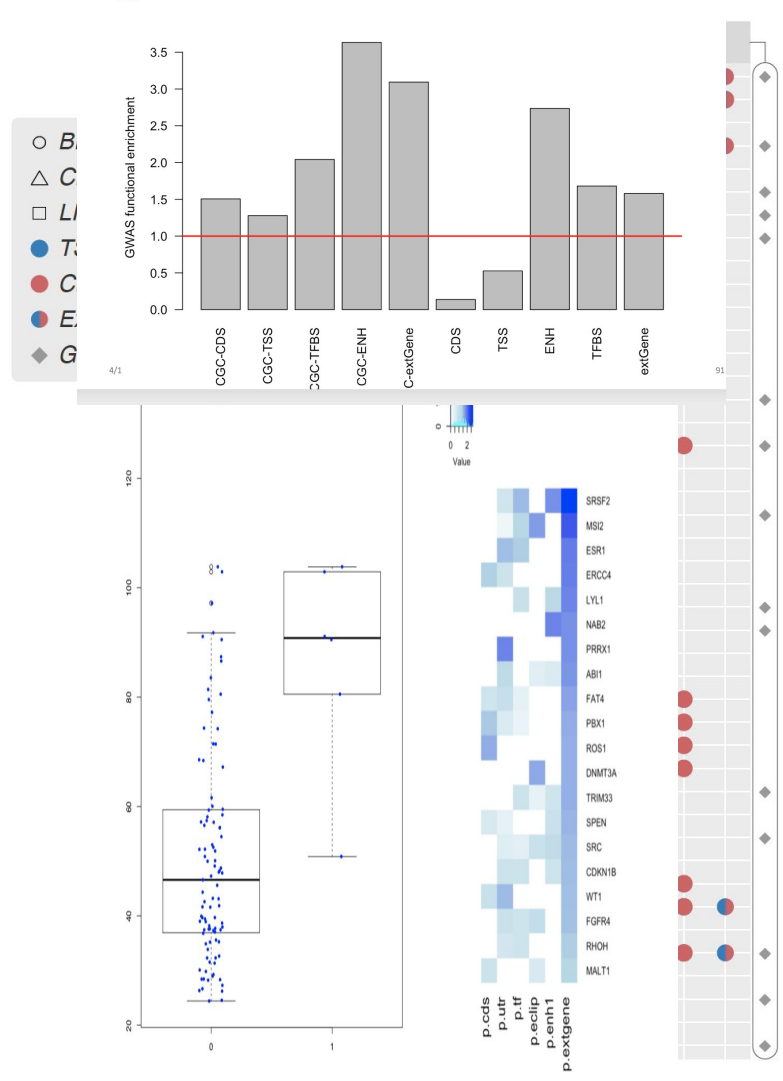
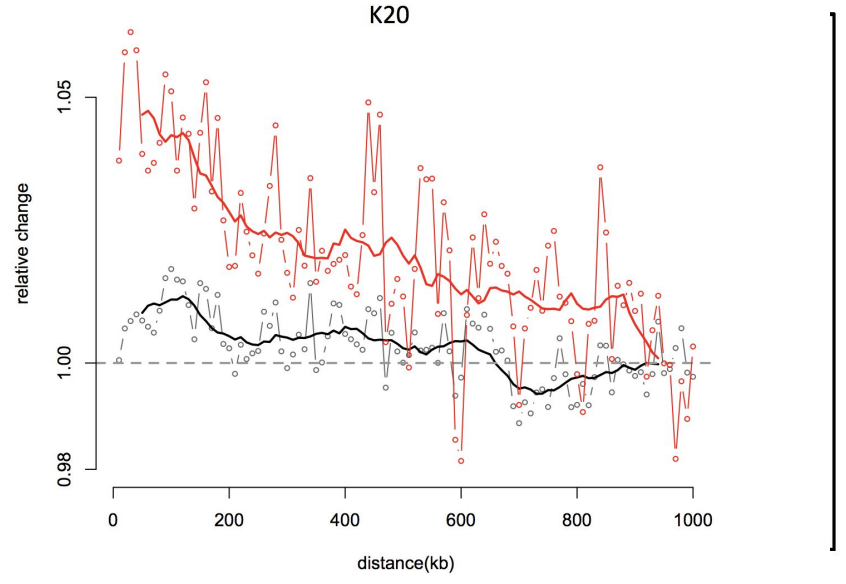
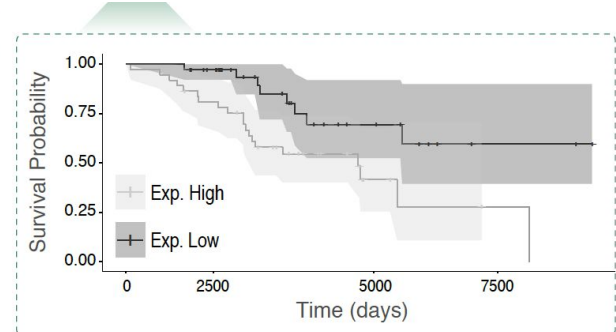


