

ADAM33	SNPs	minor allele	OR genotyping	OR imputation HM2 (HM3)	CI_L 95% genotyping	CI_L 95% imputation HM2 (HM3)	CI_U 95% genotyping	CI_U 95% imputation HM2 (HM3)	p-value genotyping	p-value imputation HM2 (HM3)	R2
	rs6084432	A	1.177	1.177 (1.177)	0.965	0.965 (0.9648)	1.436	1.436 (1.436)	0.108	0.108 (0.1082)	1
	rs12624874	T		1.093 (1.026)		0.871 (0.8133)	1.372 (1.295)		0.441 (0.827)		
	rs633330	T		2.12		0.65	6.92		0.21		
	rs630909	A		1.11		0.91	1.35		0.33		
	rs512625	A	1.037	1.046 (1.037)	0.886	0.891 (0.8856)	1.215	1.229 (1.215)	0.649	0.584 (0.6492)	0.95
	rs2787093	C		0.98		0.78	1.23		0.86		
	rs3746631	C	1.040	1.034 (1.011)	0.669	0.619 (0.5807)	1.616	1.730 (1.761)	0.862	0.898 (0.9683)	0.31
	rs2787094	C	1.122		0.937		1.342		0.211		
	rs677044	G		0.94		0.74	1.1		0.31		
	rs3918400	A	0.975	1.03	0.645	0.62	1.473	1.73	0.904	0.9	0.30
	rs43749	T	1.200		0.964		1.493		0.103		
	rs628977	-	-	-	-	-	-	-	-	-	-
	rs678881	C		0.91		0.76	1.09		0.29		
	rs2280089	A	1.161	1.160 (1.18)	0.928	0.934 (0.9475)	1.452	1.441 (1.469)	0.191	0.180 (0.1395)	0.86
	rs2280090	A	1.105		0.894		1.366		0.357		
	rs2280091	A		1.093 (1.126)		0.886 (0.9067)		1.350 (1.399)		0.407 (0.2823)	
	rs574174	T	1.149		0.957		1.379		0.138		
	rs44707	G	0.895		0.763		1.050		0.173		
	rs598418	G		0.924 (0.99)		0.787 (0.8469)		1.084 (1.158)		0.330 (0.9007)	
	rs2853209	A	1.042		0.887		1.224		0.619		
	rs528557 **	G	1.504		1.247		1.813		0.000019		
	rs3918396	T	1.291		0.996		1.672		0.054		
	rs612709	A	1.177		0.948		1.461		0.139		
	rs3918395	A	1.123	1.059 (1.118)	0.894	0.856 (0.8969)	1.411	1.310 (1.394)	0.320	0.600 (0.3211)	0.67
	rs2280092	-	-	-	-	-	-	-	-	-	-
	rs511898	T	1.127	1.103	0.961	0.9399	1.323	1.294	0.142	0.23	
	rs2787095	C		1.07		0.91	1.26		0.4		
	rs621394	A		2.12		0.65	6.92		0.21		
	rs2853210	T		0.939 (0.9078)		0.773 (0.7463)	1.140 (1.104)		0.523 (0.3336)		
	rs4987246	G		1.68		0.74	1.84		0.22		
	rs487377	A	1.054	1.071 (1.054)	0.879	0.894 (0.8794)	1.264	1.284 (1.264)	0.569	0.455 (0.5679)	0.98
	rs570269	C		0.99		0.81	1.22		0.94		
	rs7354032	T		0.774 (1.029)		0.507 (0.8771)		1.184 (1.208)		0.238 (0.7237)	
	rs554743	G	1.038	1.03	0.881	0.88	1.222	1.21	0.658	0.71	0.98
	rs2853215	A	1.085	1.085 (1.082)	0.917	0.917 (0.9141)	1.284	1.284 (1.28)	0.340	0.339 (0.3607)	1
	rs8124504	T		1.01		0.64	1.6		0.96		
	rs532448	T		1		0.86	1.17		0.97		
	rs8124665	G		1.3		0.68	2.5		0.43		
CCL5	SNPs	minor allele	OR genotyping	OR imputation HM2 (HM3)	CI_L 95% genotyping	CI_L 95% imputation HM2 (HM3)	CI_U 95% genotyping	CI_U 95% imputation HM2 (HM3)	p-value genotyping	p-value imputation HM2 (HM3)	R2
	rs16963927	G		0.649 (0.8788)		0.380 (0.6258)		1.106 (1.234)		0.112 (0.4554)	
	rs2280788	C	1.217	0.75	0.775	0.2	1.912	2.8	0.393	0.67	0.12
	rs1800825	G		1.12		0.34	3.7		0.85		
	rs2107538	T	1.190	1.088 (1.095)	0.978	0.889 (0.9086)	1.448	1.331 (1.32)	0.082	0.412 (0.3395)	0.78
	rs4239253	C		1.22		0.53	2.8		0.64		
