

Table S1. Genotyped miRNA target site SNPs and the *P* values of their association with overall survival

SNP ID	Gene	Chromosome	Position	Allele	MAF		<i>P</i> *
					Hapmap	CHB This study	
rs5876	<i>HIAT1</i>	chr1	100320581	C>T	0.34	0.45	0.199
rs10510454	<i>DAZL</i>	chr3	16604608	C>T	0.47	0.54	0.109
rs1128154	<i>HCLSI</i>	chr3	122833305	G>T	0.32	0.28	0.654
rs1058267	<i>PCDH10</i>	chr4	134332051	A>G	0.48	0.52	0.174
rs2292899	<i>ACSL1</i>	chr4	185915103	G>A	0.40	0.32	0.580
rs8488	<i>HISPPD1</i>	chr5	102566650	T>C	0.41	0.49	1.000
rs998754	<i>GABRA1</i>	chr5	161258511	A>C	0.44	0.46	0.134
rs4351800	<i>SYT9</i>	chr11	7446394	A>C	0.37	0.38	0.109
rs1062836	<i>NECAP1</i>	chr12	8140316	C>G	0.41	0.44	0.570
rs4298982	<i>LARP4</i>	chr12	49156695	A>C	0.48	0.52	0.713
rs2466551	<i>NFYB</i>	chr12	103037047	T>G	0.38	0.42	0.549
rs7158951	<i>POMT2</i>	chr14	76812334	C>T	0.50	0.44	0.458
rs11704	<i>ZNF839</i>	chr14	101878407	G>C	0.47	0.50	0.218
rs3210967	<i>ZDHHC7</i>	chr16	83566963	C>T	0.46	0.57	0.264
rs1051424	<i>RPS6KB1</i>	chr17	55379105	A>G	0.36	0.40	0.075
rs2073859	<i>LIMK2</i>	chr22	30004552	G>A	0.40	0.31	0.576

Abbreviations: SNP, single nucleotide polymorphism; MAF, minor allele frequency.

*Log-rank test for trend.