Supplemental Material

Table S1. International Statistical Classification of Diseases-10th Revision (ICD-10) codes used to define cardiovascular death.

Diagnosis	Version	Code
Ischemic Heart Disease	ICD-10	120* 121* 122* 123* 124* 125*
Heart Failure & cardiomyopathy	ICD-10	110* 111* 12* 13* 14* 15* 16* 42* 255 81 150* 1517
Valvular heart disease	ICD-10	134* 135* 136* 137*
Cardiac arrest (due to cardiac condition)	ICD-10	1462
Ventricular tachycardia and ventricular fibrillation	ICD-10	1470 1472 14901 14902
Acute stroke (ischemic, non-ischemic and hemorrhagic)	ICD-10	160*, 1161*, 163* 164* 165*, 1166*, 167* 168* 169* G46*
Cardiogenic shock	ICD-10	R570
Thromboembolism	ICD-10	126* 182*
Peripheral vascular disease	ICD-10	170* 171* 172* 173* 174* 175* 176* 178* 179* 179*
Infective endocarditis	ICD-10	I33* I38*

^{*}All digits after omitted

Table S2. Baseline characteristics of AF and HF phenotype groups according to the composite outcome (HF hospitalization or cardiovascular death) at 2.5 years.

Baseline Characteristic	Neither AF nor	HF (N=488)	AF only (N=354)	HF only (N=369)	AF plus HF (N:	=405)
	Without Event	With event	Without Event	With event	Without Event	With event	Without	With event
	(N=452)	(N=36)	(N=299)	(N=55)	(N=277)	(N=92)	Event	(N=128)
							(N=277)	
Clinical characteristics								
Age, median, (IQR)	65 (56–73)	70 (65–79)	70 (61–78)	79 (69–85)	68 (59–76)	72 (57–79)	74 (67–81)	75 (67–81)
Female sex, n (%)	205/452 (45)	17/36 (47)	120/299 (40)	30/55 (55)	98/277 (35)	30/92 ((33)	107/277 (39)	37/128 (29)
Race, n (%)								
Courseign	306/452 (68)	26/36 (72)	253/299 (85)	49/55 (89)	209/277 (75)	55/92 (59)	239/277 (86)	101/128
Caucasian								(79)
Asian	92/452 (20)	8/36 (22)	29/299 (10)	2/55 (4)	44/277 (16)	19/92 (21)	18/277 (7)	12/128 (9)

Afro-Caribbean	53/452 (12)	2/36 (6)	17/299 (6)	3/ 55 (5)	24/277 (9)	18/92 (20)	19/277 (7)	15/128 (12)
Other	1/452 (0.2)	-	-	1/55 (2)	-	-	1/277 (0.4)	-
Heart Rhythm, n (%)								
Sinus Rhythm	452/452 (100)	36/36 (100)	-	-	277/277 (100)	92/92 (100)	-	-
Paroxysmal AF	-	-	178/299 (60)	17/55 (31)	-	-	137/277 (49)	47/128 (37)
Persistent AF	-	-	57/299 (19)	19/55 (35)	-	-	66/277 (24)	34/128 (27)
Permanent AF	-	-	52/299 (17)	17/55 (31)	-	-	59/277 (21)	43/128 (34)
Atrial Flutter	-	-	12/299 (4)	2/55 (4)	-	-	15/277 (5)	4/128 (3)
BMI, kg/m², median, (IQR)) *	29 (25–33)	29 (25–33)	29 (26, 33)	27 (23–32)	28 (25–32)	28 (24–32)	29 (25–33)	29 (26–33)
Systolic BP, mmHg , median,	126 (113–140)	131 (111–145)	130 (118–145)	123 (109–135)	122 (110–137)	120 (106–130)	121 (110–	119 (106–
(IQR)							140)	134)
Heart rate/min, median, (IQR)	68 (61–78)	73 (60–89)	68 (58–80)	71 (59–88)	72 (63–81)	74 (62–83)	76 (63–90)	77 (64–89)