	rs4796123	C		1.204 (1.175)		0.970 (0.954)		1.495 (1.448)		0.093 (0.1291)	
CD14	SNPs	minor allele	OR genotyping	OR imputation HM2 (HM3)	CI_L 95% genotyping	CI_L 95% imputation HM2 (HM3)	CI_U 95% genotyping	CI_U 95% imputation HM2 (HM3)	p-value genotyping	p-value imputation HM2 (HM3)	R2
	rs778588	C		0.970 (0.9388)		0.826 (0.7982)		1.139 (1.104)		0.711 (0.446)	
	rs11574651	A		0.47		0.12	1.87		0.28		
	rs4914	G		0.92		0.75	1.13		0.43		
	rs2569190	A	1.011	0.99	0.865	0.85	1.182	1.16	0.890	0.92	0.83
	rs5744448	T		2.196 (0.6662)		0.566 (0.2104)		8.527 (2.109)		0.256 (0.4898)	
	rs5744442	C		0.86		0.6	1.23		0.4		
	rs5744441	A		1.031 (1.06)		0.864 (0.8885)		1.231 (1.264)		0.734 (0.5183)	
DPP10	SNPs	minor allele	OR genotyping	OR imputation HM2 (HM3)	CI_L 95% genotyping	CI_L 95% imputation HM2 (HM3)	CI_U 95% genotyping	CI_U 95% imputation HM2 (HM3)	p-value genotyping	p-value imputation HM2 (HM3)	R2
	rs1036022 (rs13392783 / r ² =0.88)	C	0.939	0.94	0.808	0.81	1.092	1.1	0.416	0.45	1
	rs10208402	C		0.944 (0.9554)		0.814 (0.8242)		1.095 (1.107)		0.445 (0.5443)	
	rs1367178	G		1.005 (0.9918)		0.654 (0.669)		1.542 (1.47)		0.984 (0.9675)	
	rs17048357	T		1.1		0.57	2.13		0.77		
	rs17048359	C		0.976 (1.008)		0.823 (0.8546)		1.157 (1.189)		0.778 (0.9244)	
	rs10192393	C	1.053	1.057 (1.053)	0.906	0.909 (0.9059)	1.224	1.229 (1.224)	0.500	0.475 (0.5003)	1
	rs1430092	A		0.85		0.65	1.2		0.36		
	rs12473953	C		1.005 (0.9894)		0.857 (0.8428)		1.180 (1.161)		0.948 (0.8963)	
	rs1430091	A		1.04		0.86	1.28		0.67		
	rs4849333	C		0.937 (0.8835)		0.776 (0.7329)		1.131 (1.065)		0.499 (0.1937)	
	rs1430090	G	1.049	1.040 (1.032)	0.891	0.884 (0.8776)	1.236	1.224 (1.214)	0.565	0.638 (0.7014)	1
	rs6737251	T	0.975	0.975 (0.9753)	0.833	0.833 (0.833)	1.142	1.142 (1.142)	0.756	0.756 (0.7557)	1
	rs17048372	T		1.02		0.84	1.23		0.88		
	rs6709636	A		1.41		0.4	5.01		0.6		
	rs11685217	T	1.006	1.006 (1.006)	0.833	0.833 (0.8327)	1.216	1.216 (1.216)	0.947	0.947 (0.9469)	1
	rs1430089	G		0.98		0.84	1.15		0.78		
	rs17764091	G		1.006 (0.9758)		0.802 (0.7821)		1.261 (1.218)		0.962 (0.8284)	
	rs17764121 *	C		5.21		1.15	23.61		0.03		
	rs17048408	C		0.73		0.37	1.46		0.38		
	rs7559909 (rs13011555 / r ² =0.9)	A	0.946	0.944 (0.9443)	0.811	0.809 (0.8092)	1.104	1.102 (1.102)	0.483	0.467 (0.4667)	1
	rs17048446	A		0.65		0.4	1.06		0.08		
	rs17048451	C		0.56		0.2	1.54		0.26		
	rs10496465	G	0.929	0.931 (0.9307)	0.755	0.757 (0.7565)	1.143	1.145 (1.145)	0.486	0.497 (0.4968)	1
	rs4849338	C		1		0.86	1.18		0.96		
	rs13397905	C		0.56		0.13	2.35		0.43		
	rs1820924	T		1.032 (0.9804)		0.849 (0.8099)		1.255 (1.187)		0.749 (0.8393)	
	rs6750402	C	1.006	1.012 (1.008)	0.847	0.852 (0.8486)	1.195	1.202 (1.197)	0.946	0.892 (0.9285)	1
	rs10187644	T		0.920 (0.9084)		0.792 (0.7813)		1.070 (1.056)		0.278 (0.2117)	
	rs1346762	G	1.110	1.112 (1.112)	0.947	0.950 (0.9495)	1.300	1.303 (1.303)	0.197	0.188 (0.1877)	1
