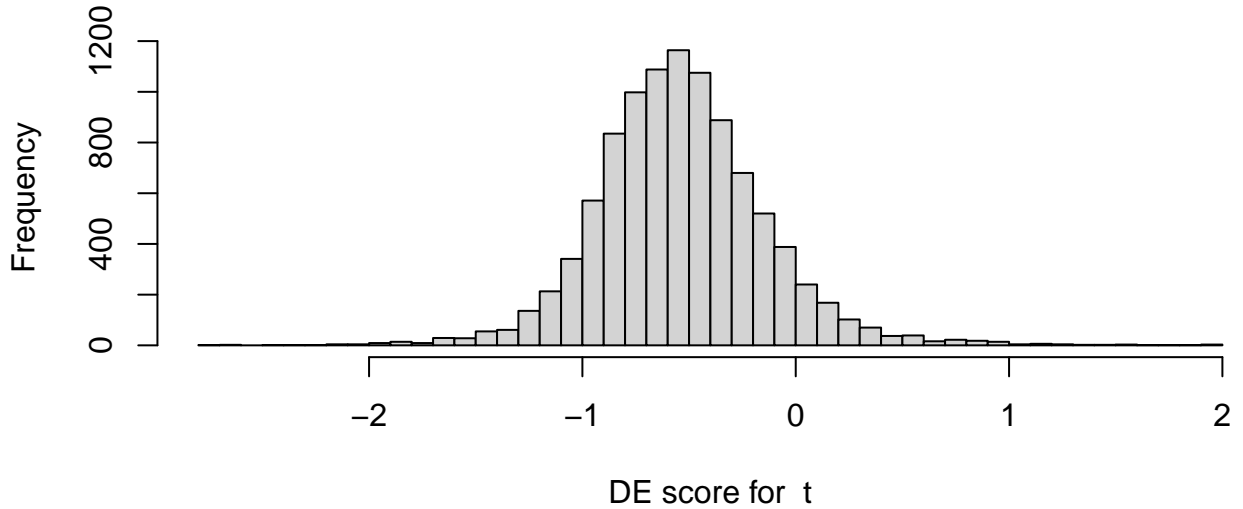
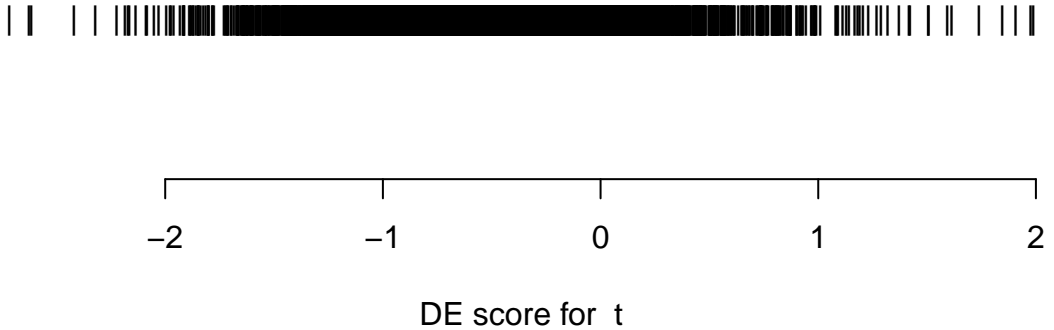


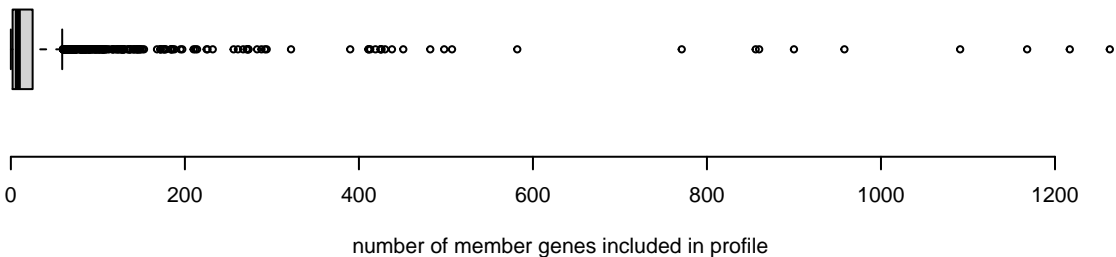
Distribution of DE scores



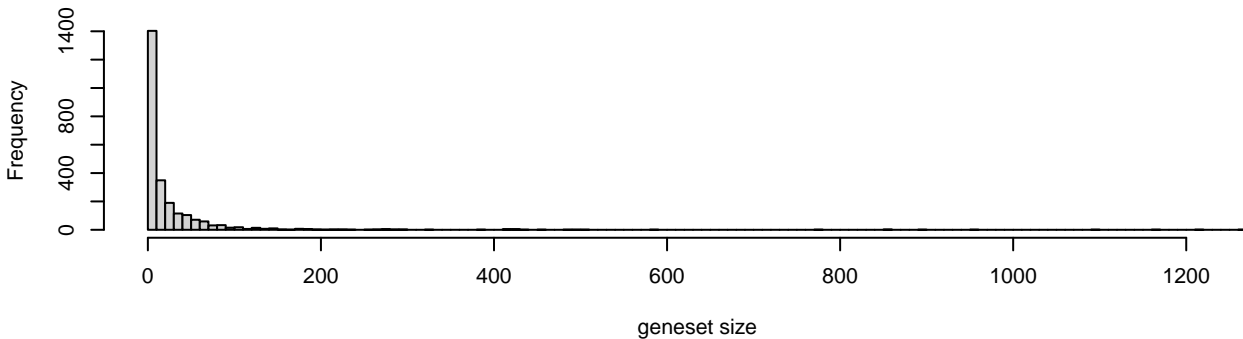
9869 genes in total, 753 trending up-regulated, 9116 trending down-regulated



