

Supplementary Material

Interaction of Alzheimer's Disease-Associated Genetic Risk with Indicators of Socioeconomic Position on Mild Cognitive Impairment in the Heinz Nixdorf Recall Study

Supplementary Table 1. Risk alleles of loci associated with Alzheimer's disease [1] used for calculating the genetic risk score (GRS_{AD}) and their age- and sex-adjusted effects on mild cognitive impairment in the study population.

Original SNP	Proxy SNP	r ²	Chr:Position Build 37	Locus	Risk Allele	Risk Allele Frequency	OR (95% CI)	p
rs4844610	rs679515	0.92	1: 207802552	<i>CRI</i>	T	0.211	0.95 (0.81; 1.11)	0.50
rs6733839			2:127892810	<i>BINI</i>	T	0.395	0.92 (8.81; 1.05)	0.20
rs10933431	rs7570061	0.75	2:233981912	<i>INPP5D</i>	G	0.166	0.93 (0.79; 1.10)	0.39
rs9271058	rs9271192	1.00	6:32575406	<i>HLA -DRB1</i>	C	0.283	1.14 (0.99; 1.31)	0.07
rs75932628			6:41129252	<i>TREM2</i>	T	NA	NA	NA
rs9473117	rs10948363	0.98	6: 47431284	<i>CD2AP</i>	G	0.264	0.88 (0.76; 1.02)	0.08
rs12539172			7: 100091795	<i>NYAPI</i>	C	0.299	0.92 (0.81; 1.06)	0.24
rs10808026	rs11767557	1.00	7: 143099133	<i>EPHA1</i>	T	0.198	0.98 (0.84; 1.15)	0.85
rs73223431	rs28834970	0.98	8 27219987	<i>PTK2B</i>	C	0.334	1.00 (0.87; 1.14)	0.99
rs9331896			8: 27467686	<i>CLU</i>	A	0.387	1.08 (0.95; 1.23)	0.24
rs7920721	rs11257240	0.87	10: 11720308	<i>ECHDC3</i>	G	0.372	1.07 (0.94; 1.23)	0.29
rs3740688	rs10769256	0.64	11: 47380340	<i>SPI1</i>	C	0.411	1.00 (0.88; 1.14)	0.99
rs7933202	rs12453	0.92	11: 59936926	<i>MS4A2</i>	A	0.389	1.08 (0.95; 1.23)	0.25
rs3851179			11: 85868640	<i>PICALM</i>	G	0.360	0.97 (0.85; 1.11)	0.63
rs11218343			11: 121435587	<i>SORL1</i>	T	0.040	1.13 (0.81; 1.64)	0.48
rs17125924	rs17125944	0.94	14: 53391680	<i>FERMT2</i>	C	0.097	1.13 (0.91; 1.39)	0.27
rs12881735	rs10498633	1.00	14: 92932828	<i>SLC24A4</i>	G	0.234	0.95 (1.03; 1.05)	0.52
rs593742	rs474875	0.78	15: 59045774	<i>ADAM10</i>	A	0.324	0.89 (0.78; 1.01)	0.08
rs7185636	rs1858973	0.94	16: 19808163	<i>IQCK</i>	A	0.171	1.06 (0.89; 1.26)	0.53
rs138190086	rs28369023	1.00	17: 61538148	<i>ACE</i>	A	0.019	1.16 (0.73; 1.78)	0.51
rs6024870	rs6064392	0.82	20: 54997568	<i>CASS4</i>	G	0.086	1.15 (0.91; 1.47)	0.26
rs2830500	rs7276338	0.85	21: 28156856	<i>ADAMTS1</i>	C	0.272	1.04 (0.90; 1.21)	0.57

[1] Kunkle BW, Grenier-Boley B, Sims R, Bis JC, Damotte V, Naj AC, Boland A, Vronskaya M, van der Lee SJ, Amlie-Wolf A, et al. (2019) Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates A β , tau, immunity and lipid processing. *Nat Genet* **51**, 414–430.

Supplementary Table 2. Age- adjusted odds ratios (OR) and corresponding 95% confidence intervals (95% CI) on mild cognitive impairment in logistic regression models including main effects of income and education as indicators of socioeconomic position, *APOE* ε4 status and an AD-associated genetic risk score (GRS_{AD}), stratified by sex.