B

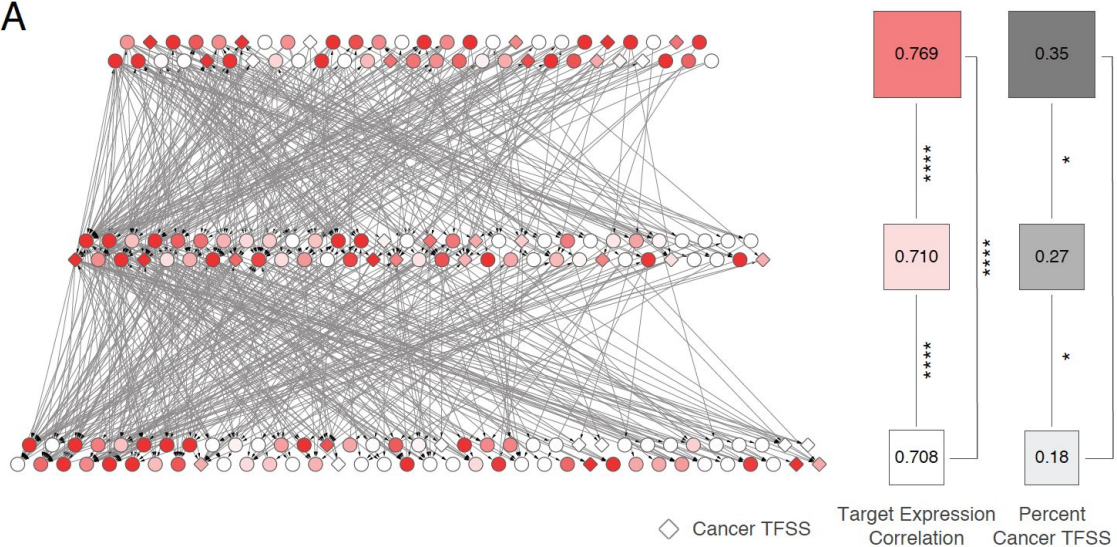


E

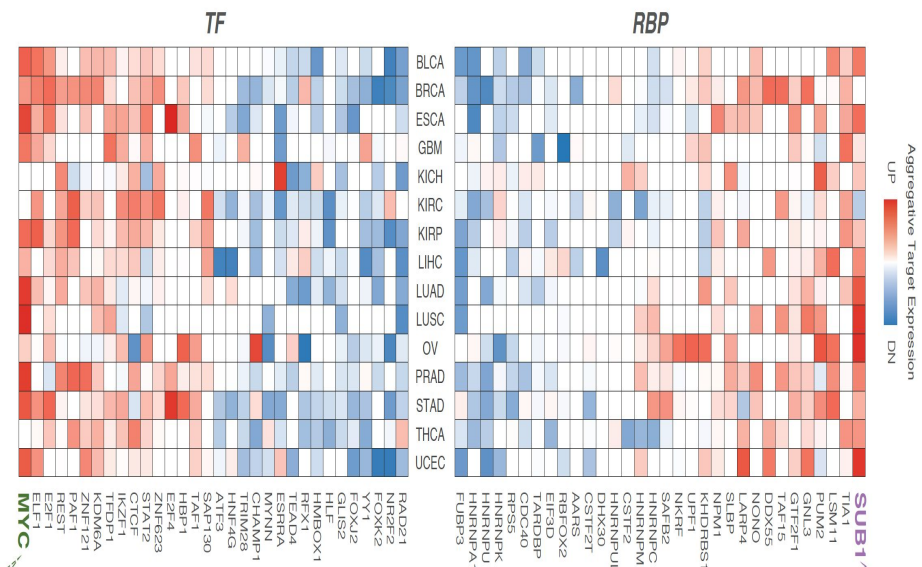


More KD
More Biolog

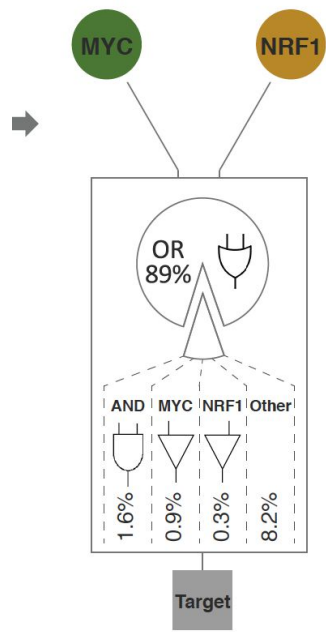
A



