

Supplementary Table 2. Association of established IBD susceptibility variants with age of onset in this study cohort

rsID	HR	<i>p</i> value	N	Beta	SE	Ethnicity
224 high-risk associations						
rs3194051	0.592	0.100	149	−0.525	0.319	European
rs2073505	1.119	0.402	341	0.112	0.134	Multi-ethnicity
rs12103	1.294	0.455	341	0.258	0.345	European
rs3197999	1.075	0.681	341	0.072	0.176	Multi-ethnicity
rs1801274	1.052	0.734	341	0.051	0.151	European
rs2227564	1.014	0.910	341	0.014	0.121	European
rs11230563	1.004	0.975	341	0.004	0.123	Multi-ethnicity
783 general associations						
rs3749171	0.705	0.025	341	−0.349	0.156	European
rs2234161	1.334	0.042	341	0.289	0.142	European
rs501916	0.804	0.106	341	−0.218	0.135	European
rs3742130	1.326	0.150	341	0.282	0.196	Multi-ethnicity
rs2641348	0.788	0.382	341	−0.238	0.272	Multi-ethnicity
rs76418789	0.882	0.526	341	−0.126	0.198	Koreans
rs3747517	0.943	0.675	341	−0.059	0.140	European
rs7076156	0.914	0.691	341	−0.090	0.226	Jewish
rs9858542	1.033	0.882	341	0.033	0.219	European
rs8049439	1.018	0.905	341	0.017	0.147	European
rs9868809	0.990	0.950	341	−0.010	0.152	Multi-ethnicity
rs13300218	1.031	0.967	341	0.030	0.725	Multi-ethnicity
rs732072	1.006	0.978	341	0.006	0.216	Multi-ethnicity
rs2476601	-	-	341	-	0	European
rs3184504	-	-	341	-	0	European

Data are presented for variants identified from established susceptibility loci. *p* values were calculated using Cox proportional hazards regression.

HR, hazard ratio; Beta, effect size (log-hazard ratio); SE, standard error; N, number of individuals; IBD, inflammatory bowel disease; rsID, reference SNP cluster ID.