	rs1346763	T		1.096 (1.056)		0.940 (0.9059)		1.278 (1.231)		0.241 (0.4862)	
	rs1430101	G	1.043	1.028 (1.03)	0.866	0.853 (0.8559)	1.256	1.239 (1.24)	0.661	0.772 (0.7535)	0.97
	rs17048470	A		0.68		0.37	1.23		0.2		
	rs2420165	T		0.993 (0.8931)		0.743 (0.6958)		1.325 (1.146)		0.960 (0.3744)	
	rs11674031	G		0.857 (0.8823)		0.709 (0.7259)		1.036 (1.072)		0.110 (0.2084)	
	rs1430103	G		1.042 (1.034)		0.859 (0.8594)		1.264 (1.245)		0.675 (0.7215)	
	rs6746254	A		1.089 (1.073)		0.863 (0.8618)		1.374 (1.335)		0.471 (0.5304)	
	rs10048644	T		0.97		0.75	1.27		0.84		
	rs11123246	T		0.944 (0.8862)		0.742 (0.7287)		1.201 (1.078)		0.638 (0.2263)	
	rs1430108	C	1.006	1.009 (1.003)	0.865	0.867 (0.8621)	1.170	1.173 (1.166)	0.940	0.912 (0.9733)	1
	rs6542214	C	0.983	0.985 (0.9856)	0.844	0.845 (0.8459)	1.146	1.148 (1.148)	0.828	0.842 (0.8521)	0.97
	rs12987334	C		0.865 (0.9043)		0.652 (0.6871)		1.147 (1.119)		0.314 (0.4731)	
	rs1835328	T		0.99		0.83	1.18		0.93		
	rs7590021	C		0.972 (1.033)		0.790 (0.8503)		1.196 (1.255)		0.786 (0.7436)	
	rs6741327	A		1.08		0.88	1.31		0.47		
	rs10514812	A	1.219	1.214 (1.221)	0.947	0.942 (0.9488)	1.569	1.564 (1.572)	0.124	0.134 (0.1207)	1
	rs10496466	A	1.007	1.002 (1.015)	0.829	0.822 (0.8359)	1.224	1.221 (1.233)	0.941	0.985 (0.879)	0.97
	rs6729737	G	1.031	1.035 (1.031)	0.880	0.883 (0.8801)	1.209	1.213 (1.208)	0.705	0.672 (0.7034)	1
	rs996135	A	1.090	1.094 (1.093)	0.917	0.921 (0.9205)	1.295	1.299 (1.299)	0.328	0.307 (0.3091)	1
	rs12999482	A		0.855 (0.8485)		0.667 (0.6613)		1.098 (1.089)		0.220 (0.1965)	
	rs1965088	G	1.002	1.000 (1.005)	0.827	0.825 (0.8291)	1.213	1.211 (1.218)	0.987	0.998 (0.9617)	1
	rs17715313	C		0.98		0.68	1.43		0.93		
	rs7582739	C	1.217	1.167 (1.169)	0.939	0.904 (0.905)	1.576	1.503 (1.509)	0.138	0.235 (0.232)	0.99
	rs1347502	C		0.974 (0.9522)		0.808 (0.7985)		1.174 (1.136)		0.781 (0.5858)	
	rs17782840	A		1.053 (0.9971)		0.815 (0.7757)		1.360 (1.282)		0.694 (0.9818)	
	rs785033	C		1.02		0.87	1.2		0.81		
	rs785041	G		1.227 (1.093)		0.885 (0.8045)		1.702 (1.484)		0.220 (0.5708)	
	rs17715765	G		0.98		0.63	1.53		0.94		
	rs13029707	T		0.94		0.4	2.17		0.88		
	rs17715867	C		1.048 (1.057)		0.732 (0.7603)		1.502 (1.47)		0.797 (0.7406)	
	rs6542217	A	0.982	0.985 (0.9781)	0.832	0.835 (0.8					

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	rs13035561	T		1.079 (1.046)		0.925 (0.9015)		1.258 (1.215)		0.333 (0.5508)	
	rs7587705	A		1.07		0.91		1.26		0.41	
	rs10201854	T		1.09		0.93		1.28		0.29	
	rs12995783	T		1.26		0.91		1.73		0.16	
	rs4277531	T	1.059 /	1.059 (1.059)	0.910 /	0.911 (0.9105)	1.232 /	1.232 (1.232)	0.459 /	0.456 (0.4561)	1
	rs10496487	T	1.111 /	1.111 (1.111)	0.943 /	0.943 (0.9431)	1.309 /	1.309 (1.309)	0.208 /	0.208 (0.2076)	1
	rs10496486	G		0.84		0.66		1.05		0.13	
	rs4459743	C	1.052 /	1.049 (1.046)	0.906 /	0.904 (0.9015)	1.222 /	1.218 (1.215)	0.507 /	0.527 (0.5508)	1
