

Table S1. APOE allele and genotype frequencies in AD, MCI and controls.

	AD		MCI		Controls	
genotypes	n	%	n	%	n	%
$\epsilon 3/\epsilon 3$	39	49.4	17	54.8	44	75.9
$\epsilon 3/\epsilon 4$	32	40.5	7	22.6	5	8.6
$\epsilon 2/\epsilon 3$	2	2.5	5	16.1	7	12.1
$\epsilon 2/\epsilon 4$	2	2.5	2	6.5	1	1.7
$\epsilon 2/\epsilon 2$	1	1.3	0	0	1	1.7
$\epsilon 4/\epsilon 4$	3	3.8	0	0	0	0
alleles						
$\epsilon 2$	6	3.8	7	11.3	10	8.6
$\epsilon 3$	112	70.9	46	74.2	100	86.2
$\epsilon 4$	40	25.3	9	14.5	6	5.2

APOE genotype distribution between groups: p=0.001; $\epsilon 4+$ between groups: p<0.001.**Table S2.** IL-1 α -IL-1 β -IL-38 estimated haplotypes (f>0.05) in AD, MCI and controls.

IL-1A-IL-1B-IL-38 haplotypes	AD		MCI		Controls	
	n	f	n	f	n	f
CCA	21	0.137	12	0.190	19	0.167
CCC	15	0.096	0	0.000	5	0.040
CTA	42	0.271	20	0.324	41	0.354
CTC	22	0.145	8	0.130	18	0.155
TCA	12	0.078	2	0.035	5	0.043
TTA	30	0.196	11	0.175	14	0.117
TTC	12	0.076	9	0.143	14	0.124

f=haplotype frequency; p=ns.

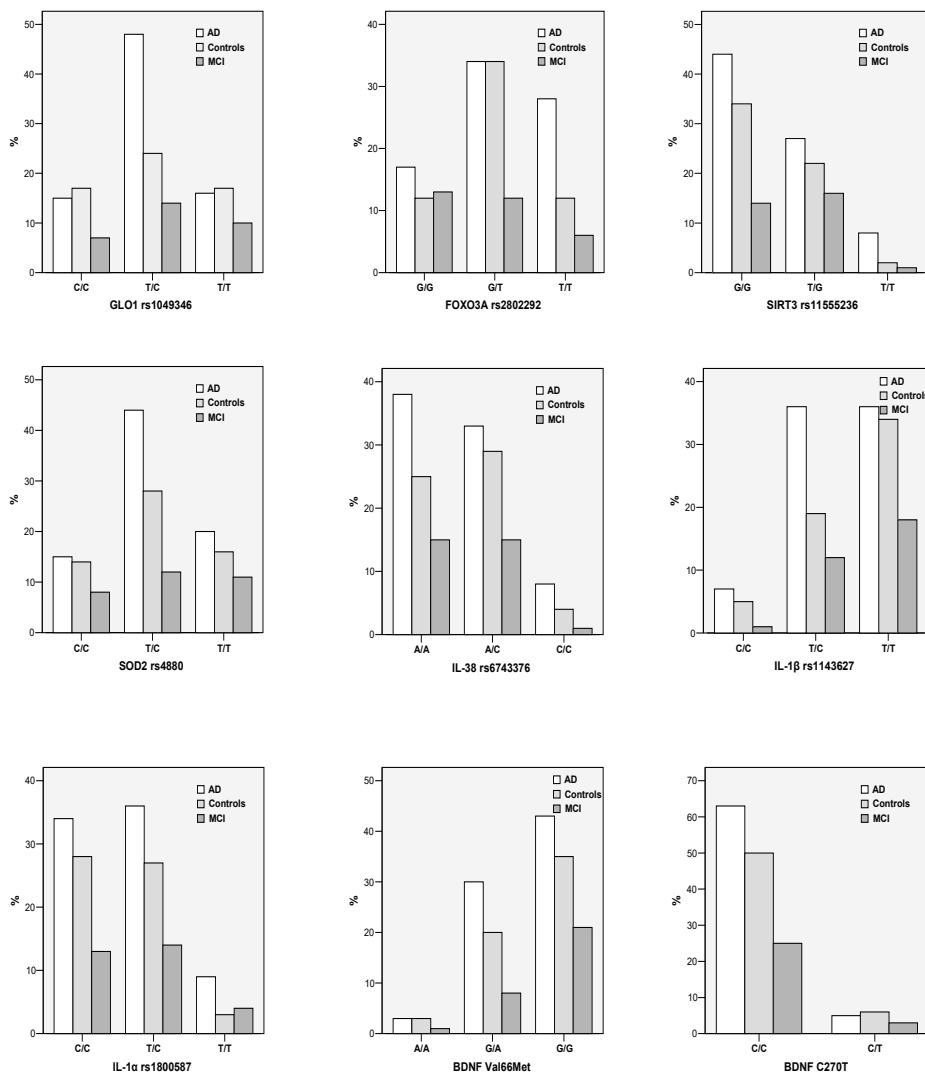


Figure S1. Genotype frequencies of the assayed polymorphisms in AD, MCI and controls ($p=ns$, Kruskal-Wallis test).