	Male			Female		
	n	OR (95%-CI)	p	n	OR (95%-CI)	p
Intercept		0.01 (0.005; 0.04)	4.6*10 ⁻¹⁵		0.02 (0.01; 0.06)	7.2*10 ⁻¹³
Income low	1,837	1.52 (1.15; 2.01)	0.003	1,772	1.26 (0.96; 1.66)	0.10
Age		1.04 (1.02; 1.06)	2.9*10 ⁻⁵		1.03 (1.01; 1.05)	0.0003
Intercept		0.01 (0.004; 0.03)	5.4*10 ⁻¹⁶		0.02 (0.01; 0.06)	9.0*10 ⁻¹⁴
Education low	1,901	1.28 (0.99; 1.67)	0.06	1,928	1.58 (1.10; 2.33)	0.02
Age		1.04 (1.02; 1.06)	2.7*10 ⁻⁶		1.03 (1.01; 1.05)	0.001
Intercept		0.01 (0.004- 0.04)	5.7*10 ⁻¹⁶		0.02 (0.01- 0.06)	9.5*10 ⁻¹⁴
<i>APOE</i> ε4	1,905	1.14 (0.85- 1.52)	0.36	1,929	1.40 (1.06- 1.84)	0.01
Age		1.04 (1.02- 1.06)	1.3*10 ⁻⁶		1.03 (1.02- 1.05)	9.1*10 ⁻⁵
Intercept		0.02 (0.004- 0.07)	8.9*10 ⁻⁸		0.02 (0.004- 0.09)	3.3*10 ⁻⁷
GRS _{AD}	1,905	0.99 (0.94- 1.04)	0.63	1,929	1.01 (0.96- 1.06)	0.75
Age		1.04 (1.03- 1.06)	1.2*10 ⁻⁶		1.03 (1.02- 1.05)	8.3*10 ⁻⁵

Supplementary Table 3. Age- adjusted effects and corresponding 95% confidence intervals (95% CI) on mild cognitive impairment in logistic regression models including main effects and interaction terms of the *APOE ε4* status and an AD-associated genetic risk score (GRS_{AD}) with indicators of socioeconomic status (years of education and income/month), stratified by sex.

	Male			Female		
	n	OR (95%-CI)	p	n	OR (95%-CI)	p
Intercept		0.01 (0.004- 0.04)	4.4*10 ⁻¹⁵		0.02 (0.01- 0.06)	6.9*10 ⁻¹³
Income low		1.57 (1.13- 2.19)	0.01		1.17 (0.84- 1.62)	0.35
<i>APOE ε4</i>	1,837	1.25 (0.75- 2.02)	0.38	1,772	1.18 (0.75- 1.83)	0.47
Age		1.04 (1.02- 1.06)	3.5*10 ⁻⁵		1.03 (1.01- 1.05)	0.0003
<i>APOE ε4</i> *Income low		0.90 (0.49- 1.69)	0.75		1.32 (0.74- 2.38)	0.35
Intercept		0.01 (0.004- 0.03)	7.1*10 ⁻¹⁶		0.02 (0.01- 0.06)	7.1*10 ⁻¹³
Education low		1.26 (0.92- 1.71)	0.14		1.28 (0.85- 1.99)	0.25
<i>APOE ε4</i>	1,901	1.10 (0.71- 1.69)	0.65	1,928	0.72 (0.30- 1.56)	0.43
Age		1.04 (1.02- 1.06)	3.1*10 ⁻⁶		1.03 (1.01- 1.05)	0.001
<i>APOE ε4</i> *Education low		1.08 (0.60- 1.94)	0.80		2.17 (0.94- 5.48)	0.08
Intercept		0.03 (0.003- 0.22)	0.001		0.02 (0.002- 0.14)	0.0002
Income low		0.53 (0.05- 5.68)	0.60		1.63 (0.16- 16.52)	0.68
GRS _{AD}	1,837	0.97 (0.90- 1.05)	0.48	1,772	1.01 (0.94- 1.09)	0.78
Age		1.04 (1.02- 1.06)	2.8*10 ⁻⁵		1.03 (1.01- 1.05)	0.0003
GRS _{AD} *Income low		1.05 (0.95; 1.16)	0.38		0.99 (0.90; 1.09)	0.82
Intercept		0.01 (0.001; 0.06)	1.74*10 ⁻⁶		0.003 (0.001; 0.07)	0.0003
Education low		3.46 (0.38; 31.80)	0.27		13.25 (0.54; 366.34)	0.12
GRS _{AD}	1,901	1.01 (0.95; 1.09)	0.71	1,928	1.09 (0.96; 1.24)	0.19
Age		1.04 (1.02; 1.06)	2.6*10 ⁻⁶		1.03 (1.01; 1.05)	0.001
GRS _{AD} *Education low		0.96 (0.87; 1.05)	0.38		0.91 (0.80; 1.05)	0.19

Supplementary Table 4. Age-adjusted relative excess risk due to interaction (RERI) and corresponding 95% CI as a measure of interaction between *APOE ε4xSES* (socioeconomic status) and the GRS_{ADXSES} on additive scale, stratified by sex.

