

Table e-1. SNPs with uncorrected  $p$  values between  $10^{-5}$  and  $10^{-6}$ . Position is based on Genome Build 36.3.

SNP	Chr	Position	Closest RefSeq Gene	Gene Name	Location relative to gene
rs10191411	2	149104438	<i>EPC2</i>	enhancer of polycomb homolog 2	14.6kb upstream
rs6759083	2	149154681	<i>EPC2</i>		Intron
rs7564433	2	149163108	<i>EPC2</i>		Intron
rs10197267	2	149203379	<i>EPC2</i>		Intron
rs11724427	4	89322196	<i>ABCG2</i>	ATP-binding cassette, sub-family G (WHITE), member 2	23.2kb upstream
rs2583693	8	20710259	<i>LOC724059</i>		124.1kb downstream
rs10974334	9	4092339	<i>GLIS3</i>	GLIS family	Intron
rs2208558	9	4093345	<i>GLIS3</i>	zinc finger 3	Intron
rs17157052	10	1853551	<i>LOC100129465</i>		48.0kb upstream
rs7123160	11	79796236	<i>LOC729790</i>		245.9kb downstream
rs2448281	11	79803240	<i>LOC729790</i>		238.9kb downstream
rs1408888	13	71326648	<i>DACHI</i>	dachshund	Intron

				homolog 1	
rs11628929	14	23913894	<i>NFATC4</i>	nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 4	Intron
rs6573766	14	23921886	<i>NFATC4</i>		3.2kb downstream
rs3936367	16	81106786	<i>CDH13</i>	cadherin 13, H-cadherin (heart)	111.3kb upstream
rs3813089	18	46600022	<i>MRO</i>	maestro	Exon
rs1885082	20	17541984	<i>RRBP1</i>	ribosome binding protein 1 homolog 180kDa	0.3kb downstream
rs6089510	20	59810819	<i>CDH4</i>	cadherin 4, type 1, R-cadherin (retinal)	Intron
rs4444	22	29535334	<i>OSBP2</i>	oxysterol binding protein 2	Intron
rs136245	22	29538544	<i>OSBP2</i>		Intron
rs136243	22	29539135	<i>OSBP2</i>		Intron
rs2269657	22	40594215	<i>SREBF2</i>	sterol regulatory element binding transcription factor 2	Intron