	rs7579207	A	1.078 /	1.074 (1.074)	0.914 /	0.911 (0.9107)	1.271 /	1.266 (1.266)	0.372 /	0.397 (0.3968)	1
	rs10496485	T		0.94		0.78		1.14		0.54	
	rs7424115	C	0.974 /	0.97	0.728 /	0.73	1.302 /	1.3	0.858 /	0.86	1
	rs9677654	C		0.791 (0.8382)		0.410 (0.4394)		1.523 (1.599)		0.482 (0.5923)	
	rs983829	T	0.956 /	0.94	0.729 /	0.72	1.255 /	1.23	0.747 /	0.65	0.99
	rs958457	G	1.036 /	1.042 (0.9381)	0.874 /	0.878 (0.7147)	1.229 /	1.235 (1.231)	0.684 /	0.639 (0.645)	1
	rs13416481	G		0.895 (0.8833)		0.715 (0.7054)		1.121 (1.106)		0.335 (0.2794)	
	rs2280193	A		0.793 (0.7599)		0.557 (0.5361)		1.130 (1.077)		0.199 (0.123)	
	rs12469615	G		1.03		0.81		1.31		0.8	
	rs17687793	C		1.097 (1.127)		0.741 (0.7702)		1.624 (1.648)		0.644 (0.5387)	
	rs17044138	G		0.8		0.57		1.13		0.21	
	rs17687884	C		0.96		0.59		1.57		0.88	
	rs2420884	A		1.01		0.76		1.35		0.93	
	rs4131302	G	1.027 /	1.020 (1.024)	0.877 /	0.872 (0.8746)	1.202 /	1.194 (1.198)	0.742 /	0.802 (0.772)	1
	rs10445863	C	1.037 /	1.032 (1.036)	0.890 /	0.887 (0.8892)	1.207 /	1.202 (1.206)	0.645 /	0.682 (0.6525)	0.98
	rs10195710	A	1.085 /	1.088 (1.085)	0.932 /	0.935 (0.932)	1.262 /	1.263 (1.262)	0.294 /	0.276 (0.2941)	1
	rs2176250	G	1.040 /	1.045 (1.067)	0.861 /	0.865 (0.8818)	1.257 /	1.265 (1.29)	0.684 /	0.649 (0.5065)	1
	rs11693764	T		1.028 (1.019)		0.883 (0.8752)		1.197 (1.186)		0.719 (0.8088)	
	rs11675397	G		1.07		0.91		1.25		0.43	
	rs7558702	A	1.119 /	1.108 (1.108)	0.924 /	0.916 (0.9158)	1.356 /	1.340 (1.34)	0.250 /	0.292 (0.2923)	1
	rs13005061	G		1.08		0.82		1.41		0.58	
	rs7566462	C		0.93		0.78		1.11		0.44	
	rs2067699	G		0.94		0.71		1.26		0.69	
	rs17044170	G		0.903 (0.9275)		0.736 (0.7534)		1.108 (1.142)		0.328 (0.478)	
	rs11123287	A		0.96		0.83		1.12		0.6	
	rs7593121	G	1.031 /	1.032 (1.032)	0.880 /	0.882 (0.8815)	1.208 /	1.209 (1.209)	0.703 /	0.692 (0.6921)	1
	rs11896538	G		0.87		0.72		1.07		0.18	
	rs956534	A		1		0.85		1.17		0.98	
	rs9308710	T	1.017 /	0.932 (0.9315)	0.741 /	0.683 (0.6834)	1.396 /	1.270 (1.27)	0.917 /	0.653 (0.6532)	0.99
	rs1123288	A		0.941 (0.949)		0.790 (0.7967)		1.121 (1.13)		0.497 (0.5572)	
	rs10496483	G	1.155 /	1.155 (1.136)	0.894 /	0.895 (0.8804)	1.492 /	1.490 (1.465)	0.269 /	0.268 (0.3277)	1
	rs17355679	T		1.03		0.88		1.21		0.72	
	rs4610066	G		1.03		0.84		1.25		0.8	
	rs4949387	G	1.143 /	1.146 (1.137)	0.885 /	0.888 (0.8809)	1.478 /	1.479 (1.467)	0.306 /	0.295 (0.3244)	0.99
	rs10204084	A		1		0.86		1.17		0.96	
	rs6542253	G		0.83		0.32		2.16		0.7	
	rs11123289	G		0.99		0.85		1.15		0.89	
	rs12991531	T		1.002 (0.9699)		0.751 (0.7288)		1.338 (1.291)		0.987 (0.8338)	
	rs6542254	G	0.931 /	0.928 (0.9277)	0.756 /	0.754 (0.7535)	1.147 /	1.142 (1.142)	0.502 /	0.480 (0.4797)	1
	rs10206444	G		1.012 (1.097)		0.708 (0.7554)		1.446 (1.594)		0.947 (0.6261)	
	rs13001269	G	1.061 /	1.061 (1.061)	0.808 /	0.808 (0.8081)	1.392 /	1.392 (1.392)	0.671 /	0.671 (0.6709)	1