	Male		Female	
	n	RERI (95% CI)	n	RERI (95% CI)
<i>APOE*Income</i>	1,837	-0.04 (-0.91; 0.82)	1,772	0.05 (-0.23; 1.24)
<i>APOE*Education</i>	1,901	0.14 (-0.58; 0.86)	1,928	1.00 (0.26; 1.75)*
<i>GRS_{AD}*Income</i>	1,837	0.06 (-0.66; 0.78)	1,772	-0.42 (-1.23; 0.38)
<i>GRS_{AD}*Education</i>	1,901	-0.18 (-0.86; 0.50)	1,928	-1.17 (-3.02; 0.68)

*p < 0.05

Supplementary Table 5. Age- and sex-adjusted effects and corresponding 95% confidence interval (95% CI) for the interaction between each Alzheimer's disease associated single nucleotide polymorphism (SNP) and income (per 1000€/month) in separate linear regression models, sorted by descending interaction effect size estimates.

Chr:Position Build 37	Locus	SNP	Risk Allele	OR _{SNPxinc} (95%CI)	RERI _{SNPxinc} (95%CI)
17: 61538148	<i>ACE</i>	rs28369023	A	1.53 (0.57; 4.56)	0.62 (-0.66; 1.90)
11: 59936926	<i>MS4A2</i>	rs12453	T	1.19 (0.90; 1.57)	0.20 (-0.04; 0.45)
2:127892810	<i>BIN1</i>	rs6733839	T	1.16 (0.88; 1.54)	0.13 (-0.15; 0.42)
20: 54997568	<i>CASS4</i>	rs6064392	G	1.14 (0.67; 1.90)	0.18 (-0.18; 0.54)
15: 59045774	<i>ADAM10</i>	rs474875	A	1.13 (0.85; 1.49)	0.09 (-0.21; 0.39)
2:233981912	<i>INPP5D</i>	rs7570061	G	1.12 (0.78; 1.60)	0.10 (-0.26; 0.47)
14: 53391680	<i>FERMT2</i>	rs17125944	C	1.12 (0.71; 1.76)	0.21 (-0.37; 0.79)
8:27219987	<i>PTK2B</i>	rs28834970	C	1.08 (0.81; 1.45)	0.08 (-0.24; 0.39)
21: 28156856	<i>ADAMTS1</i>	rs7276338	C	1.03 (0.76; 1.40)	0.05 (-0.29; 0.38)
6:32575406	<i>HLA -DRB1</i>	rs9271192	C	1.02 (0.76; 1.37)	0.09 (-0.27; 0.45)
11: 85868640	<i>PICALM</i>	rs3851179	G	1.00 (0.75; 1.34)	0.003 (-0.33; 0.33)
14: 92932828	<i>SLC24A4</i>	rs10498633	G	0.99 (0.72; 1.36)	-0.05 (-0.45; 0.35)
1: 207802552	<i>CRI</i>	rs679515	T	0.99 (0.71; 1.39)	-0.03 (-0.41; 0.36)
8: 27467686	<i>CLU</i>	rs9331896	A	0.97 (0.73; 1.29)	0.0004 (-0.32; 0.33)
11: 47380340	<i>SPI1</i>	rs10769256	C	0.96 (0.73; 1.27)	-0.04 (-0.38; 0.29)
16: 19808163	<i>IQCK</i>	rs1858973	A	0.96 (0.66; 1.40)	-0.02 (-0.26; 0.42)
6: 47431284	<i>CD2AP</i>	rs10948363	G	0.95 (0.69; 1.30)	-0.10 (-0.46; 0.26)
7: 143099133	<i>EPHA1</i>	rs11767557	T	0.89 (0.64; 1.25)	-0.14 (-0.61; 0.33)
10: 11720308	<i>ECHDC3</i>	rs11257240	G	0.86 (0.65; 1.14)	-0.17 (-0.54; 0.21)
7: 100091795	<i>NYAP1</i>	rs12539172	C	0.82 (0.61; 1.10)	-0.30 (-0.79; 0.18)
6:41129252	<i>TREM2</i>	rs75932628	T	NA	NA

Supplementary Table 6. Age- and sex-adjusted effects and corresponding 95% confidence interval (95% CI) for the interaction between each Alzheimer's disease associated single nucleotide polymorphism (SNP) and education (per year) in separate linear regression models, sorted by descending interaction effect size estimates.