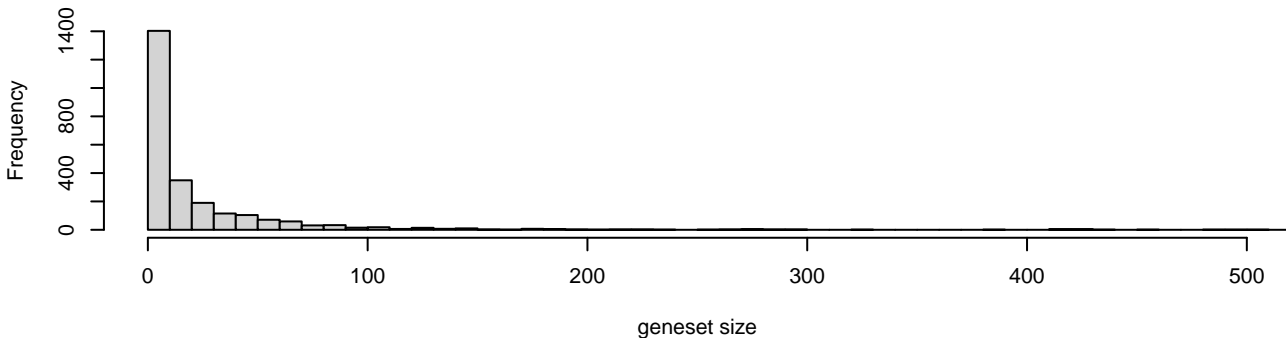
Gene set size



Histogram of geneset size

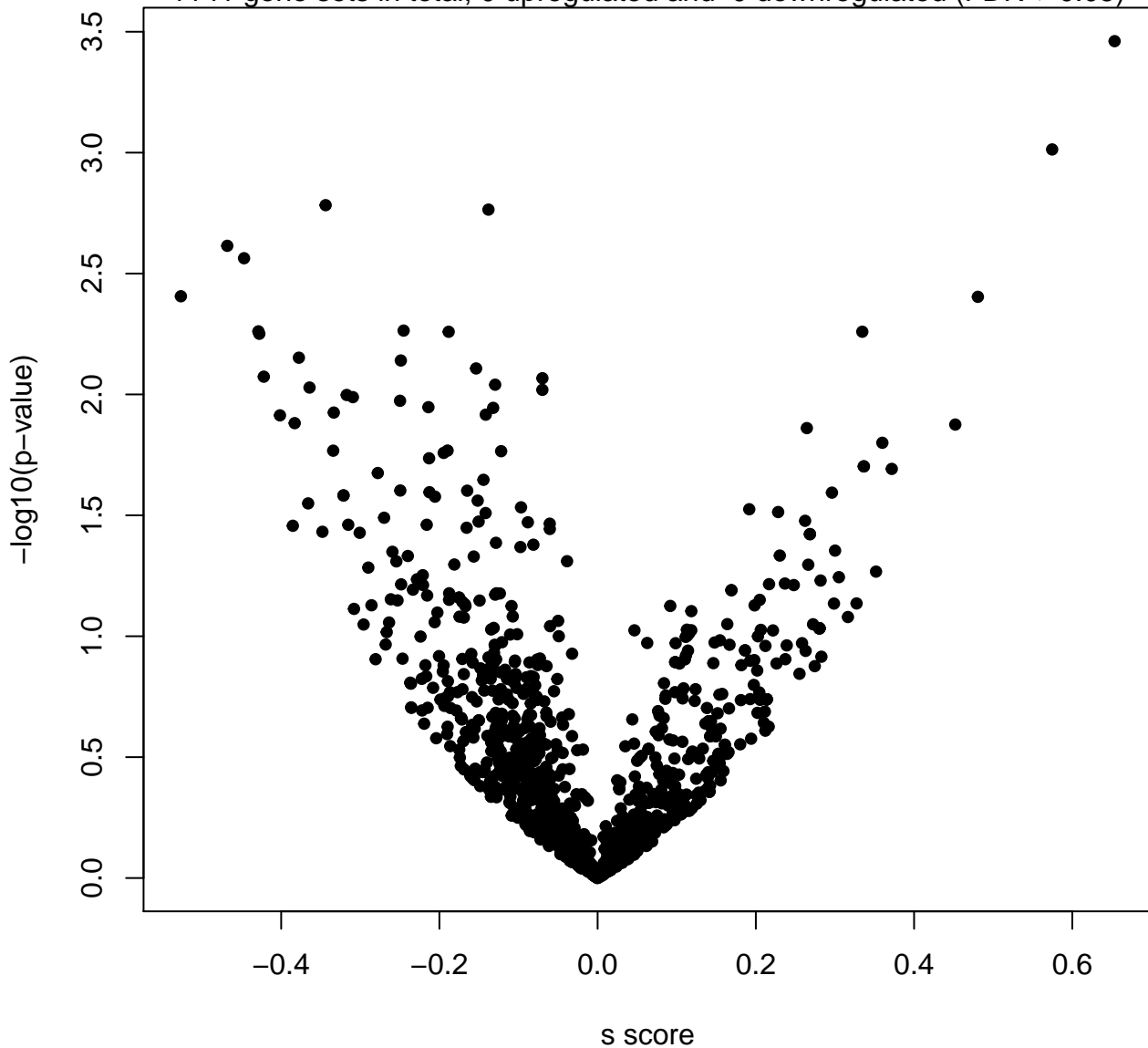


Trimmed histogram of geneset size



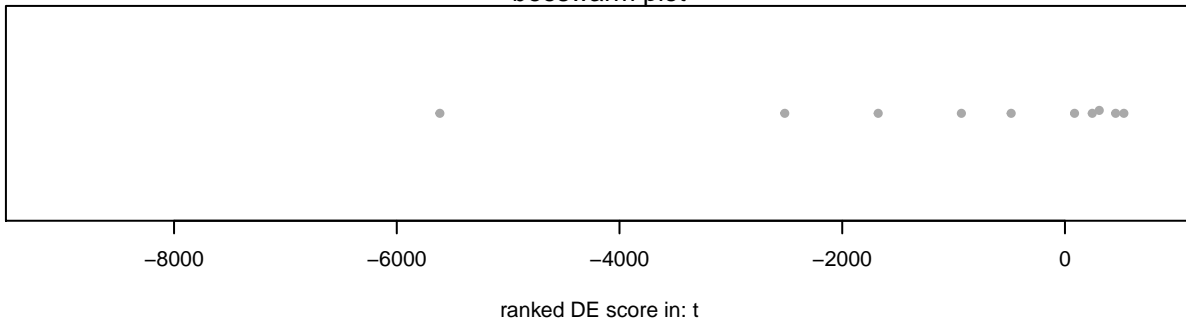
volcano plot of gene set enrichments

1141 gene sets in total, 0 upregulated and 0 downregulated (FDR<=0.05)

