Genetic predisposition, modifiable risk factor profile and long-term dementia risk in the general population

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Appendix A. Supplementary figure	.3
Appendix B. Supplementary tables	4

Appendix A. Supplementary figure

Figure 1. Flow-chart of study population



Appendix B. Supplementary tables

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Chr	BP	SNP	Gene	Risk	Locus discovered in	Effect size	Minor	Maf_RS1	R ² -RS1	R ² -RS2
				allele			Allele			
1	207802552	rs4844610	CR1	А	Lambert et al. (2009)	0.157003749	А	0.19039	0.9968	0.99498
2	127892810	rs6733839	BIN1	Т	Seshadri et al.	0.182321557	Т	0.39584	0.95965	0.91998
2	233981912	rs10933431	INPP5D	С	Lambert et al.	0.094310679	G	0.21024	0.93279	0.92668
5	88223420	rs190982	MEF2C	А	Lambert et al. (2013)	0.061875404	G	0.395	0.97929	0.93602
6	41129252	rs75932628	TREM2	Т	Guerreiro et al., Jonsson et al.	0.732367894	Т	0.00197	0.7976	0.77763
6	47431284	rs9473117	CD2AP	С	Hollingworth et al. , Naj et al.	0.086177696	С	0.27668	0.99424	0.99517
6	41129207	rs143332484	TREM2	Т	Sims et al., (2017)	0.457424847	Т	0.01288	0.83663	0.8099
7	100091795	rs12539172	ZCWPW1	С	Lambert et al., Kunkle et al.	0.083381609	Т	0.34739	0.99958	0.99947
7	143099133	rs10808026	EPHA1	С	Hollingworth et al. , Naj et al.	0.105360516	А	0.20154	0.98093	0.98031
7	37844263	rs4723711	NME8	А	Lambert et al. (2013)	0.061875404	Т	0.35445	0.99933	0.99924
8	27219987	rs73223431	PTK2B	Т	Lambert et al. (2013)	0.09531018	Т	0.38498	0.99562	0.99678
8	27467686	rs9331896	CLU	Т	Harold et al., Lambert et al. (2009)	0.127833372	С	0.39809	0.90325	0.97633
11	47380340	rs3740688	CELF1	Т	Lambert et al. (2013)	0.083381609	G	0.4464	0.98617	0.98504
11	59936926	rs7933202	MS4A2	А	Hollingworth et al. , Naj et al.	0.116533816	С	0.40726	0.99544	0.99719
11	85868640	rs3851179	PICALM	С	Harold et al.	0.127833372	Т	0.37041	0.9992	0.99951
11	121435587	rs11218343	SORL1	Т	Lambert et al. (2013)	0.223143551	С	0.04651	0.9979	0.99592
14	53391680	rs17125924	FERMT2	G	Lambert et al. (2013)	0.131028262	G	0.09948	0.99862	0.99842
14	92932828	rs12881735	SLC24A4	Т	Lambert et al. (2013)	0.083381609	С	0.24057	0.99526	0.99591
15	59045774	rs593742	ADAM10	А	Kunkle et al. (2019)	0.072570693	G	0.33247	0.99626	0.99625
16	19808163	rs7185636	IQCK	Т	Kunkle et al. (2019)	0.083381609	С	0.17569	0.98562	0.98756
16	79355857	rs62039712	WWOX	А	Kunkle et al. (2019)	0.148420005	А	0.12707	0.77987	0.80096
16	81942028	rs72824905	PLCG2	С	Sims et al., (2017)	0.431782416	С	0.01013	0.94589	0.94491
17	61538148	rs138190086	ACE	А	Kunkle et al. (2019)	0.262364264	А	0.01703	0.83116	0.86548
17	47297297	rs616338	ABI3	Т	Sims et al., (2017)	0.350656872	Т	0.01137	0.64079	0.62751
19	1056492	rs3752246	ABCA7	G	Hollingworth et al. , Naj et al.	0.139761942	G	0.16336	0.91902	0.93084
20	54997568	rs6024870	CASS4	G	Lambert et al. (2013)	0.127833372	А	0.0815	0.9842	0.98621
21	28156856	rs2830500	ADAMTS1	С	Kunkle et al. (2019)	0.072570693	А	0.30571	0.97586	0.97907