Chr:Position Build 37	Locus	SNP	Risk Allele	OR _{SNPxedu} (95%CI)	RERI _{SNPxedu} (95%CI)
11: 47380340	<i>SPII</i>	rs10769256	C	1.31 (0.99; 1.73)	0.25 (0.02; 0.48)*
11: 59936926	<i>MS4A2</i>	rs12453	T	1.22 (0.91; 1.63)	0.22 (-0.03; 0.46)
17: 61538148	<i>ACE</i>	rs28369023	A	1.18 (0.46; 3.31)	0.27 (-0.96; 1.50)
11: 121435587	<i>SORL1</i>	rs11218343	T	1.09 (0.44; 2.48)	0.14 (-0.47; 0.75)
11: 85868640	<i>PICALM</i>	rs3851179	G	1.06 (0.79; 1.42)	0.05 (-0.26; 0.36)
10: 11720308	<i>ECHDC3</i>	rs11257240	G	1.02 (0.76; 1.37)	0.04 (-0.28; 0.37)
14: 92932828	<i>SLC24A4</i>	rs10498633	G	1.02 (0.73; 1.42)	0.009 (-0.37; 0.38)
2: 127892810	<i>BIN1</i>	rs6733839	T	0.98 (0.74; 1.32)	-0.05 (-0.38; 0.28)
6: 47431284	<i>CD2AP</i>	rs10948363	G	0.97 (0.70; 1.35)	-0.07 (-0.43; 0.28)
21: 28156856	<i>ADAMTS1</i>	rs7276338	C	0.97 (0.70; 1.33)	-0.02 (-0.38; 0.34)
15: 59045774	<i>ADAM10</i>	rs474875	A	0.95 (0.71; 1.28)	-0.10 (-0.48; 0.28)
16: 19808163	<i>IQCK</i>	rs1858973	A	0.92 (0.61; 1.36)	-0.07 (-0.54; 0.40)
8: 27219987	<i>PTK2B</i>	rs28834970	C	0.91 (0.68; 1.22)	-0.11 (-0.47; 0.24)
6: 32575406	<i>HLA -DRBI</i>	rs9271192	C	0.91 (0.67; 1.24)	-0.05 (-0.44; 0.33)
2: 233981912	<i>INPP5D</i>	rs7570061	G	0.90 (0.61; 1.31)	-0.16 (-0.7; 0.37)
14: 53391680	<i>FERMT2</i>	rs17125944	C	0.89 (0.57; 1.41)	-0.09 (-0.69; 0.50)
20: 54997568	<i>CASS4</i>	rs6064392	G	0.86 (0.49; 1.47)	-0.09 (-0.73; 0.54)
7: 143099133	<i>EPHA1</i>	rs11767557	T	0.81 (0.56; 1.15)	-0.29 (-0.85; 0.27)
7: 100091795	<i>NYAPI</i>	rs12539172	C	0.79 (0.57; 1.08)	-0.35 (-0.87; 0.16)
8: 27467686	<i>CLU</i>	rs9331896	A	0.78 (0.58; 1.04)	-0.28 (-0.71; 0.15)
1: 207802552	<i>CRI</i>	rs679515	T	0.73 (0.52; 1.03)	-0.40 (-0.85; 0.05)
6: 41129252	<i>TREM2</i>	rs75932628	T	NA	NA

*p < 0.05

Supplementary Table 7. Sex- and age- adjusted odds ratios (OR) and corresponding 95% confidence intervals (95% CI) for the association of income, education, *APOE* ε4 genotype and the AD-associated genetic risk score (GRS_{AD}) with mild cognitive impairment (MCI) using a more rigorously defined non-MCI group by including only those participants who did not report a cognitive complaint and did not show any impairment in the applied cognitive tests calculated using separate logistic regression models.

	n	OR (95% CI)	p
Intercept	1,819	0.02(0.01; 0.05)	<2*10 ⁻¹⁶
Low income		1.65 (1.33; 2.04)	5.1*10 ⁻⁶
Sex		1.07 (0.87; 1.31)	0.53
Age		1.04 (1.03; 1.06)	4.6*10 ⁻¹⁰
Intercept	1,939	0.02 (0.01; 0.05)	<2*10 ⁻¹⁶
Low education		1.51 (1.20; 1.90)	0.0005
Sex		0.91 (0.74; 1.13)	0.40
Age		1.05 (1.03; 1.06)	1*10 ⁻¹¹
Intercept	1,940	0.02 (0.01; 0.04)	<2*10 ⁻¹⁶
<i>APOE</i> ε4		1.31 (1.04; 1.63)	0.02
Sex		1.03 (0.84; 1.25)	0.79
Age		1.05 (1.04; 1.06)	1.6*10 ⁻¹³
Intercept	1,940	0.03 (0.01; 0.10)	2.4*10 ⁻⁹
GRS _{AD}		0.99 (0.95; 1.02)	0.43
Sex		1.03 (0.85; 1.26)	0.75
Age		1.05 (1.04; 1.06)	1.2*10 ⁻¹³

Supplementary Table 8. Sex- and age- adjusted odds ratios (OR) and corresponding 95% confidence intervals (95% CI) in logistic regression models including main effects and interaction terms for the interaction of *APOE* ε4 status and AD-associated genetic risk score (GRS_{AD}) with education and income on mild cognitive impairment (MCI) using a more rigorously defined non-MCI group by including only those participants who did not report a cognitive complaint and did not show any impairment in the applied cognitive tests.

