migraine (%)	8913 (4.4)	2194 (1.5)	75 (4.2)	64 (1.4)	237 (4.6)	116 (1.4)
Severe mental health (%)	883 (0.4)	735 (0.5)	9 (0.5)	28 (0.6)	40 (0.8)	51 (0.6)
Chronic kindey disease(%)	123 (0.1)	136 (0.1)	5 (0.3)	13 (0.3)	14 (0.3)	30 (0.4)
Blood pressure lowering medication (%)	34994 (17.1)	30513 (20.8)	611 (34.6)	1399 (31.3)	1761 (34.4)	2883 (33.8)
Systemic steroid medications (%)	1794 (0.9)	1152 (0.8)	31 (1.8)	72 (1.6)	99 (1.9)	126 (1.5)
Atypical antipsychotic medications (%)	443 (0.2)	423 (0.3)	4 (0.2)	10 (0.2)	19 (0.4)	24 (0.3)
First degree relative with CHD (%)	90542 (44.3)	55506 (37.9)	1023 (57.9)	2249 (50.3)	2873 (56.2)	4088 (47.9)
Cholesterol / HDL ratio (mean (SD))	4.0 (1.1)	4.8 (1.3)	4.8 (1.3)	5.4 (1.4)	4.6 (1.3)	5.2 (1.4)
Diabetes medication (%)	4427 (2.2)	5860 (4.0)	114 (6.4)	335 (7.5)	324 (6.3)	698 (8.2)
Cholesterol medication (%)	21242 (10.4)	24645 (16.8)	364 (20.6)	1106 (24.7)	1124 (22.0)	2258 (26.5)
Person-years of observation (mean (SD))	8.0 (1.0)	7.9 (1.3)	4.6 (2.4)	4.3 (2.4)	4.5 (2.4)	4.3 (2.4)

**eTable 3E.** Descriptive Characteristics of the QRISK3 Prospective Cohort and Excluded Participants

		ive cohort		Excluded	
		50,730) Name	14/2	(N=71,703)	Baissins sound
N	<b>Women</b> 204157	<b>Men</b> 146573	Women	27902	Missing count
Age (years) (mean (SD))		55.8 (8.3)		55.98 (8.37)	
Mean SBP (mmHg) (mean (SD))	1 1	140.9 (17.3)		140.55 (17.56)	1230
		5.3 (4.3)			1655
SD of SBP (mmHg) (mean (SD))				5.25 (4.37) 27.75 (4.35)	
BMI (kg/m²) (mean (SD))		27.7 (4.2)			2665
Townsend deprivation score (mean (SD))	-1.4 (3.0)	-1.3 (3.1)	-1.19 (3.11)	-1.06 (3.25)	520
Reported Ethnicity (%)	1179 (0.6)	1154 (0.0)	267 (0.9)	272 (1.0)	
		1154 (0.8)	367 (0.8)		
Bangladeshi		97 (0.1)	14 (0.0)		
		1110 (0.8)	537 (1.2)	, ,	
		391 (0.3)	186 (0.4)		
		1751 (1.2)	597 (1.4)		
		2066 (1.4)	839 (1.9)		
Other Asian		631 (0.4)	156 (0.4)		
		682 (0.5)	142 (0.3)		
White/Not stated	193635 (94.8)	138691 (94.6)	40963 (93.5)	26036 (93.3)	2054
Smoking category (%)*	121000 (50 7)	74007 (50.5)	25.454./64.0\	10500 (51.0)	3854
	121888 (59.7)	` '		13522 (51.8)	
	1	55521 (37.9)		10114 (38.8)	
Light Smoker (0-9/day)		4762 (3.2)		692 (2.7)	
Moderate Smoker (10-19/day)		6025 (4.1)		825 (3.2)	
Heavy Smoker (20+/day)		6258 (4.3)		934 (3.6)	
Type 1 diabetes (%)		262 (0.2)	1813 (4.1)		
Type 2 diabetes (%)		9310 (6.4)	6 (0.0)		
Erectile disfunction (%)		971 (0.7)	243 (0.6)		
Atrial fibrillation (%)		1825 (1.2)	649 (1.5)		
Rheumatoid arthritis (%)		1027 (0.7)	90 (0.2)	, ,	
Systemic lupus erythematosus (%)	1 1	43 (0.0)	1840 (4.2)		
Migraine (%)		2194 (1.5)	215 (0.5)		
Severe mental health (%)	883 (0.4)	735 (0.5)	45 (0.1)		
Chronic kidney disease (%)		136 (0.1)		6254 (22.4)	
Blood pressure lowering medication (%)	` ,	30513 (20.8)	418 (1.0)		
Systemic steroid medications (%)		1152 (0.8)		111 (0.4)	
Atypical antipsychotic medications (%)	443 (0.2)	423 (0.3)		10486 (37.6)	
First degree relative with CHD (%)		55506 (37.9)		225.96 (44.06)	
Cholesterol / HDL ratio (mean (SD))		4.8 (1.3)		48.35 (12.37)	62712
Diabetes medication (%)		5860 (4.0)		1327 (4.8)	
Cholesterol medication (%)	21242 (10.4)	24645 (16.8)	4869 (11.1)	5136 (18.4)	
Person-years of observation (mean (SD))	8.0 (1.0)	7.9 (1.3)	8.05 (1.23)	7.93 (1.52)	

