Supplemental Figure 1. Z-scores for heritability estimates.



Note: Phenotypes are presented in ascending order of heritability estimates (i.e., with the smallest Z-scores all the way to the left and the largest Z-scores all the way to the right). Actual Z-scores for each dataset are presented in each bar. As noted in the key, orange bars correspond to African Ancestry (AA) samples and blue bars correspond to European Ancestry (EA) samples. Confidence intervals (95%) bars are shown for each estimate. h2=Heritability, SNP=Single Nucleotide Polymorphism, Re-Exp=reexperiencing, MVP= Million Veteran Program, AD=Alcohol Dependence, AUD=Alcohol Use Disorder, Max. Alc.=Maximum Alcohol Intake, AUDIT=Alcohol Use Disorder Identification Test, T=total, P=problems, C=consumption, DPW=Drinks per Week, UKB=UK Biobank, GSCAN= GWAS & Sequencing Consortium of Alcohol and Nicotine Use, PGC=Psychiatric Genomics Consortium.