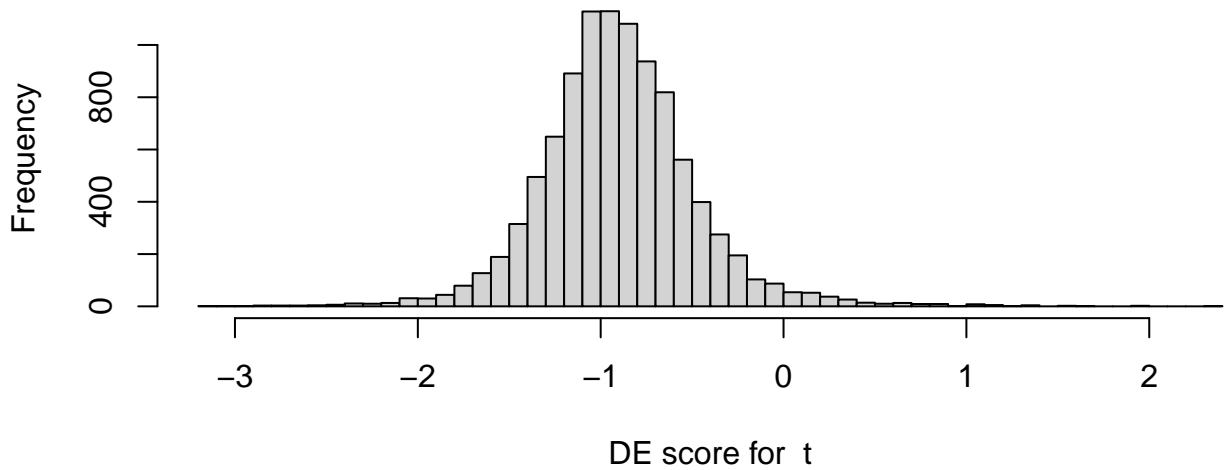
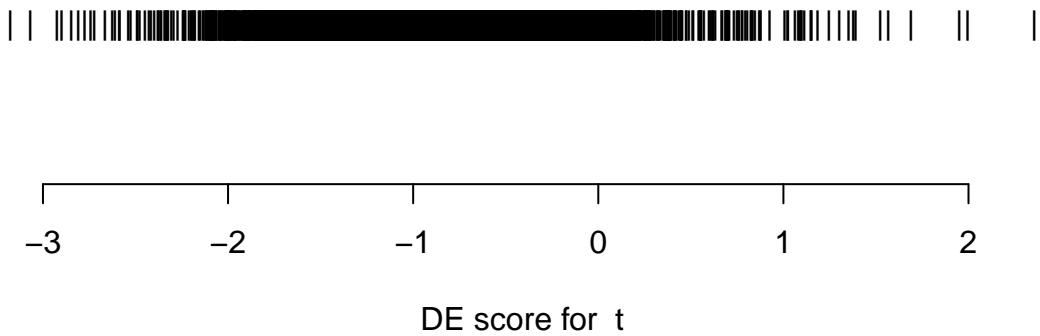


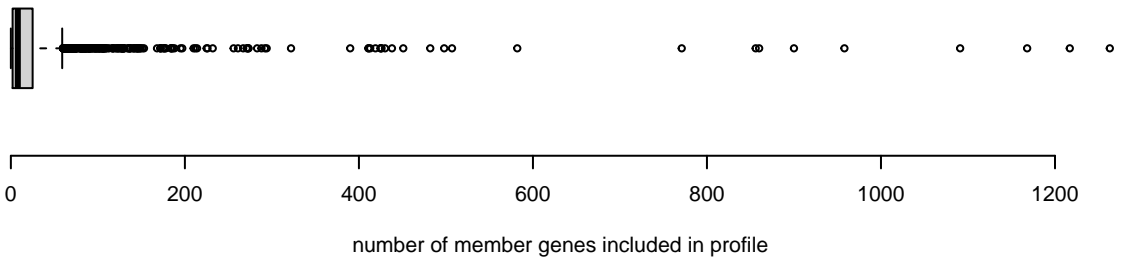
Distribution of DE scores



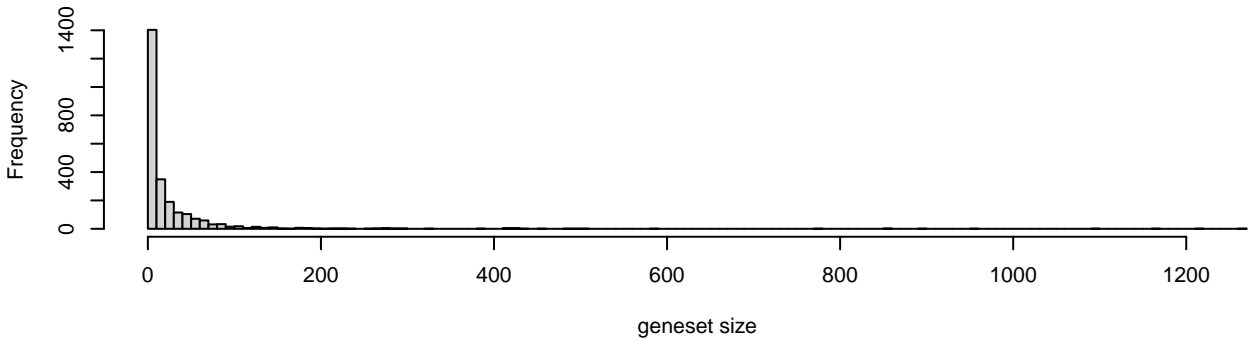
9869 genes in total, 249 trending up-regulated, 9620 trending down-regulated