Ejection fraction, %, median,	61 (57–68)	60 (54–71)	62 (57–68)	58 (54–68)	47 (38–59)	39 (29–52)	49 (40–58)	40 (27–52)
(IQR)								
Ejection fraction <50%, n (%)	-	-	-	-	158/267 (59)	66/90 (73)	141/262 (54)	91/126 (72)
Previous diagnosis of stable HF	-	-	-	-	99/277 (36)	53/92 (58)	123/277 (44)	80/128 (63)
Symptomatic HF								
NYHA II HF, n (%)	-	-	-	-	118/277 (43)	25/92 (25)	120/275 (44)	39/126 (31)
NYHA III HF, n (%)	-	-	-	-	52/277 (19)	32/92 (32)	65/275 (24)	46/126 (37)
NYHA IV HF, n (%)	-	-	-	-	11/277 (4)	9/92 (9)	16/275 (6)	15/126 (12)
LBBB, n (%)	6/452 (1)	0/36 (0)	5/299 (2)	1/55 (2)	15/277 (4)	11/92 (12)	12/277 (4)	10/128 (8)
Medical history, n (%)								
Diabetes	195/452 (43)	17/36 (47)	59/299 (20)	16/55 (29)	113/277 (41)	53/92 (58)	66/277 (24)	46/128 (36)
Hypertension	299/452 (66)	23/36 (64)	167/299 (56)	38/55 (69)	166/277 (60)	54/92 (59)	130/277 (47)	69/128 (54)
Coronary artery disease	206/452 (46)	18/36 (50)	50/299 (17)	8/55 (15)	155/277 (56)	48/92 (52)	88/277 (32)	56/128 (44)

Hyponatremia (Na <135 mmol/L)*	71/445 (16)	6/36 (17)	30/272 (11)	13/55 (24)	46/274 (17)	25/92 (27)	28/270 (11)	29/127 (23)
Severe valvular heart disease	9/452 (2)	0/36 (0)	8/299 (3)	9/55 (16)	7/277 (3)	5/92 (5)	21/277 (8)	20/128 (16)
HF hospitalization at presentation	-	-	-	-	15/277 (5)	8/92 (9)	4/277 (1)	12/128 (9)
Laboratory measurements								
eGFR mL/min/1.73m², (CKD-EPI),	81 (64–95)	66 (43–86)	75 (62–88)	56 (42–75)	75 (57–91)	59 (44–77)	66 (47–82)	56 (40–78)
median, (IQR)								
NT-proBNP pg/mL, median,	192 (68–558)	968 (250–	485 (167–	2580 (1204–	659 (218–	1801 (522–	1279 (461–	2793
(IQR)		2478)	1191)	6465)	2067)	4752)	3330)	(1220–
								6314)
NT-proBNP ≥125pg/ml, n (%)	266/452 (59)	32/36 (89)	240/299 (80)	55/55 (100)	225/277 (81)	87/92 (95)	256/277 (92)	126/128
W problem 2123pg/mi, m (70)								(98)
NT-proBNP groups, n (%)								
<300pg/mL	276/452 (61)	10/36 (28)	119/299 (40)	2/55 (4)	88/277 (32)	11/91 (12)	50/277 (18)	6/128 (5)

