Table S4. Coefficients of determination (R<sup>2</sup>) on the observed scale (Nagelkerke's) and the liability scale for model variables in our adjusted GRS models for AFR and EUR Veterans in the MVP. We calculated Nagelkerke's R<sup>2</sup> and liability R<sup>2</sup> for each set of model variables to account for covariates in our adjusted models. To do this, we first entered age and sex into the model, then we added 10 PCs to account for population structure. We then added each GRS approach (unweighted, cross-ancestry meta-analysis weighted, and ancestry-specific weighted) into the model and calculated increases in R<sup>2</sup> by each GRS type. In addition, we include the single genome-wide significant AFR-specific variant in *IQGAP1* (rs16944405) to each GRS. For the liability scale calculations, we used a single fixed prevalence value for POAG (2.4%; cited in Zhang et al. 2021) for the sake of comparison.

AFR		EUR	
Nagelkerke's R <sup>2</sup> (Observed Scale)	Liability R <sup>2</sup>	Nagelkerke's R <sup>2</sup> (Observed Scale)	Liability R <sup>2</sup>
0.08645841	0.05212212	0.001087428	0.001253605
0.1031472	0.06273584	0.002002607	0.002308977
0.1271559	0.07834008	0.04673353	0.05427347
0.1328011	0.08206801	0.05788033	0.06734026
0.117113	0.07176384	0.05691808	0.06621041
0.1273081	0.0784403	0.04647621	0.05397238
0.1317377	0.08136405	0.05788689	0.06734797
0.1171453	0.07178492	0.05692639	0.06622016
0.0240087	0.01560424	0.044730923	0.051964493
0.0296539	0.01933217	0.055877723	0.065031283
0.0139658	0.009028	0.054915473	0.063901433
0.0241609	0.01570446	0.044473603	0.051663403
0.0285905	0.01862821	0.055884283	0.065038993
0.0139981	0.00904908	0.054923783	0.063911183
	Nagelkerke's R <sup>2</sup> (Observed Scale)  0.08645841  0.1031472  0.1271559  0.1328011  0.117113  0.1273081  0.1317377  0.1171453  0.0240087  0.0296539  0.0139658  0.0241609  0.0285905	Nagelkerke's R² (Observed Scale)         Liability R²           0.08645841         0.05212212           0.1031472         0.06273584           0.1271559         0.07834008           0.1328011         0.08206801           0.117113         0.07176384           0.1273081         0.0784403           0.1317377         0.08136405           0.1171453         0.07178492           0.0240087         0.01560424           0.0296539         0.01933217           0.0139658         0.009028           0.0241609         0.01570446           0.0285905         0.01862821	Nagelkerke's R² (Observed Scale)Liability R²Nagelkerke's R² (Observed Scale)0.086458410.052122120.0010874280.10314720.062735840.0020026070.12715590.078340080.046733530.13280110.082068010.057880330.1171130.071763840.056918080.12730810.07844030.046476210.13173770.081364050.057886890.11714530.071784920.056926390.02400870.015604240.0447309230.02965390.019332170.0558777230.01396580.0090280.0549154730.02416090.015704460.0444736030.02859050.018628210.055884283

**Abbreviations:** POAG = primary open-angle glaucoma, PCs = principal components, GRS = genetic risk scores, AFR = non-Hispanic African ancestry, EUR = non-Hispanic European ancestry, MVP = Million Veteran Program.