S2 Table. Previously identified colorectal cancer susceptibility single-nucleotide polymorphisms by genome-wide association studies[a)]

SNP	Reported gene	Chr.	Functional Consequence	Allele[a)]		Reference
				M1	M2	
rs6687758	*Intergenic*	1	-	A	G	[1]
rs10936599	*MYNN*	3	-	C	T	[1]
rs647161	*C5orf66*	5	Intron variant	A	C	[2]
rs7758229	*SLC22A3*	6	-	G	T	[3]
rs6983267	*CASC8, CCAT2*	8	Intron variant, nc transcript variant	G	T	[4]
rs7014346	*CASC8*	8	Intron variant	G	A	[5]
rs10505477	*CASC8*	8	Intron variant	A	G	[6]
rs10795668	*LOC105376400*	10	Intron variant	G	A	[7]
rs704017	*ZMIZ1-AS1*	10	Intron variant	A	G	[8]
rs11196172	*TCF7L2*	10	Intron variant	G	A	[8]
rs1665650	*HSPA12A*	10	Intron variant	C	T	[2]
rs174537	*MYRF*	11	Intron variant	G	T	[8]
rs174550	*FADS1*	11	Intron variant	T	C	[8]
rs1535	*FADS2*	11	Intron variant	A	G	[8]
rs3802842	*COLCA1, COLCA2*	11	Intron variant, upstream variant 2KB	A	C	[5]
rs10849432	*LOC105369625*	12	Intron variant	C	T	[8]
rs10774214	*CCND2-AS1*	12	Intron variant	C	T	[2]
rs11169552	*ATF1*	12	Upstream variant 2KB	C	T	[1]
rs7136702	*Intergenic*	12	-	C	T	[9]
rs4444235	*Intergenic*	14	-	T	C	[10]
rs1957636	*LOC105370507*	14	Intron variant	T	C	[8]
rs4779584	*Intergenic*	15	-	C	T	[7]
rs9929218	*CDH1*	16	Intron variant	G	A	[10]
rs12603526	*NXN*	17	Intron variant, utr variant 5 prime	T	C	[8]
rs10411210	*RHPN2*	19	Intron variant	C	T	[10]
rs1800469	*TGFB1, B9D2*	19	Downstream variant 500B, upstream variant 2KB	G	A	[8]
rs2241714	*B9D2, TMEM91*	19	Intron variant, missense, upstream variant 2KB	C	T	[8]
rs961253	*Intergenic*	20	-	C	A	[10]
rs2423279	*Intergenic*	20	-	T	C	[2]
rs4813802	*Intergenic*	20	-	T	G	[11]

SNP, single-nucleotide polymorphism; Chr, chromosome; M1, major allele; M2, minor allele. [a)]Based on NCBI dbSNP (National Center for Biotechnology Information Database of Single Nucleotide Polymorphisms).