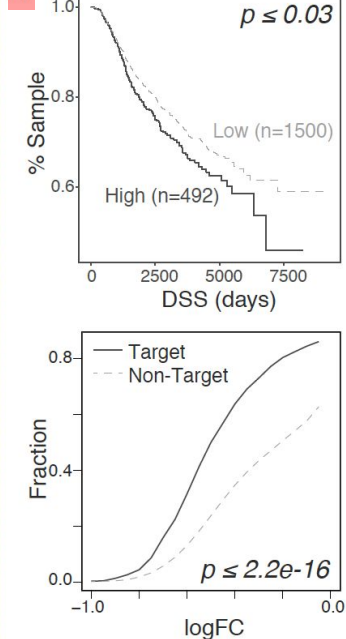
A



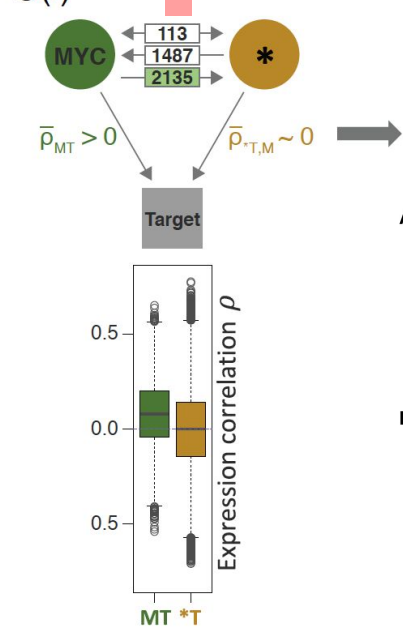
C(iii)



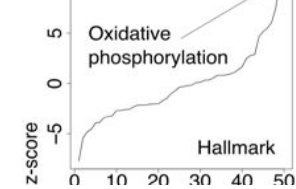
B



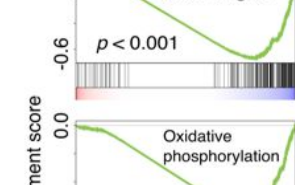
C(i)