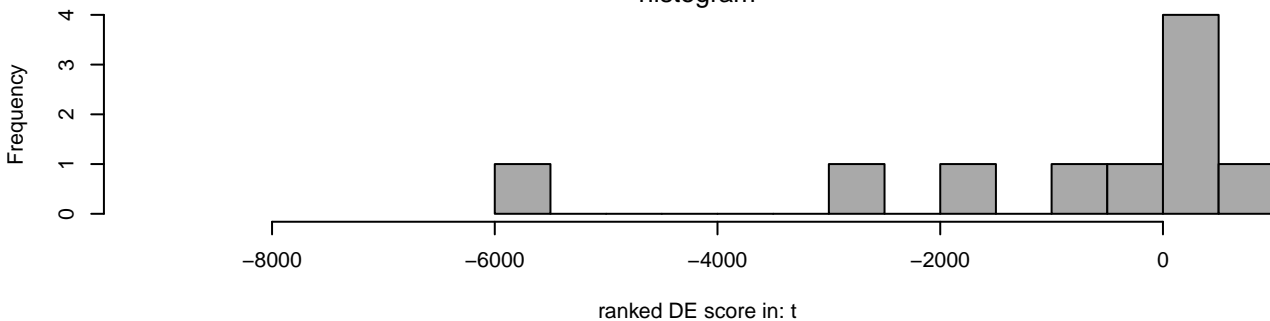


Assembly of collagen fibrils and other multimeric structures

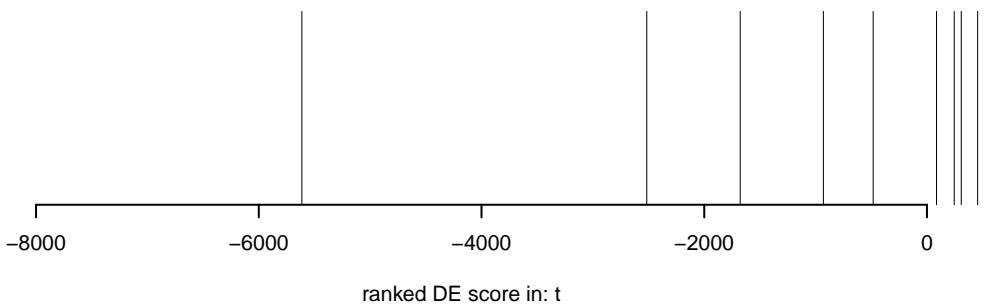
beeswarm plot



histogram

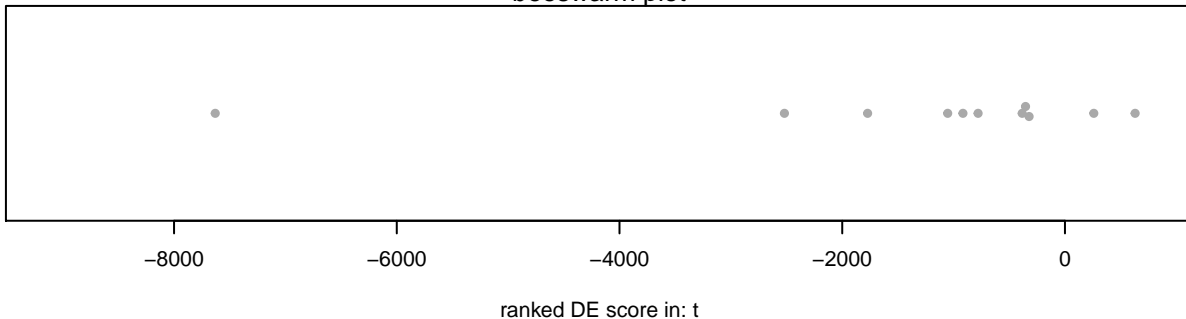


rugplot

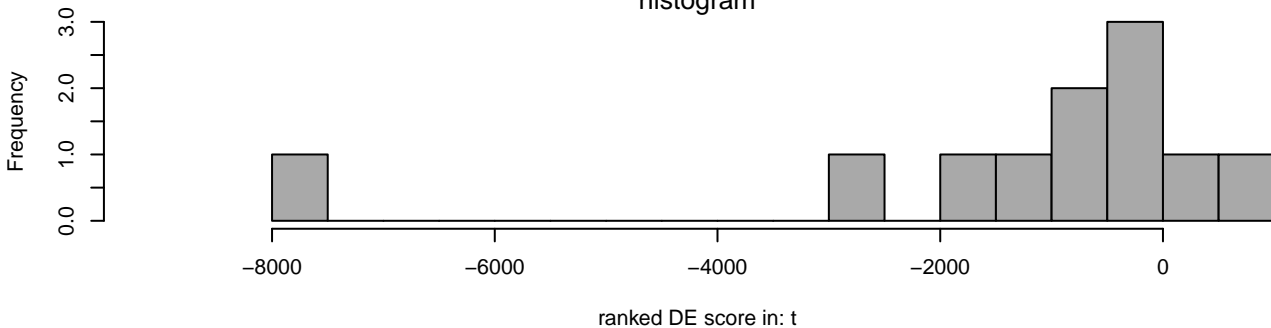


Negative regulation of NMDA receptor-mediated neuronal transmission

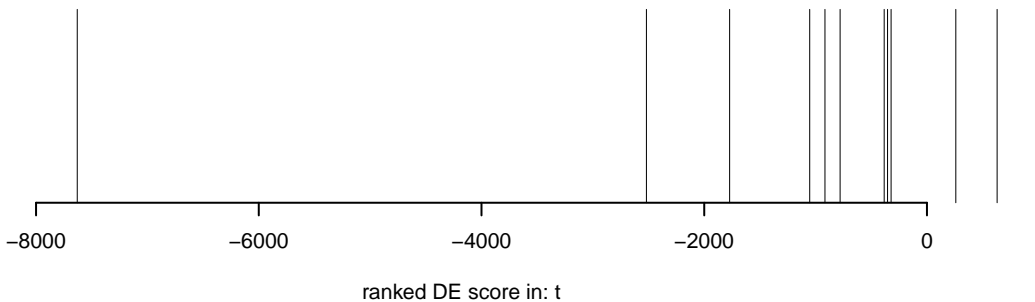
beeswarm plot



histogram

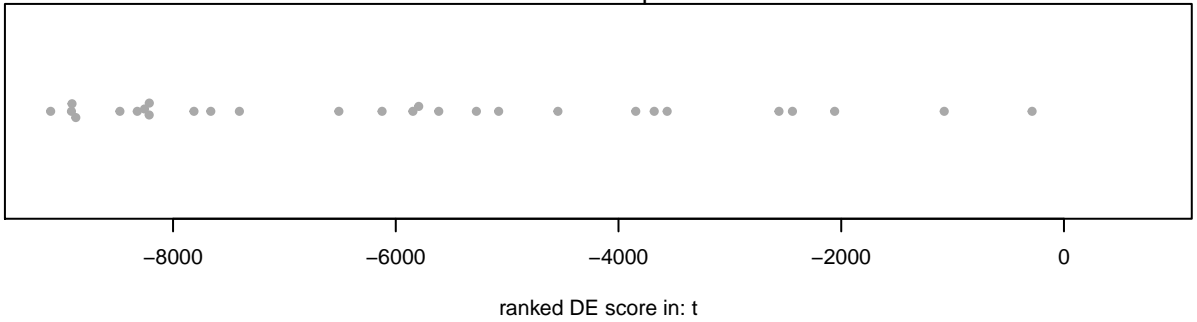


rugplot

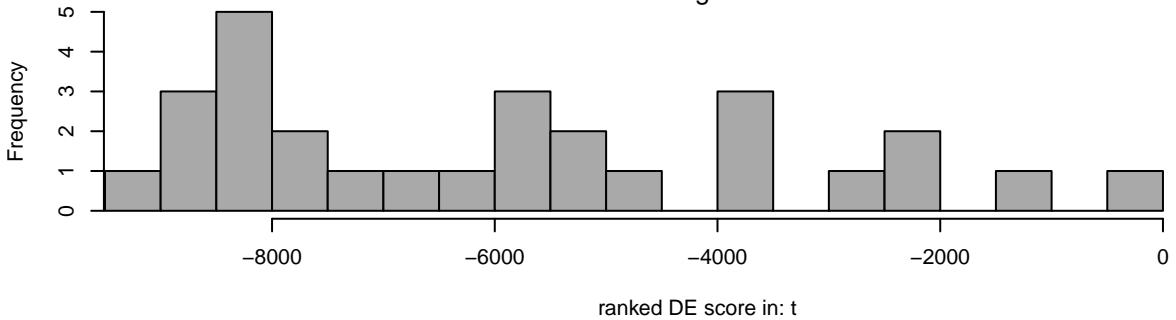


mRNA 3'-end processing

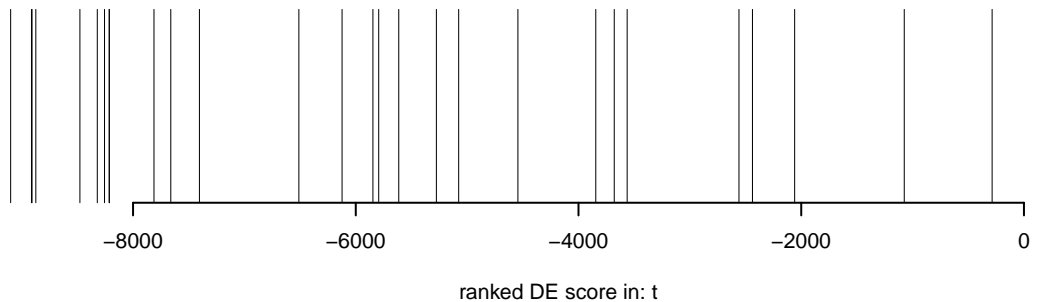
beeswarm plot



histogram

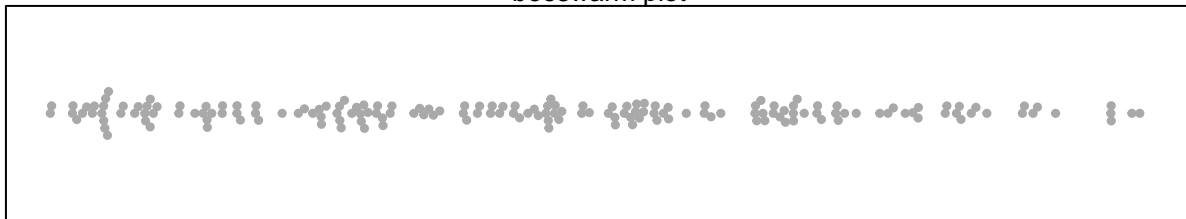


rugplot



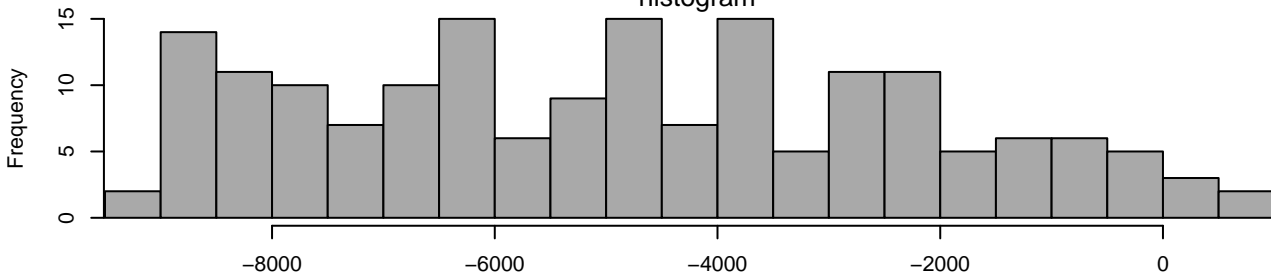
Processing of Capped Intron-Containing Pre-mRNA

