Guide to the R markdown file

Attie\_eQTL\_paper\_circlePlots\_and\_gwasEnrichment.Rmd contains the generation of circle plots and enrichment analysis, including:

* counting number of diabetes GWAS SNPs in 1Mbp windows across the genome,
* getting nearest mouse gene to clinical and module QTLs,
* mapping mouse genes to orthologous human genes,
* enrichment analysis of QTLs in human GWAS diabetes SNPs.