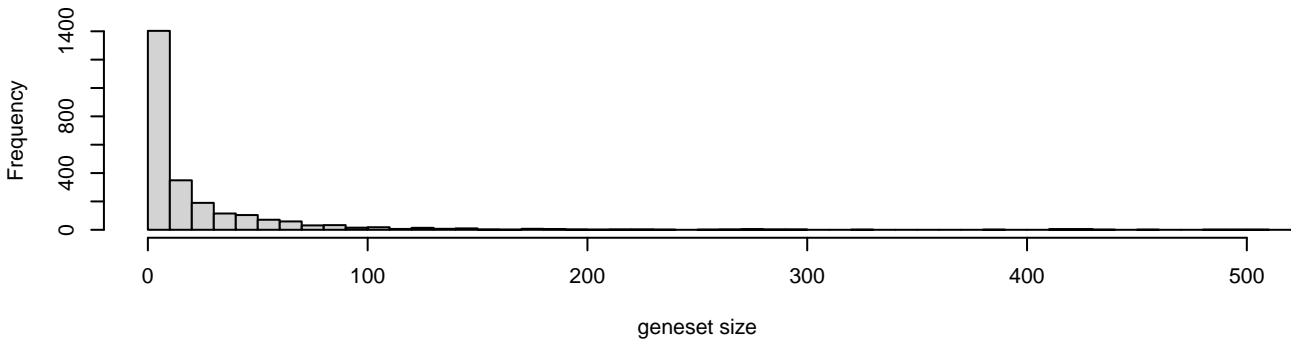
Gene set size



Histogram of geneset size

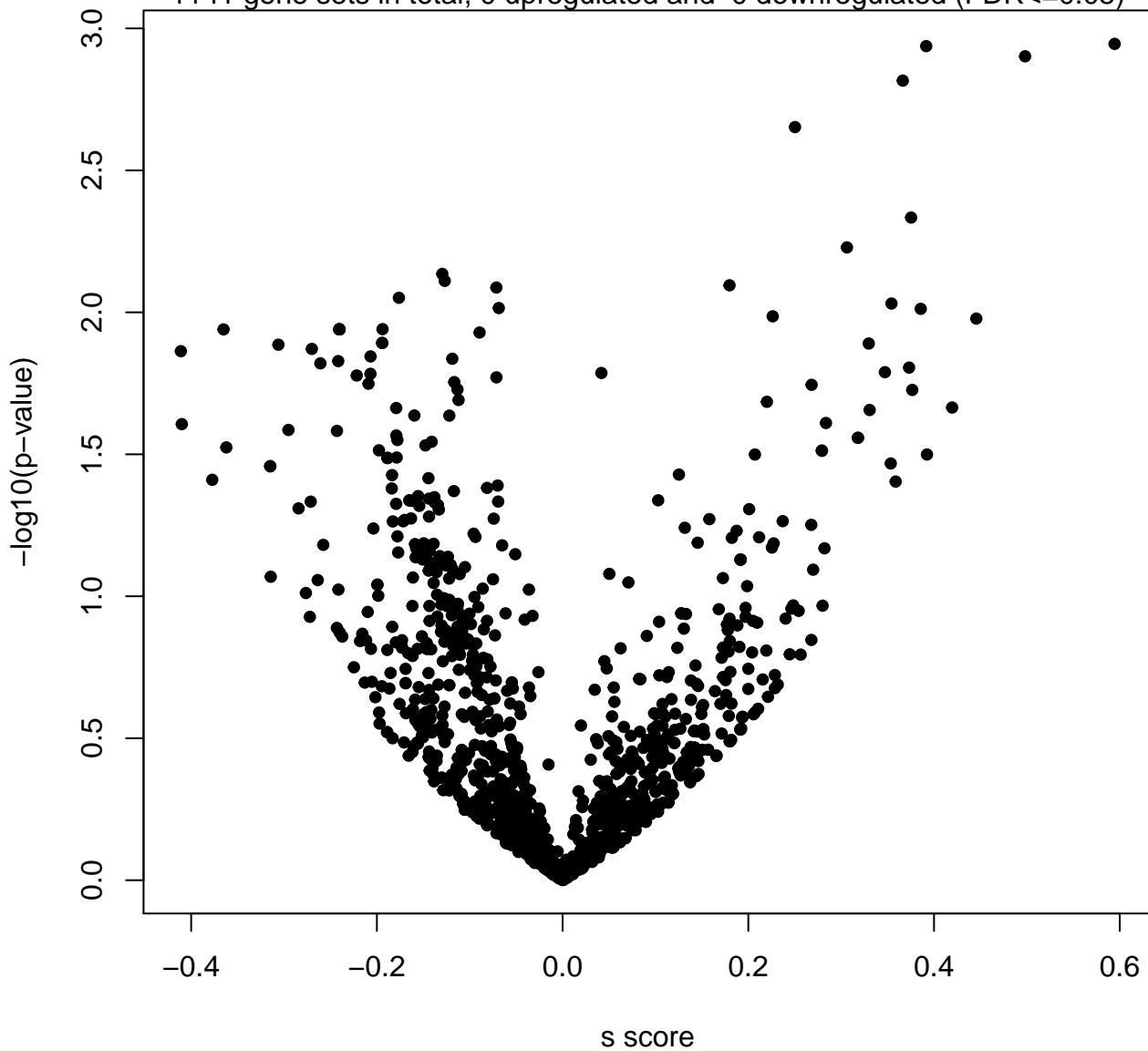


Trimmed histogram of geneset size



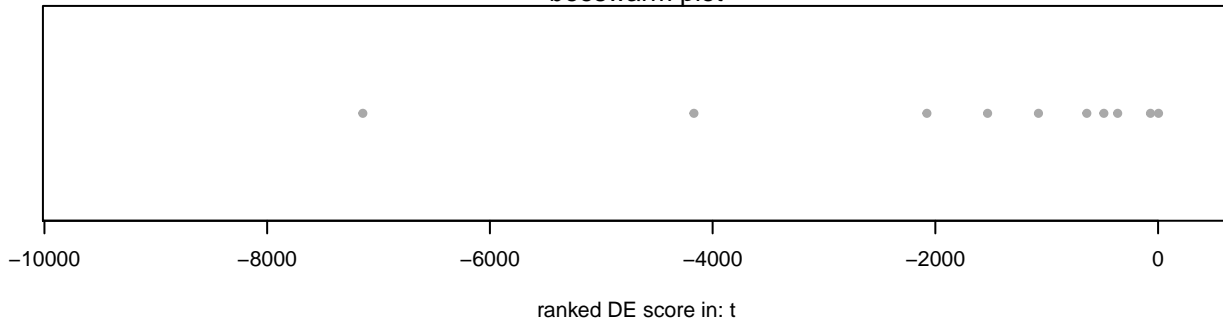
volcano plot of gene set enrichments

1141 gene sets in total, 0 upregulated and 0 downregulated (FDR \leq 0.05)