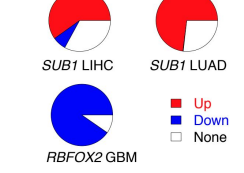
a



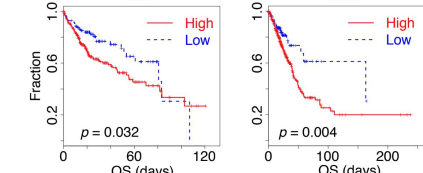
b



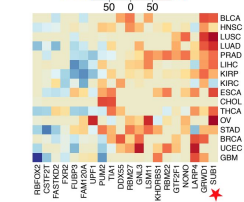
A



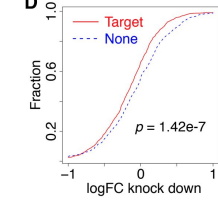
C



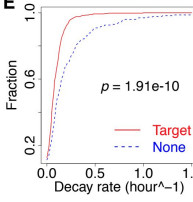
B



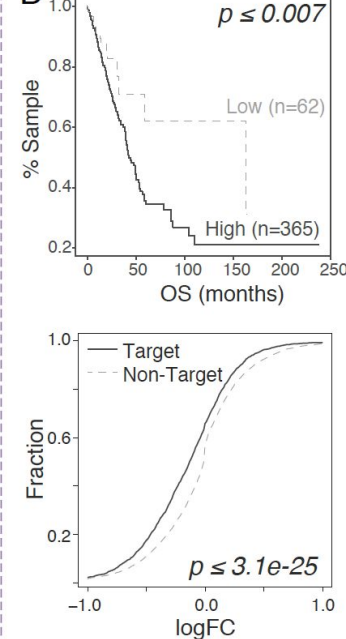
D



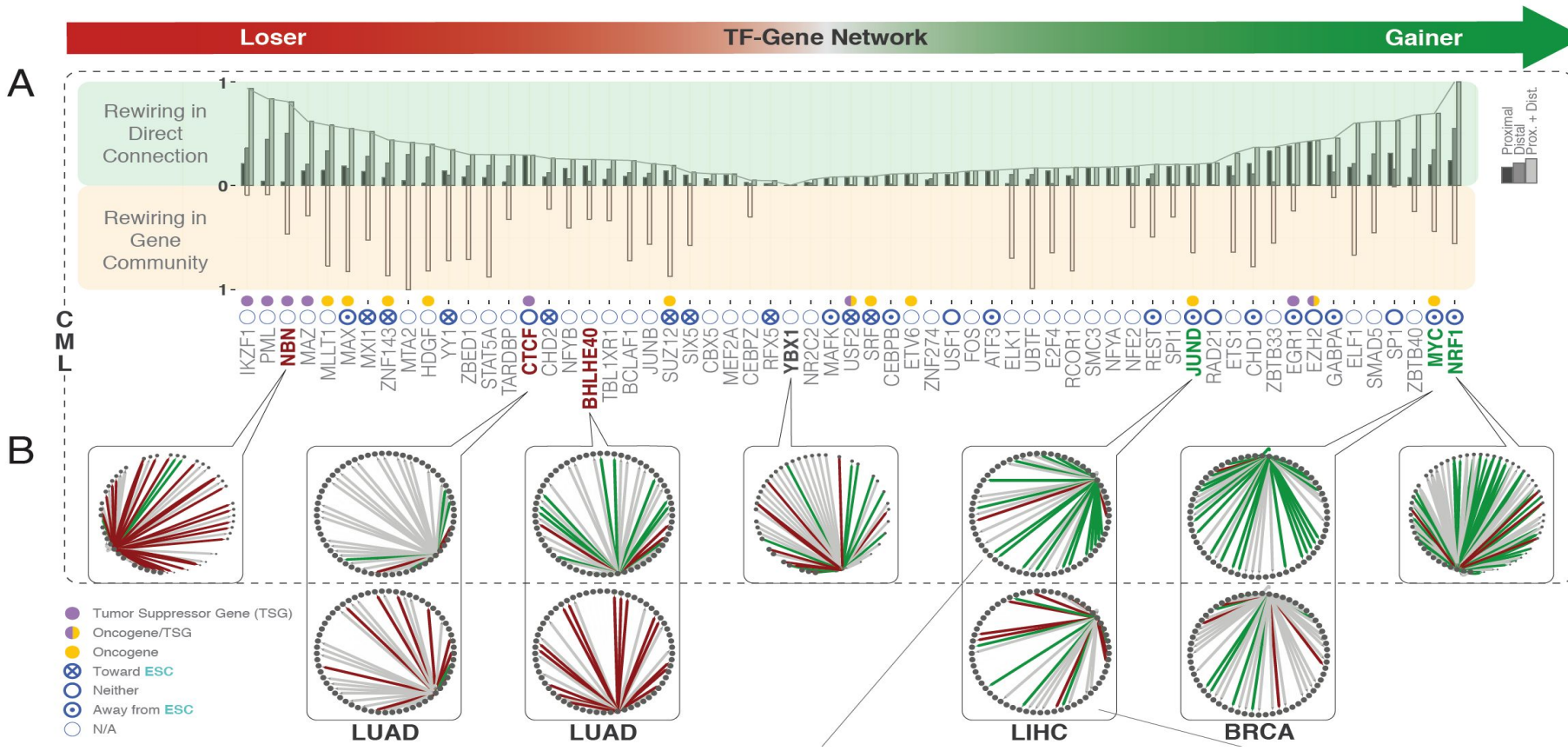
E



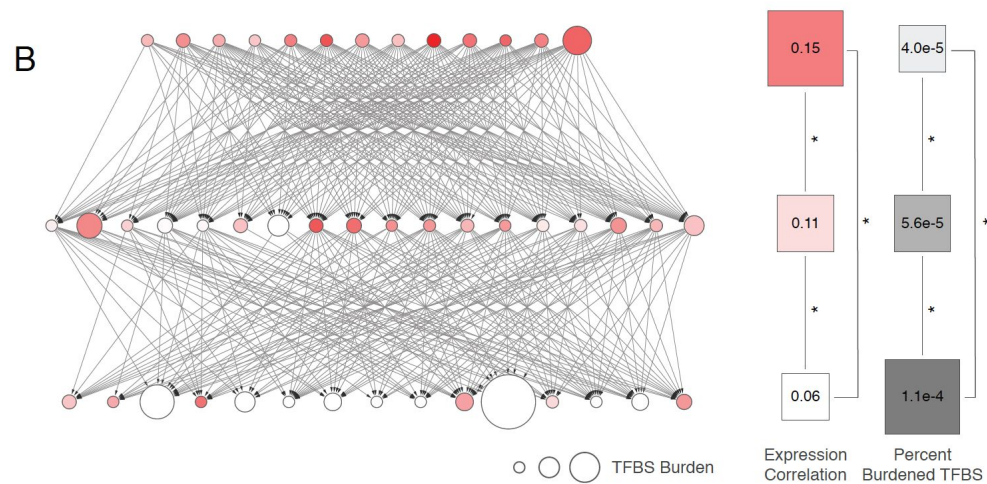
D



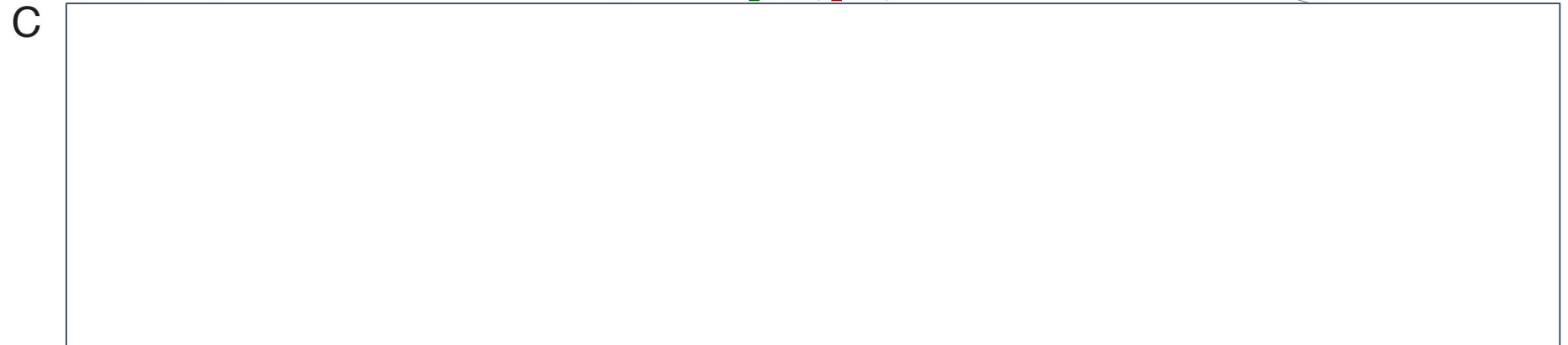