beeswarm plot



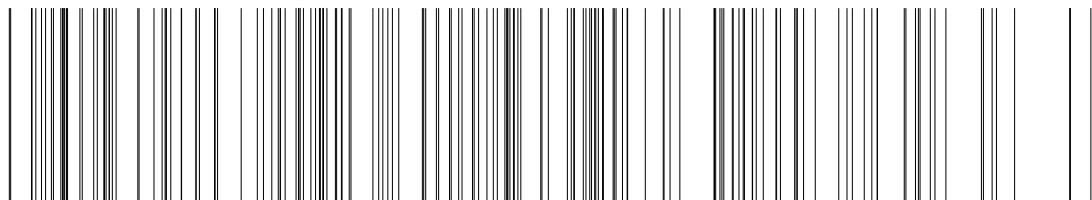
ranked DE score in: t

histogram



ranked DE score in: t

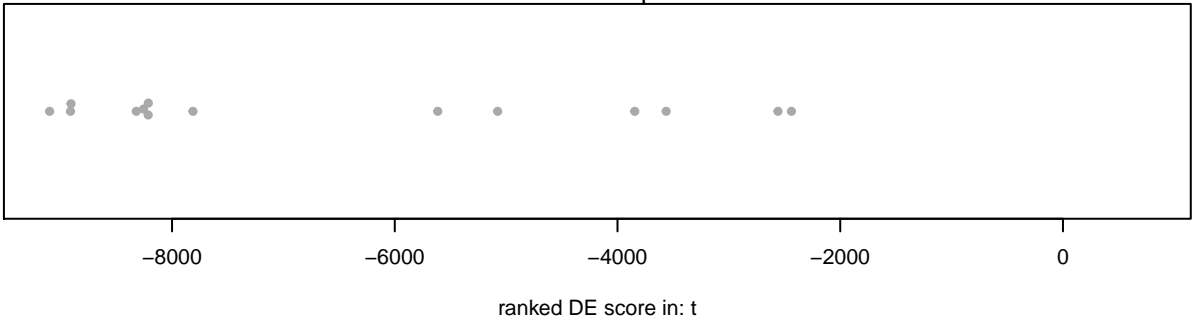
rugplot



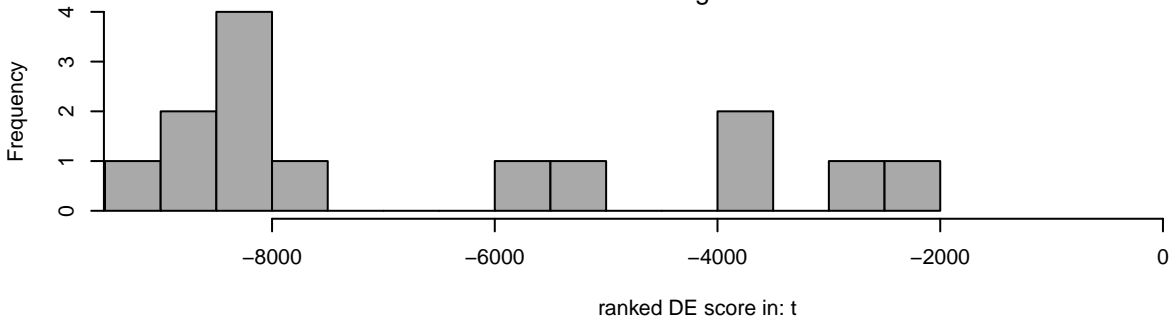
ranked DE score in: t

Processing of Intronless Pre-mRNAs

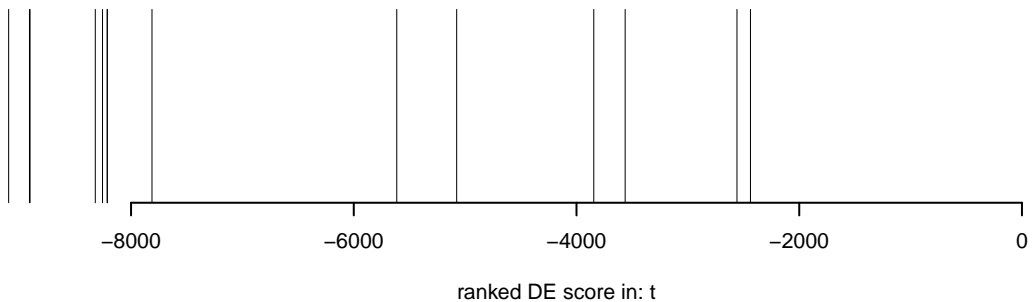
beeswarm plot



histogram

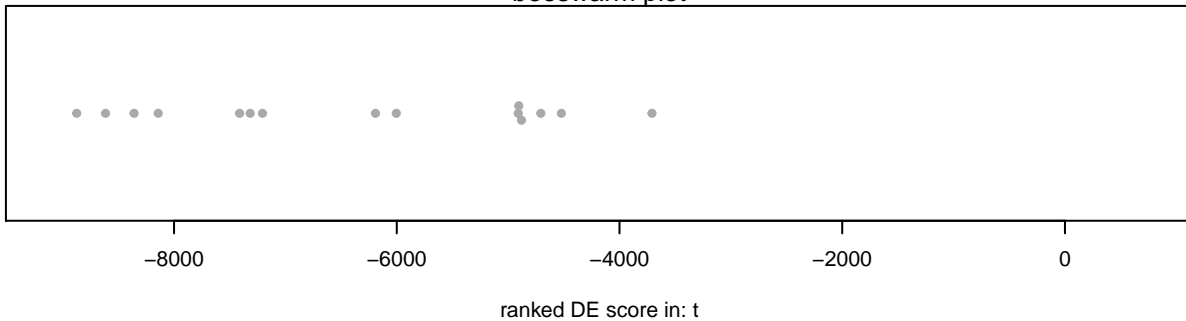


rugplot

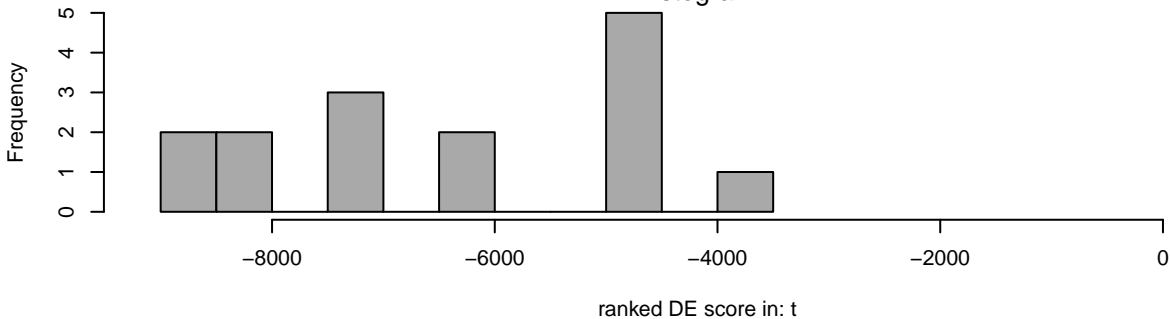


Removal of the Flap Intermediate from the C-strand

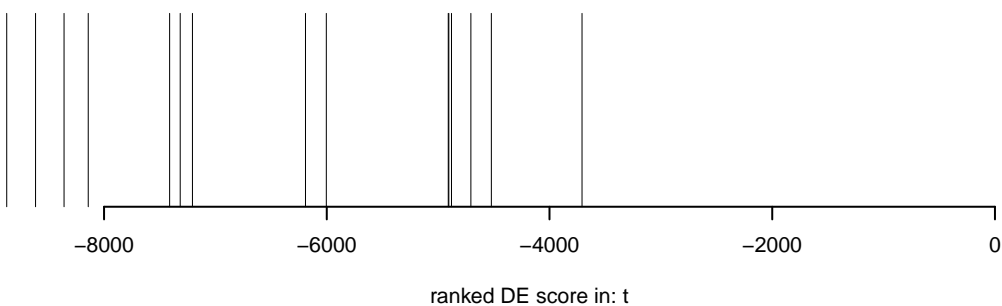
beeswarm plot



histogram

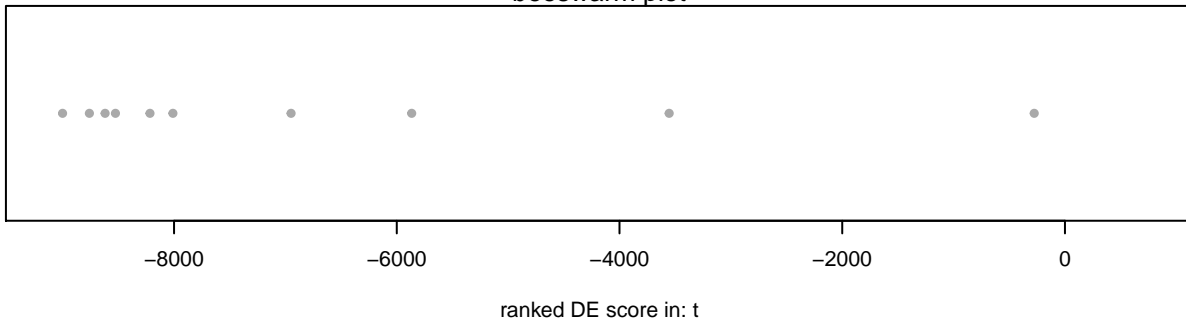


rugplot

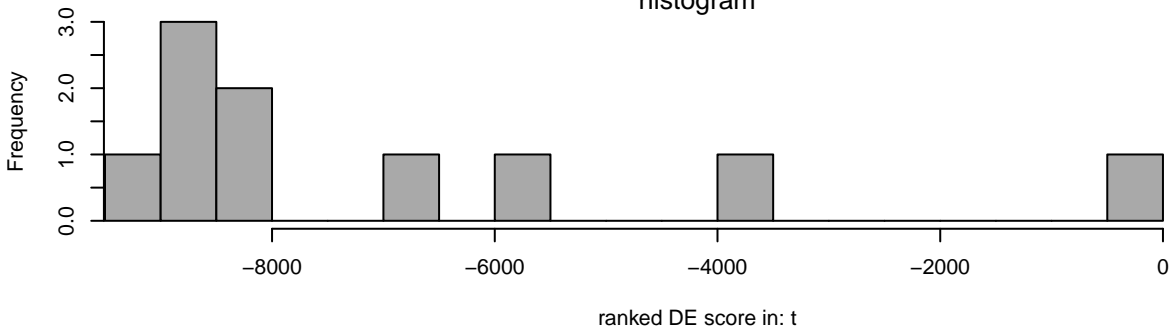


Condensation of Prometaphase Chromosomes

