

Supplementary Table 2. Association between unweighted polygenic risk scores and intraocular pressure across five cross-validation runs

Model	Number of SNPs selected	β	SE	P
Iteration 1	1,253	0.0238	0.0008	2.9×10^{-194}
Iteration 2	1,262	0.0232	0.0008	2.6×10^{-189}
Iteration 3	1,242	0.0236	0.0008	1.2×10^{-194}
Iteration 4	1,274	0.0249	0.0008	3.0×10^{-208}
Iteration 5	1,250	0.0244	0.0008	1.5×10^{-206}

SNP = single nucleotide polymorphisms; SE = standard error.