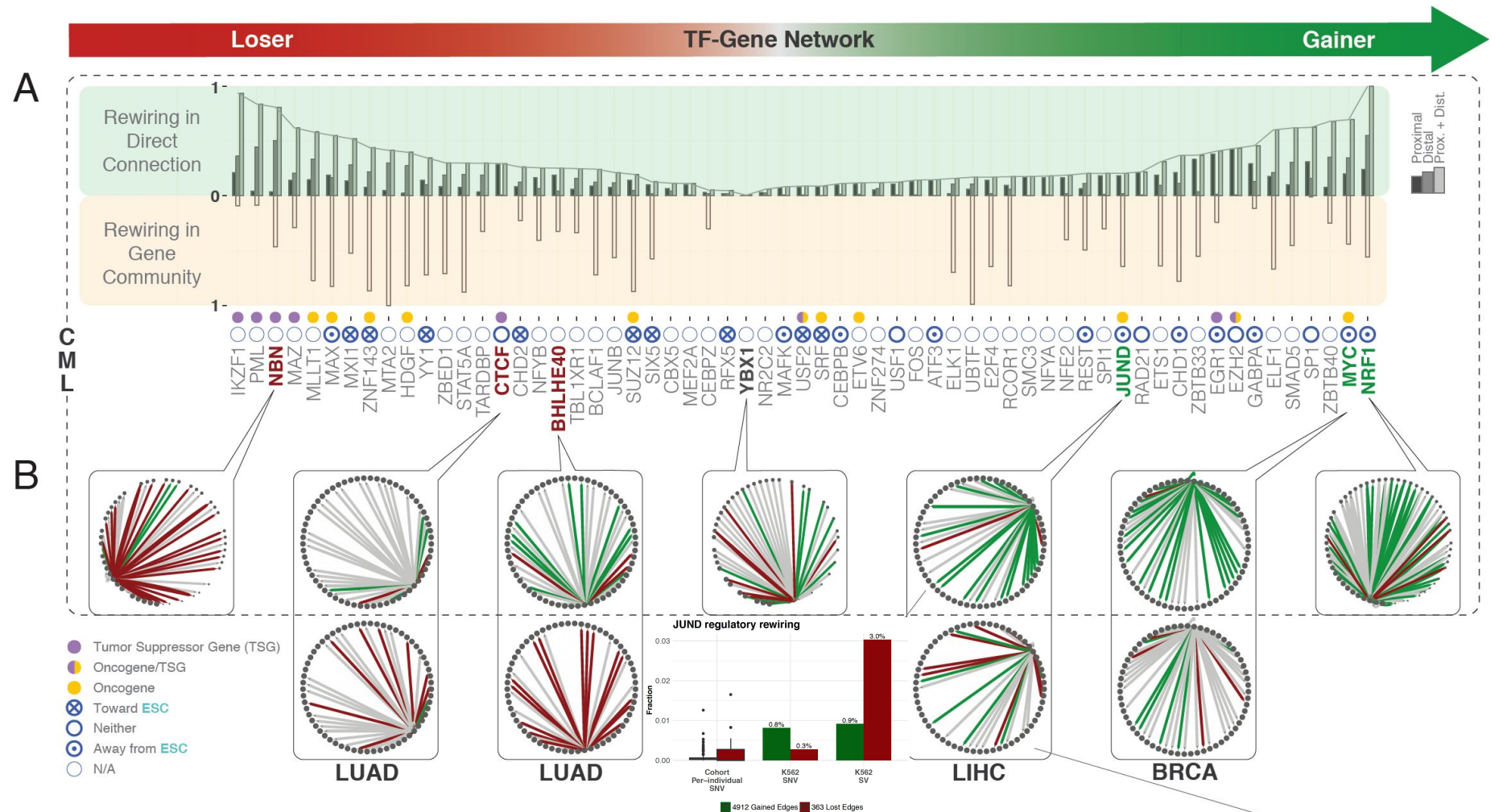
Inference of RNA binding proteins that drive tumor specific expression patterns. Based on



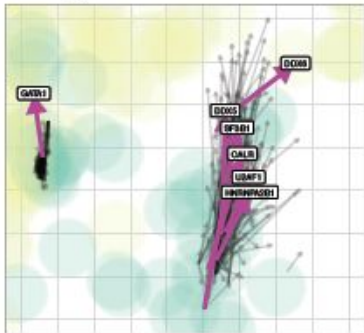
Order shuffling results on cell lines?