	n	OR (95% CI)	p
Intercept	1,819	0.02 (0.01; 0.05)	<2*10 ⁻¹⁶
Income low		1.64 (1.27; 2.11)	0.0001
<i>APOE</i> ε4		1.30 (0.90; 1.85)	0.15
Sex		1.06 (0.86; 1.31)	0.55
Age		1.04 (1.03; 1.06)	7.3*10 ⁻¹⁰
<i>APOE</i> ε4*Income low		1.04 (0.65; 1.67)	0.87
Intercept	1,939	0.02 (0.01; 0.05)	<2*10 ⁻¹⁶
Education low		1.37 (1.05; 1.80)	0.02
<i>APOE</i> ε4		1.03 (0.68; 1.55)	0.88
Sex		0.91 (0.74; 1.13)	0.39
Age		1.05 (1.03; 1.06)	1.5*10 ⁻¹¹
<i>APOE</i> ε4*Education low		1.41 (0.87; 2.32)	0.17
Intercept	1,819	0.03 (0.01; 0.14)	1.6*10 ⁻⁵
Income low		1.39 (0.23; 8.37)	0.72
GRS _{AD}		0.99 (0.93; 1.05)	0.70
Sex		1.07 (0.87; 1.32)	0.53
Age		1.04 (1.03; 1.06)	4.8*10 ⁻¹⁰
GRS _{AD} *Income low		1.01 (0.93; 1.09)	0.85
Intercept	1,939	0.01 (0.001; 0.05)	1.0*10 ⁻⁷
Education low		8.22 (1.29; 53.35)	0.03
GRS _{AD}		1.04 (0.97; 1.11)	0.26
Sex		0.92 (0.74; 1.14)	0.44
Age		1.05 (1.03; 1.06)	9.4*10 ⁻¹²
GRS _{AD} *Education low		0.93 (0.86; 1.01)	0.07

Supplementary Table 9. Age- and sex-adjusted relative excess risk due to interaction (RERI) and corresponding 95% confidence intervals as a measure of interaction between *APOE ε4xSEP* and the GRS_{ADXSEP} on the additive scale using a more rigorously defined non-MCI group by including only those participants who did not report a cognitive complaint and did not show any impairment in the applied cognitive tests.

	n	RERI (95% CI)
<i>APOE ε4*Income</i>	1819	0.24 (-0.12; 0.61)
<i>APOE ε4*Education</i>	1939	0.31 (0.12; 0.51)*
GRS _{AD} *Income	1819	0.006 (-0.09; 0.11)
GRS _{AD} *Education	1939	-0.32 (-1.14; 0.49)

*p < 0.05

Supplementary Table 10. Sex- and age- adjusted odds ratios (OR) and corresponding 95% confidence intervals (95% CI) for the association of income, education, *APOE* ε4 genotype and the AD-associated genetic risk score (GRS_{AD}) with amnestic mild cognitive impairment calculated using separate logistic regression models.

