

Supplementary table 1. Sequence variants identified by DNA sequencing in ADIPOQ coding sequence

SNP	amino acid coding change	Position	MAF	location	p-value for association
C-186T	-	188043005	0.013	promoter	0.20
rs2241766	G15G	188053586	0.14	exon 2	0.32
G133C	G45R	188053674	0.039	exon 2	<0.0001
C75T	P25P	188053616	0.0087	exon 2	0.79
rs17366743	Y111H	188054783	0.022	exon 3	0.84

*position in build NCBI36/hb18