

Total participants  
genotyped that passed  
QC  
N = 43,060

Total participants with  
phenotype data (age,  
sex, platelet traits)  
N = 35,994

Participants with both  
genotype and phenotype  
data  
N = 33,462

Participants with genetic  
risk score and remained  
consented  
N = 33,458

Participants with body  
mass index measured  
N = 33,388

Study population for  
linear regression and  
Mendelian randomization  
N = 33,388

