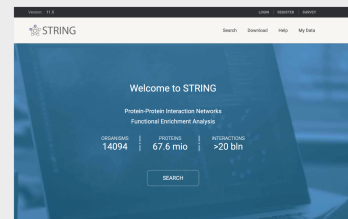
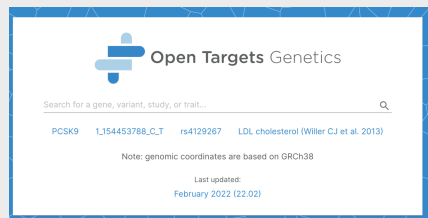
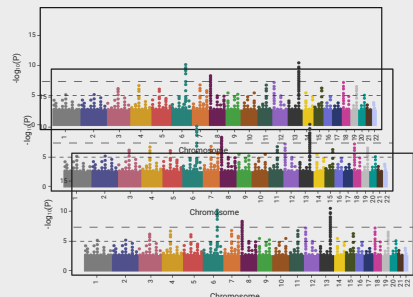
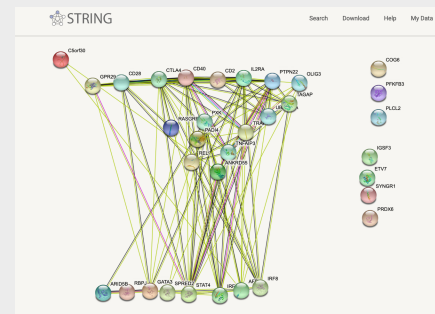
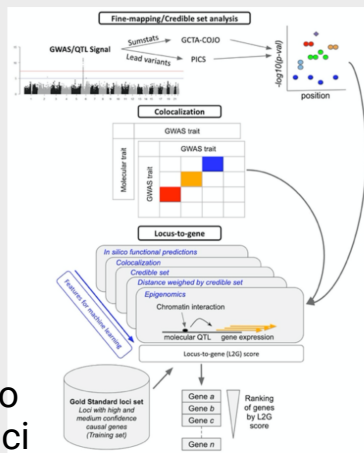


2. Annotate with predicted Causal Gene from Open Targets Genetics L2G

3. Get the predicted protein-protein interaction network from STRING using putative Causal Gene List

1. Gather Regions from GWAS catalog



Overview of Locus-to-gene model to prioritise genes across all GWAS loci

4. Load STRING networks into cytoscape to arrange and merge