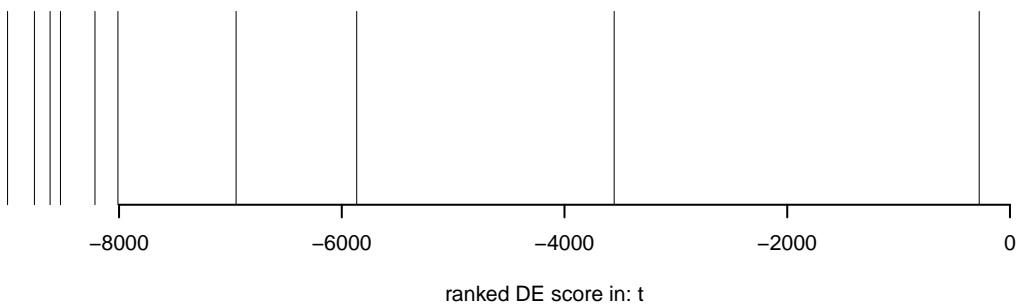
beeswarm plot



histogram

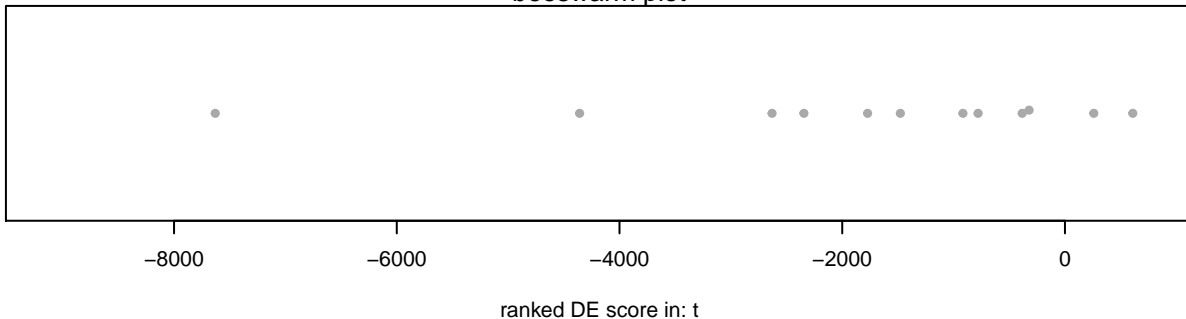


rugplot

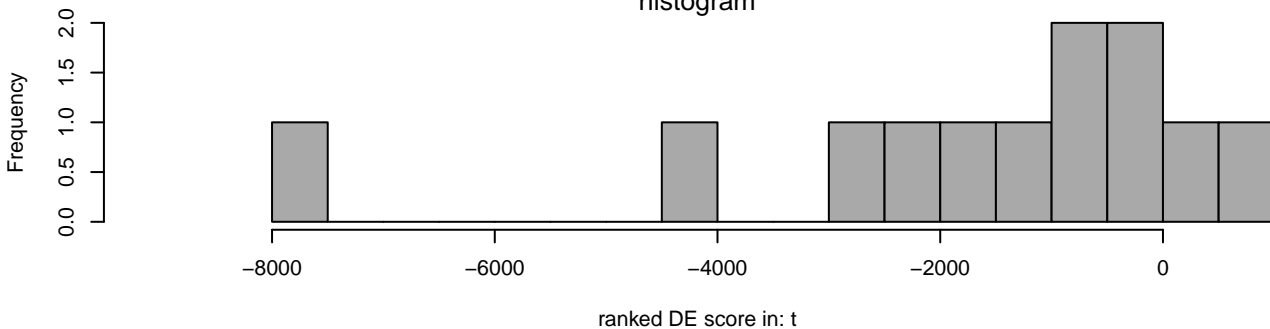


Assembly and cell surface presentation of NMDA receptors

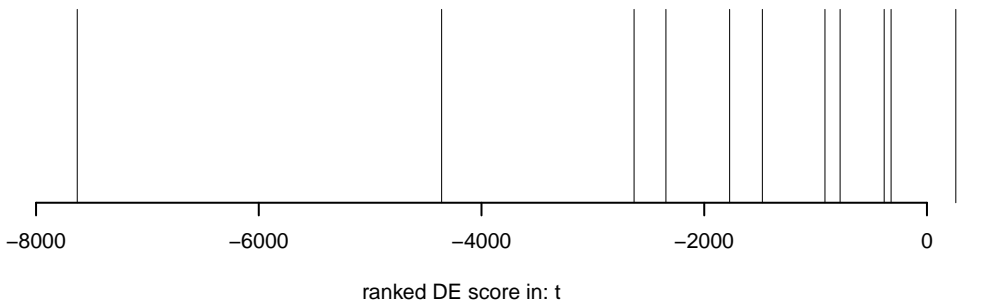
beeswarm plot



histogram

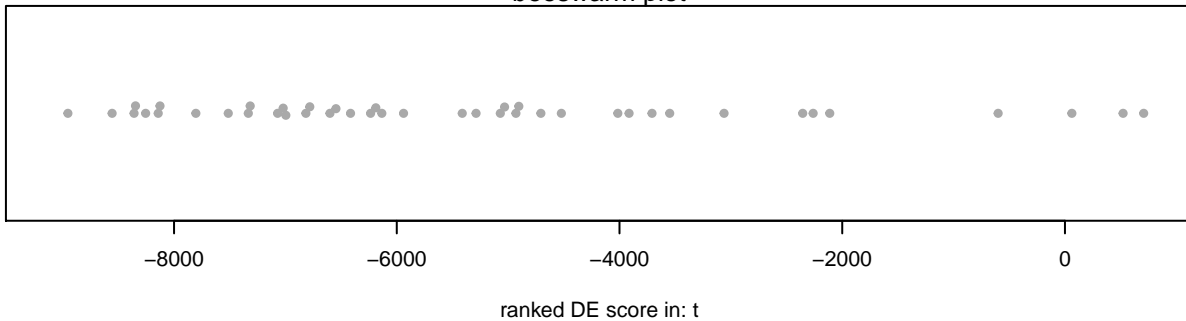


rugplot

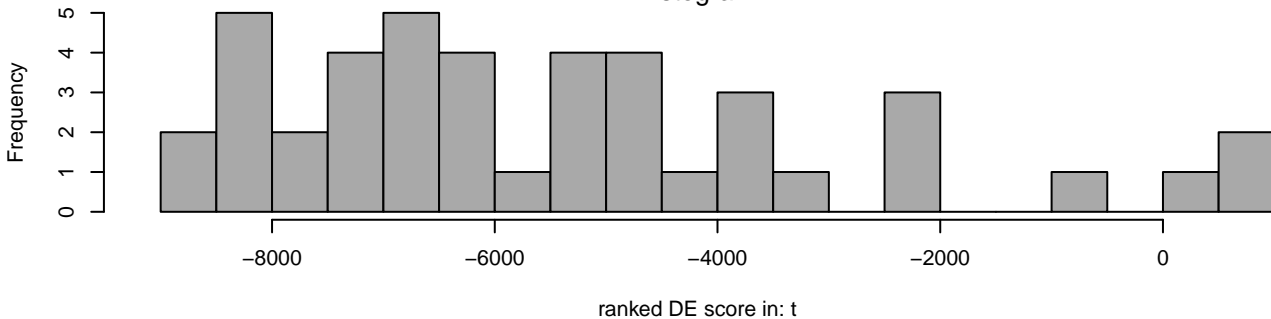


Reproduction

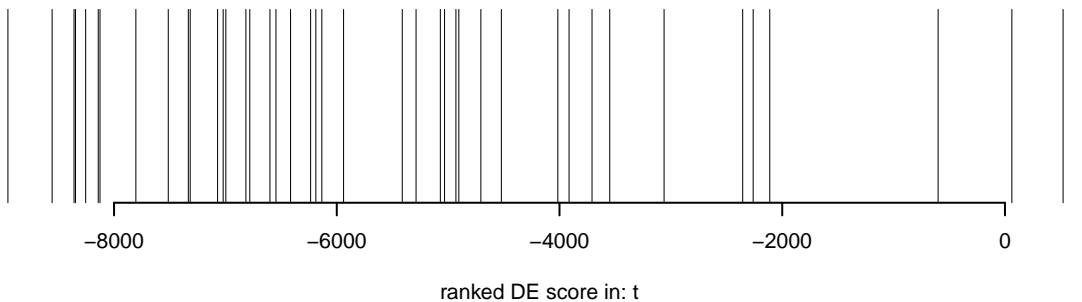
beeswarm plot



histogram

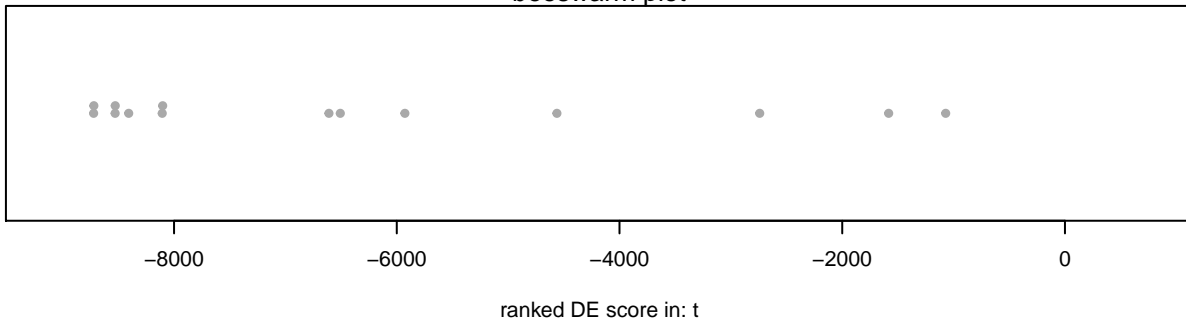


rugplot

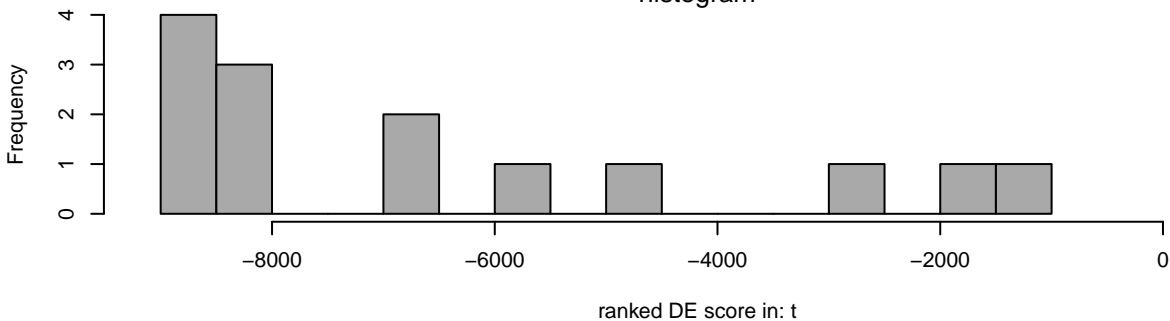


Butyrate Response Factor 1 (BRF1) binds and destabilizes mRNA

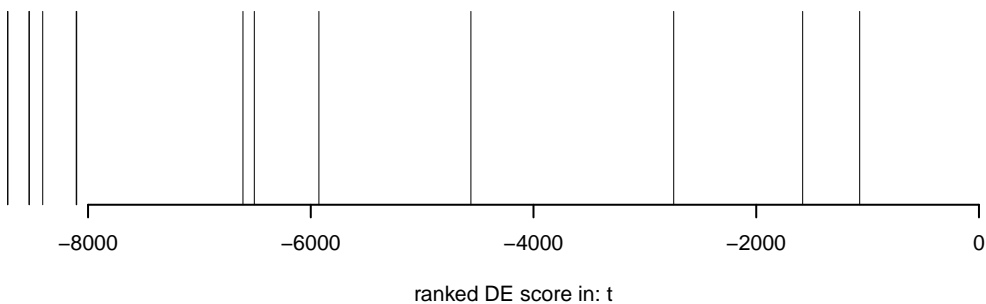
beeswarm plot



histogram

