## SUPPLEMENTARY TABLES

Supplementary Table 1. Effects of genetic markers on the age at the midpoint of cognitive function decline (M) and the rate of cognitive function decline (R) parameters based on the MMSE after adjusting for sex and education years (n = 283).

Covariates	Minor allele	Regression coefficients on M		<b>Regression coefficients on</b> <i>R</i>	
Covariates	Minor allele	Posterior mean	95% CI	Posterior mean	95% CI
Sex (female vs. male)	_	-1.02	(-2.93, 0.92)	0.64	(0.23, 1.11)*
Education years	—	0.39	(0.18, 0.61)*	-0.04	(-0.09, -0.003)*
APOE	ε4	-2.10	(-3.79, -0.34)*	0.90	(0.37, 1.42)*
ABCA7 rs3764650	G	-1.74	(-3.16, -0.29)*	-0.20	(-0.51, 0.13)
SORL1 rs3737529	Т	2.65	(0.41, 4.90)*	0.56	(0.01, 1.07)*
SORL1 rs1784933	G	1.16	(-0.91, 3.22)	0.12	(-0.33, 0.52)
SORL1 rs2298813	А	0.88	(-1.78, 3.61)	-0.005	(-0.60, 0.71)
BIN1 rs744373	G	-0.47	(-1.79, 0.86)	-0.16	(-0.43, 0.11)
GAB2 rs2373115	А	-0.35	(-1.69, 0.99)	0.09	(-0.23, 0.40)
CD33 rs3865444	А	1.32	(-0.34, 2.95)	-0.05	(-0.41, 0.36)

Note: The subset of 255 AD and 28 MCI progression to AD patients were analyzed. Abbreviations: MMSE, Mini-Mental State Examination; CI, credible interval.

\*Statistically significant with 95% credible interval.

Supplementary Table 2. Model selection among the eight genetic markers after adjusting for sex and education
years ( <i>n</i> = 283).

Models	Variables in the model	Posterior model probabilities
Model 1	APOE, SORL1 rs3737529	0.8988
Model 2	APOE, ABCA7 rs3764650, SORL1 rs3737529, SORL1 rs1784933	0.0565
Model 3	APOE, SORL1 rs3737529, CD33 rs3865444	0.0216
Model 4	APOE, SORL1 rs3737529, SORL1 rs1784933	0.0131
Model 5	APOE, SORL1 rs3737529, SORL1 rs2298813	0.0066
Model 6	APOE, SORL1 rs3737529, GAB2 rs2373115	0.0024
Model 7	APOE, SORL1 rs3737529, BIN1 rs744373, GAB2 rs2373115	0.0009
Model 8	APOE, SORL1 rs3737529, BIN1 rs744373	0.0002

Note: The subset of 255 AD and 28 MCI progression to AD patients were analyzed.

Supplementary Table 3. Effects of *APOE* and *SORL1* rs3737529 on the age at the midpoint of cognitive function decline (M) and the rate of cognitive function decline (R) parameters based on the MMSE after Bayesian variable selection (n = 283).

Compristor	Minor allele	Regression coefficients on M		Regression coefficients on R	
Covariates		Posterior mean	95% CI	Posterior mean	95% CI
Sex (female vs. male)		-0.89	(-2.88, 1.12)	0.59	(0.12, 1.12)*
Education years		0.40	(0.19, 0.62)*	-0.04	(-0.09, 0.003)
APOE	ε4	-2.50	(-4.25, -0.73)*	0.82	(0.37, 1.30)*
SORL1 rs3737529	Т	1.58	(0.01, 3.23)*	0.43	(0.06, 0.80)*

Note: The subset of 255 AD and 28 MCI progression to AD patients were analyzed.

Abbreviations: MMSE, Mini-Mental State Examination; CI, credible interval.

\*Statistically significant with 95% credible interval.

Supplementary Table 4. Effects of genetic markers on the age at the midpoint of cognitive function decline (M) and the rate of cognitive function decline (R) parameters based on the MMSE after adjusting for sex and education years using the ADNI data (n = 385).

Conomiator	Minon allala	Regression coefficients on M		Regression coefficients on R	
Covariates Minor allele		Posterior mean	95% CI	Posterior mean	95% CI
Sex (female vs. male)	_	-1.47	(-3.34, 0.39)	-0.05	(-0.29, 0.18)
Education years		-0.04	(-0.36, 0.28)	-0.07	(-0.10, -0.03)*
APOE	ε4	-3.82	(-5.14, -2.44)*	-0.05	(-0.23, 0.13)
ABCA7 rs3764650	G	1.35	(-0.55, 3.22)	0.09	(-0.18, 0.37)
SORL1 rs3737529	Т	-1.67	(-5.06, 1.92)	0.16	(-0.44, 1.02)
SORL1 rs1784933	G	1.68	(-1.24, 4.57)	0.11	(-0.30, 0.56)
SORL1 rs2298813	А	2.19	(-0.62, 4.97)	-0.14	(-0.44, 0.18)
BIN1 rs744373	G	-0.13	(-1.50, 1.26)	-0.14	(-0.32, 0.04)
GAB2 rs2373115	А	-0.39	(-1.94, 1.16)	-0.07	(-0.26, 0.14)
CD33 rs3865444	А	0.57	(-0.79, 1.95)	0.12	(-0.06, 0.31)

Note: The sample of 150 AD and 235 MCI progression to AD patients with at least four MMSE measurements were analyzed. Abbreviations: MMSE, Mini-Mental State Examination; CI, credible interval.

\*Statistically significant with 95% credible interval.

Gene	SNP	TaqMan genotyping assay ID		
APOE_e2/3/4	rs429358	C_3084793_20		
APOE_e2/3/4	rs7412	C_904973_10		
ABCA7	rs3764650	C_27478162_20		
SORL1	rs3737529	C_25803133_10		
SORL1	rs1784933	C_8799397_10		
SORL1	rs2298813	C_16190780_10		
BIN1	rs744373	C_1042213_10		
GAB2	rs2373115	C_12033202_20		
CD33	rs3865444	C_1487395_40		

Supplementary Table 5. The TaqMan genotyping assay ID for the selected single nucleotide polymorphism (SNP).