

Table S8. Functional information for SNPs in LD with rs11191659

No. rs	Chromosome	Position	Allele	LDsnp	Pop/LD	TFBS	Splicing(site)	Splicing(ESE or ESS)	Splicing(abolish domain)	miRNA(miRanda)	miRNA(Sanger)	nsSNP	Stop Codon	Polyphen	SNPs3D(svm profile)	SNPs3D(svm structure)	RegPotential	Conservation	Nearby Gene	Distance (bp)	Allele	GIH
1 rs10883841	10	104924699	C/T	rs11191659	GIH/0.828	--	--	--	--	--	--	Y	--	benign	--	--	0.17821	1	NT5C2	86797 18307	T	0.972
2 rs11191606	10	104946025	C/T	rs11191659	GIH/1.000	Y	--	--	--	--	--	--	--	--	--	--	0	0.02	LOC729081 LOC401648	-19983 -19431	T	0.966
3 rs11191655	10	105068783	C/T	rs11191659	GIH/1.000	--	--	--	--	--	--	--	--	--	--	--	0	0.014	PCGF6	16240 32098	T	0.966
4 rs11191659	10	105091691	C/T	rs11191659	1	--	--	--	--	--	--	--	--	--	--	--	NA	0	PCGF6	39148 9190	C	0.966