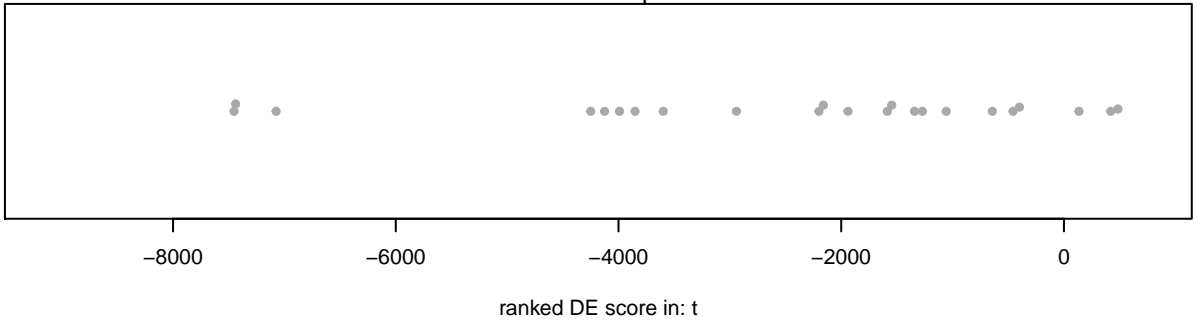


rugplot

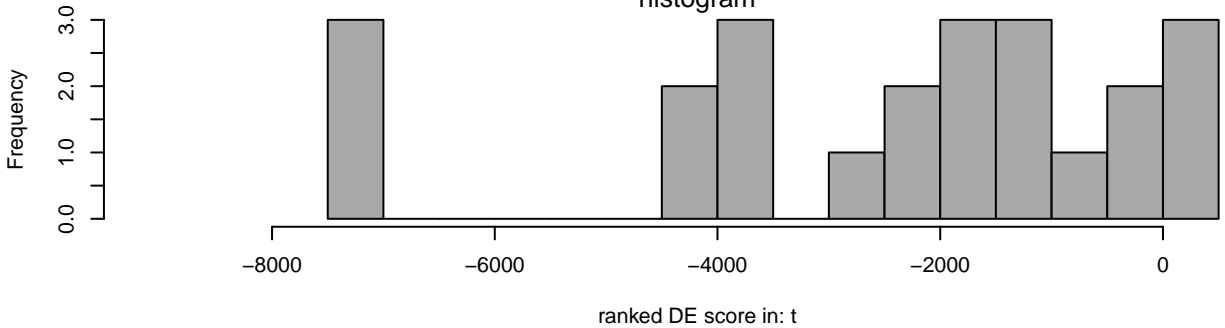


Signaling by WNT in cancer

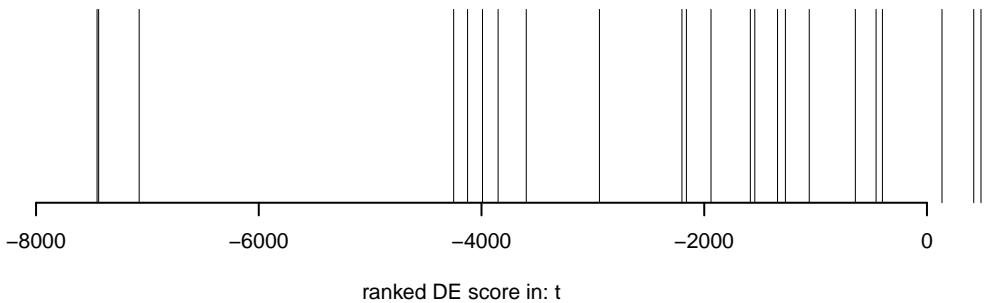
beeswarm plot



histogram

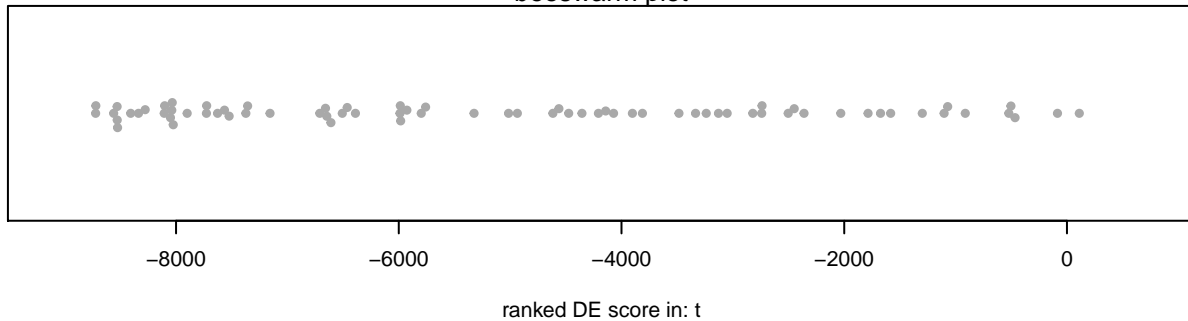


rugplot

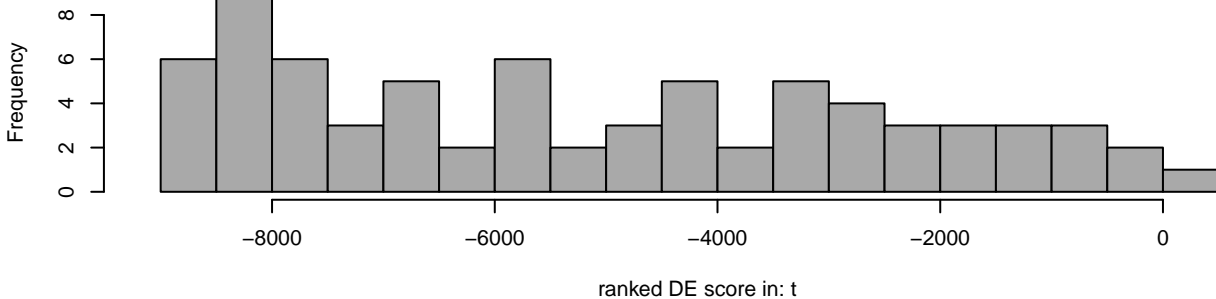


Regulation of mRNA stability by proteins that bind AU-rich elements

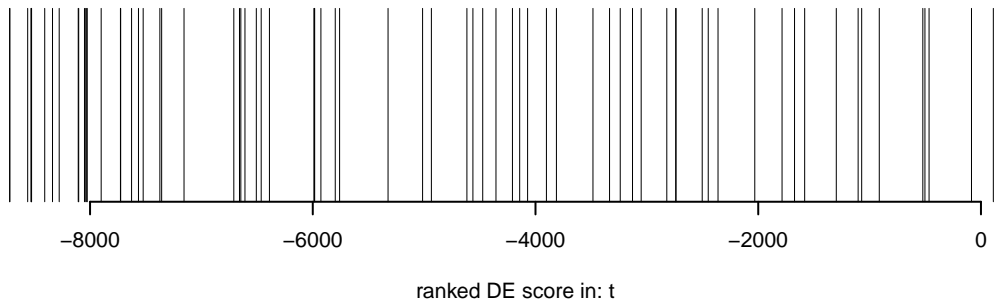
beeswarm plot



histogram

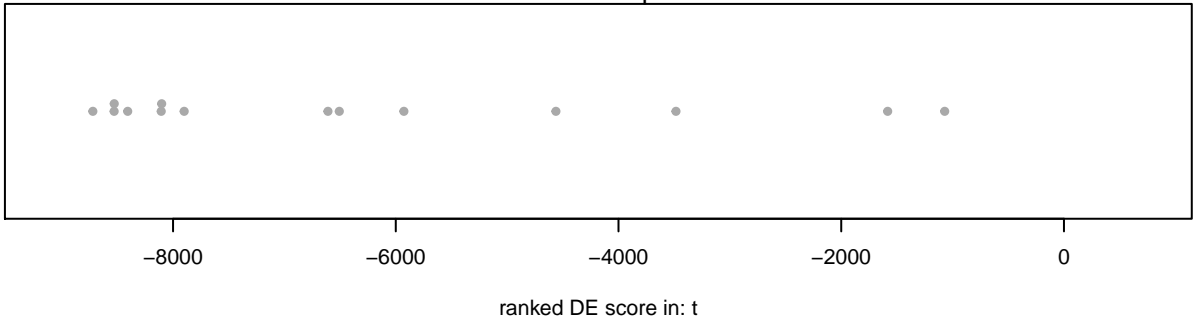


rugplot

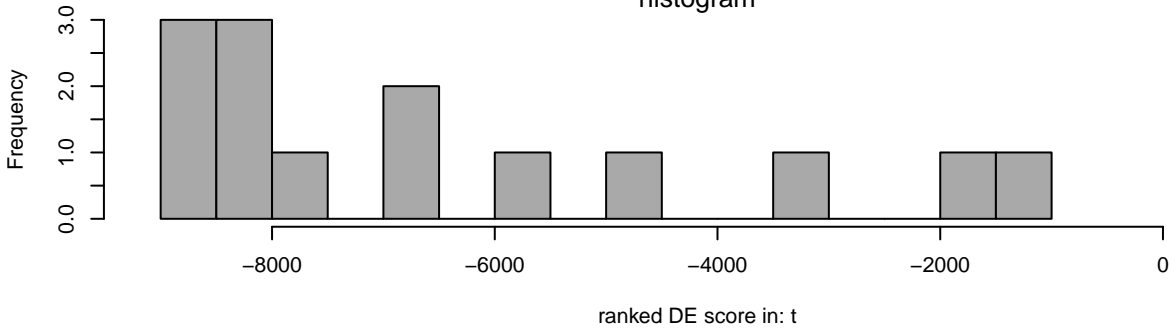


Tristetraprolin (TTP, ZFP36) binds and destabilizes mRNA

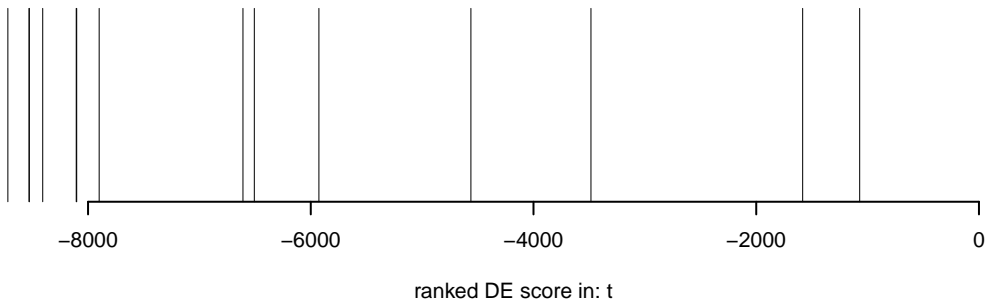
beeswarm plot



histogram

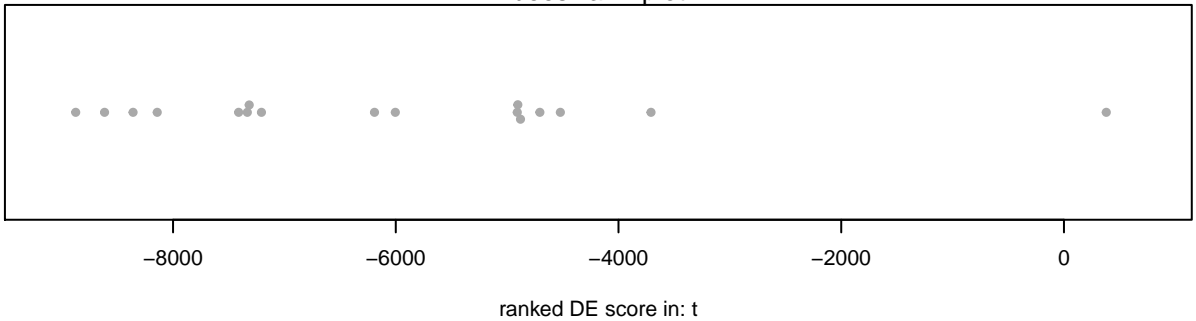


rugplot