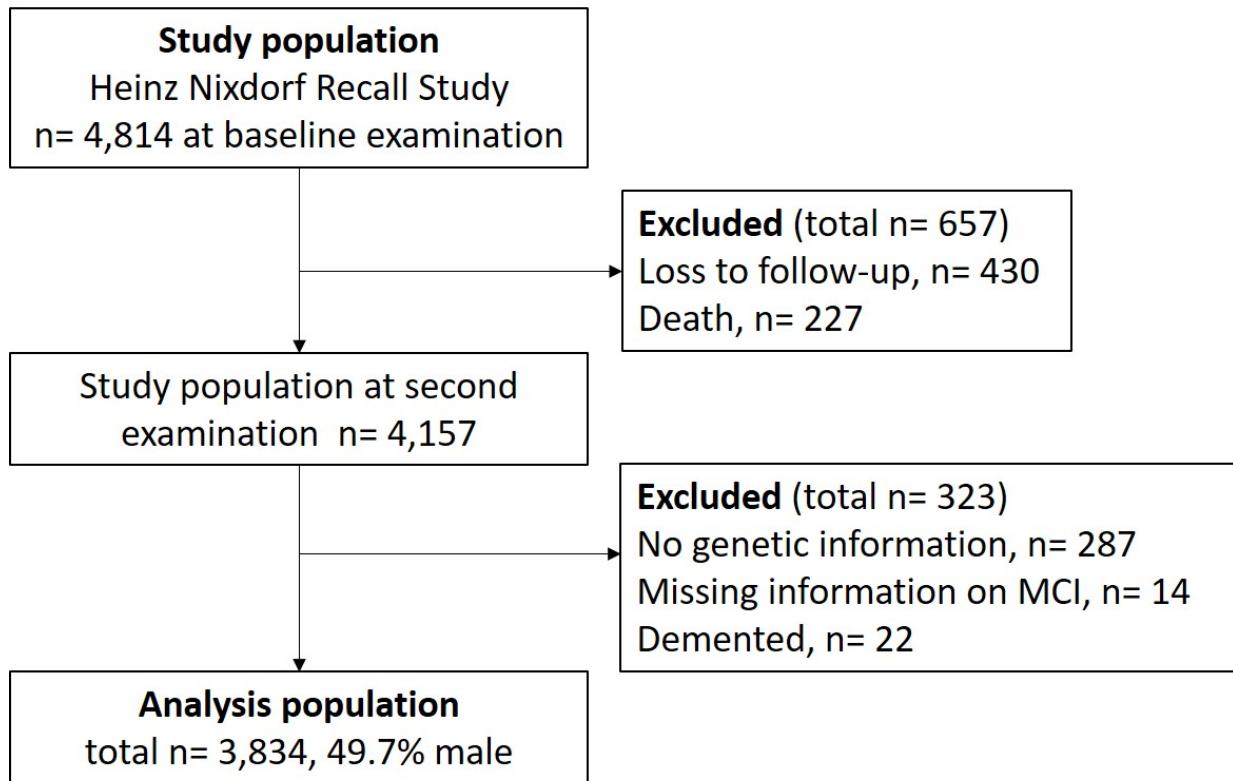
	n	OR (95% CI)	p
Intercept	3,361	0.01 (0.005; 0.04)	<2*10 ⁻¹⁶
Low income		1.65 (1.27; 2.16)	0.0002
Sex		0.85 (0.66; 1.09)	0.20
Age		1.03 (1.01; 1.05)	0.001
Intercept	3,562	0.01 (0.005; 0.03)	<2*10 ⁻¹⁶
Low education		1.53 (1.16; 2.03)	0.003
Sex		0.72 (0.56; 0.93)	0.01
Age		1.03 (1.01; 1.05)	0.0002
Intercept	3,567	0.01 (0.05; 0.03)	<2*10 ⁻¹⁶
<i>APOE</i> ε4		1.36 (1.04; 1.76)	0.02
Sex		0.81 (0.64; 1.03)	0.09
Age		1.03 (1.02; 1.05)	3.0*10 ⁻⁵
Intercept	3,567	0.01 (0.002; 0.04)	3.9*10 ⁻¹¹
GRS _{AD}		1.02 (0.97; 1.06)	0.48
Sex		0.81 (0.64; 1.03)	0.09
Age		1.03 (1.02; 1.05)	2.6*10 ⁻⁵

Supplementary Table 11. Sex- and age- adjusted odds ratios (OR) and corresponding 95% confidence intervals (95% CI) in logistic regression models including main effects and interaction terms for the interaction of *APOE ε4* status and AD-associated genetic risk score (GRS_{AD}) with education and income on amnestic mild cognitive impairment.

	n	OR (95% CI)	p
Intercept	3,361	0.01 (0.004; 0.03)	<2*10 ⁻¹⁶
Income low		1.86 (1.35; 2.59)	0.0002
<i>APOE ε4</i>		1.69 (1.08; 2.62)	0.02
Sex		0.85 (0.66; 1.08)	0.19
Age		1.03 (1.01; 1.04)	0.001
<i>APOE ε4</i> *Income low		0.71 (0.41; 1.25)	0.23
Intercept	3,562	0.01 (0.004; 0.03)	<2*10 ⁻¹⁶
Education low		1.52 (1.10; 2.14)	0.01
<i>APOE ε4</i>		1.34 (0.81; 2.16)	0.24
Sex		0.72 (0.56; 0.92)	0.01
Age		1.03 (1.01; 1.05)	0.0002
<i>APOE ε4</i> *Education low		1.04 (0.59; 1.87)	0.89
Intercept	3,361	0.01 (0.001; 0.05)	1.6*10-6
Income low		1.91 (0.21; 17.49)	0.56
GRS _{AD}		1.03 (0.96; 1.11)	0.41
Sex		0.85 (0.66; 1.09)	0.19
Age		1.03 (1.01; 1.05)	0.001
GRS _{AD} *Income low		0.99 (0.90; 1.09)	0.90
Intercept	3,562	0.002 (0.0002; 0.02)	2.7*10-8
Education low		10.15 (1.07; 100.57)	0.05
GRS _{AD}		1.08 (0.99; 1.17)	0.07
Sex		0.72 (0.56; 0.93)	0.01
Age		1.03 (1.01; 1.05)	0.0002
GRS _{AD} *Education low		0.92 (0.84; 1.02)	0.10