300–999pg/mL	99/452 (22)	8/36 (22)	91/299 (30)	10/55 (18)	78//277 (28)	21/91 (23)	71/277 (26)	16/128 (13)
1000–1999pg/mL	38/452 (8)	6/36 (17)	48/299 (16)	10/55 (18)	40//277 (14)	16/91 (18)	51/277 (18)	28/128 (22)
≥2000pg/mL	39/452 (9)	12/36 (33)	41/299 (14)	33/55 (60)	71//277 (26)	43/91 (47)	105/277 (38)	78/128 (61)
Sodium mmol/L, median, (IQR) *	138 (136–140)	138 (136–140)	139 (137–141)	139 (135–141)	138 (136–140)	137 (134–140)	139 (137–	138 (135–
							141)	140)
Urea mmol/L, median, (IQR) *	5.5 (4.4–7.0)	6.4 (5.0–10.3)	5.6 (4.6–6.9)	6.9 (5.8–9.1)	5.8 (4.5–7.2)	7.8 (5.7–11.8)	6.3 (4.9–8.8)	9.2 (6.1–
orea minor, E, median, (really								13.1)
Hemoglobin g/L, median, (IQR) *	134 (121–145)	119 (109–134)	137 (124–148)	124 (105–137)	131 (118–144)	122 (105–139)	130 (116–	121 (110–
Hemoglobin g/ L, median, (iQN)							142)	136)
Pharmacotherapy, n (%)								
Beta-blocker	247/452 (55)	18/36 (50)	153/299 (51)	29/55 (53)	173 /277 (63)	59/92 (64)	161/277 (58)	68/128 (53)
ACE-inhibitors or ARB	223/452 (49)	18/36 (50)	137/299 (46)	24/55 (44)	164/277 (59)	51/92 (55)	138/277 (50)	61/128 (48)
NOAC	9/452 (2)	0/36 (0)	133/299 (44)	25/55 (45)	6/277 (2)	3/92 (3)	133/277 (48)	57/128 (45)

Warfarin	4/452 (1)	1/36 (3)	65/299 (22)	13/55 (24)	8/277 (3)	5/92 (5)	72/277 (26)	38/128 (30)
Diuretic	83/452 (18)	14/36 (39)	59/299 (20)	24/55 (44)	97/277 (35)	30/92 (33)	135/277 (49)	41/128 (32)
MRA	6/452 (1)	0/36 (0)	5/299 (2)	4/55 (7)	23/277 (7)	22/92 (24)	20/277 (7)	24/128 (19)
Complex device (ICD or CRT)	4/452 (1)	1/36 (3)	2/299 (1)	3/55 (5)	14/277 (5)	12/92 (13)	20/277 (7)	18/128 (14)

ACE, Angiotensin converting enzyme; AF, atrial fibrillation; ARB, angiotensin receptor blocker; BMI, body mass index; BP, blood pressure; CKD-EPI,

Chronic Kidney Disease Epidemiology Collaboration; CRT, cardiac resynchronization therapy; eGFR, estimate glomerular filtration rate; ICD, implantable

cardioverter defibrillator; IQR, interquartile range; LBBB, left bundle branch block; MRA, mineralocorticoid receptor antagonist; Na, sodium; NOAC,

Novel oral anticoagulant; NT-proBNP, N-terminal pro-B-type natriuretic peptide; VKA, vitamin K antagonist.

^{*} Baseline data were missing in 3.6% of the study population for BMI, 2.4% for hemoglobin, 2.8% for urea and sodium.

Table S3. Outcomes stratified according to AF and HF phenotype groups.

Patient Group	Composite outcome	HF Hospitalization	Cardiovascular Death	All-Cause Mortality
	Events/person-yrs	Events/person-yrs	Events/person-yrs	Events/person-yrs
	(incidence/100 person-yrs)	(incidence/100 person-yrs)	(incidence/100 person-yrs)	(incidence/100 person-yrs)
Entire Cohort	310/3381 (9.2)	202/3686 (5.5)	168/3657 (4.6)	254/3657 (7.0)
Neither AF nor HF	36/1135 (3.2))	18/1190 (1.5)	22/1160 (1.9)	40/1159 (3.4)
AF only	55/775 (7.1)	34/826 (4.1)	32/819 (3.9)	47/819 (5.7)
HF only	91/759 (12.1)	59/824 (7.2)	52/828 (6.3)	66/828 (8.0)
AF plus HF	128/722 (17.7)	91/846 (10.8)	62/850 (7.3)	101/850 (11.9)

Table S4. C-Statistic of NT-proBNP as a continuous variable in each patient group for the composite outcome, its individual components i.e., HF hospitalization and cardiovascular death, and all-cause mortality.