<sup>\*</sup>Smoking intensity was imputed for current smokers in the prospective cohort.

**eTable 4.** Association of Different Polygenic Risk Scores (PRS) With Coronary Artery Disease (CAD) and Cardiovascular Disease (CVD) in Tuning Case-Control Studies and in Prospective Cohort Study

				AUC (95	% CI) PRS		
			CA	AD	C	/D	
Method	D value / tuning narameter	N SNPs in the	Case control study	Prospective study	Case control study	Prospective study	
Wethod	P value/ tuning parameter	PRS	(N=15,947)	(N=6,272)	(N=33,458)	(N=13,753)	
Clumping & Thresholding	0.000001	109	0.597 (0.591-0.603)	0.588 (0.581-0.595)	0.549 (0.545-0.554)	0.549 (0.544-0.554)	
Clumping & Thresholding	0.000005	156	0.600 (0.594-0.606)	0.590 (0.583-0.598)	0.552 (0.547-0.556)	0.551 (0.546-0.556)	
Clumping & Thresholding	0.00001	177	0.603 (0.597-0.609)	0.593 (0.585-0.600)	0.553 (0.549-0.558)	0.552 (0.547-0.557)	
Clumping & Thresholding	0.00005	274	0.609 (0.603-0.615)	0.598 (0.591-0.606)	0.557 (0.553-0.561)	0.555 (0.550-0.560)	
Clumping & Thresholding	0.0001	342	0.607 (0.600-0.613)	0.598 (0.591-0.605)	0.559 (0.554-0.563)	0.554 (0.550-0.559)	
Clumping & Thresholding	0.0002	459	0.609 (0.602-0.615)	0.598 (0.591-0.605)	0.560 (0.556-0.565)	0.556 (0.551-0.561)	
Clumping & Thresholding	0.0003	539	0.610 (0.603-0.616)	0.599 (0.592-0.606)	0.561 (0.557-0.566)	0.557 (0.552-0.562)	
Clumping & Thresholding	0.0004	614	0.611 (0.605-0.617)	0.600 (0.593-0.607)	0.562 (0.557-0.566)	0.558 (0.553-0.563)	
Clumping & Thresholding	0.0005	671	0.611 (0.605-0.618)	0.600 (0.593-0.607)	0.562 (0.558-0.567)	0.558 (0.554-0.563)	
Clumping & Thresholding	0.0006	740	0.612 (0.606-0.618)	0.600 (0.593-0.608)	0.563 (0.558-0.567)	0.559 (0.554-0.564)	
Clumping & Thresholding	0.0007	812	0.612 (0.606-0.618)	0.600 (0.593-0.607)	0.563 (0.559-0.567)	0.559 (0.554-0.564)	
Clumping & Thresholding	0.0008	869	0.612 (0.606-0.619)	0.600 (0.593-0.607)	0.563 (0.559-0.568)	0.558 (0.553-0.563)	
Clumping & Thresholding	0.0009	930	0.612 (0.605-0.618)	0.599 (0.592-0.606)	0.563 (0.558-0.567)	0.558 (0.553-0.563)	
Clumping & Thresholding	0.001	987	0.612 (0.606-0.618)	0.599 (0.592-0.606)	0.563 (0.559-0.567)	0.558 (0.553-0.563)	