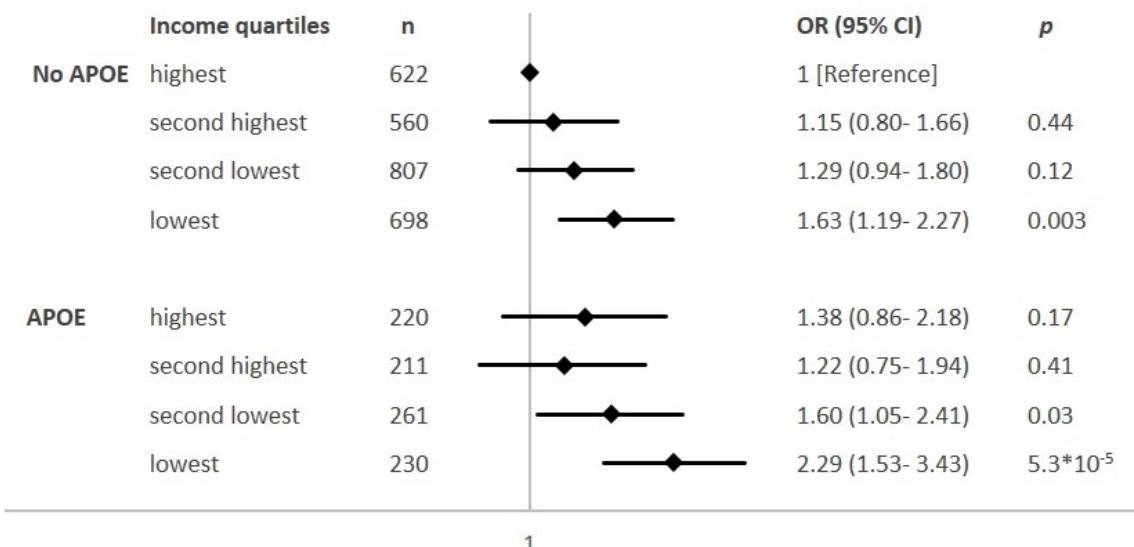
Supplementary Table 12. Age- and sex-adjusted relative excess risk due to interaction (RERI) and corresponding 95% confidence intervals as a measure of interaction between *APOE ε4*xSEP and the GRS_{AD}xSEP on the additive scale (amnestic mild cognitive impairment).

	n	RERI (95% CI)
<i>APOE ε4</i> *Income	3,361	-0.09 (-1.01; 0.84)
<i>APOE ε4</i> *Education	3,562	0.23 (-0.19; 0.65)
GRS _{AD} *Income	3,361	0.02 (-0.03; 0.06)
GRS _{AD} *Education	3,562	-0.14 (-0.75; 0.47)



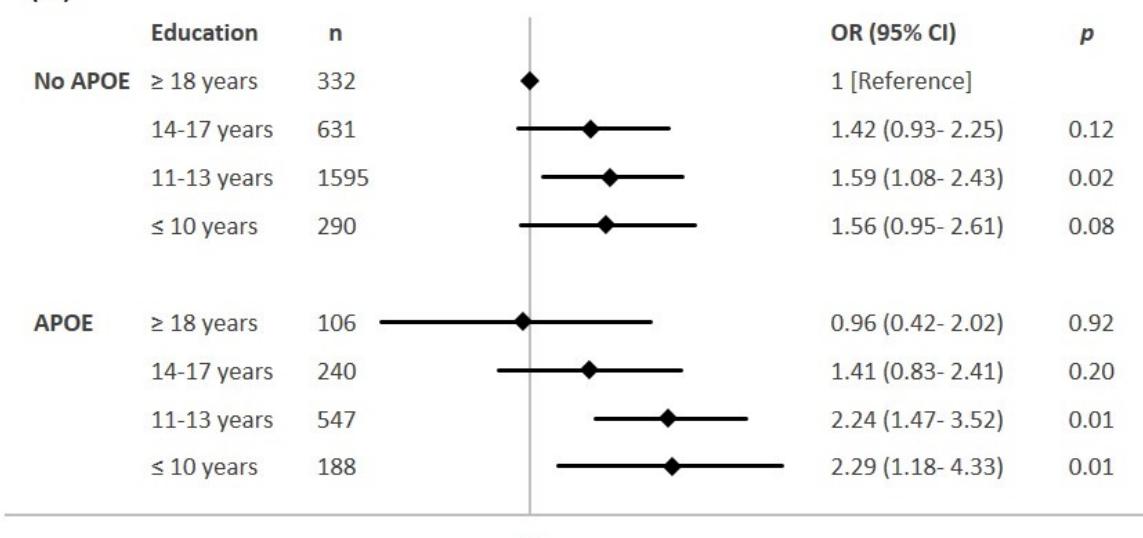
Supplementary Figure 1. Flow-chart of participants of the Heinz Nixdorf Recall Study cohort included in the analysis.