Patient Group	Composite Outcome	P Value	HF Hospitalization	P Value	Cardiovascular	P Value	All-Cause Mortality	P Value
	C-Statistic (95% CI)		C-Statistic (95% CI)		Death		C-Statistic (95% CI)	
					C-Statistic (95% CI)			
Entire Cohort	0.74 (0.72 to 0.77)	<0.001	0.72 (0.69 to 0.75)	<0.001	0.76 (0.73 to 0.80)	<0.001	0.72 (0.69 to 0.75)	<0.001
Neither AF nor HF	0.73 (0.65 to 0.81)	<0.001	0.74 (0.64 to 0.84)	<0.001	0.74 (0.63 to 0.85)	<0.001	0.67 (0.59 to 0.76)	<0.001
AF only	0.82 (0.77 to 0.87)	<0.001	0.79 (0.72 to 0.85)	<0.001	0.86 (0.80 to 0.91)	<0.001	0.79 (0.73 to 0.85)	<0.001
HF only	0.66 (0.60 to 0.72)	<0.001	0.64 (0.57 to 0.71)	<0.001	0.71 (0.64 to 0.78)	<0.001	0.70 (0.63 to 0.77)	<0.001
AF plus HF	0.66 (0.61 to 0.70)	<0.001	0.61 (0.55 to 0.66)	<0.001	0.68 (0.62 to 0.75)	<0.001	0.65 (0.59 to 0.71)	<0.001

Table S5. Optimal cut-point in the entire cohort and each patient group (Youden index) and important cut-offs with associated area under the ROC curve, sensitivity, specificity, positive predictive value and negative predictive value of NT-proBNP for the composite outcome (heart failure hospitalization or cardiovascular death).

NT-proBNP cut-off	Patient Group	AUC (95% CI)	Sensitivity (95% CI)	Specificity (95% CI)	Positive predictive	Negative predictive
					value (95% CI)	value (95% CI)
Optimal cut-point						
(Youden index)						
1079pg/ml	Entire Cohort	0.70 (0.68 to 0.73)	71% (66% to 76%)	69% (67% to 72%)	36% (32% to 39%)	91% (89% to 93%)
229pg/ml	Neither AF nor HF	0.68 (0.61 to 0.75)	81% (64% to 92%)	55% (51% to 60%)	13% (9% to 18%)	97% (95% to 99%)
1182pg/ml	AF only	0.77 (0.71 to 0.83)	78% (65% to 88%)	75% (70% to 80%)	36% (28% to 46%)	95% (91% to 97%)
1407pg/ml	HF only	0.64 (0.58 to 0.70)	60% (49% to 70%)	68% (62% to 74%)	39% (31% to 47%)	84% (78% to 88%)
2128pg/ml	AF plus HF	0.62 (0.57 to 0.67)	60% (51% to 69%)	64% (58% to 69%)	43% (36% to 51%)	78% (72% to 83%)
Important cut-offs						

125pg/ml	Entire Cohort	0.60 (0.59 to 0.62)	97% (94% to 98%)	24% (22% to 27%)	23% (21% to 26%)	97% (94% to 98%)
	Neither AF nor HF	0.65 (0.59 to 0.71)	89% (74% to 97%)	41% (37% to 46%)	11% (7% to 15%)	98% (95% to 99%)
	AF only	0.60 (0.58 to 0.62)	100% (94 to 100%)	20% (15% to 25%)	19% (14% to 24%)	100% (94% to 100%)
	HF only	0.57 (0.53 to 0.60)	95% (88 to 98%)	19% (14% to 24%)	28% (23% to 33%)	91% (81% to 97%)
	AF plus HF	0.53 (0.51 to 0.55)	98% (95% to 100%)	8% (5% to 11%)	33% (28% to 38%)	91% (72% to 99%)
300pg/ml	Entire Cohort	0.66 (0.64 to 0.68)	90% (0.87 to 0.93)	41% (38% to 44%)	27% (24% to 30%)	95% (93% to 96%)
	Neither AF nor HF	0.67 (0.59 to 0.74)	72% (55% to 86%)	61% (56% to 66%)	13% (9% to 18%)	97% (94% to 98%)
	AF only	0.68 (0.64 to 0.72)	96% (88% to 100%)	40% (34% to 46%)	23% (18% to 29%)	98% (94% to 100%)
	HF only	0.59 (55 to 0.64)	87% (78% to 93%)	32% (26% to 38%)	30% (24% to 36%)	88% (80% to 94%)
	AF plus HF	0.57 (0.54 to 0.60)	95% (90% to 98%)	18% (14% to 23%)	35% (30% to 40%)	89% (78% to 96%)
1000pg/ml	Entire Cohort	0.70 (0.67 to 0.73)	73% (67% to 78%)	67% (64% to 69%)	34% (31% to 38%)	91% (89% to 93%)
	Neither AF nor HF	0.67 (0.58 to 0.75)	50% (33% to 67%)	83% (79% to 86%)	19% (12% to 28%)	95% (93% to 97%)