Ordered by chromosome number. Effect estimates taken from Kunkle et al.¹ and Sims et al.² (TREM2- rs143332484, PLCG2 and ABI3) Additional references: Harold et al.³, Seshadri et al.⁴, Hollingworth et al.⁵, Naj et

al.⁶, Lambert et al. $(2009, 2013)^{7.8}$, Jonsson et al.⁹, and Ruiz et al.¹⁰ Minor allele Frequency (Maf) of Rotterdam Study (RS) 1 is shown and is representative of the MAF in RS2. R²⁼ imputation quality. RS1= initial Rotterdam Study cohort and RS2=first extension Rotterdam Study. Rotterdam Study cohorts were imputed separately. Gene names are ncbi gene names assigned to the loci in the corresponding references

Table 2. Absolute 15-year anticipated risk of dementia according to modifiable risk factor profiles, across different levels of genetically determined risk based on *APOE* carrier status

		N/n	Low genetic risk	N/n	Intermediate	N/n	High genetic risk
			Mean (95% CI)		genetic risk		Mean (95% CI)
					Mean (95% CI)		
Overall	Overall	887/85	15.8% (4.6;32.0)	3718/456	16.1% (10.0;19.2)	1747/374	18.6% (12.8;23.8)
Modifiable	Favorable	568/44	12.6% (4.5;26.8)	2453/253	13.5% (8.9;15.6)	1132/241	18.2% (12.7;22.2)
risk factor profile	Intermediate	224/23	16.7% (8.0;36.5)	884/139	20.6% (12.6;22.1)	443/97	19.2% (12.6;21.7)
1	Unfavorable	95/18	32.1% (0.0;59.9)	381/64	22.0% (8.3;39.2)	172/36	19.5% (2.5;27.3)

Based on age, sex, education, and modifiable risk factor profile. Confidence intervals computed based on 1000 bootstrap samples. Since baseline characteristics of participants across strata of genetic risk are not identical, anticipated absolute risks may vary beyond the contribution of genetic and modifiable risk factors. Given that these estimations also take into account the contribution of other risk factors for dementia risk, namely age, sex and educational level, predicted risks are not directly comparable across genetic and modifiable risk strata.

Abbreviations: N=number of people at risk, n=number of cases.

Table 3. Risk of incident dementia stratified by modifiable risk factor profiles and genetic risk, separately for participants below and above the median age (68.2 years)

Genetic risk			Low		Intermediate		High
		N/n	HR (95% CI)	N/n	HR (95% CI)	N/n	HR (95% CI)
	Favorable	311/7	Ref.	1421/110	Ref.	641/79	Ref.
<68.2 years	Intermediate	84/2	1.21 (0.24;6.19)	371/19	1.71 (1.00;2.93)	207/27	1.23 (0.80;1.92)
	Unfavorable	33/3	7.50 (1.73;32.4)	133/10	2.69 (1.33;5.44)	64/6	0.96 (0.42;2.23)
	Favorable	257/37	Ref.	1319/46	Ref.	491/162	Ref.
≥68.2 years	Intermediate	140/21	1.14 (0.96;1.36)	513/120	1.26 (1.00;1.60)	236/70	0.97 (0.73;1.30)
	Unfavorable	62/15	1.31 (1.04;1.63)	248/54	1.33 (0.97;1.82)	118/30	1.05 (0.70;1.57)
<68.2 years ≥68.2 years	Intermediate Unfavorable Favorable Intermediate Unfavorable	84/2 33/3 257/37 140/21 62/15	1.21 (0.24;6.19) 7.50 (1.73;32.4) Ref. 1.14 (0.96;1.36) 1.31 (1.04;1.63)	371/19 133/10 1319/46 513/120 248/54	1.71 (1.00;2.93) 2.69 (1.33;5.44) Ref. 1.26 (1.00;1.60) 1.33 (0.97;1.82)	207/27 64/6 491/162 236/70 118/30	1.23 (0.80;1.92) 0.96 (0.42;2.23) Ref. 0.97 (0.73;1.30) 1.05 (0.70;1.57)

Adjusted for: age, sex and education Abbreviations: n=number of cases, N=number of people at risk, HR=hazard ratio, CI=confidence interval.

Table 4. Risk of incident dementia stratified by modifiable risk factor profiles and genetic risk, separately for participants below and above age

 of 70 years

Genetic risk			Low		Intermediate		High
		N/n	HR (95% CI)	N/n	HR (95% CI)	N/n	HR (95% CI)
	Favorable	351/10	Ref.	1515/65	Ref.	716/90	Ref.
<70.0 years	Intermediate	98/3	1.29 (0.35;4.81)	423/25	1.68 (1.06;2.68)	239/3	1.19 (0.79;1.78)
	Unfavorable	39/4	3.93 (1.19;12.94)	157/10	1.80 (0.92;3.53)	73/11	1.65 (0.87;3.13)
	Favorable	217/34	Ref.	938/188	Ref.	416/151	Ref.
≥70.0 years	Intermediate	126/20	1.14 (0.63;2.05)	462/114	1.25 (0.98;1.59)	204/64	0.95 (0.70;1.29)
	Unfavorable	56/14	2.18 (1.12;4.24)	224/54	1.38 (1.00;1.89)	99/25	0.85 (0.55;1.31)

Adjusted for: age, sex and education Abbreviations: n=number of cases, N=number of people at risk, HR=hazard ratio, CI=confidence interval.

