

rsID	Gene	"-" variant	"+" variant
rs4343	ACE	A	G
rs699	AGT	A	G
rs26538	ATG12	T	C
rs10210302	ATG16L1	C	T
rs510432	ATG5	C	T
rs4680	COMT V158M	G	A
rs4646	CYP19A1	C	A
rs1048943	CYP1A1 L4889G	T	C
rs1800440	CYP1B1	T	C
rs225014	DIO2	T	C
rs6025	FACTOR V	C	T
rs3211719	FACTOR X	A	G
rs2071010	FOLR1	G	A
rs651933	FOLR2	A	G
rs1867277	FOXE1	G	A

rsID	Gene	"-" variant	"+" variant
rs6165	FSHR	C	T
rs2282679	GC or DBP	T	G
rs1695	GSTP1:I105V	A	G
rs268	LPL	A	G
rs1076991	MTHFD1	C	T
rs1801131	MTHFR:A1298C	T	G
rs1801133	MTHFR:C677T	G	A
rs1051266	SLC19A1	T	C
rs166050	SRD5A1	A	G
rs526934	TCN1	A	G
rs1801198	TCN2	C	G
rs731236	VDR Taq	A	G