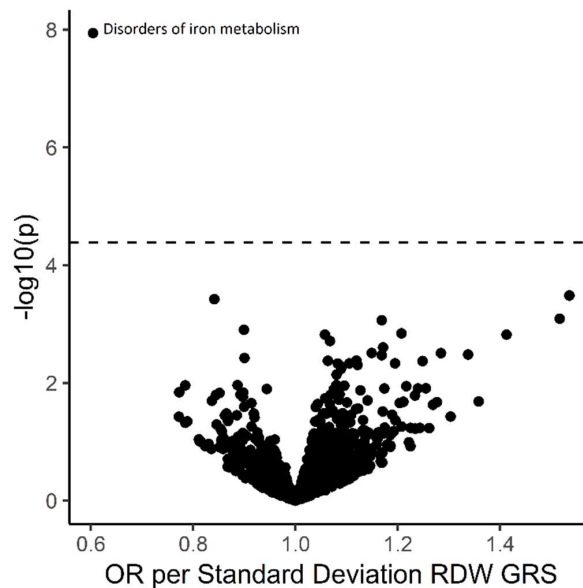


Supplemental Table 2

Validation of each genetic risk score for a cardiovascular disease or cardiovascular disease risk factor against predicted phenotype

Genetic risk score (GRS)	Number of genetic variants	PheWAS Phenotype	OR (per standard deviation GRS)	P
Coronary artery disease	24	“Coronary atherosclerosis”	1.25	2×10^{-23}
Heart failure with reduced ejection fraction	7	“Congestive heart failure (CHF) NOS”	1.16	4×10^{-11}
Peripheral arterial disease	12	“Atherosclerosis of the extremities”	1.24	2×10^{-7}
Venous thromboembolism	20	“Deep vein thrombosis [DVT]”	1.36	4×10^{-18}
Atrial fibrillation	63	“Atrial fibrillation and flutter”	1.26	1×10^{-20}
Systolic blood pressure	127	“Hypertension”	1.12	3×10^{-9}
Diastolic blood pressure	184	“Hypertension”	1.09	1×10^{-6}
Low-density lipoprotein	64	“Hypercholesterolemia”	1.24	3×10^{-20}
High-density lipoprotein	76	“Hypercholesterolemia”	0.9	2×10^{-5}
Triglycerides	75	“Hyperglyceridemia”	1.32	6×10^{-5}
Body mass index	1027	“Obesity”	1.42	1×10^{-59}
Type 2 diabetes mellitus	92	“Type 2 diabetes”	1.38	2×10^{-63}
Estimated glomerular filtration rate	65	“Chronic renal failure [CKD]”	0.87	1×10^{-9}



Supplemental Figure S1: Full phenome-wide association studies of RDW GRS revealed no significant non-hematologic phenotype associations with RDW GRS. Dashed line represents Bonferroni corrected P value threshold determined by logistic regression.