

rsID	Gene	"-" variant	"+" variant
rs819147	AHCY	T	C
rs429358	APOE	T	C
rs7412	APOE	C	T
rs26538	ATG12	T	C
rs10210302	ATG16L1	C	T
rs510432	ATG5	C	T
rs6265	BDNF	C	T
rs10402876	C3	G	C
rs2569191	CD14	T	C
rs4680	COMT V158M	G	A
rs1021737	CTH	G	T
rs231775	CTLA4	A	G
rs1108580	DBH	A	G
rs1076560	DRD2	C	A
rs769407	GAD1	G	C
rs3828275	GAD1	C	T
rs7483	GSTM3	C	T

rsID	Gene	"-" variant	"+" variant
rs6313	HTR2	G	A
rs1800925	IL13	C	T
rs1143634	IL1B	G	A
rs2069812	IL5	A	G
rs1800795	IL6	G	C
rs6323	MAO-A	T	G
rs1799836	MAO-B	T	C
rs3798963	PARK2 (Parkin)	A	T
rs1042173	SLC6A4	A	C
rs10181656	STAT4	C	G
rs1142636	SYN1	A	G
rs1800629	TNF	G	A
rs361525	TNF	G	A
rs4570625	TPH2	G	T