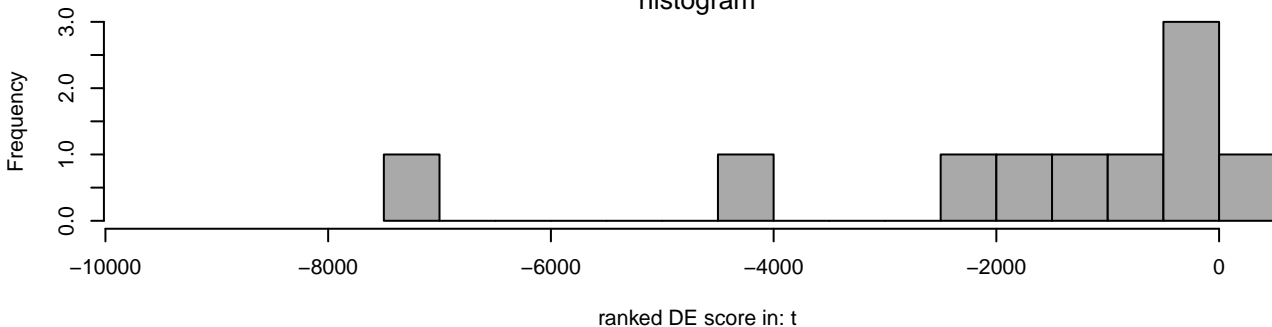


RORA activates gene expression

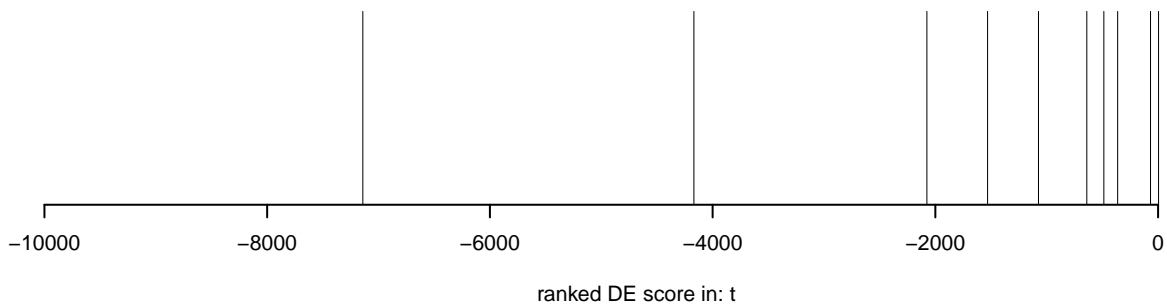
beeswarm plot



histogram

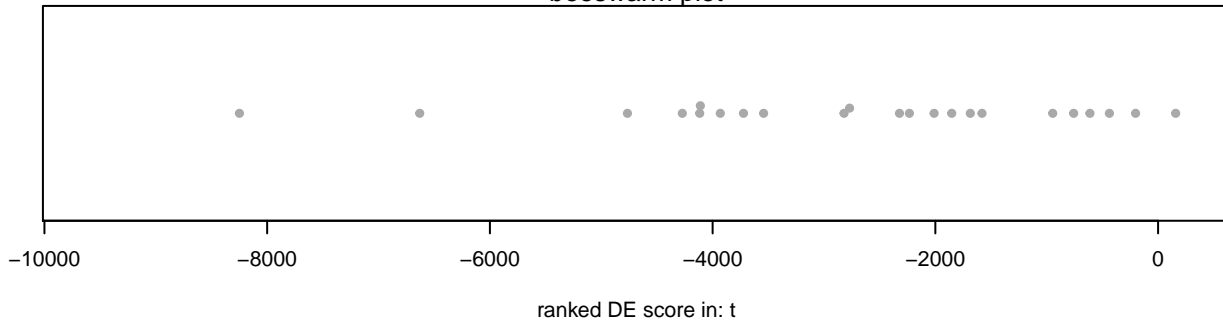


rugplot

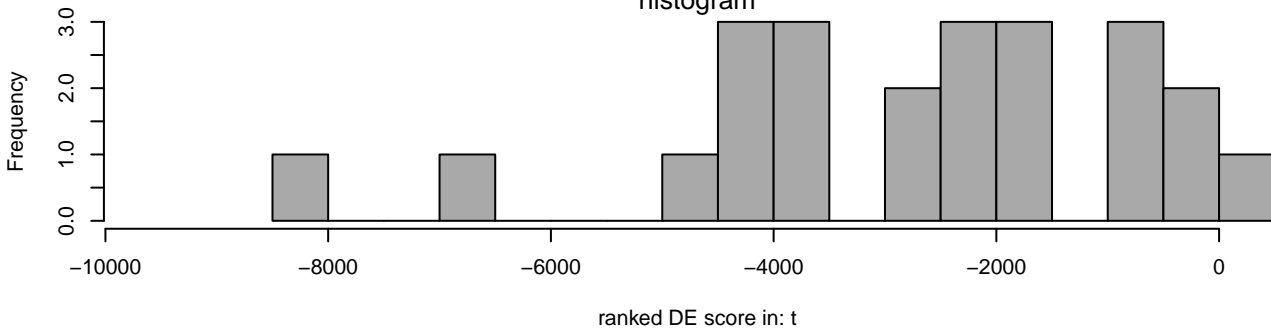


GPVI-mediated activation cascade

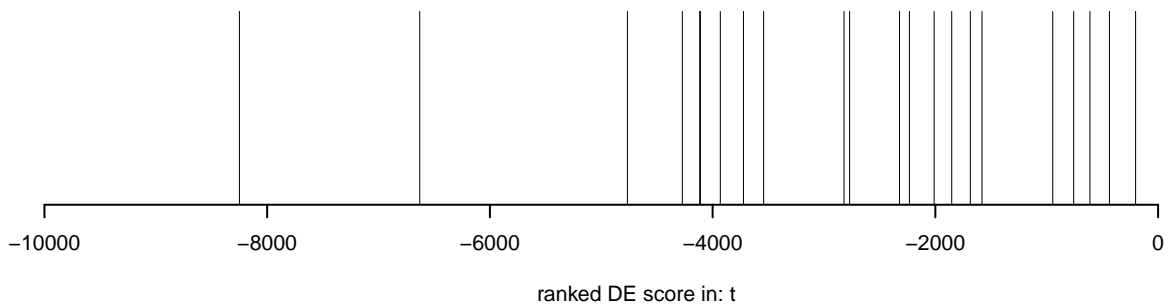
beeswarm plot



histogram

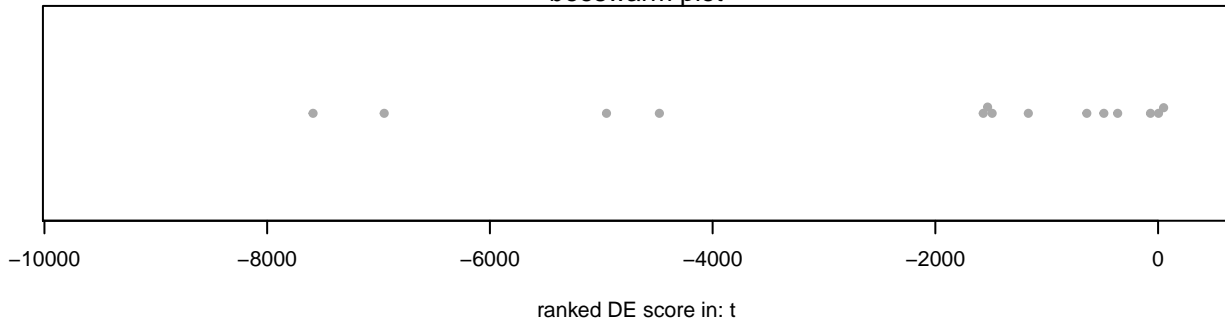


