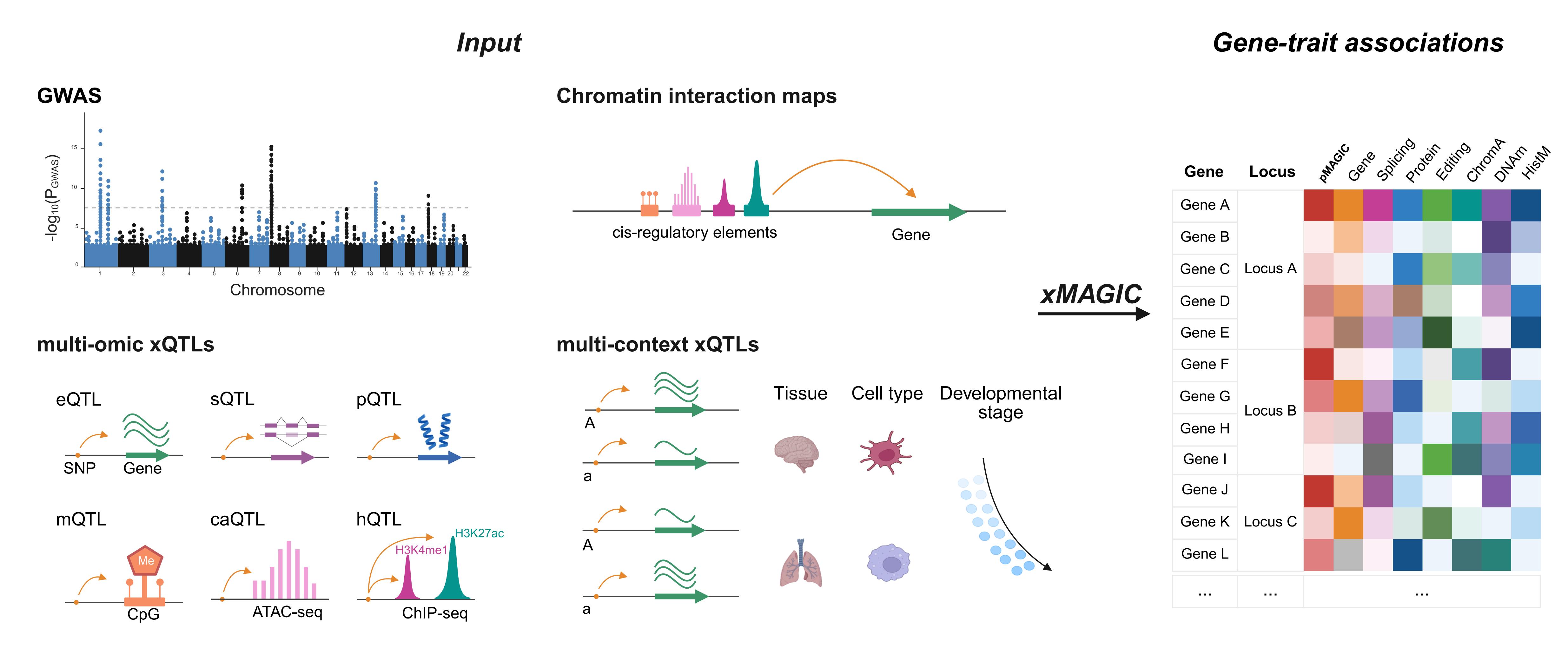
a Online analysis



b Database

25 diseases 20 quantitative traits **Associates** controls 🚱 cases 🚱 Overall P TPI Layer-specific P Dataset-specific P/ 19K Genes **45 Traits** Omics layers Diseases "ve oatabase contex **GWAS** data for Contexts Quantitative / 45 traits Loops 11 mQTL 286 eQTL SNP Gene **xMAGIC** 428 xQTL datasets Portal Search **Wisualization** By trait, gene GWAS Advanced 49 edQTL xQTLs 56 sQTL search **Download** Loops H3K27ac Gene H3K4me1 Associations by trait Significant xQTLs 5 caQTL 14 pQTL 7 hQTL

c Locus visualization