Clumping & Thresholding	0.002	1,488	0.612 (0.606-0.618)	0.599 (0.592-0.606)	0.563 (0.559-0.567)	0.559 (0.554-0.563)	
Clumping & Thresholding	0.003	1,934	0.612 (0.606-0.618)	0.597 (0.590-0.604)	0.564 (0.560-0.569)	0.558 (0.553-0.563)	
Clumping & Thresholding	0.004	2,363	0.613 (0.606-0.619)	0.595 (0.588-0.602)	0.565 (0.561-0.570)	0.557 (0.552-0.562)	
Clumping & Thresholding	0.005	2,795	0.611 (0.605-0.617)	0.594 (0.587-0.602)	0.565 (0.561-0.569)	0.557 (0.552-0.562)	
Genome-wide significant	5.00E-08	455	0.551 (0.545-0.557)	0.552 (0.545-0.560)	0.524 (0.519-0.528)	0.527 (0.522-0.532)	
Lassosum (INFO score>0.999	s=0.5; lambda=0.00428 (CAD) s=0.9; lambda=0.00207 (CVD)	40,079 (CAD) 297,862 (CVD)	0.629 (0.623-0.635)	0.608 (0.601-0.615)			
(N=1,037,385 variants))		,()			0.582 (0.577-0.586)	0.564 (0.559-0.569)	
Lassosum (INFO score>0.3 (N=6,750,670 variants))	s=0.2; lambda=0.00546 (CAD) s=0.2; lambda=0.00886 (CVD)	39,219 (CAD) 1,227 (CVD)	0.618 (0.612-0.624)	0.605 (0.598-0.612)	0.548 (0.543-0.552)	0.544 (0.539-0.549)	
(IN-0,750,070 Varialls))		1,227 (CVD)			0.546 (0.545-0.552)	0.544 (0.559-0.549)	

Clumping parameter: R2=0.6 (no clumping for genome wide)

To allow recalculation of the PRS, Lassosum coefficients for CAD and CVD models can be downloaded from 10.6084/m9.figshare.11778534 (CAD) and 10.6084/m9.figshare.11778543 (CVD).

eTable 5. Recalibration Coefficients for CAD and CVD Analyses From Cox Regression (PCE)

		PF	RS	P	CE	Se	ex	Age	1**	Age	2**	Age	3**	Age	4**
	S0	β	SE												
CAD															
PCE	0.984			0.58	0.01	1.06	0.03								
PRS	0.978	2.79	0.09												
Age + sex	0.984					1.29	0.03	2.13	0.14	2.13	0.11	4.36	0.33	2.09	0.11
PRS + age + sex	0.985	2.99	0.09			1.31	0.03	2.15	0.14	2.15	0.11	4.41	0.33	2.15	0.11
PCE + PRS	0.986	2.79	0.09	0.57	0.01	1.08	0.03								
						C	CVD								
PCE	0.963			0.57	0.01	0.66	0.02								
PRS	0.952	0.29	0.01												
Age + sex	0.961					0.88	0.02	1.95	0.09	2.07	0.07	4.21	0.22	2.27	0.07
PRS + age + sex	0.962	0.32	0.01			0.89	0.02	1.97	0.09	2.09	0.07	4.24	0.22	2.3	0.07
PCE + PRS	0.963	0.28	0.01	0.56	0.01	0.67	0.02								