Genetic risk		Low		Intermediate		High	
	N/n	HR (95% CI)	N/n	HR (95% CI)	N/n	HR (95% CI)	
Favorable	252/15	Ref.	1185/110	Ref.	557/108	Ref.	
Intermediate	74/3	0.88 (0.25;3.09)	350/41	1.39 (0.96;2.00)	175/22	0.84 (0.52;1.33)	
Unfavorable	32/5	3.72 (1.24;11.1)	109/8	1.04 (0.50;2.17)	45/5	0.67 (0.27;1.68)	
1							
Favorable	316/29	Ref.	1268/143	Ref.	575/133	Ref.	
Intermediate	150/20	1.28 (0.70;2.34)	534/98	1.24 (0.95;1.63)	268/75	1.09 (0.82;1.47)	
Unfavorable	62/13	2.33 (1.17;4.62)	272/56	1.48 (1.07;2.04)	127/31	1.15 (0.77;1.71)	
	Favorable Intermediate Unfavorable Favorable Intermediate Unfavorable	N/nFavorable252/15Intermediate74/3Unfavorable32/5Favorable316/29Intermediate150/20Unfavorable62/13	N/n HR (95% CI) Favorable 252/15 Ref. Intermediate 74/3 0.88 (0.25;3.09) Unfavorable 32/5 3.72 (1.24;11.1) Favorable 316/29 Ref. Intermediate 150/20 1.28 (0.70;2.34) Unfavorable 62/13 2.33 (1.17;4.62)	LowN/nHR (95% CI)N/nFavorable252/15Ref.1185/110Intermediate74/30.88 (0.25;3.09)350/41Unfavorable32/53.72 (1.24;11.1)109/8Favorable316/29Ref.1268/143Intermediate150/201.28 (0.70;2.34)534/98Unfavorable62/132.33 (1.17;4.62)272/56	Low Intermediate N/n HR (95% CI) N/n HR (95% CI) Favorable 252/15 Ref. 1185/110 Ref. Intermediate 74/3 0.88 (0.25;3.09) 350/41 1.39 (0.96;2.00) Unfavorable 32/5 3.72 (1.24;11.1) 109/8 1.04 (0.50;2.17) Favorable 316/29 Ref. 1268/143 Ref. Intermediate 150/20 1.28 (0.70;2.34) 534/98 1.24 (0.95;1.63) Unfavorable 62/13 2.33 (1.17;4.62) 272/56 1.48 (1.07;2.04)	Low Intermediate N/n HR (95% CI) N/n HR (95% CI) N/n Favorable 252/15 Ref. 1185/110 Ref. 557/108 Intermediate 74/3 0.88 (0.25;3.09) 350/41 1.39 (0.96;2.00) 175/22 Unfavorable 32/5 3.72 (1.24;11.1) 109/8 1.04 (0.50;2.17) 45/5 Favorable 316/29 Ref. 1268/143 Ref. 575/133 Intermediate 150/20 1.28 (0.70;2.34) 534/98 1.24 (0.95;1.63) 268/75 Unfavorable 62/13 2.33 (1.17;4.62) 272/56 1.48 (1.07;2.04) 127/31	

Table 5. Risk of incident dementia stratified by modifiable risk factor profiles and genetic risk, separately for men and women

Adjusted for: age, sex and education Abbreviations: n=number of cases, N=number of people at risk, HR=hazard ratio, CI=confidence interval.

Polygenic risk score tertiles	N/n	Model 1	Model 2
		HR (95% CI)	HR (95% CI)
Low risk (first tertile)	1864/189	Reference	Reference
Intermediate risk (second	1885/213	1.16 (0.96;1.42)	1.15 (0.95;1.40)
tertile)			
High risk (third tertile)	1811//251	1.53 (1.27;1.85)	1.54 (1.27;1.86)
P for trend		7.82×10 ⁻⁶	7.46×10 ⁻⁶

Table 6. Risk of incident Alzheimer's disease according to tertiles of the polygenic risk score

Model 1 - adjusted for: age, sex and education

Model 2 - additionally adjusted for: parental history of dementia, history of stroke, systolic blood pressure, total and high-density lipoprotein cholesterol

Abbreviations: N=number of individuals at risk, n=number of Alzheimer's disease cases during follow-up,

HR=hazard ratio, CI=confidence interval. The log rank test was used to calculate P values. Two-sided P values were uncorrected for multiple testing.