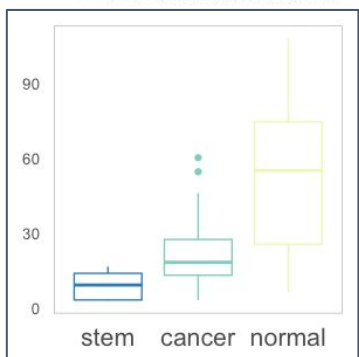
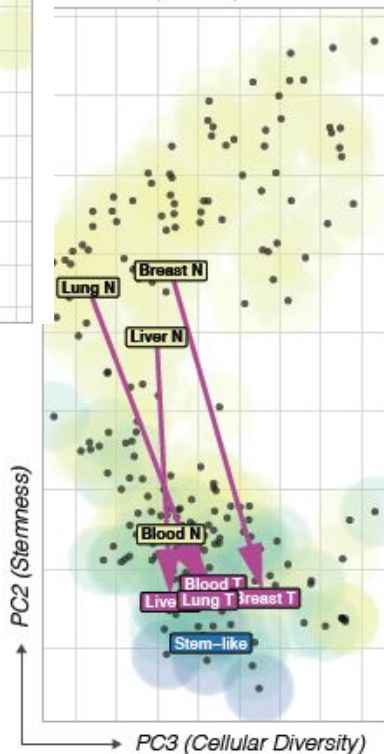
KD for some bars in the examples
 H1: 1~2 panels
 SV added in D



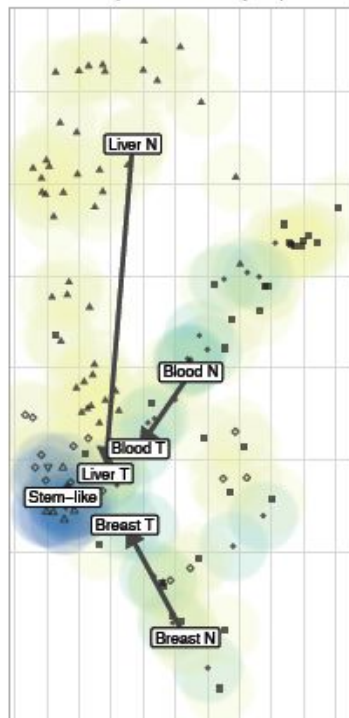
Oncogene KD



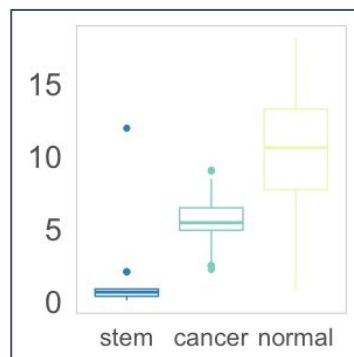
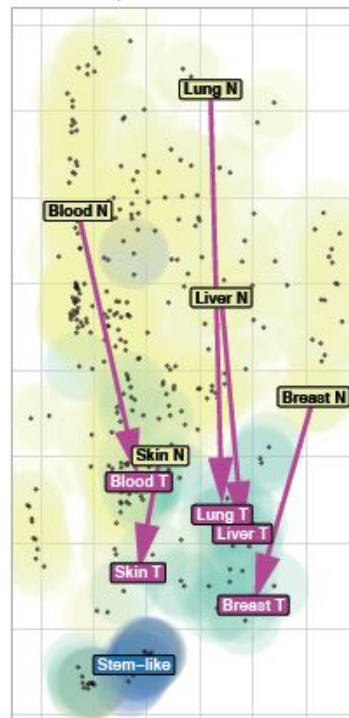
Prox Net (CTCF)