<sup>\*</sup>S0: recalibrated baseline survival at 10 years

<sup>\*\*</sup>Non-linear effects of age are modelled with cubic splines (breakpoints at quartiles of age)

**eTable 6.** C Statistics (Derived From Cox Regression) for CVD Using Recalibrated Models in the PCE Prospective Cohort. Results are presented for the full population and stratified by gender and age class (above or below 55 years of age).

# A. All participants

	All	< 55 years old	>= 55 years old	Men	Women
	(N=352,660	(N=147,985	(N=204,675	(N=147,363	(N=205,297
	13,753 events)	2,854 events)	10,899 events)	8,595 events)	5,158 events)
PRS	0.56 (0.56-0.57)	0.59 (0.58-0.60)	0.56 (0.55-0.56)	0.57 (0.56-0.57)	0.56 (0.56-0.57)
Age and sex	0.70 (0.70-0.71)	0.68 (0.67-0.69)	0.65 (0.64-0.65)	0.65 (0.65-0.66)	0.68 (0.67-0.68)
PRS + age and sex	0.71 (0.71-0.72)	0.70 (0.69-0.71)	0.66 (0.65-0.66)	0.67 (0.66-0.67)	0.69 (0.68-0.69)
PCE	0.72 (0.72-0.73)	0.73 (0.72-0.74)	0.67 (0.67-0.68)	0.68 (0.67-0.69)	0.71 (0.70-0.72)
PRS + PCE	0.73 (0.73-0.74)	0.74 (0.73-0.75)	0.68 (0.68-0.69)	0.69 (0.68-0.70)	0.72 (0.71-0.72)
B. Participants not rec	eiving lipid lowerin	g treatment at base	eline		
	All	< 55 years old	>= 55 years old	Men	Women
	(N=306,421	(N=140,266	(N=166,155	(N=122,546	(N=183,875
	10,332 events)	2,432 events)	7,900 events)	6,313 events)	4,019 events)
PRS	0.57 (0.56-0.57)	0.59 (0.58-0.61)	0.56 (0.56-0.57)	0.57 (0.56-0.58)	0.56 (0.55-0.57)
Age and sex	0.71 (0.70-0.71)	0.68 (0.67-0.69)	0.65 (0.65-0.66)	0.66 (0.65-0.67)	0.68 (0.67-0.69)
PRS + age and sex	0.72 (0.71-0.72)	0.70 (0.69-0.71)	0.66 (0.66-0.67)	0.68 (0.67-0.68)	0.69 (0.68-0.70)
PCE	0.73 (0.72-0.73)	0.72 (0.71-0.73)	0.68 (0.67-0.68)	0.69 (0.68-0.69)	0.71 (0.70-0.72)
PRS + PCE	0.74 (0.73-0.74)	0.73 (0.72-0.74)	0.68 (0.68-0.69)	0.70 (0.69-0.70)	0.72 (0.71-0.72)

eTable 7. Recalibration Coefficients for CAD and CVD Analysis From Cox Regression (QRISK3)

		PR	S	PO	CE	Se	ex	Age	1**	Age	2**	Age	3**	Age	4**
	S0	β	SE												
CAD															
PCE	0.987			1.01	0.02	1.3	0.03								
PRS	0.978	2.79	0.09												
Age + sex	0.984					1.29	0.03	2.12	0.14	2.12	0.11	4.31	0.33	2.09	0.11
PRS + age + sex	0.985	2.99	0.09			1.31	0.03	2.14	0.14	2.14	0.11	4.36	0.33	2.15	0.11
PCE + PRS	0.987	2.68	0.09	0.99	0.02	1.32	0.03								
						C	VD								
PCE	0.967			0.97	0.01	0.89	0.02								
PRS	0.952	0.29	0.01												
Age + sex	0.961					0.88	0.02	1.97	0.09	2.08	0.07	4.24	0.22	2.28	0.07
PRS + age + sex	0.962	0.32	0.01			0.89	0.02	1.98	0.09	2.1	0.07	4.28	0.22	2.31	0.07
PCE + PRS	0.968	0.26	0.01	0.96	0.01	0.9	0.02								