	AF only	0.74 (0.68 to 0.80)	78% (65% to 88%)	70% (65% to 75%)	33% (25% to 41%)	95% (91% to 97%)
	HF only	0.62 (0.56 to 0.68)	64% (54% to 74%)	60% (54% to 66%)	35% (28% to 42%)	83% (78% to 88%)
	AF plus HF	0.63 (0.59 to 0.68)	83% (75% to 89%)	44% (38% to 50%)	41% (35% to 47%)	85% (78% to 90%)
2000pg/ml	Entire Cohort	0.67 (0.64 to 0.70))	53% (48% to 59%)	80% (78% to 83%)	39% (35% to 44%)	88% (86% to 90%)
	Neither AF nor HF	0.62 (0.54 to 0.70)	33% (19% to 51%)	91% (88% to 94%)	24% (13% to 38%)	95% (92% to 96%)
	AF only	0.73 (0.66 to 0.80)	60% (46% to 73%)	86% (82% to 90%)	45% (33% to 57%)	92% (88% to 95%)
	HF only	0.61 (0.55 to 0.66)	47% (36% to 57%)	74% (69% to 79%)	38% (29% to 47%)	81% (75% to 85%)
	AF plus HF	0.62 (0.56 to 0.67)	61% (52% to 69%)	62% (56% to 68%)	43% (35% to 50%)	78% (71% to 83%)

Table S6. Performance of important cut-offs at predicting the composite outcome evaluated using discrimination, calibration, and reclassification. An NT-proBNP cut-off of 300pg/ml was used as a reference for NRI with two risk levels were selected: >20% and <20% risk of the composite outcome.

NT-proBNP		Discrimination	Calibration				Reclassification	
Cut-off								
		C-Statistic	Brier Score	AIC	BIC	Likelihood Ratio	NRI (20%)	IDI
125pg/ml	Entire Cohort	0.60 (0.58 to 0.61)	0.12 (0.10 to 0.13)	4425	4430	p=0.013	No change	0.001 (p=0.040)
	Neither AF nor HF	0.65 (0.60 to 0.70)	0.06 (0.04 to 0.08)	428	432	p=0.076	No change	0.003 (<i>p</i> =0.120)
	AF only	0.59 (0.57 to 0.61)	0.11 (0.08 to 0.14)	615	619	p=0.101	No change	-0.003 (p<0.001)
	HF only	0.56 (0.53 to 0.59)	0.13 (0.11 to 0.16)	1050	1054	p=0.270	No change	0.002 (<i>p</i> =0.298)
	AF plus HF	0.53 (0.51 to 0.54)	0.14 (0.12 to 0.16)	1472	1476	p=0.679	No change	0.0001 (<i>p</i> =0.734)
300pg/ml	Entire Cohort	0.65 (0.63 to 0.66)	0.11 (0.99 to 0.12)	4364	4370	Reference	Reference	Reference
	Neither AF nor HF	0.66 (0.59 to 0.74)	0.06 (0.41 to 0.08)	428	432	Reference	Reference	Reference
	AF only	0.67 (0.64 to 0.70)	0.90 (0.72 to 0.12)	601	605	Reference	Reference	Reference