Table 7. Risk of incident Alzheimer's disease while stratifying participants both on tertiles of	a
polygenic risk score and different modifiable risk factor profiles	

Polygenic risk score	Risk factor profile	N/n	HR (95% CI)
	Favorable	1191/99	Reference
Low (Lowest tertile)	Intermediate	478/60	1.11 (0.79;1.56)
	Unfavorable	195/30	1.46 (0.95;2.25)
	P for trend		0.0547
Intermediate (Middle)	Favorable	1230/125	Reference
	Intermediate	454/63	1.22 (0.89;1.68)
	Unfavorable	201/25	1.08 (0.69;1.67)
	P for trend		0.3974
	Favorable	1204/159	Reference
High (Highest tertile)	Intermediate	429/60	0.97 (0.71;1.31)
	Unfavorable	178/32	1.27 (0.85;1.89)
	P for trend		0.9574

Adjusted for: age, sex and education Abbreviations: n=number of cases, N=number of people at risk, HR=hazard ratio, CI=confidence interval. The log rank test was used to calculate P values. Two-sided P values were uncorrected for multiple testing.

APOE-related risk	Risk factor profile	N/n	HR (95% CI)
	Favorable	18/1	Reference
	Intermediate	426/30	1.31 (0.18;9.70)
Low ($\epsilon 2\epsilon 2/\epsilon 2\epsilon 3$)	Unfavorable	443/54	1.93 (2.64;14.11)
	P for trend		0.0942
	Favorable	35/4	Reference
	Intermediate	1871/199	0.80 (0.28;2.13)
Intermediate (ɛ3ɛ3)	Unfavorable	1812/253	0.85 (0.31;2.33)
	P for trend		0.2944
	Favorable	20/5	Reference
	Intermediate	849/189	0.72 (0.30;1.77)
High ($\epsilon 2\epsilon 4/\epsilon 3\epsilon 4/\epsilon 4\epsilon 4$)	Unfavorable	778/180	0.55 (0.23;1.36)
	P for trend		0.0085

Table 8. Risk of incident dementia while stratifying participants both on *APOE* and on levels of modifiable risk based on the Ideal Cardiovascular Health metric

Adjusted for: age, sex and education

Abbreviations: n=number of cases, N=number of people at risk, HR=hazard ratio, CI=confidence interval. The log rank test was used to calculate P values. Two-sided P values were uncorrected for multiple testing.

APOE-related risk	Risk factor profile	N/n	HR (95% CI)
	Favorable	386/15	Reference
	Intermediate	225/20	1.22 (0.58;2.55)
Low ($\epsilon 2\epsilon 2/\epsilon 2\epsilon 3$)	Unfavorable	256/45	1.73 (0.73;4.06)
	P for trend		0.1866
	Favorable	1539/83	Reference
	Intermediate	991/123	1.34 (0.98;1.83)
Intermediate (ɛ3ɛ3)	Unfavorable	1061/237	1.66 (1.16;2.38)
	P for trend		0.0061
	Favorable	734/108	Reference
	Intermediate	480/109	1.01 (0.58;1.28)
High ($\epsilon 2\epsilon 4/\epsilon 3\epsilon 4/\epsilon 4\epsilon 4$)	Unfavorable	469/139	0.86 (0.74;1.37)
	P for trend		0.4528

Table 9. Risk of incident dementia while stratifying participants both on APOE and on their 10year predicted risk of fatal cardiovascular diseases

Adjusted for: age, sex and education Abbreviations: n=number of cases, N=number of people at risk, HR=hazard ratio, CI=confidence interval. The log rank test was used to calculate P values. Two-sided P values were uncorrected for multiple testing.

	N/n	Model 1	Model 2
		HR (95% CI)	HR (95% CI)
Abstaining from smoking	5093/790	0.97 (0.80;1.17)	0.97 (0.84;1.18)
Absence of depression	5717/792	0.65 (0.54;0.79)	0.66 (0.54;0.80)
Absence of diabetes	5618/796	0.74 (0.61;0.90)	0.74 (0.61;0.91)
Regular physical activity	3598/434	0.90 (0.78;1.05)	0.91 (0.78;1.06)
Avoiding social isolation	4526/559	1.04 (0.89;1.21)	1.05 (0.90;1.22)
Adherence to a healthy dietary pattern	952/181	1.09 (0.92;1.29)	1.10 (0.93;1.30)

Table 10. Health and lifestyle-specific associations with dementia risk

Model 1 - adjusted for: age, sex and education.