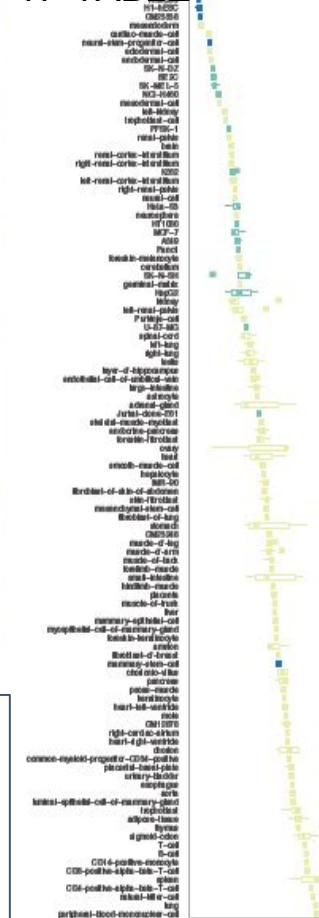
Dist Net (ccRE hotspot)



Gene Expression

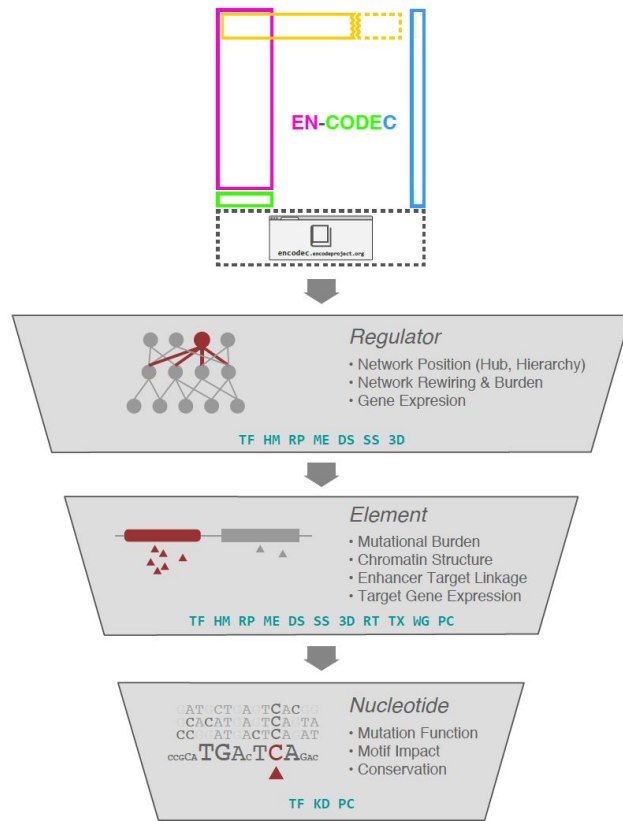


TF TABLE

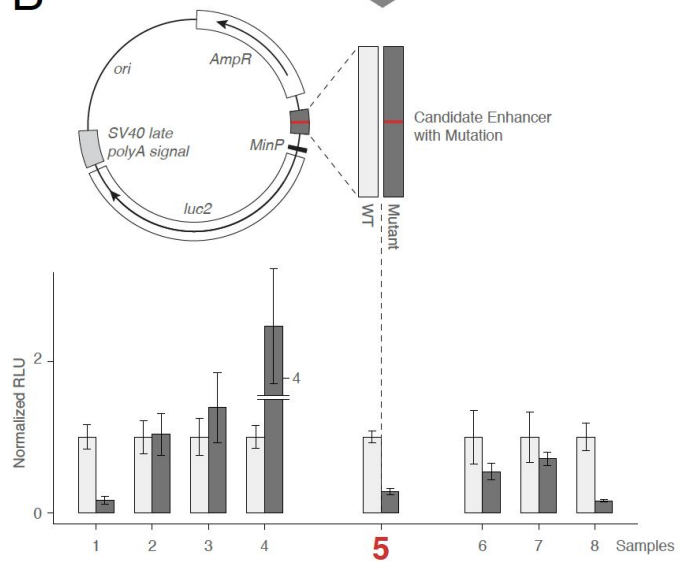


L2 distance to Stem-like cluster

A



B



C

