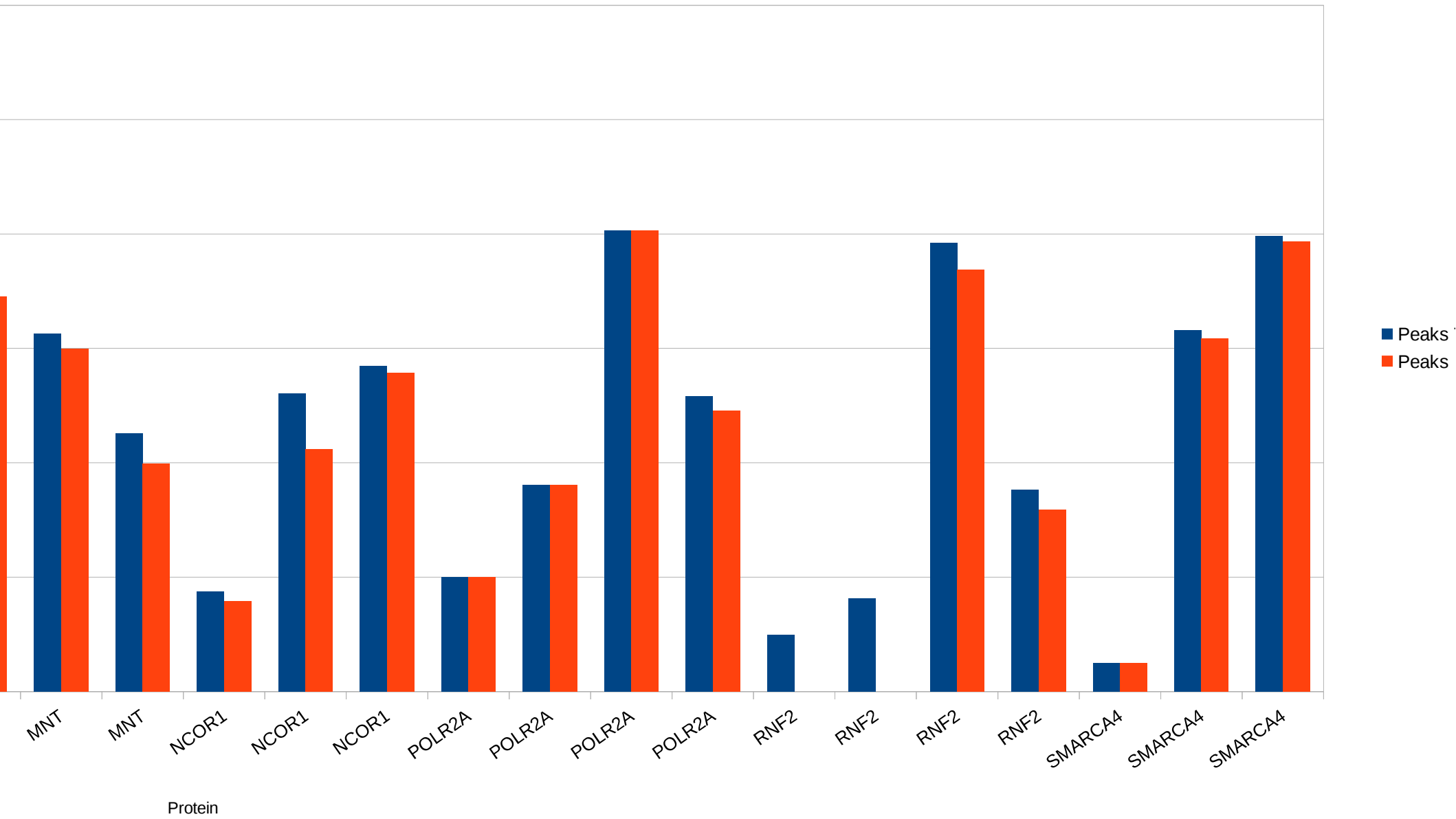


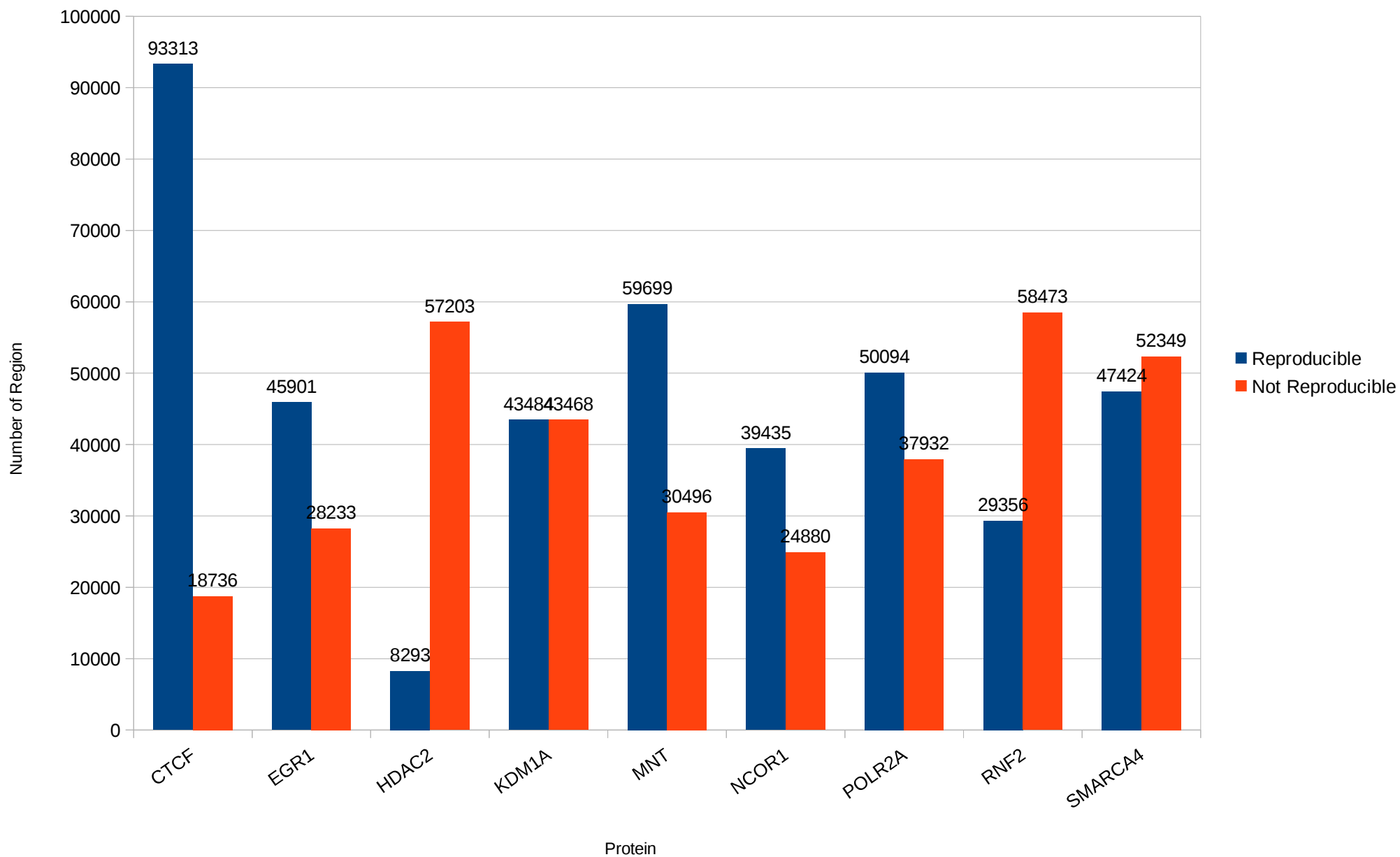