rugplot

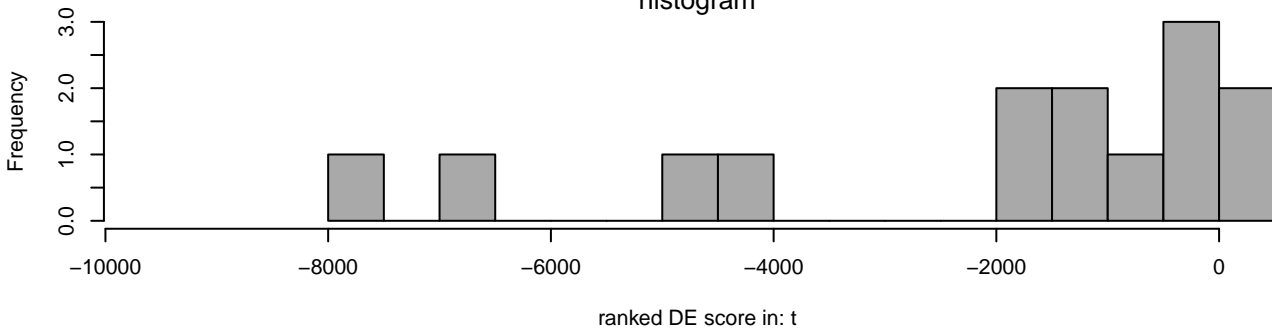


BMAL1:CLOCK,NPAS2 activates circadian gene expression

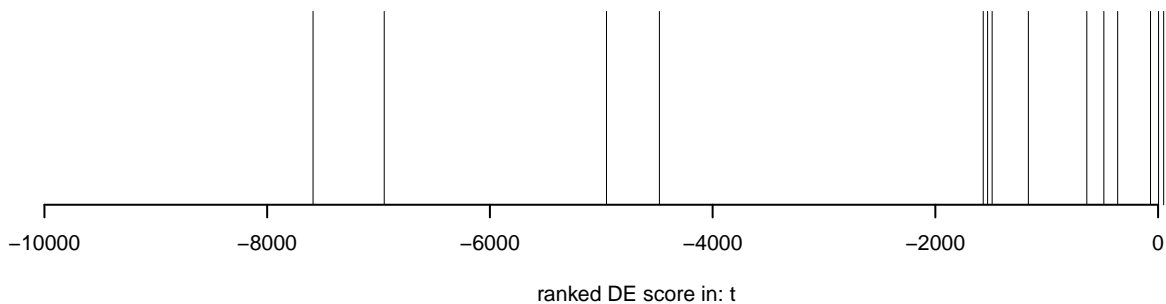
beeswarm plot



histogram

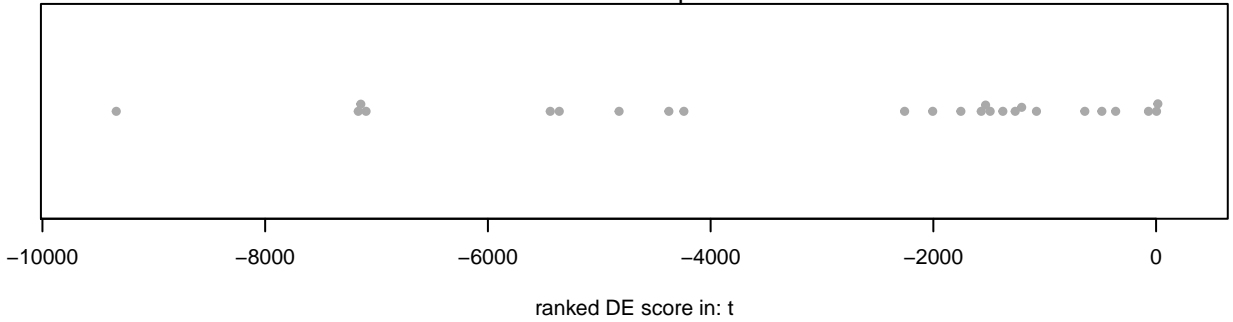


rugplot

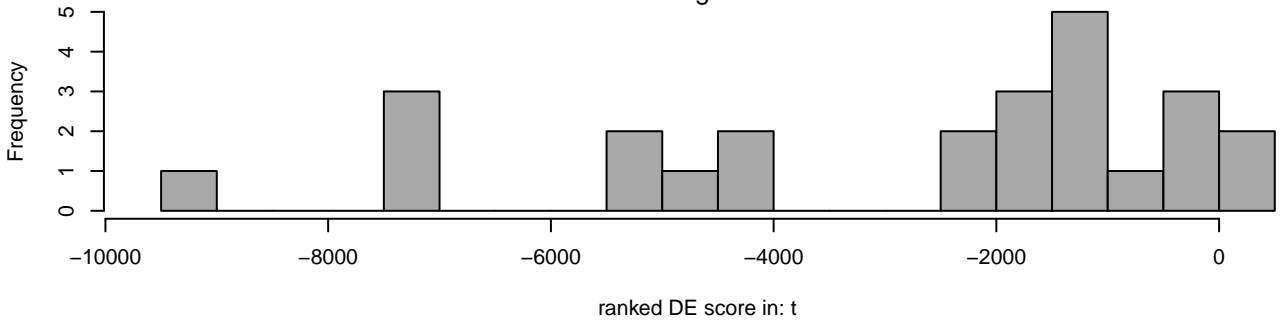


Heme signaling

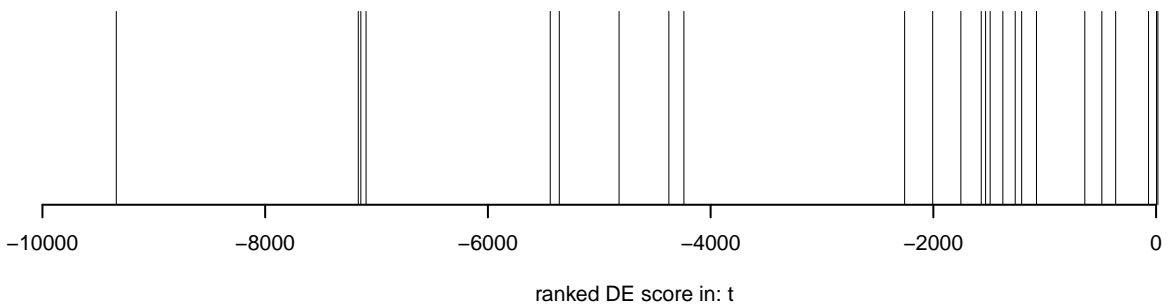
beeswarm plot



histogram

