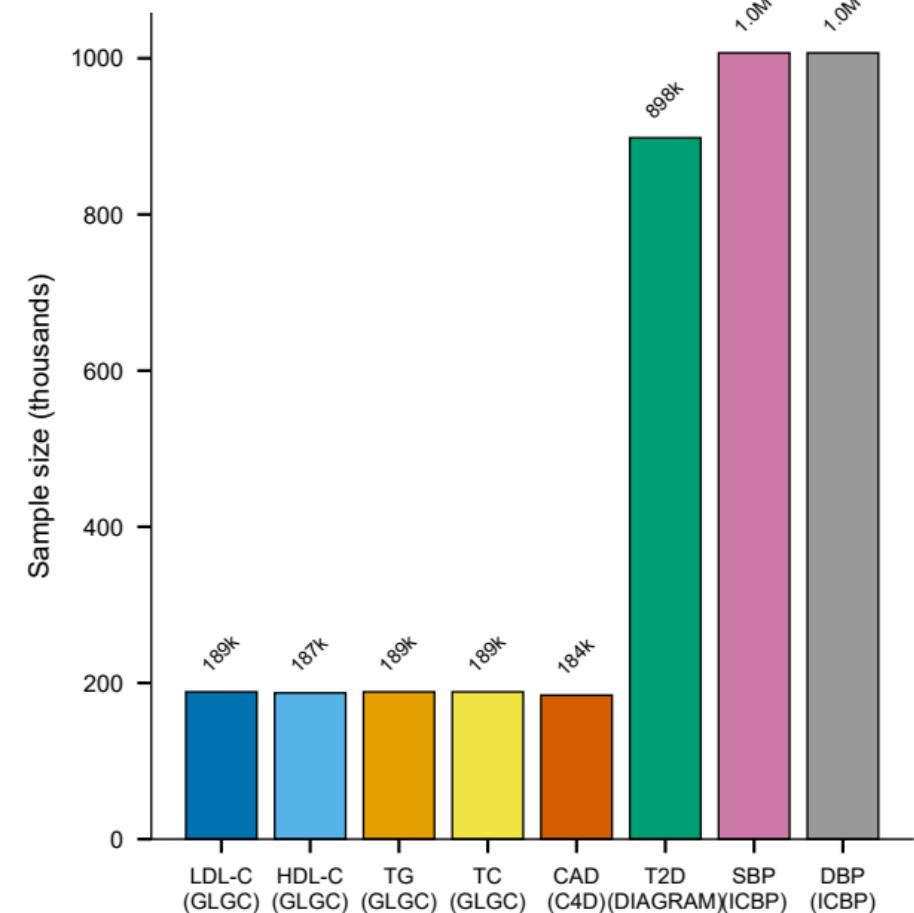


**a**

GWAS datasets analyzed

**b**

Loci per trait