(A)



1

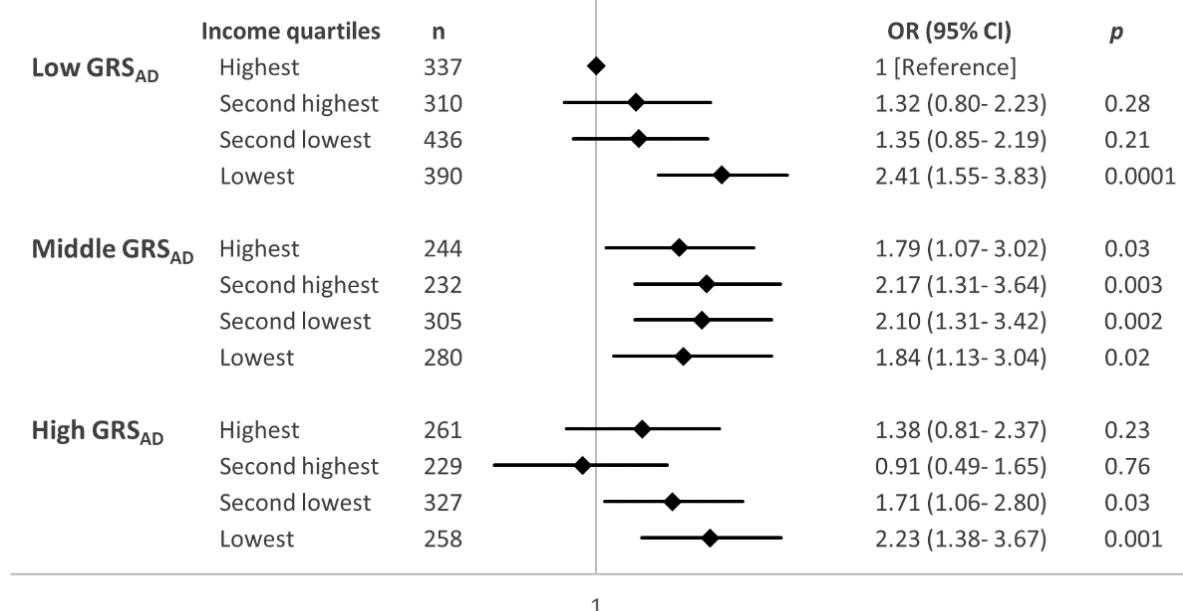
(B)



1

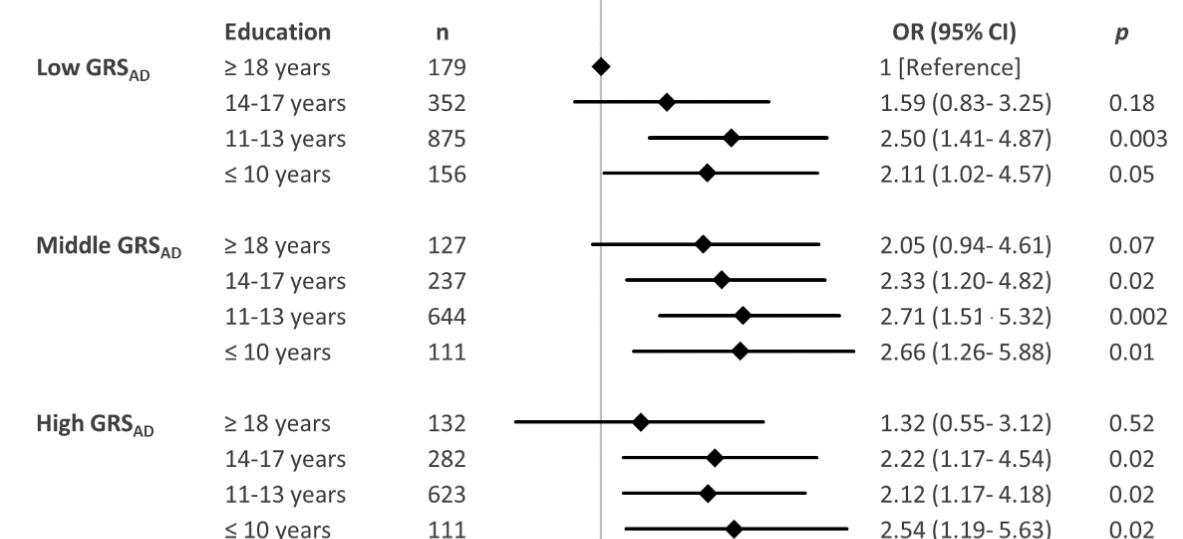
Supplementary Figure 2. Sex- and age-adjusted odds ratios (OR) and corresponding 95% confidence intervals (95% CI) for single reference joint effects of *APOE* ε4 genotype and socioeconomic position indicators on mild cognitive impairment calculated in separate logistic regression models for (A) income quartiles and (B) education categories, with the group of a negative *APOE* ε4 status and a high socioeconomic position as reference.

(A)



1

(B)



1

Supplementary Figure 3. Sex- and age-adjusted odds ratios (OR) and corresponding 95% confidence intervals (95% CI) for single reference joint effects of tertiles of an AD-associated genetic risk score (GRS_{AD}) and socioeconomic position indicators on mild cognitive impairment calculated in separate logistic regression models for (A) income quartiles and (B) education categories, with the group of a low GRS and a high socioeconomic position as reference.