<sup>\*</sup>S0: recalibrated baseline survival at 10 years

<sup>\*\*</sup>Non-linear effects of age are modelled with cubic splines (breakpoints at quartiles of age)

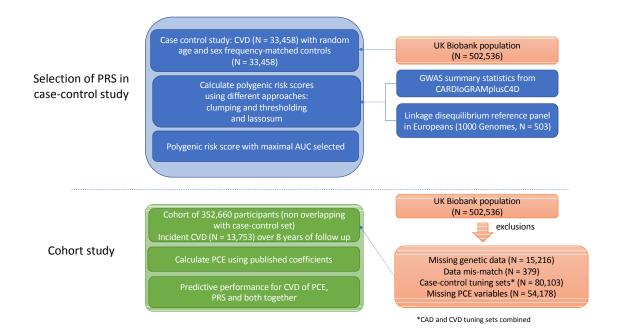
**eTable 8.** C Statistics (Derived From Cox Regression) for CAD and CVD Using Recalibrated Models in the QRISK3 Prospective Cohort. Results are presented for the full population and stratified by gender and age class (above or below 55 years of age).

CAD									
	Full population	< 55 years old	>= 55 years old	Men	Women				
	(N=350,730	(N=147,110	(N=203,620	(N=146,573	(N=204,157				
	6,239 events)	1,339 events)	4,900 events)	4,471 events)	1,768 events)				
QRISK3	0.61 (0.60-0.62)	0.64 (0.63-0.66)	0.60 (0.59-0.61)	0.61 (0.60-0.62)	0.61 (0.60-0.63)				
Age (and sex*)	0.73 (0.72-0.74)	74) 0.73 (0.72-0.75) 0.68 (0.68-0.69) 0.64 (0.63-0.65		0.64 (0.63-0.65)	0.68 (0.67-0.70)				
PRS + age (and sex*)	0.76 (0.75-0.76)	0.76 (0.75-0.78)	0.71 (0.70-0.72)	0.68 (0.67-0.69)	0.71 (0.70-0.73)				
QRISK3*	0.78 (0.77-0.79)	0.81 (0.80-0.83)	0.73 (0.73-0.74)	0.71 (0.70-0.72)	.70-0.72) 0.76 (0.75-0.78)				
PRS + QRISK3*	0.79 (0.79-0.80)		0.73 (0.72-0.74)	0.78 (0.76-0.79)					
Participants not receiving lipid lowering treatment at baseline									
	All	< 55 years old	>= 55 years old	Men	Women				
	(N=304,843	(N=139,475	(N=165,368	(N=121,928	(N=182,915				
	4,769 events)	1,139 events)	3,630 events)	3,365 events)	1,404 events)				
QRISK3	0.61 (0.60-0.62)		0.62 (0.61-0.63)	0.61 (0.60-0.63)					
Age (and sex*)	(and sex*) 0.74 (0.73-0.75) 0.74 (0.7		0.69 (0.68-0.70)	.69 (0.68-0.70)					
PRS + age (and sex*)	0.76 (0.76-0.77)	0.77 (0.75-0.79)	0.72 (0.71-0.73)	0.70 (0.69-0.71)	0.72 (0.70-0.73)				
QRISK3* 0.79 (0.78-0.80)		0.81 (0.80-0.83)	0.74 (0.73-0.75)	0.72 (0.71-0.73)	0.77 (0.75-0.78)				