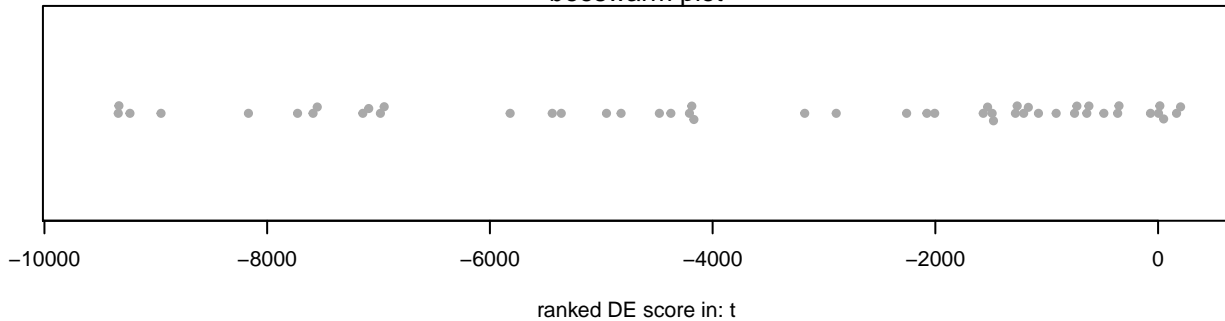


rugplot

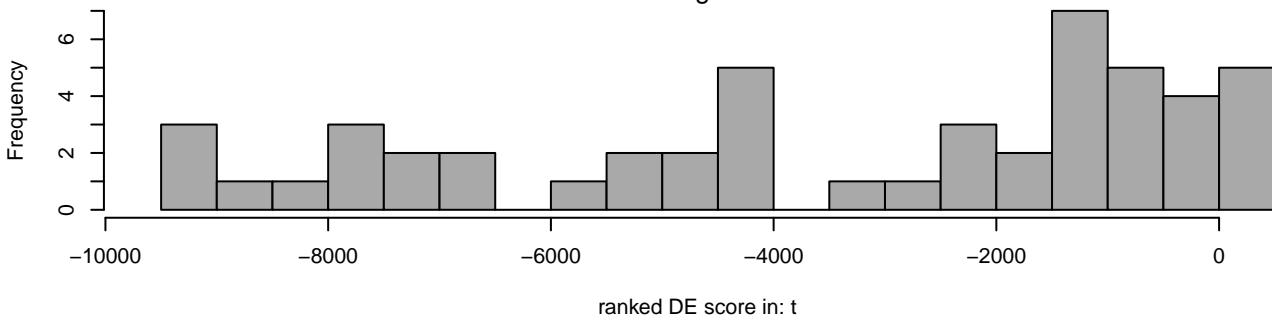


Circadian Clock

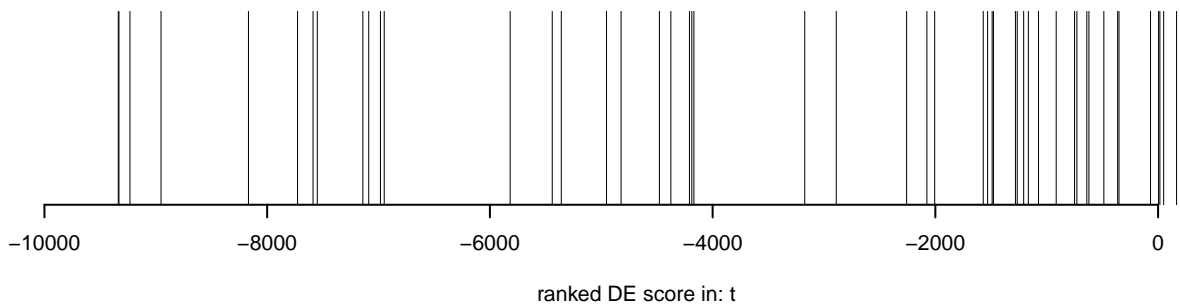
beeswarm plot



histogram

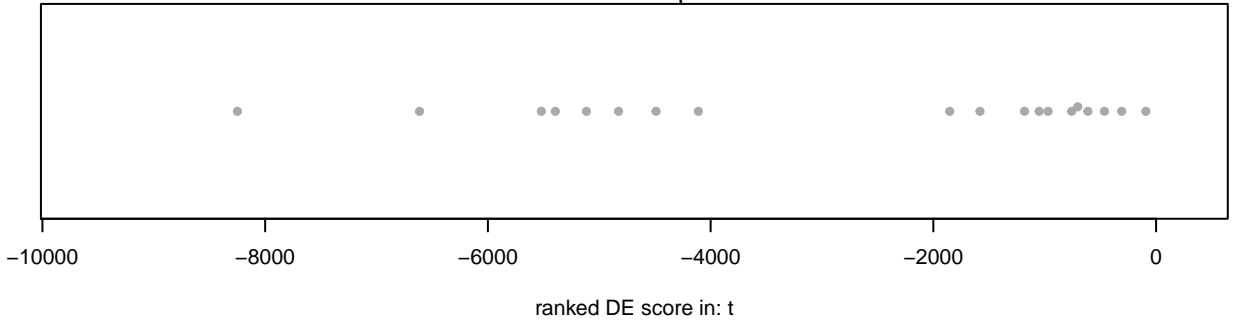


rugplot

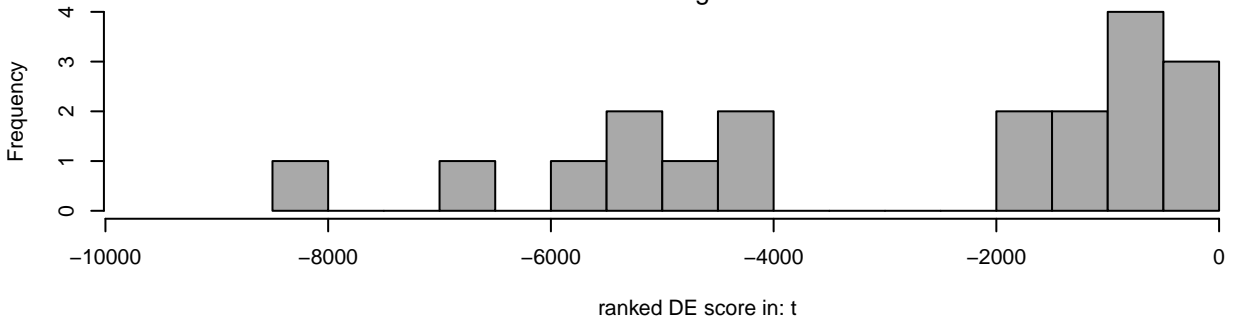


Signaling by Erythropoietin

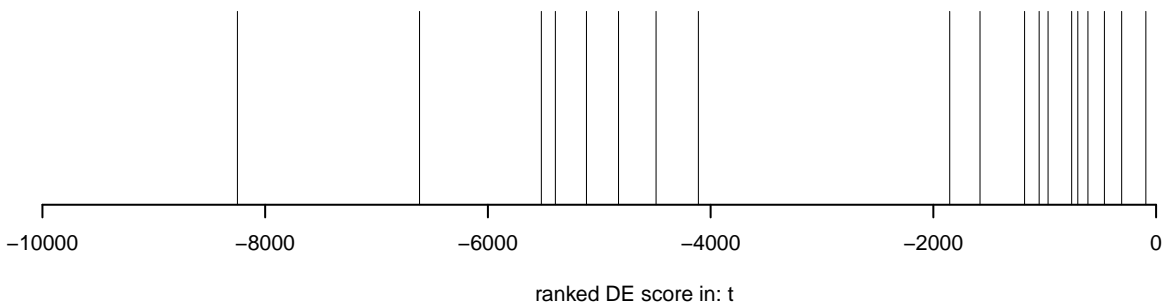
beeswarm plot



histogram