	rs4849389	A	1.051 /	1.03	0.819 /	0.81	1.348 /	1.31	0.697 /	0.84	1
	rs13009552	G		1.03		0.82		1.29		0.82	
	rs10496482	G		0.956 (0.9637)		0.823 (0.8287)		1.111 (1.121)		0.558 (0.6311)	
	rs12991665	A	0.959 /	0.979 (0.9785)	0.633 /	0.647 (0.6473)	1.452 /	1.479 (1.479)	0.842 /	0.918 (0.9181)	0.99
	rs6759005	C		0.99		0.75		1.31		0.95	
	rs7577833	G	0.973 /	0.976 (0.9787)	0.837 /	0.839 (0.8422)	1.130 /	1.134 (1.137)	0.716 /	0.747 (0.7788)	1
	rs13386690	G		0.970 (0.9773)		0.822 (0.8274)		1.145 (1.154)		0.719 (0.787)	
	rs13033618	G		1.1		0.77		1.58		0.59	
	rs4849390	T	1.003 /	1.001 (0.9917)	0.834 /	0.832 (0.8249)	1.206 /	1.203 (1.192)	0.973 /	0.993 (0.9296)	1
	rs12619497	A	1.001 /	0.979 (0.9785)	0.664 /	0.647 (0.6473)	1.511 /	1.479 (1.479)	0.995 /	0.918 (0.9181)	0.99
	rs11690705	A	1.080 /	1	0.890 /	0.83	1.310 /	1.2	0.437 /	0.99	0.98
	rs6741354	G		0.31		0.03		3		0.31	
	rs9646900	C		1.18		0.68		2.04		0.56	
	rs6542256	C	0.986 /	0.977 (0.9774)	0.764 /	0.758 (0.7583)	1.271 /	1.260 (1.26)	0.911 /	0.860 (0.8598)	1
	rs908599	G		0.97		0.82		1.15		0.73	
	rs1356383	T		0.93		0.66		1.32		0.7	
	rs2420815	C	1.009 /	0.997 (0.9927)	0.853 /	0.843 (0.8397)	1.193 /	1.178 (1.174)	0.921 /	0.967 (0.9318)	0.99
	rs10176449	T		1.01		0.86		1.19		0.91	
	rs17044234	A		0.84		0.56		1.26		0.39	
	rs2420819	T	0.992 /	0.990 (0.9902)	0.843 /	0.841 (0.8413)	1.168 /	1.165 (1.165)	0.925 /	0.906 (0.9058)	1
	rs17623623 *	T		1.62		1.05		2.52		0.03	
	rs17044254	C		0.94		0.62		1.42		0.76	
	rs2421100	A	0.960 /	0.960 (0.9595)	0.814 /	0.814 (0.8143)	1.131 /	1.131 (1.131)	0.622 /	0.622 (0.6216)	1
	rs70925	C	0.937 /	0.937 (0.9369)	0.806 /	0.806 (0.8057)	1.089 /	1.089 (1.089)	0.397 /	0.397 (0.3972)	1
	rs9308712	G	1.016 /	1.016 (1.016)	0.833 /	0.833 (0.8328)	1.240 /	1.240 (1.24)	0.875 /	0.875 (0.8752)	1
	rs13022199	A		1.007 (0.9992)		0.797 (0.7908)		1.271 (1.262)		0.957 (0.9944)	
	rs2421101	C	0.964 /	0.962 (0.9615)	0.829 /	0.827 (0.8268)	1.122 /	1.118 (1.118)	0.638 /	0.610 (0.6097)	1
	rs17044296	A		1.18		0.71		1.97		0.52	
	rs1155664	G	0.877 /	0.873 (0.8727)	0.715 /	0.711 (0.7112)	1.076 /	1.071 (1.071)	0.210 /	0.192 (0.1919)	1
	rs17453235	C		1.130 (1.058)		0.909 (0.8464)		1.404 (1.322)		0.272 (0.6214)	
	rs11683441	T	1.002 /	0.993 (0.9925)	0.857 /	0.849 (0.8488)	1.172 /	1.161 (1.161)	0.979 /	0.925 (0.9249)	1
	rs7561977	A		0.93		0.69		1.27		0.65	
	rs843399	T	0.950 /	0.950 (0.95)	0.803 /	0.803 (0.8025)	1.125 /	1.125 (1.125)	0.551 /	0.551 (0.5511)	1
	rs952747	C	0.950 /	0.950 (0.95)	0.803 /	0.803 (0.8025)	1.125 /	1.125 (1.125)	0.551 /	0.551 (0.5511)	1
	rs1717054 (rs843385 / r ² =1)	T	1.066 /	1.066 (1.066)	0.821 /	0.821 (0.8205)	1.384 /	1.384 (1.384)	0.633 /	0.633 (0.6334)	1
	rs2030440	T		0.98		0.84		1.14		0.79	
	rs1517363	G		1.19		0.65		2.19		0.58	
	rs843408	G		0.974 (0.9712)		0.836 (0.834)		1.134 (1.131)		0.735 (0.7072)	
	rs843447	T		0.72		0.47		1.09		0.12	
	rs6732466	T		0.98		0.84		1.15		0.79	
