

Supplemental Table 4. Single-variant association stratified by self-reported ancestry group. SNV: single nucleotide variant, GROUP: self-reported ancestry group, N: sample size, MAF: minor allele frequency, OR: odds ratio, CI: confidence interval, Z: Z-score, *P*: *P* value, NA: association not tested due to low allele count. Nominally significant results are highlighted in bold ($P < 0.05$).

SNV	GROUP	N	MAF	OR	95% CI	Z	<i>P</i>
<i>FZD3</i> rs352214	European-American	8,203	0.425	0.79	0.72-0.87	-5.15	< 1E-05
	African-American	1,315	0.313	0.97	0.75-1.27	-0.20	0.8420
	Hispanic-American	276	0.422	0.80	0.48-1.36	-0.82	0.4130
	Other	228	0.381	0.47	0.23-0.99	-1.99	0.0460
<i>MYH11</i> rs10852375	European-American	8,203	0.406	1.27	1.16-1.39	5.33	< 1E-05
	African-American	1,318	0.390	1.29	1.02-1.63	2.10	0.0360
	Hispanic-American	276	0.411	2.16	1.22-3.83	2.63	0.0090
	Other	228	0.318	1.95	0.94-4.00	1.81	0.0710