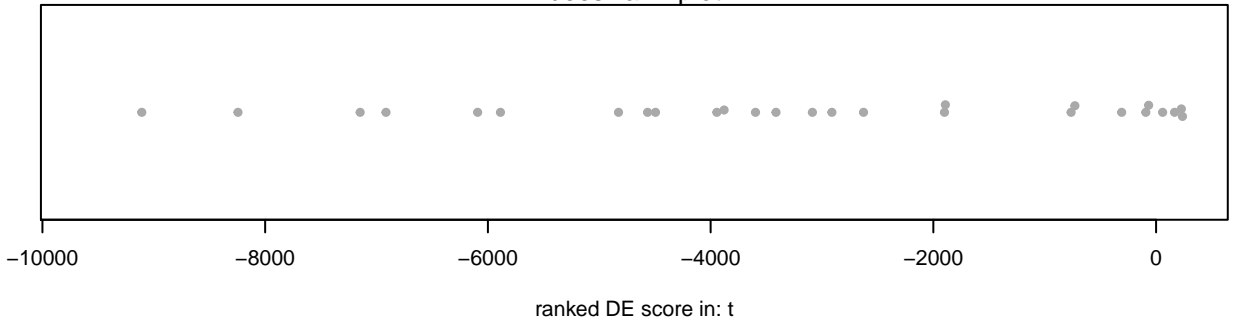


rugplot

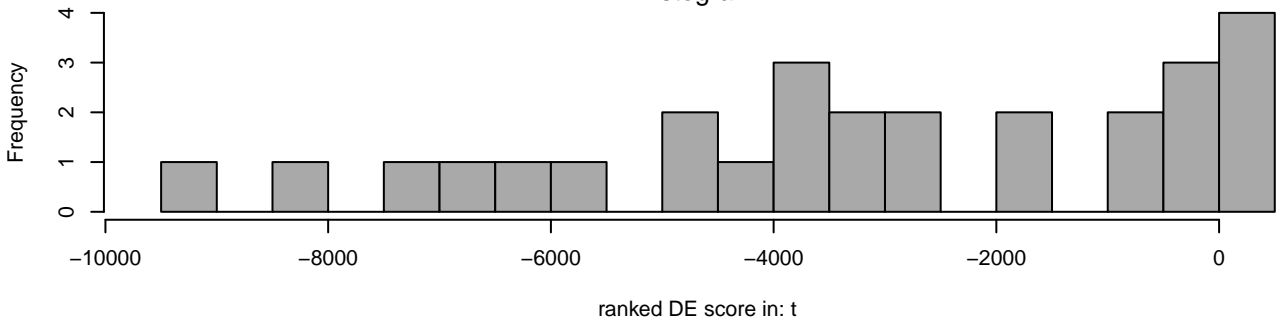


RAF activation

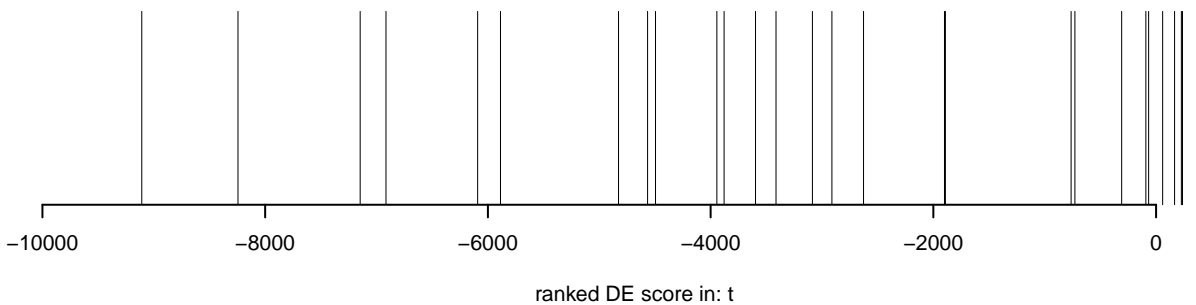
beeswarm plot



histogram

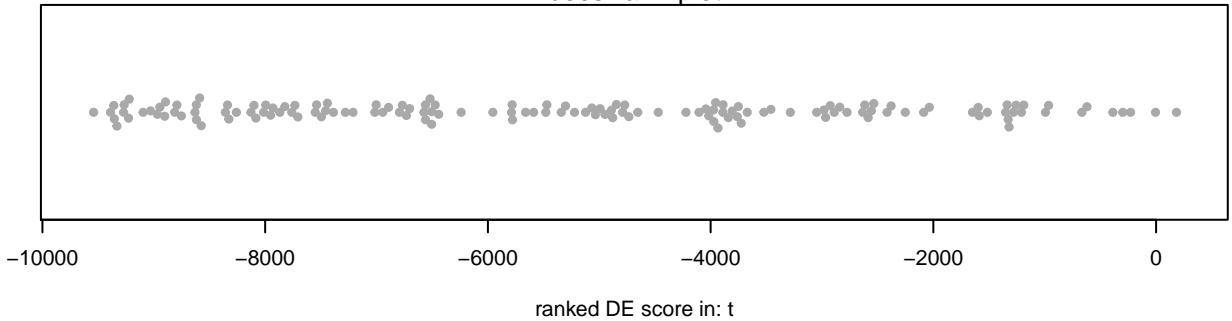


rugplot

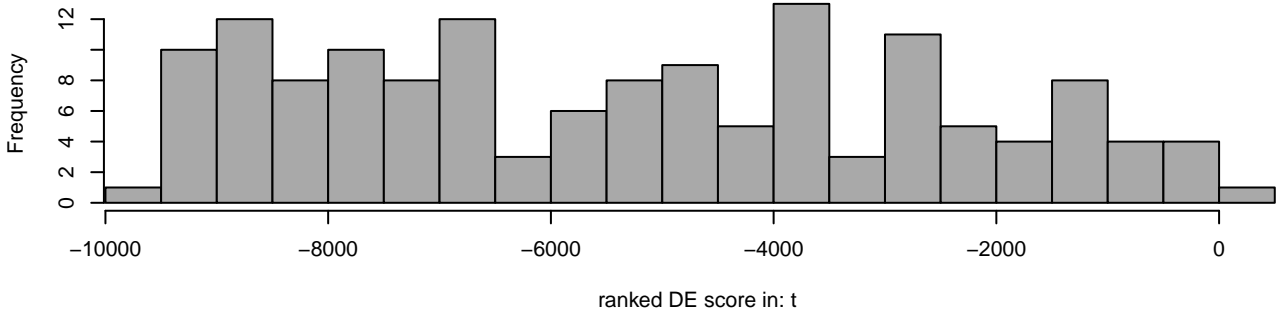


rRNA processing in the nucleus and cytosol

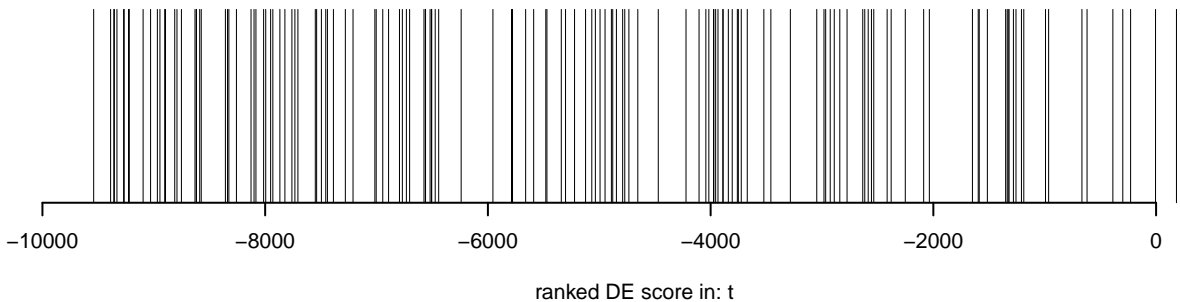
beeswarm plot