PRS + QRISK3*	0.81 (0.80-0.81)	0.83 (0.81-0.85)	0.76 (0.75-0.77)	0.75 (0.74-0.76)	0.78 (0.77-0.80)				
CVD									
	Full population	< 55 years old	>= 55 years old	Men	Women				
	(N=350,730	(N=147,110	(N=203,620	(N=146,573	(N=204,157				
	13,650 events)	2,826 events)	10,824 events)	8,536 events)	5,114 events)				
QRISK3	0.56 (0.56-0.57)	0.59 (0.58-0.61)	.61) 0.56 (0.55-0.56) 0.57 (0.56-0.57)		0.56 (0.56-0.57)				
Age (and sex*)	e (and sex*) 0.70 (0.70-0.71) 0.68 (0.6		0.65 (0.64-0.65)	0.65 (0.65-0.66)	0.68 (0.67-0.68)				
PRS + age (and sex*)	d sex*) 0.71 (0.71-0.72) 0.70 (0.69-0.71) 0.66 (0.65-0.66) 0.67 (0.6		0.67 (0.66-0.67)	0.69 (0.68-0.69)					
QRISK3*	SK3* 0.75 (0.74-0.75) 0.76 (0.75-0.77) 0.70 (0.69		0.70 (0.69-0.70)	0.71 (0.70-0.71)	0.74 (0.73-0.74)				
PRS + QRISK3*	K3* 0.75 (0.75-0.76) 0.77 (0.76-0.78) 0.70 (0.70-0.71) 0.71 (0.71-0.72)		0.71 (0.71-0.72)	0.74 (0.73-0.75)					
	Participants	not receiving lipid lo	wering treatment at	baseline					
	All	< 55 years old	>= 55 years old	Men	Women				
	(N=304,843	(N=139,475	(N=165,368	(N=121,928	(N=182,915				
	10,267 events)	2,408 events)	7,859 events)	6,277 events)	3,990 events)				
QRISK3	0.57 (0.56-0.57)	0.59 (0.58-0.61)	0.56 (0.56-0.57)	0.57 (0.56-0.58)	0.56 (0.55-0.57)				
Age (and sex*)	ge (and sex*) 0.71 (0.70-0.71) 0.68 (0.67-0.69) (		0.65 (0.65-0.66)	0.66 (0.65-0.67)	0.68 (0.67-0.69)				
PRS + age (and sex*)	S + age (and sex*) 0.72 (0.71-0.72) 0.70 (0.69-0.71) 0.66 (0.66-0.67)		0.68 (0.67-0.68)	0.69 (0.68-0.70)					
QRISK3* 0.75 (0.75-0.76) 0.76 (0.75-0.77) 0.70		0.70 (0.69-0.71)	0.71 (0.71-0.72)	0.74 (0.73-0.74)					
PRS + QRISK3*	0.76 (0.75-0.76)	0.76 (0.75-0.78)	0.71 (0.70-0.71)	0.72 (0.72-0.73)	0.74 (0.73-0.75)				

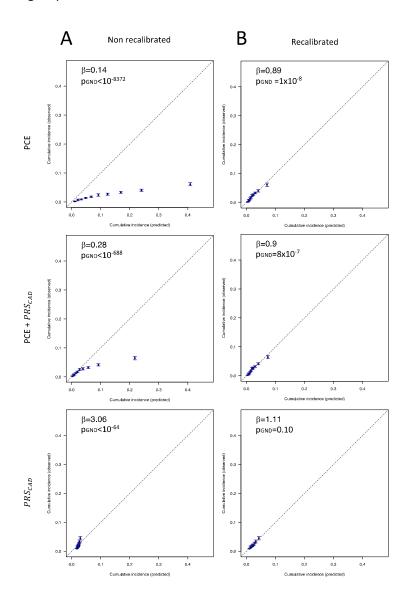
**eTable 9.** Risk Reclassification at 7.5% and 10% Thresholds for CAD and CVD Using Recalibrated Models: QRISK3 and QRISK3 Plus PRS.