Model 2 - additionally adjusted for: parental history of dementia, history of stroke, systolic blood pressure, total and high-density lipoprotein cholesterol

Abbreviations: n=number of cases, N=number of people at risk, HR=hazard ratio, CI=confidence interval.

	N/n	Model 1	Model 2
		HR (95% CI)	HR (95% CI)
1. Abstaining from smoking	5093/790	0.97 (0.80;1.17)	0.97 (0.84;1.18)
2. Ideal body mass index	2071/306	1.09 (0.95;1.26)	1.08 (0.94;1.24)
3. Ideal physical activity	3598/434	0.74 (0.61;0.90)	0.74 (0.61;0.91)
4. Adherence to a healthy dietary pattern	952/181	1.09 (0.92;1.29)	1.10 (0.93;1.30)
5. Untreated total cholesterol <5.2 mmol/L	1565/209	0.97 (0.83;1.14)	1.04 (0.88;1.22)
6. Untreated systolic blood pressure <120	737/95	1.10 (0.90;1.37)	1.09 (0.88;1.35)
mmHg & diastolic blood pressure <80 mmHg			
7. Fasting plasma glucose <5.5 mmol/L	3013/447	1.00 (0.88;1.14)	1.01 (0.89;1.15)

Table 11. Associations of individual Ideal Cardiovascular Health components with the risk of dementia

Model 1 - adjusted for: age, sex and education.

Model 2 - additionally adjusted for: parental history of dementia and history of stroke.

Abbreviations: n=number of cases, N=number of people at risk, HR=hazard ratio, CI=confidence interval.

	N/n	Model 1	Model 2
		HR (95% CI)	HR (95% CI)
Age, per year	6352/915	1.13 (1.12;1.14)	1.13 (1.12;1.14)
Current smoking	1259/125	1.00 (0.81;1.24)	0.99 (0.80;1.23)
Systolic blood pressure, per	6352/915	1.01 (0.98;1.04)	1.01 (0.98;1.04)
10 mmHg			
Total cholesterol, per	6352/915	1.04 (0.97;1.11)	1.04 (0.97;1.12)
mmol/L			

Table 12. Associations of the individual SCORE risk equation components with the risk of dementia

Model 1 - adjusted for: sex.

Model 2 - additionally adjusted for: parental history of dementia and history of stroke.

Abbreviations: n=number of cases, N=number of people at risk, HR=hazard ratio, CI=confidence interval.

	Included participants	Excluded participants	
	N=6352	N=1433	
Age, years	69.1 (8.2)	70.7 (10.6)	
Women	3572 (56.2)	956 (66.7)	
Educational years, median (IQR)	10 (7-13)	10 (7-13)	
Parental history of dementia	427 (6.7)	13 (0.9)	
History of stroke	231 (3.6)	67 (4.7)	
Body mass index, kg/m ²	27.4 (4.0)	26.9 (3.9)	
Systolic blood pressure, mmHg	143 (21)	139 (19)	
Diastolic blood pressure, mmHg	77 (11)	80 (13)	
Total cholesterol, mmol/L	5.8 (1.0)	5.3 (1.0)	
High density lipoprotein cholesterol,	1.4 (0.4)	1.3 (0.3)	
mmol/L			
Fasting glucose, mmol/L	6.0 (1.6)	6.1 (2.1)	
Age of dementia diagnosis	83.1 (6.5)	84.2 (6.5)	
Modifiable health and lifestyle factor	'S		
No current smoking	5064 (79.7)	92 (79.3)*	
Absence of depression	5717 (90.0)	96 (85.0)*	
Absence of diabetes	5301 (83.5)	38 (82.6)*	
Regular physical activity	3598 (56.6)	1 (7.7)*	
Absence of social isolation	4508 (71.0)	79 (66.9)*	
Healthy diet score	749 (11.8)	136 (17.2)*	

Table 13. Baseline characteristics for included and excluded participants of the current study

Abbreviations: N=number of people at risk, SD=standard deviation, IQR=interquartile range. Data are shown for non-imputed data. * Percentages shown for available, non-missing data. Data presented as frequency (percent) for categorical values and mean \pm SD for continuous variables unless indicated otherwise.

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