histogram

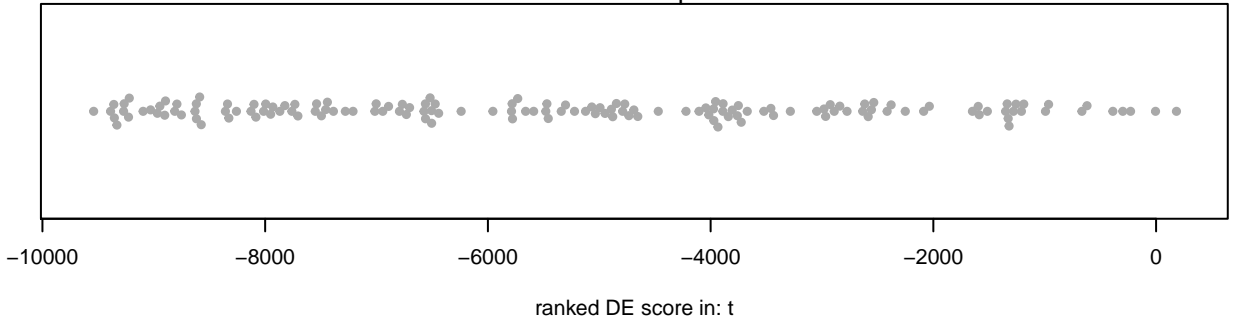


rugplot

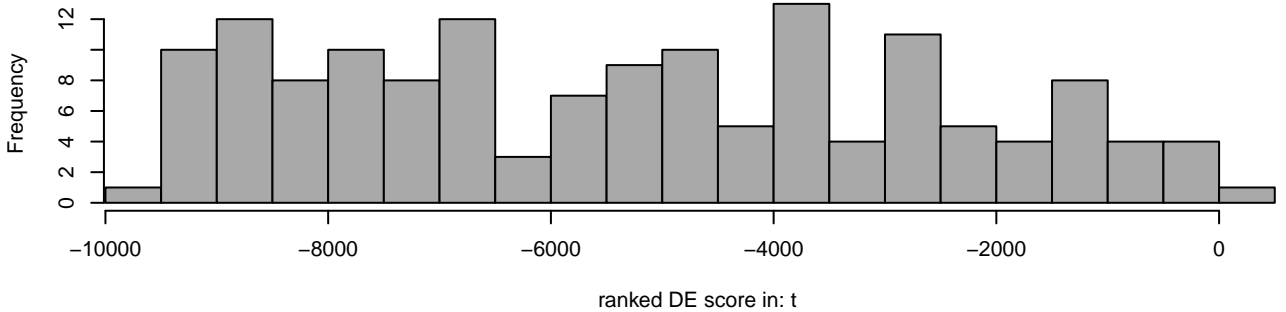


rRNA processing

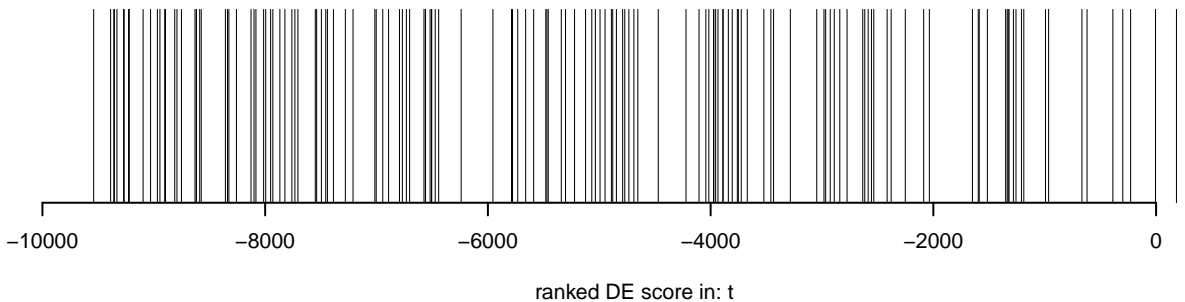
beeswarm plot



histogram

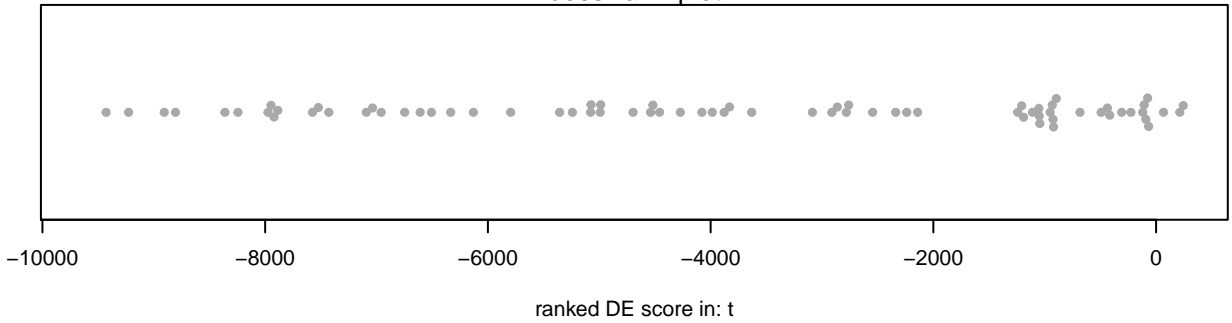


rugplot

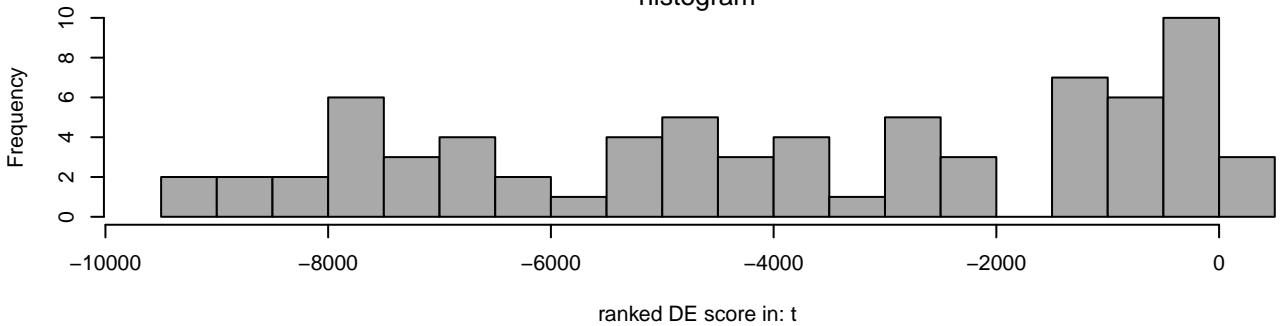


Neurotransmitter receptors and postsynaptic signal transmission