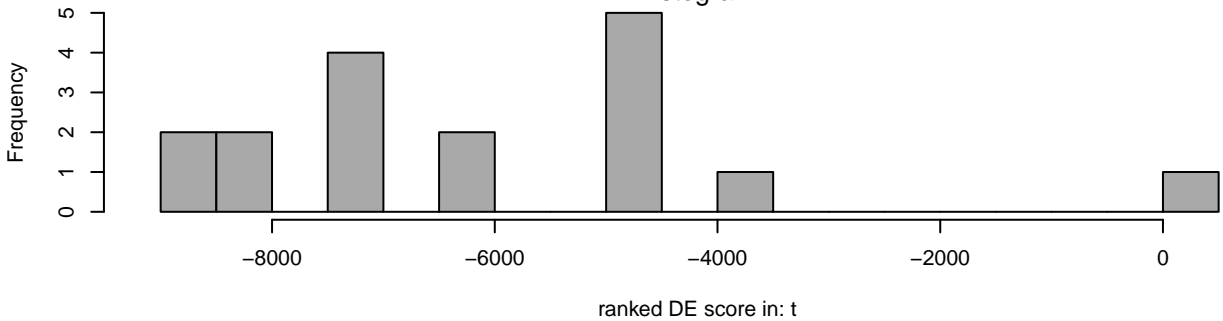


Processive synthesis on the C-strand of the telomere

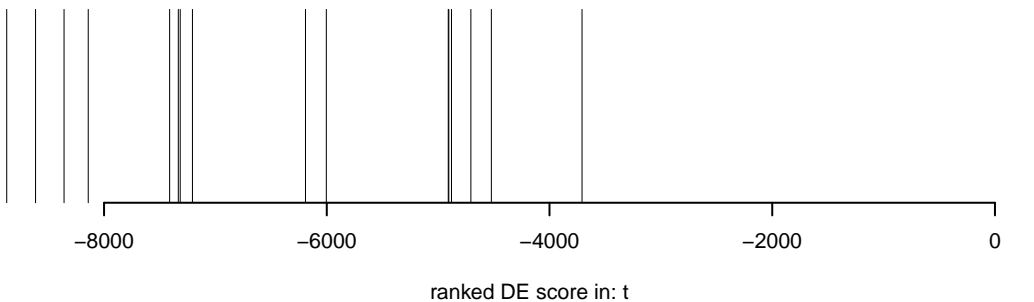
beeswarm plot



histogram

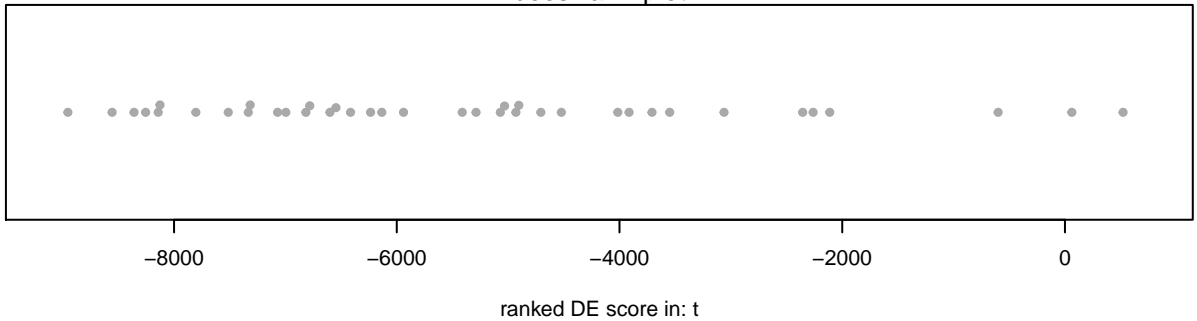


rugplot

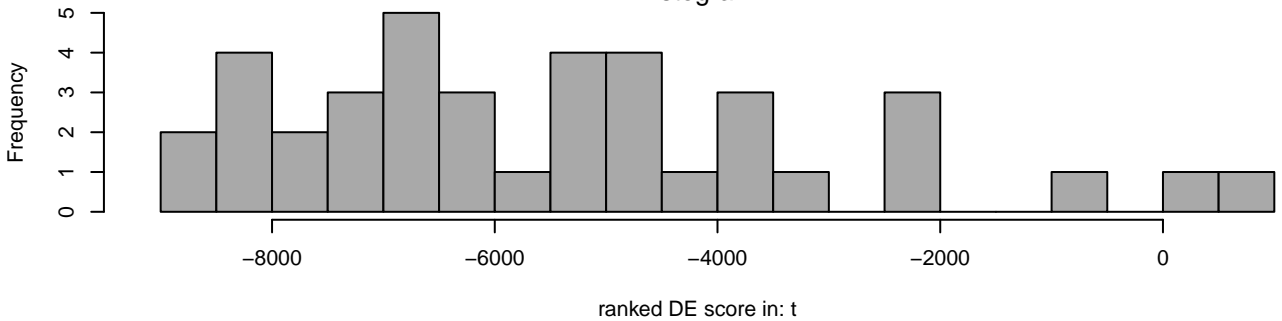


Meiosis

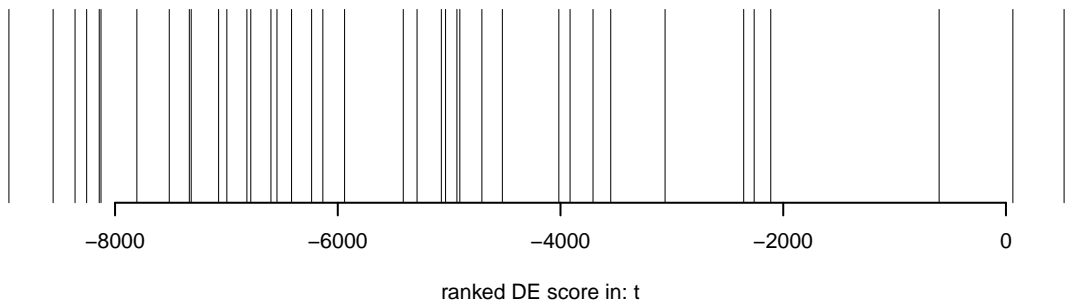
beeswarm plot



histogram

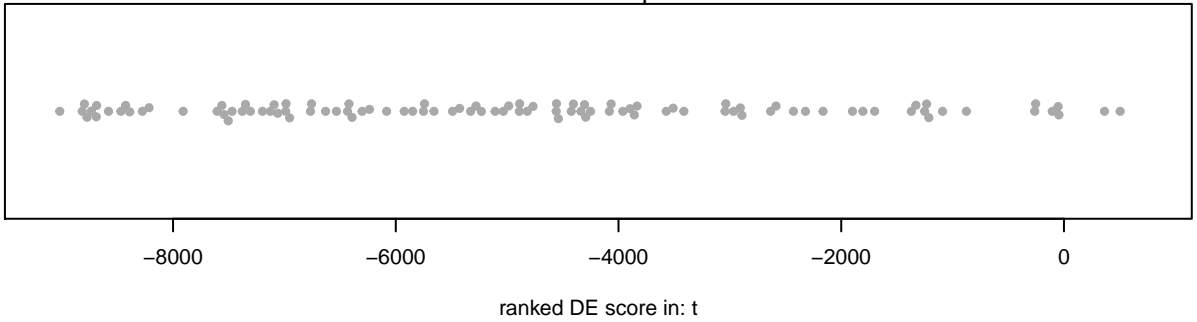


rugplot

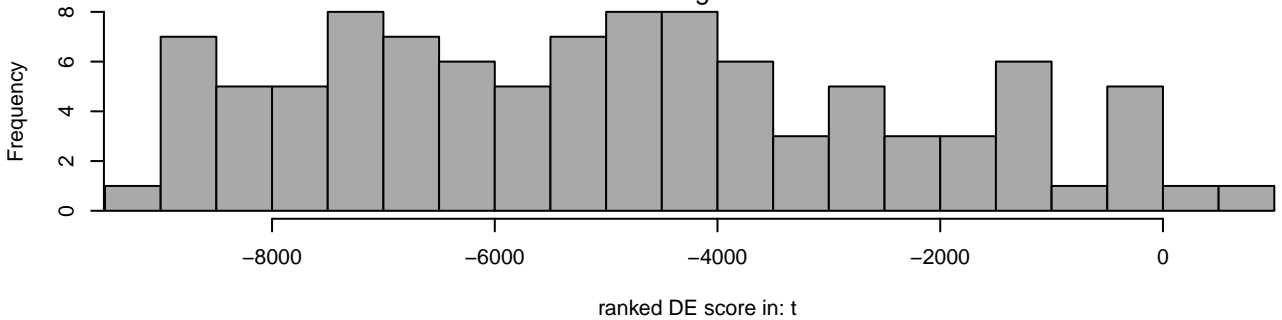


ER to Golgi Anterograde Transport

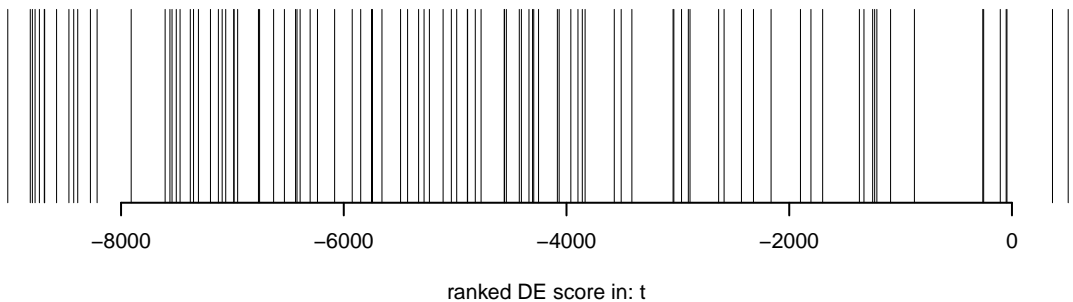
beeswarm plot



histogram

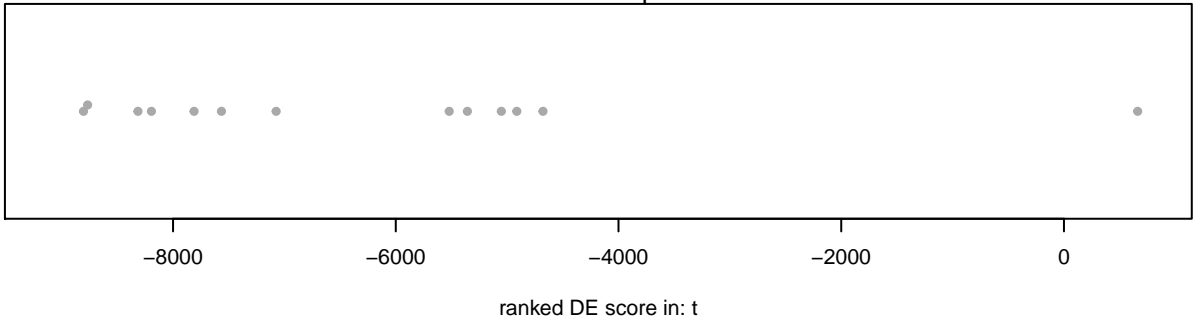


rugplot

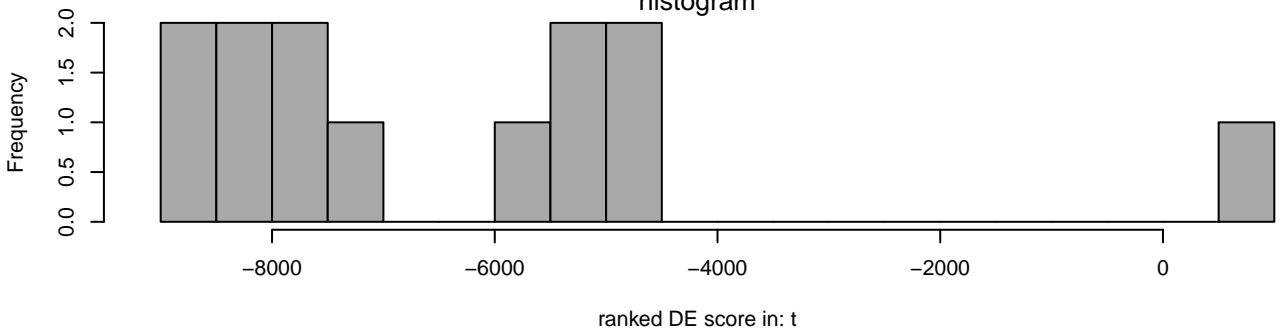


Myogenesis

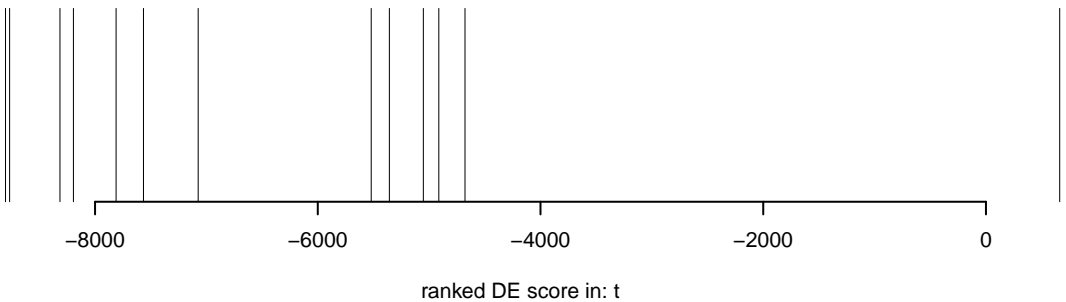
beeswarm plot



histogram