	HF only	0.59 (0.55 to 0.62)	0.13 (0.11 to 0.16)	1047	1051	Reference	Reference	Reference
	AF plus HF	0.56 (0.54 to 0.58)	0.14 (0.12 to 0.16)	1463	1467	Reference	Reference	Reference
1000pg/ml	Entire Cohort	0.69 (0.66 to 0.71)	0.10 (0.09 to 0.12)	4345	4350	P<0.001	0.08 (<i>p</i> =0.003)	0.04 (<i>p</i> <0.001)
	Neither AF nor HF	0.66 (0.58 to 0.74)	0.06 (0.04 to 0.08)	425	429	P=0.017	No change	0.02 (<i>p</i> =0.013)
	AF only	0.73 (0.68 to 0.78)	0.09 (0.06 to 0.11)	590	594	P<0.001	0.12 (<i>p</i> =0.062)	0.06 (<i>p</i> <0.001)
	HF only	0.62 (0.57 to 0.67)	0.13 (0.11 to 0.15)	1043	1047	P=0.008	No change	0.02 (<i>p</i> =0.014)
	AF plus HF	0.62 (0.59 to 0.65)	0.13 (0.11 to 0.15)	1448	1452	<i>P</i> <0.001	0.13 (<i>p</i> =0.002)	0.04 (<i>p</i> <0.001)
2000pg/ml	Entire Cohort	0.66 (0.63 to 0.69	0.11 (0.10 to 0.12)	4372	4377	<i>P</i> <0.001	0.03 (<i>p</i> =0.507)	0.05 (<i>p</i> <0.001)
	Neither AF nor HF	0.62 (0.54 to 0.70)	0.06 (0.04 to 0.08)	428	432	P=0.014	No change	0.02 (<i>p</i> =0.028)
	AF only	0.72 (0.66 to 0.78)	0.08 (0.06 to 0.10)	586	590	<i>P</i> <0.001	0.10 (<i>p</i> =0.263)	0.11 (<i>p</i> <0.001)
	HF only	0.60 (0.55 to 0.65)	0.13 (0.11 to 0.16)	1046	1050	P=0.008	No Change	0.02 (<i>p</i> =0.016)
	AF plus HF	0.61 (0.56 to 0.65)	0.14 (0.11 to 0.16)	1458	1462	P<0.001	No Change	0.03 (<i>p</i> =0.001)

Table S7. Univariate Cox proportional hazards models based on cut-offs with B coefficient and baseline hazard in the entire cohort.

NT-proBNP cut-off	B coefficient	95% Confidence	Standard Error	Z score	<i>p</i> value	Baseline Hazard
		Interval				
125pg/ml	2.08	1.48 to 2.69	0.31	6.79	<0.001	0.03
300pg/ml	1.78	1.39 to 2.15	0.19	9.20	<0.001	0.05
1000pg/ml	1.55	1.30 to 1.80	0.13	12.17	<0.001	0.08
2000pg/ml	1.40	1.18 to 1.63	0.11	12.32	<0.001	0.11

Transparent reporting of a multivariable prediction model for individual prognosis or diagnosis (TRIPOD): The TRIPOD Checklist for Prediction Model Development (Collins GS, Reitsma JB, Altman DG, Moons KG. Transparent reporting of a multivariable prediction model for individual prognosis or diagnosis (TRIPOD): The TRIPOD statement).