	rs1037205	G	0.979 /	0.979 (0.9859)	0.835 /	0.835 (0.8407)	1.148 /	1.148 (1.156)	0.792 /	0.792 (0.8616)	1
	rs17362720	A		1.094 (1.076)		0.848 (0.8275)		1.411 (1.4)		0.489 (0.583)	
	rs13033419	A	1.018 /	1.020 (1.019)	0.849 /	0.851 (0.8504)	1.220 /	1.222 (1.222)	0.849 /	0.833 (0.8355)	1
	rs1868327	C	1.015 /	1.020 (1.02)	0.846 /	0.851 (0.8506)	1.218 /	1.222 (1.222)	0.871 /	0.833 (0.833)	1
	rs17362881	T		0.92		0.72		1.18		0.52	
	rs17453808	T		1.05		0.4		2.75		0.92	
	rs13429901	T	0.962 /	0.962 (0.9619)	0.765 /	0.765 (0.7648)	1.210 /	1.210 (1.21)	0.740 /	0.740 (0.7396)	1
	rs7584395	A	0.996 /	0.996 (1.003)	0.846 /	0.846 (0.8518)	1.172 /	1.172 (1.18)	0.960 /	0.960 (0.9758)	1
	rs1971190	C		1.029 (1.032)		0.860 (0.8636)		1.231 (1.234)		0.755 (0.7269)	
	rs1006342	G	0.968 /	0.968 (0.9616)	0.770 /	0.770 (0.7648)	1.217 /	1.217 (1.209)	0.780 /	0.780 (0.7372)	1
	rs12471022	T		2.531 (1.752)		0.984 (0.6947)		6.507 (4.419)		0.054 (0.2347)	
	rs17044686	A		0.82		0.55		1.23		0.34	
	rs1545396 (rs756583 / r ² =1)	T	0.972 /	0.986 (0.9864)	0.739 /	0.750 (0.7501)	1.280 /	1.297 (1.297)	0.841 /	0.922 (0.9217)	1
	rs1550985	T	1.046 /	1.035 (1.042)	0.887 /	0.878 (0.884)	1.233 /	1.221 (1.228)	0.594 /	0.680 (0.6246)	1
	rs6738130	C	1.058 /	1.042 (1.045)	0.889 /	0.884 (0.8871)	1.259 /	1.229 (1.232)	0.524 /	0.622 (0.5968)	0.99
	rs921211	C	1.019 /	1.019 (1.019)	0.876 /	0.875 (0.8752)	1.187 /	1.187 (1.187)	0.805 /	0.808 (0.8081)	1
	rs17044762	G	0.826 /	0.835 (0.8354)	0.595 /	0.603 (0.6029)	1.146 /	1.158 (1.158)	0.252 /	0.280 (0.2798)	

Gene	SNPs	minor allele	OR genotyping	OR imputation HM2 (HM3)	CI_L 95% genotyping	CI_L 95% imputation HM2 (HM3)	CI_U 95% genotyping	CI_U 95% imputation HM2 (HM3)	p-value genotyping	p-value imputation HM2 (HM3)	R2	
GSTP1	rs7927381	T		1.273 (1.242)		0.981 (0.9596)		1.654 (1.608)		0.070 (0.09953)		
	rs614080 **	A	0.778 /	0.778 (0.7779)	0.668 /	0.668 (0.6679)	0.906 /	0.906 (0.906)	0.0013 /	0.0012 (0.0012)	1	
	rs6591255	A	1.159 /	(1.192)	0.990 /	(1.024)	1.356 /	(1.388)	0.066 /	(0.02357)		
	rs6591256 *	G	1.190 /	1.18	1.022 /	1.01	1.387 /	1.37	0.026 /	0.03	0.99	
	rs1695	G	1.125 /	(1.165 (1.212))	0.962 /	(0.996 (1.034))	1.316 /	(1.362 (1.421))	0.141 /	(0.056 (0.01765))	0.94	
	rs1799811	T	0.985 /	1.04	0.758 /	0.81	1.279 /	1.35	0.908 /	0.76	1	
	rs1871042	T		1.161 (1.227)		0.993 (1.047)		1.358 (1.439)		0.061 (0.0116)		
IL12B	rs6870828	C		0.94		0.81		1.09		0.4		
	rs1965014	T		0.92		0.76		1.12		0.43		
	rs3181224	C	1.013 /	1.01	0.797 /	0.8	1.289 /	1.29	0.914 /	0.91	1	
	rs3212227	G		1.09		0.9		1.32		0.4		
	rs3213120	T		1.21		0.77		1.9		0.41		
	rs3213102 *	T		3.04		1.11		8.35		0.03		
	rs2853694	T		1.028 (1.028)		0.884 (0.8844)		1.195 (1.195)		0.718 (0.7184)		
	rs3213096	T		5.656 (2.12)		0.679 (0.6498)		47.100 (6.918)		0.109 (0.2129)		
	rs3213094	T		1.08		0.89		1.31		0.43		
	rs2569253	T	1.012 /	0.988 (0.9913)	0.868 /	0.847 (0.8496)	1.181 /	1.153 (1.157)	0.877 /	0.879 (0.9112)	1	
