

Barrett et al SNP	WTCCC SNP	r2	Chr	LD region	Gene or Num of genes
rs2269241	rs2269241	1	1p31.3	63.87–63.94	<i>PGM1</i>
rs2476601	rs6679677	1	1p13.2	113.62–114.46	<i>PTPN22</i>
rs2816316	rs1323296	1	1q31.2	190.73–190.82	<i>RGS1</i>
rs3024505	rs3024505	1	1q32.1	204.87–205.12	<i>IL10</i>
rs1534422	rs10175906	0.72	2p25.1	12.53–12.60	(0)
rs917997	rs2041756	1	2q12.1	102.22–102.58	<i>IL18RAP</i>
rs1990760	rs7608315	0.59	2q24.2	162.67–163.10	<i>IFIH1</i>
rs3087243	rs3087243	1	2q33.2	204.38–204.53	<i>CTLA4</i>
rs11711054	rs6441961	1	3p21.31	45.96–46.63	<i>CCR5</i>
rs10517086	rs17698094	0.54	4p15.2	25.64–25.75	(0)
rs4505848	rs17388568	0.83	4q27	123.13–123.83	<i>IL2</i>
rs6897932	rs6897932	1	5p13.2	35.84–36.07	<i>IL7R</i>
rs9268645	rs9268645	1	6p21.32	24.70–34.00	<i>MHC</i>
rs11755527	rs11755527	1	6q15	90.86–91.10	<i>BACH2</i>
rs9388489	rs9388489	1	6q22.32	126.48–127.46	<i>C6orf173</i>
rs2327832	rs2327832	1	6q23.3	137.80–138.40	<i>TNFAIP3</i>
rs1738074	rs2016588	0.57	6q25.3	159.13–159.62	<i>TAGAP</i>
rs7804356	rs10486478	1	7p15.2	26.62–27.17	(10)
rs4948088	rs10486748	0.41	7p12.1	50.87–51.64	<i>COBL</i>
rs7020673	rs7020673	1	9p24.2	4.22–4.31	<i>GLIS3</i>
rs12251307	rs12251307	1	10p15.1	6.07–6.24	<i>IL2RA</i>
rs11258747	rs11258747	1	10p15.1	6.48–6.59	<i>PRKCQ</i>
rs10509540	rs11816865	0.66	10q23.31	90.00–90.27	<i>C10orf59</i>
rs7111341 ¹	rs6578252	0.07	11p15.5	2.02–2.26	<i>INS</i>
rs4763879	rs4763879	1	12p13.31	9.51–9.87	<i>CD69</i>
rs2292239	rs2292239	1	12q13.2	54.64–55.09	<i>ERBB3</i>
rs1678536	rs1678542	0.62	12q13.3	55.27–56.82	<i>Multiple</i>
rs3184504	rs1265566	0.25	12q24.12	109.77–111.72	<i>SH2B3</i>
rs1465788	rs1465788	1	14q24.1	68.24–68.39	(2)
rs4900384	rs4900384	1	14q32.2	97.43–97.60	(0)
rs3825932	rs3825932	1	15q25.1	76.77–77.05	<i>CTSH</i>
rs12708716	rs12708716	1	16p13.13	10.92–11.56	<i>CLEC16A</i>
rs12444268	rs12444268	1	16p12.3	20.17–20.28	(2)
rs4788084	rs151181	0.9	16p11.2	28.19–28.94	<i>IL27</i>
rs7202877	rs4536500	0.64	16q23.1	73.76–74.09	(7)
rs16956936	rs16956936	1	17p13.1	7.56–7.66	(2)
rs2290400	rs1008723	1	17q12	34.63–35.51	<i>ORMDL3</i>
rs7221109	rs7221109	1	17q21.2	35.95–36.13	(3)
rs1893217	rs16939895	0.69	18p11.21	12.73–12.92	<i>PTPN2</i>
rs763361	rs1788101	0.18	18q22.2	65.63–65.72	<i>CD226</i>

rs425105	rs4804000	0.69	19q13.32	51.84–52.02	(5)
rs2281808	rs2281808	1	20p13	1.44–1.71	(3)
rs11203203	rs11203203	1	21q22.3	42.68–42.76	<i>UBASH3A</i>
rs229541	rs229541	1	22q13.1	35.90–36.00	<i>C1QTNF6</i>
rs5753037	rs41176	1	22q12.2	28.14–29.00	(14)
rs2664170 ²			Xq28	153.48–154.10	(16)

1: The *INS* locus is not well covered by the Affymetrix array.

2: The chromosome X marker is not used in our study.