## Imputing gene expression

## **Testing associations**

## eQTLGen Consortium



31,684 blood examples



cis-SNPs

**1000 Genomes Project** 



LD reference (R)

**GTEx Portal** 



Gene expression reference

Constructing

Tuning & validation

Individual-level GWAS data

Phenotype ~ Imputed expression

Summary-level GWAS data

$$Z^* pprox rac{w^T Z}{\sqrt{w^T R w}}$$

where **Z** is the SNP-trait standardized effect.

cis-eQTL effect sizes (w)