Section/Topic			Checklist Item	Page	
Title and abstra	act				
Title	1	D;V	Identify the study as developing and/or validating a multivariable prediction model, the target population, and the outcome to be predicted.	1	
Abstract	2	D;V	Provide a summary of objectives, study design, setting, participants, sample size, predictors, outcome, statistical analysis, results, and conclusions.	3-4	
			Introduction		
Background and	3a	D;V	Explain the medical context (including whether diagnostic or prognostic) and rationale for developing or validating the multivariable prediction model, including references to existing models.	6-7	
objectives	3b	D;V	Specify the objectives, including whether the study describes the development or validation of the model or both.	6-7	
		•	Methods		
Source of	4a	D;V	Describe the study design or source of data (e.g., randomized trial, cohort, or registry data), separately for the development and validation data sets, if applicable.	7-8	
data	4b	D;V	Specify the key study dates, including start of accrual; end of accrual; and, if applicable, end of follow-up.	7-8 Figure 1	
D	5a	D;V	Specify key elements of the study setting (e.g., primary care, secondary care, general population) including number and location of centers.		
Participants	5b	D;V	Describe eligibility criteria for participants.	7-8	
	5c	D;V	Give details of treatments received, if relevant.	n/a	
Outcome	6a	D;V	Clearly define the outcome that is predicted by the prediction model, including how and when assessed.		
	6b	D;V	Report any actions to blind assessment of the outcome to be predicted.	n/a	
Predictors	7a	D;V	Clearly define all predictors used in developing the multivariable prediction model, including how and when they were measured.	11-13	
	7b	D;V	Report any actions to blind assessment of predictors for the outcome and other predictors.	n/a	
Sample size	8	D;V	Explain how the study size was arrived at.	7-8	
Missing data	9	D;V	Describe how missing data were handled (e.g., complete-case analysis, single imputation, multiple imputation) with details of any imputation method.	11-13	
analysis methods	10a	D	Describe how predictors were handled in the analyses.	11-13	
	10b	D	Specify type of model, all model-building procedures (including any predictor selection), and method for internal validation.	11-13	
	10c	V	For validation, describe how the predictions were calculated.	n/a	

	10d	D;V	Specify all measures used to assess model performance and, if relevant, to compare multiple models.	11-13
	10e	V	Describe any model updating (e.g., recalibration) arising from the validation, if done.	n/a
Risk groups	11	D;V	Provide details on how risk groups were created, if done.	11
Development vs. validation	12	V	For validation, identify any differences from the development data in setting, eligibility criteria, outcome, and predictors.	n/a
	ı		Results	T
	13a	D;V	Describe the flow of participants through the study, including the number of participants with and without the outcome and, if applicable, a summary of the follow-up time. A diagram may be helpful.	Figure 1
Participants	13b	D;V	Describe the characteristics of the participants (basic demographics, clinical features, available predictors), including the number of participants with missing data for predictors and outcome.	Table 1
	13c	V	For validation, show a comparison with the development data of the distribution of important variables (demographics, predictors and outcome).	n/a
Model	14a	D	Specify the number of participants and outcome events in each analysis.	14 Figure 1
development	14b	D	If done, report the unadjusted association between each candidate predictor and outcome.	Table 2
Model specification	15a	D	Present the full prediction model to allow predictions for individuals (i.e., all regression coefficients, and model intercept or baseline survival at a given time point).	Table 2 Supplementa Materials Table 7
	15b	D	Explain how to use the prediction model.	15-18
Model performance	16	D;V	Report performance measures (with CIs) for the prediction model.	Supplementa Materials Table 5, Supplementa Materials Table 6
Model- updating	17	V	If done, report the results from any model updating (i.e., model specification, model performance).	n/a
upuuting			Discussion	
Limitations	18	D;V	Discuss any limitations of the study (such as nonrepresentative sample, few events per predictor, missing data).	22
Interpretatio	19a	V	For validation, discuss the results with reference to performance in the development data, and any other validation data.	n/a
n	19b	D;V	Give an overall interpretation of the results, considering objectives, limitations, results from similar studies, and other relevant evidence.	18
Implications	20	D;V	Discuss the potential clinical use of the model and implications for future research.	20-21
	ı		Other information	T
Supplementar y information	21	D;V	Provide information about the availability of supplementary resources, such as study protocol, Web calculator, and data sets.	n/a
Funding	22	D;V	Give the source of funding and the role of the funders for the present study.	23