	rs2569254	T	0.953 /	(0.955 (0.9603))	0.784 /	(0.787 (0.7906))	1.157 /	(1.160 (1.166))	0.624 /	(0.646 (0.6827))	1	
	rs2546891	A		3.78		0.8		17.84		0.09		
	rs3181216	T	1.023 /	(1.02)	0.866 /	(0.87)	1.209 /	(1.2)	0.789 /	(0.79)	0.97	
	rs3213117	C		1.05		0.69		1.61		0.82		
	rs1003199	T		1.013 (0.9847)		0.868 (0.8438)		1.181 (1.149)		0.872 (0.8449)		
	rs1433048	G	0.884 /	0.872 (0.8816)	0.724 /	0.713 (0.7221)	1.080 /	1.065 (1.076)	0.227 /	0.180 (0.2159)	0.98	
	rs2546893	A		1.01		0.86		1.17		0.94		
rs730691	T	0.983 /	0.982 (0.9818)	0.837 /	0.836 (0.836)	1.154 /	1.153 (1.153)	0.831 /	0.821 (0.8228)	1		
rs2546890	G	0.980 /	(0.96 (0.9685))	0.839 /	(0.828 (0.8308))	1.144 /	(1.126 (1.129))	0.795 /	(0.655 (0.6831))	1		
rs10052709	G		0.98		0.79		1.23		0.89			
rs1422876	C		0.94		0.8		1.09		0.41			
IL13	rs3091307	G		1.11		0.93		1.33		0.24		
	rs1881457	C	1.076 /	(1.12)	0.898 /	(0.9393)	1.291 /	(1.334)	0.427 /	(0.2074)		
	rs1800925	T	1.085		0.904		1.303		0.380			
	rs1295686 *	A	1.220 /	1.230 (1.23)	1.016 /	1.024 (1.024)	1.464 /	1.478 (1.478)	0.033 /	0.027 (0.02681)	0.99	
	rs20541 *	T	1.227 /	(1.225 (1.221))	1.021 /	(1.020 (1.016))	1.475 /	(1.472 (1.468))	0.029 /	(0.030 (0.0319))	1	
	rs1295683 *	A		1.26		1		1.59		0.05		
	rs2243297	A		1.39		0.92		2.11		0.12		
	rs2243204	T	1.129 /	1.129 (1.129)	0.870 /	0.870 (0.8701)	1.464 /	1.464 (1.464)	0.362 /	0.362 (0.3616)	1	
	rs2243298	A		2.15		0.7		6.56		0.18		
	IL4	rs762534	A		1.14		0.86		1.51		0.37	
rs2243240		T		1.62		0.63		4.13		0.32		
rs2243248		G		0.988 (0.9442)		0.772 (0.6999)		1.291 (1.274)		0.990 (0.7071)		
rs2243250		T	1.225 /	(1.23)	0.997 /	(1.006)	1.505 /	(1.505)	0.054 /	(0.04348)		
rs2070874		T	1.221 /	(1.356 (1.23))	0.997 /	(1.098 (1.006))	1.494 /	(1.675 (1.505))	0.054 /	(0.005 (0.04348))	0.77	
rs2243263		C		0.98		0.76		1.25		0.84		
rs2243274 *		A		1.24		1.01		1.52		0.04		
rs2406539		T		1.16		0.98		1.37		0.09		
IL4R	rs8057585	C		1.104 (1.097)		0.946 (0.9404)		1.288 (1.28)		0.209 (0.2388)		
	rs2057768 *	C	1.267 /	(1.119 (1.106))	1.061 /	(0.944 (0.9335))	1.512 /	(1.326 (1.31))	0.0089 /	(0.195 (0.2443))	0.89	
	rs6498012	T		1.080 (1.071)		0.922 (0.9145)		1.263 (1.253)		0.340 (0.3959)		
	rs1110470	A		0.89		0.76		1.03		0.12		
	rs4787948	G	1.110 /	1.111 (1.115)	0.940 /	0.940 (0.9434)	1.312 /	1.312 (1.317)	0.218 /	0.217 (0.2022)	1	
	rs2283563	A	0.896 /	0.893 (0.8928)	0.761 /	0.759 (0.7588)	1.054 /	1.050 (1.05)	0.185 /	0.172 (0.1719)	1	
	rs3024530	G	1.134 /	1.134 (1.134)	0.975 /	0.975 (0.9748)	1.319 /	1.319 (1.319)	0.103 /	0.103 (0.1033)	1	
	rs3024536	T	1.091 /	1.091 (1.085)	0.889 /	0.889 (0.8842)	1.338 /	1.338 (1.331)	0.406 /	0.406 (0.4348)	1	
	rs1805010 *	G	1.234 /	(1.149)	1.053 /	(0.9865)	1.446 /	(1.337)	0.0093 /	(0.07421)		
	rs3024560	G		1.151 (1.157)		0.985 (0.9903)		1.344 (1.352)		0.078 (0.06614)		
	rs2301807	A		0.85		0.6		1.2		0.36		