CAD (7.5% threshold)											
			QRISK								
			<7.5%	>=7.5%	% reclassified	Categorical NRI:	Continuous NRI:				
QRISK3	<7.5%	Cases	4347	585	11.86	0.053 (0.043.0.064)	0.149				
	>=7.5%	Cases	260	1047	19.89	0.052 (0.043;0.061)	(0.125;0.174)				
	<7.5%	Non-cases	319802	7154	2.19	-0.005 (-0.006;-	0.147				
	>=7.5%	Non-cases	5365	12170	30.6	0.005)	(0.143;0.150)				
NRI in the full population:						0.047 (0.038;0.056)	0.296 (0.271;0.321)				
	IDI: 0.0064 (0.0058-0.007)										
CAD (10% threshold)											
QRISK3 + PRS											
			<10%	>=10%	% reclassified	Categorical NRI:	Continuous NRI:				
QRISK3	<10%	Cases	5175	411	7.36	0.043 (0.036;0.051)	0.149				
	>=10%	Cases	141	512	21.59	0.043 (0.030,0.031)	(0.125;0.174)				
	<10%	Non-cases	332698	4521	1.34	-0.006 (-0.007;-	0.147				
	>=10%	Non-cases	2359	4913	32.44	0.006)	(0.143;0.150)				
				0.037 (0.030;0.044)	0.296						
	IDI	0.0064/0.00		NRI in the	full population:		(0.271;0.321)				
IDI: 0.0064 (0.0058-0.007)						1					
			QRISK3		(7.5% threshold	)					
			<7.5%	>=7.5%	% reclassified	Categorical NRI:	Continuous NRI:				
	<7.5%	Cases	5831	628	9.72	Categorical INTI.	0.079				
3	>=7.5%	Cases	521	6670	7.25	0.008 (0.003;0.013)	(0.062;0.096)				
QRISK3	<7.5%	Non-cases	260120	8104	3.02		0.075				
ď	>=7.5%	Non-cases	9171	59685	13.32	0.003 (0.002;0.004)	(0.072;0.078)				
	7-7.570	14011 cases	31/1	33003	15.52	0.011 (0.006;0.016)	0.154				
			ı	NRI in the	full population:		(0.137;0.171)				
	IDI:	0.0032 (0.00	29-0.0036	)							
		-		CVD	(10% threshold)						
QRISK3 + PRS											
			<10%	>=10%	% reclassified	Categorical NRI:	Continuous NRI:				
QRISK3	<10%	Cases	7967	682	7.89	0.014 (0.009;0.018)	0.079				
	>=10%	Cases	497	4504	9.94	0.014 (0.009;0.018)	(0.062;0.096)				
	<10%	Non-cases	290884	6570	2.21	-0.0005 (-	0.075				
	>=10%	Non-cases	6404	33222	16.16	0.001;0.000)	(0.072;0.078)				
			ı	0.013 (0.008;0.018)	0.154 (0.137;0.171)						
	IDI: 0.0032 (0.0029-0.0036)										

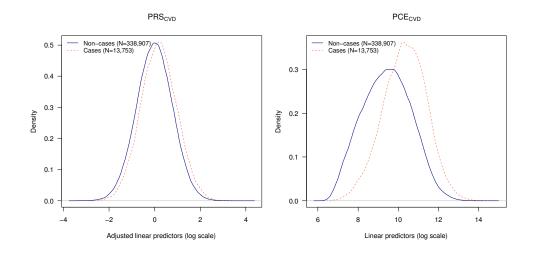
eFigure 1. Study Design and Flowchart for Cardiovascular Disease With PCE



