

Supplementary Table 3. Imputed results of the ImmunoChip study with Crohn's disease in Koreans

SNP	Chr	Position (hg19)	Gene	Risk allele	Non-risk allele	RAF in cases	RAF in controls	OR	95% CI	<i>P</i> trend
rs75323844	4q24	103,774,293	<i>UBE2D3</i>	T	TC	0.60	0.52	1.42	(1.19-1.69)	8.51E-05
rs73771645	6q22	106,016,785	<i>PREP</i>	T	C	0.05	0.02	3.13	(1.77-5.54)	3.90E-05
rs58761034	11p15	524,864	(<i>PTDSS2, RNHI</i>)	G	A	0.86	0.80	1.55	(1.24-1.94)	9.36E-05
rs637304	11q14	85,685,166	<i>PICALM</i>	A	G	0.30	0.23	1.51	(1.24-1.83)	4.31E-05
rs600743	11q14	85,686,223	<i>PICALM</i>	A	T	0.30	0.23	1.50	(1.23-1.83)	4.87E-05
rs585820	11q14	85,696,670	<i>PICALM</i>	G	A	0.30	0.22	1.51	(1.24-1.84)	3.94E-05
rs670606	11q14	85,698,598	<i>PICALM</i>	G	A	0.30	0.23	1.49	(1.22-1.81)	7.10E-05
rs654194	11q14	85,700,000	<i>PICALM</i>	C	A	0.30	0.23	1.49	(1.22-1.81)	7.10E-05
rs595972	11q14	85,705,885	<i>PICALM</i>	G	T	0.30	0.23	1.48	(1.22-1.81)	7.61E-05
rs616555	11q14	85,710,180	<i>PICALM</i>	G	A	0.30	0.23	1.48	(1.22-1.80)	8.10E-05
rs28645497	17p13	5,042,241	<i>USP6</i>	A	G	0.55	0.46	1.42	(1.19-1.69)	9.49E-05
rs140200174	17p13	5,047,599	<i>USP6</i>	A	ATATT	0.54	0.46	1.44	(1.20-1.72)	7.48E-05
rs200642633	21q11	16,790,941	(<i>USP25</i>)	C	CA	0.78	0.70	1.49	(1.22-1.82)	9.98E-05
rs72635015	21q21	27,451,908	<i>APP</i>	A	T	0.22	0.15	1.59	(1.27-1.99)	4.95E-05
rs11910285	21q21	27,458,446	<i>APP</i>	C	T	0.22	0.15	1.59	(1.27-1.99)	4.95E-05

Note: Chr, chromosome; CI, confidence interval; OR, odds ratio; RAF, risk allele frequency.