Peaks K562 Cell Type

Before and After FDR <= 0.01 selection

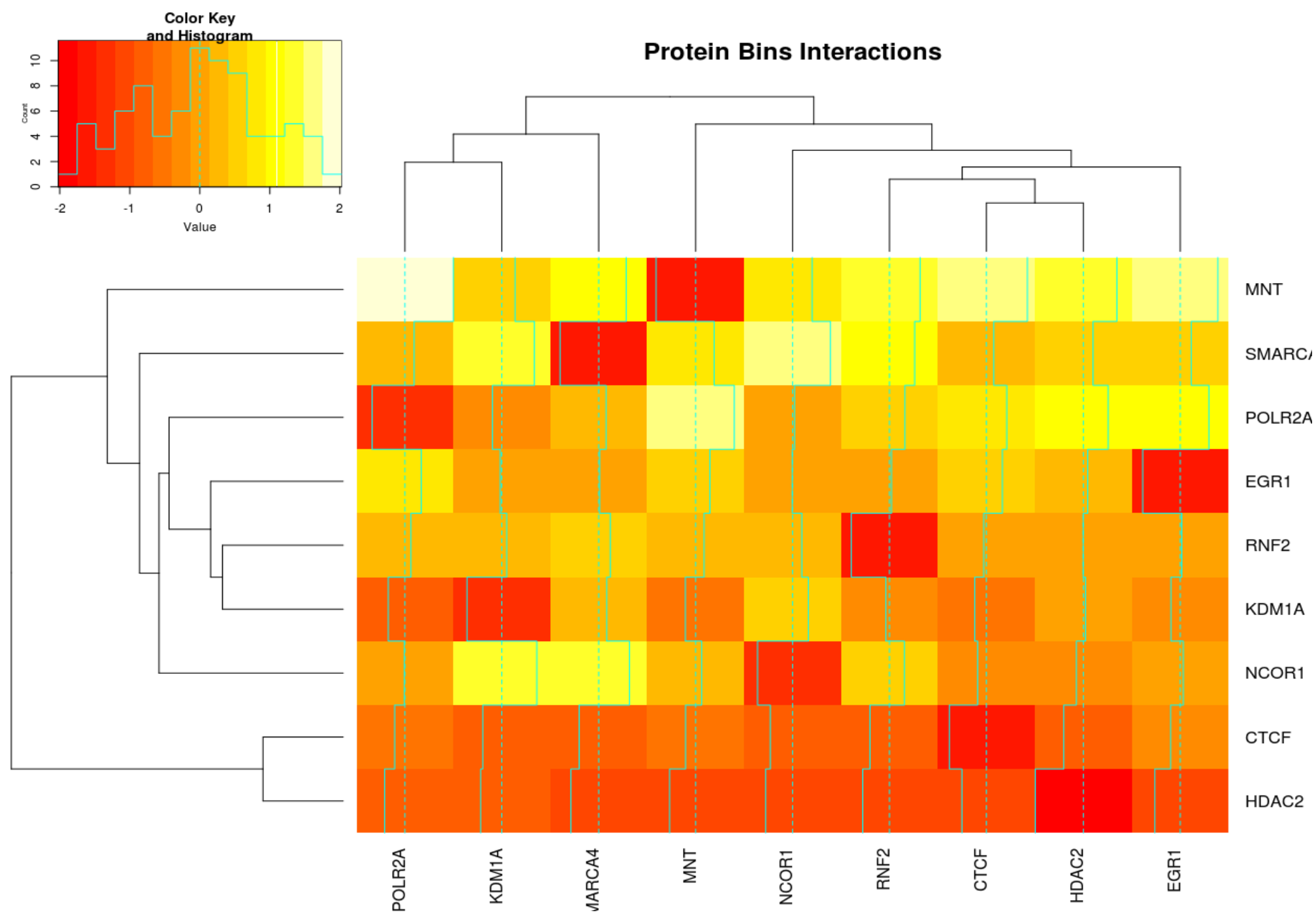