	rs2239349	A		0.933 (0.8096)		0.542 (0.5906)		1.609 (1.11)		0.804 (0.1893)		
	rs2239346 *	A		0.84		0.72		0.98		0.02		
	rs3116578	T		1.9		0.85		4.25		0.12		
	rs3024585 *	A	1.180 /	1.179 (1.175)	1.014 /	1.013 (1.01)	1.373 /	1.371 (1.366)	0.032 /	0.033 (0.03679)	1	
	rs3024613	T		0.869 (0.8786)		0.747 (0.756)		1.010 (1.021)		0.067 (0.09138)		
	rs3024620	G		0.86		0.6		1.23		0.41		
	rs3024622 *	G		1.22		1.04		1.42		0.01		
	rs3024630	G	0.932 /	0.946 (0.9454)	0.724 /	0.736 (0.7348)	1.199 /	1.215 (1.216)	0.582 /	0.661 (0.6621)	0.99	
	rs3024691	A		0.56		0.13		2.35		0.43		
	rs3024647	G		0.94		0.76		1.16		0.55		
	rs3024660	C	0.910 /	0.943 (0.912)	0.741 /	0.765 (0.742)	1.118 /	1.163 (1.121)	0.371 /	0.584 (0.3811)	0.96	
	rs2234897	C		0.85		0.52		1.39		0.51		
	rs1805011	C	1.114 /	(1.103 (1.103))	0.870 /	(0.862 (0.8624))	1.428 /	(1.411 (1.411))	0.392 /	(0.435 (0.4346))	1	
rs1805011 (rs2234898 / r ² =0.95)	C	1.114 /	(1.103 (1.104))	0.870 /	(0.862 (0.8628))	1.428 /	(1.411 (1.413))	0.392 /	(0.435 (0.4311))	1		
rs1805012	C	1.111 /	(1.104 (1.104))	0.865 /	(0.863 (0.8628))	1.427 /	(1.413 (1.413))	0.411 /	(0.431 (0.4311))	1		
rs2234900	C	- /	(1.104)	- /	(0.8628)	- /	(1.413)	- /	(0.4311)			
rs1805015	C	1.035 /	(1.030 (1.024))	0.842 /	(0.838 (0.8293))	1.272 /	(1.265 (1.264))	0.744 /	(0.782 (0.828))	1		
rs1801275	G	1.052 /	(1.05)	0.874 /	(0.87)	1.267 /	(1.27)	0.589 /	(0.59)	1		
rs1805016	G		0.848 (0.9305)		0.602 (0.6517)		1.195 (1.328)		0.347 (0.6917)			
rs2074570	C		1.13		0.77		1.67		0.53			
rs1049631	G	1.083 /	1.084 (1.084)	0.931 /	0.932 (0.9315)	1.260 /	1.260 (1.26)	0.301 /	0.298 (0.2983)	1		
rs3024698	A		0.85		0.46		1.57		0.6			
rs3024685	C		1.020 (1.027)		0.874 (0.8805)		1.190 (1.198)		0.802 (0.7353)			
rs12102586	T		1.04		0.8		1.36		0.78			
rs4787956	G		1.037 (1.049)		0.886 (0.8969)		1.214 (1.227)		0.650 (0.5495)			
PTGDR	rs803014	A		1.07		0.86		1.33		0.56		
	rs803012 (rs708487 / r ² =1)	C	0.939 /	(0.941 (0.9412))	0.784 /	(0.786 (0.7863))	1.124 /	(1.127 (1.127))	0.493 /	(0.509 (0.5087))	1	
	rs803011	T		0.945		0.786		1.137		0.52		
	rs8004654	T	0.913 /	(0.9531)	0.782 /	(0.8204)	1.066 /	(1.107)	0.250 /	(0.5302)		
	rs11157907	C	0.975		0.788		1.207		0.816			
	rs11851957	T	0.957 /	0.957 (0.9516)	0.775 /	0.775 (0.7703)	1.182 /	1.182 (1.176)	0.684 /	0.684 (0.6457)	1	
	rs17831669	C		1.134 (1.111)		0.866 (0.8503)		1.484 (1.453)		0.361 (0.4392)		
	rs708486	C	0.987 /	(0.99)	0.847 /	(0.85)	1.150 /	(1.15)	0.866 /	(0.87)	1	
	rs810633 (rs803010 / r ² =0.85)	T	1.042 /	(1.042 (1.042))	0.879 /	(0.879 (0.8785))	1.235 /	(1.235 (1.235))	0.639 /	(0.639 (0.6391))	1	
	rs12895027	T		1.155 (1.581)		0.651 (0.8259)		2.048 (3.025)		0.623 (0.1669)		
	rs1953367	C	1.012 /	1.014 (1.005)	0.848 /	0.849 (0.8428)	1.207 /	1.210 (1.199)	0.898 /	0.881 (0.9552)	0.99	
	TNF	rs915654	A		1.12		0.95		1.31		0.18	
		rs2844482 (rs1800630 / r ² =0.89)	A	1.145 /	(1.170 (1.17))	0.940 /	(0.962 (0					