(A)				
Weighted PRS Model	PRS built using SNPs with $P < 5 \times 10^{-5}$		PRS built using SNPs with $P < 5 \times 10^{-8}$	
	Number of SNPs selected	Р	Number of SNPs selected	Р
Iteration 1	1,253	1.0×10^{-190}	220	8.4×10^{-121}
Iteration 2	1,262	5.2×10^{-201}	228	4.9×10^{-138}
Iteration 3	1,242	3.2×10^{-202}	223	3.6×10^{-144}
Iteration 4	1,274	5.1×10^{-205}	234	4.7×10^{-141}
Iteration 5	1,250	3.5×10^{-209}	219	2.1×10^{-150}

Supplementary Table 3. Associations between polygenic risk scores and intraocular pressure across five cross-validation runs

(B)

Unweighted PRS Model	PRS built using SNPs with $P < 5 \times 10^{-5}$		PRS built using SNPs with $P < 5 \times 10^{-8}$	
	Number of SNPs selected	Р	Number of SNPs selected	Р
Iteration 1	1,253	2.9×10^{-194}	220	3.7×10^{-117}
Iteration 2	1,262	2.6×10^{-189}	228	5.2×10^{-129}
Iteration 3	1,242	1.2×10^{-194}	223	1.4×10^{-142}
Iteration 4	1,274	3.0×10^{-208}	234	2.5×10^{-136}
Iteration 5	1,250	1.5×10^{-206}	219	8.1×10^{-143}

(A) The association results between IOP and weighted PRSs using SNPs with $P < 5 \times 10^{-5}$ and 5×10^{-8} ; (B) The association results between IOP and unweighted PRSs. The association results of all runs in the 5 $\times 10^{-5}$ threshold category are more significant than those in the 5 $\times 10^{-8}$ category.

Abbreviations: SNP = single nucleotide polymorphism; IOP = intraocular pressure; PRS = polygenic risk score.