Reproducible and Not Reproducible Regions in bins of 400 base pairs

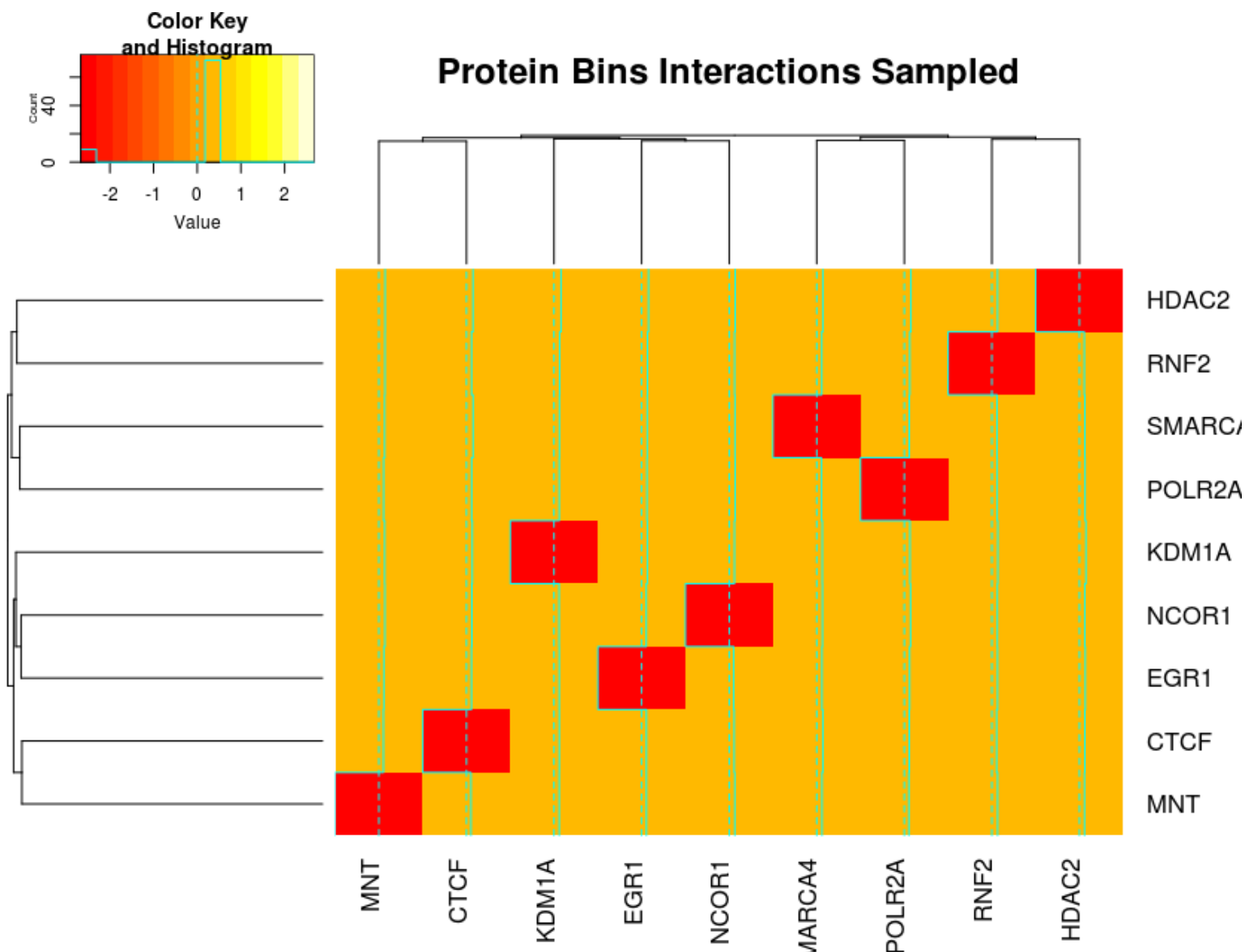
Cell Type K562



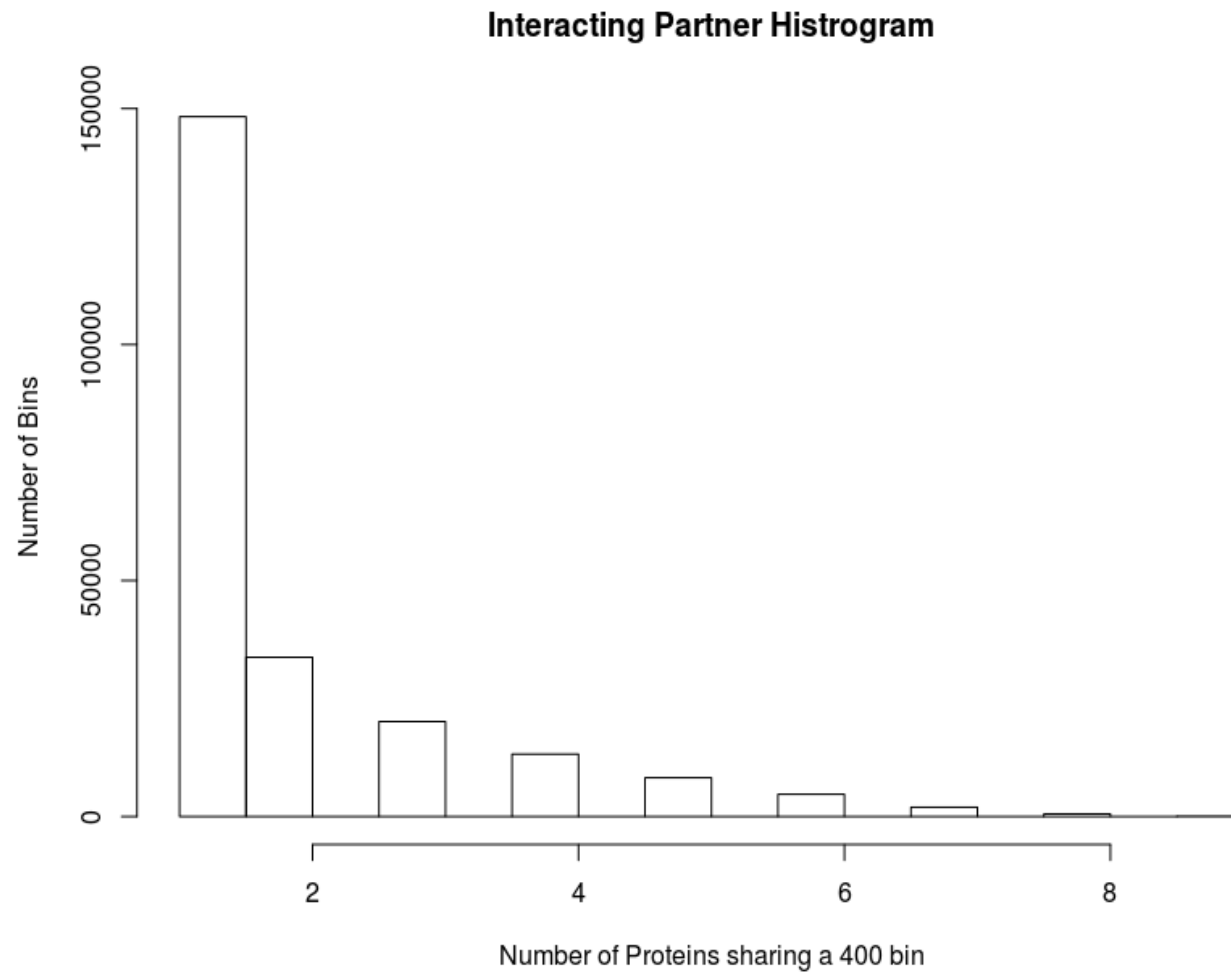
Heat-map of Protein/bins interaction (scaled)



Heat-map of Protein/bins interaction sampled (scaled)