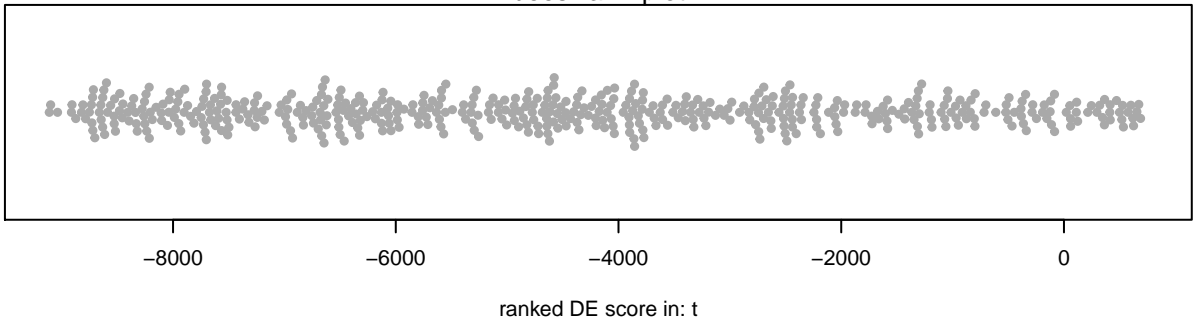


rugplot

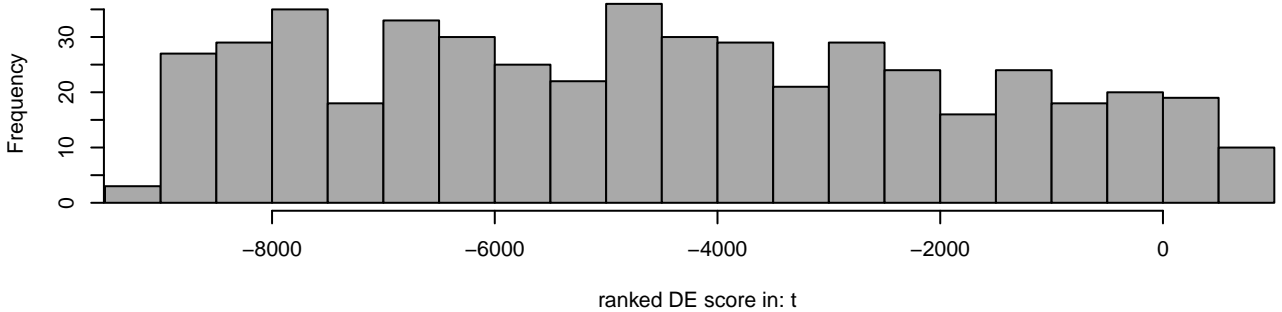


Metabolism of RNA

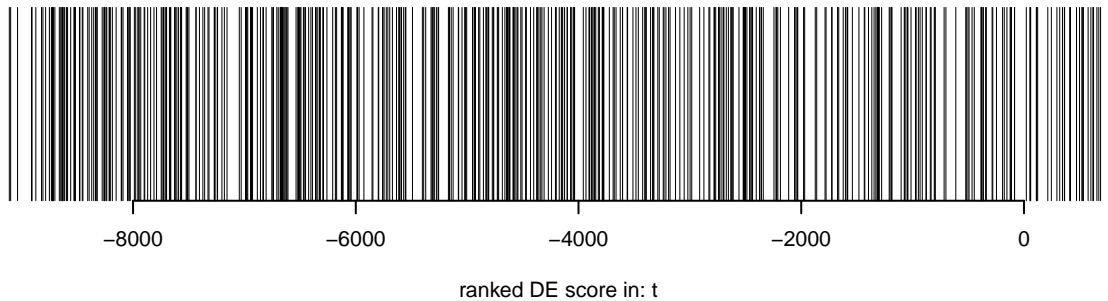
beeswarm plot



histogram

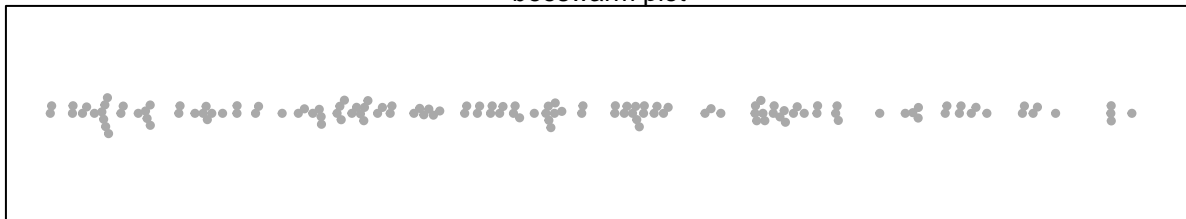


rugplot



mRNA Splicing

beeswarm plot



-8000

-6000

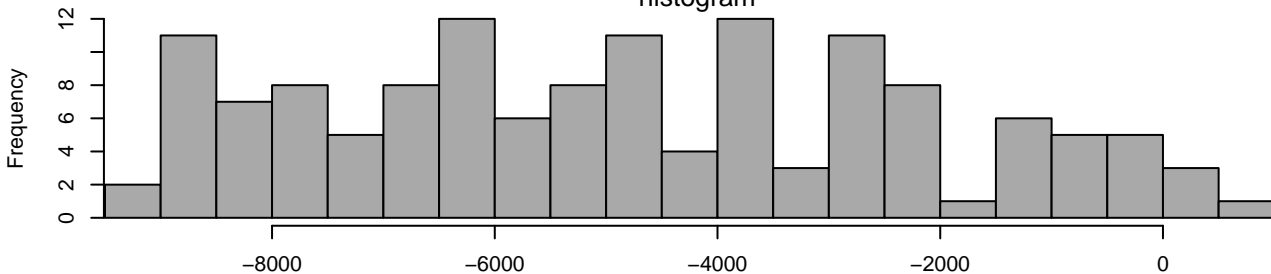
-4000

-2000

0

ranked DE score in: t

histogram



-8000

-6000

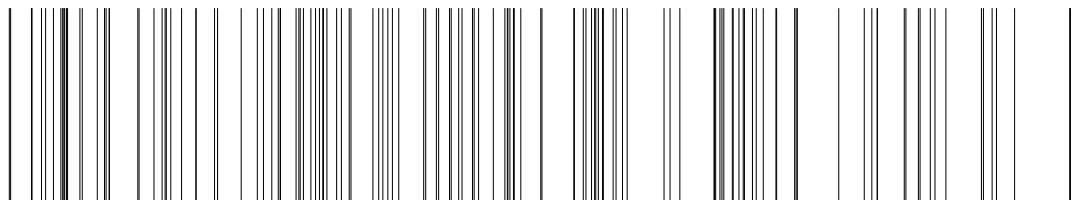
-4000

-2000

0

ranked DE score in: t

rugplot



-8000

-6000

-4000

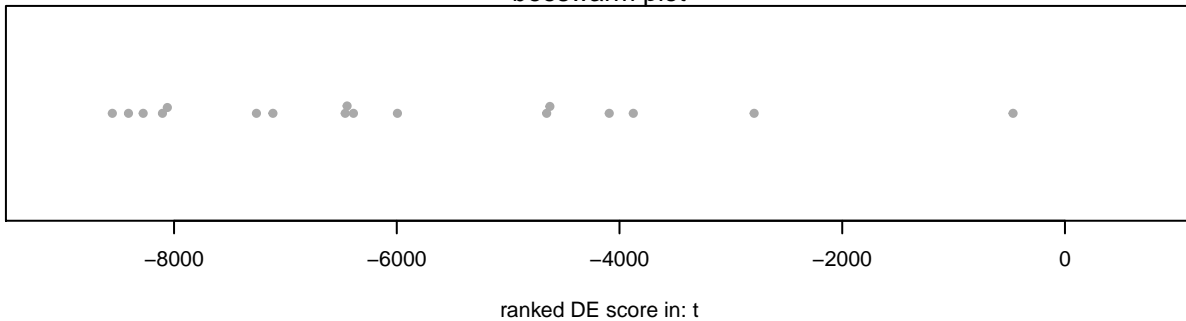
-2000

0

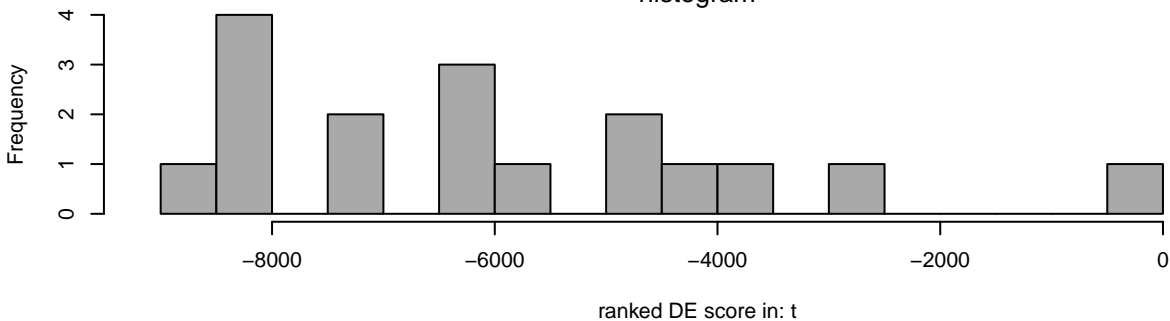
ranked DE score in: t

IKK complex recruitment mediated by RIP1

beeswarm plot



histogram



rugplot

