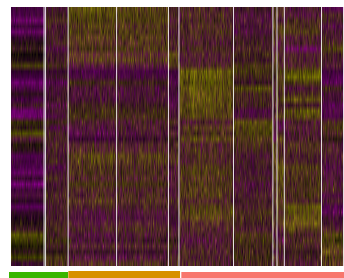


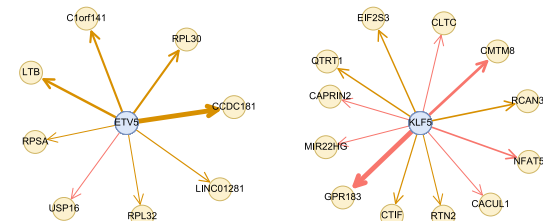
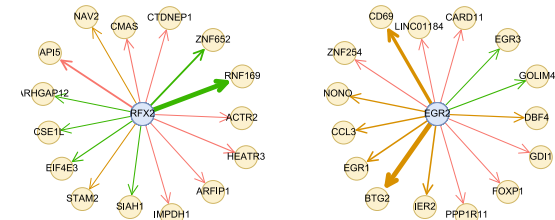
scRNA-seq data



GRNBoost2  
+  
Motif enrichment

TF-target networks

Quantile  
normalization



Differential analysis of  
link importances between  
groups  
considering only  
simultaneously activated TFs