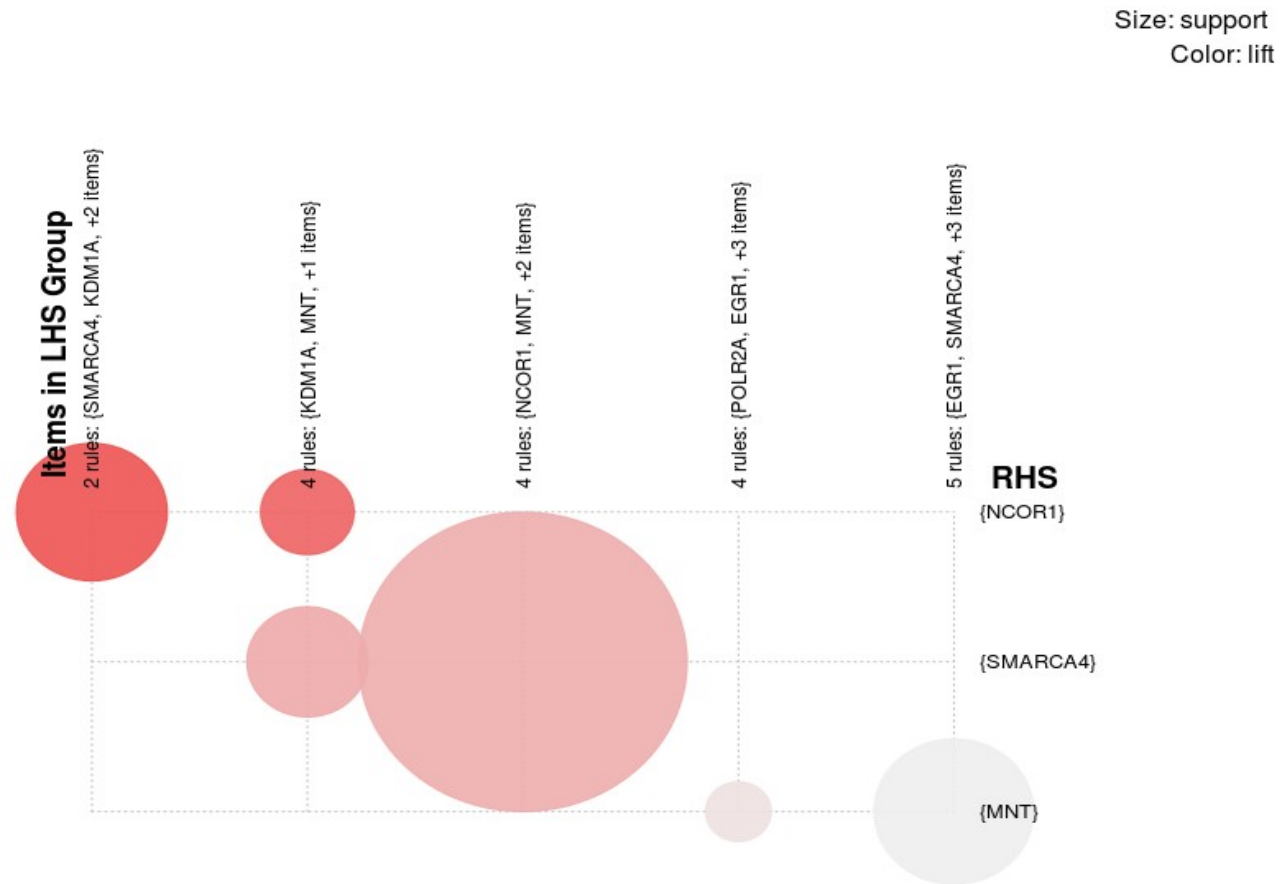


Histogram of Protein sharing the same bin



Top 19 Rules generated

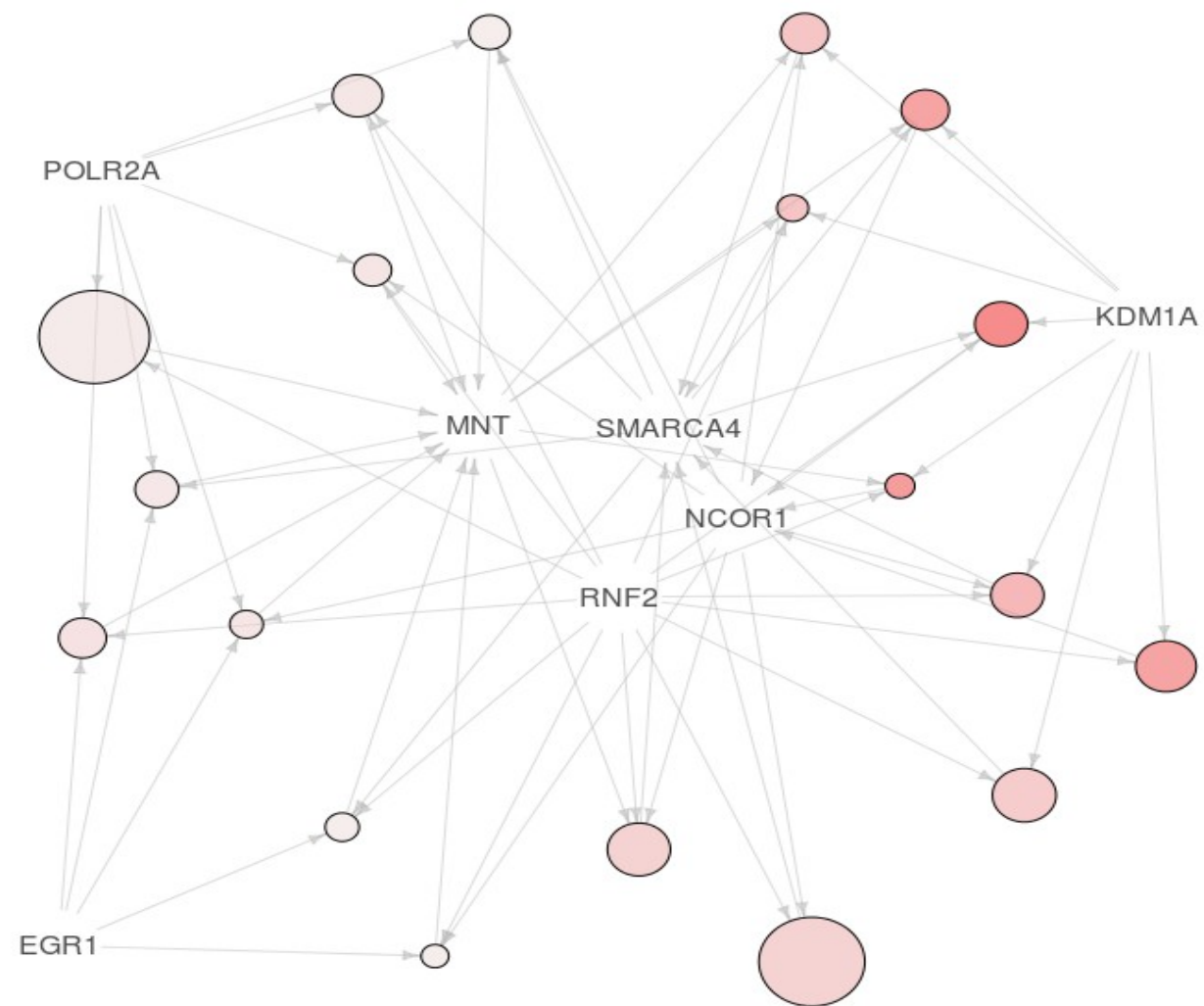
Grouped Matrix for 19 Rules



Support > 0.02
Confidence > 0.80

Graph for 19 rules

size: support (0.02 - 0.05)
color: lift (3.109 - 5.184)



K562 Methylome