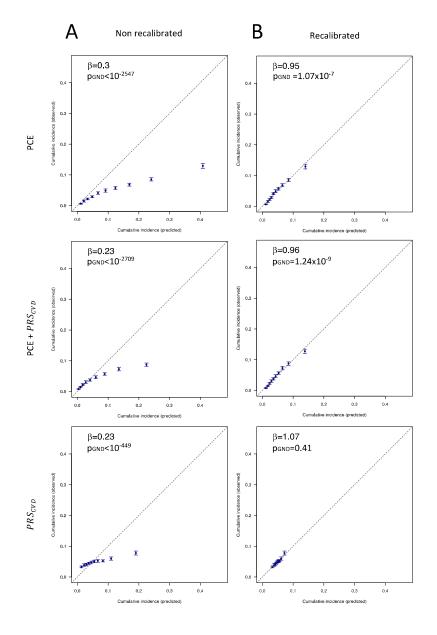
eFigure 2. Calibration Plots for PCE, Polygenic Risk Score for Coronary Artery Disease (CAD) (PRS<sub>CAD</sub>) and Both Combined, Using a UK Biobank Prospective Cohort Sample (N = 352,660) A) Non-recalibrated model performance. B) Recalibrated model performance. Recalibration was undertaken by fitting the predicted log hazard from original models as a covariate in a Cox survival model. P<sub>GND</sub> is the P-value associated with the Greenwood-Nam-D'Agostino test statistic and tests the null hypothesis that the observed and expected probabilities are identical in each group.



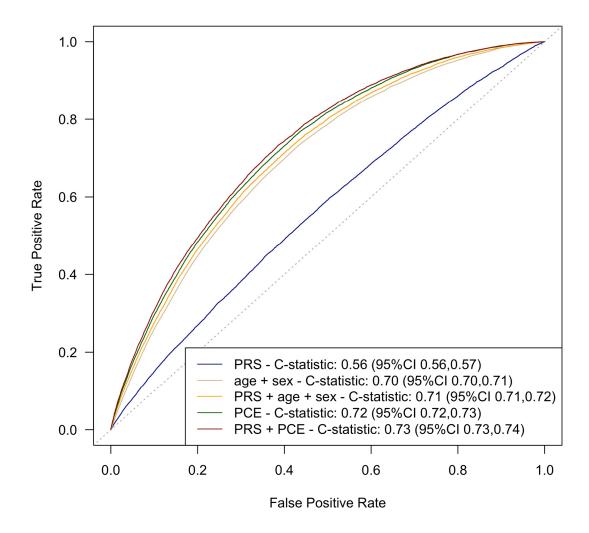
eFigure 3. Distributions of PRS and PCE for CVD in the Prospective Cohort



**eFigure 4.** Calibration Plots for PCE, Polygenic Risk Score for CVD (PRS<sub>CVD</sub>) and Both Combined, Using a UK Biobank Prospective Cohort Sample (N=352,662). A. Non-recalibrated model performance. B. Recalibrated model performance. Recalibration was undertaken by fitting the predicted log hazard from original models as a covariate in a Cox survival model. P<sub>GND</sub> is the P-value associated with the Greenwood-Nam-D'Agostino test statistic.



**eFigure 5.** ROC Curves and C Statistics for Different Models in Prospective Cohort Analyses for CVD



eFigure 6. Change in the Predicted Probabilities (Expressed as a Percentage) of the Recalibrated PCE Model After the Addition of the Polygenic Risk Score for Cardiovascular Disease (CVD) (PRS<sub>CvD</sub>). The x-axis is the predicted probability from the original PCE model, and the y-axis is the difference in predicted probabilities between the PRS<sub>CvD</sub>-augmented models and PCE. The associated table shows the percentage of participants whose predicted probabilities change by less than given thresholds. B. Predicted probabilities by PCE and PCE plus PRS<sub>CvD</sub> with dotted lines showing the 7.5% threshold. The associated table shows the reclassification percentages for 7.5% threshold as well as the Net Reclassification Improvement (IDI).