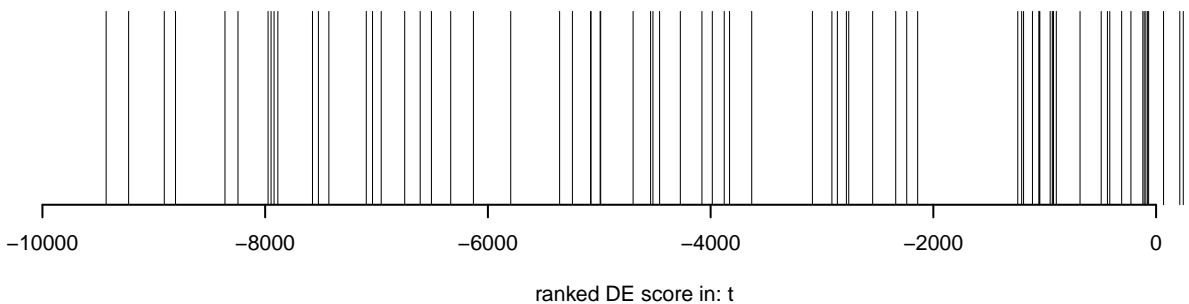
beeswarm plot



histogram

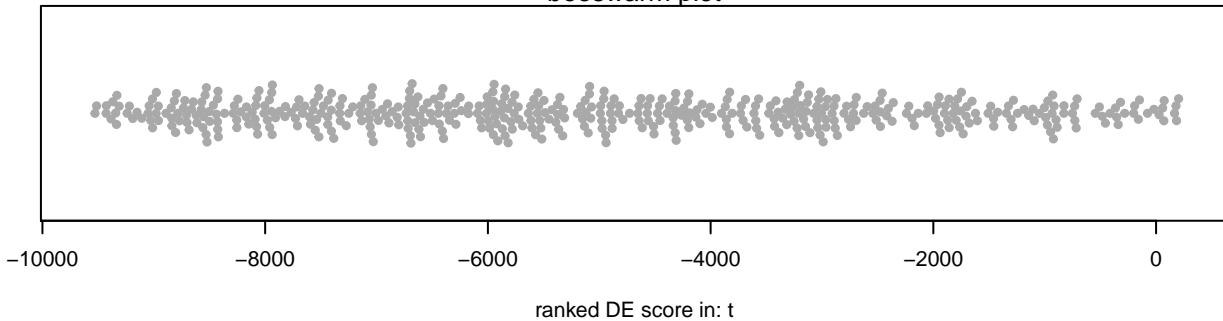


rugplot

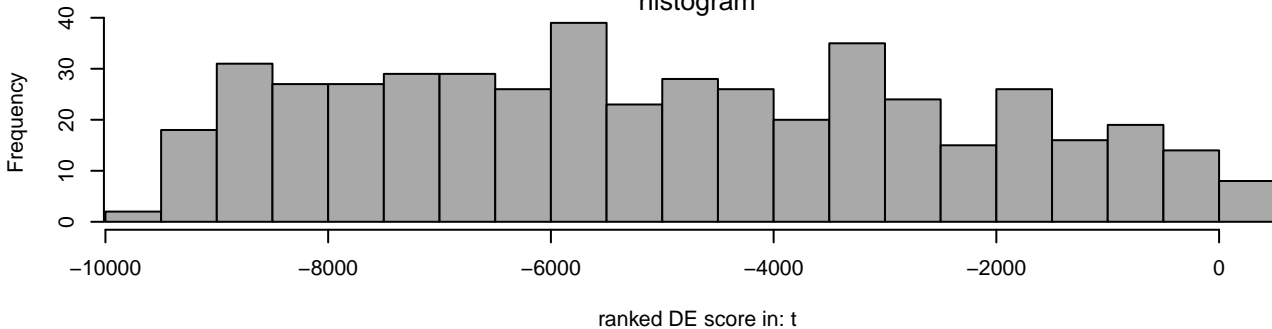


Cell Cycle

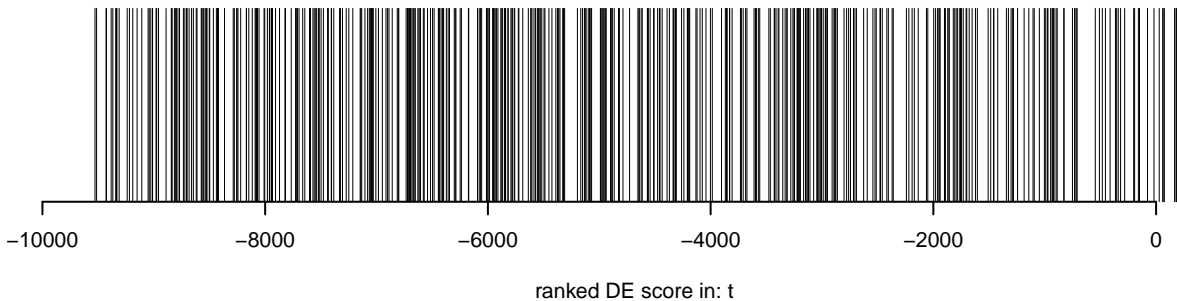
beeswarm plot



histogram

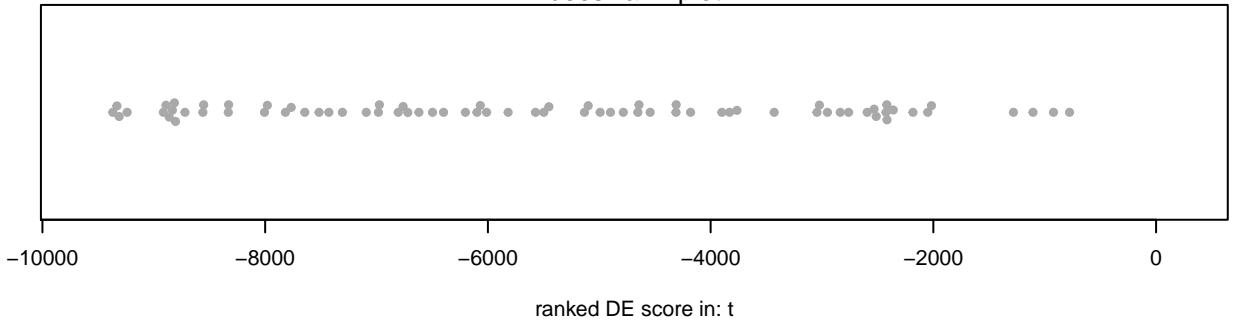


rugplot

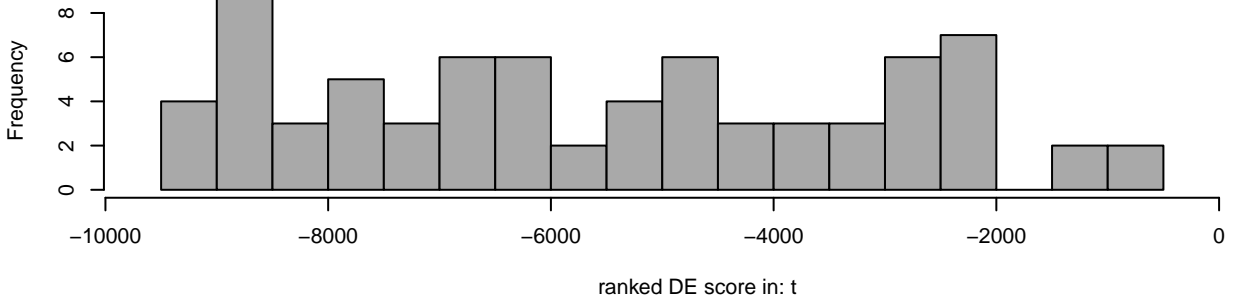


Regulation of TP53 Activity through Phosphorylation

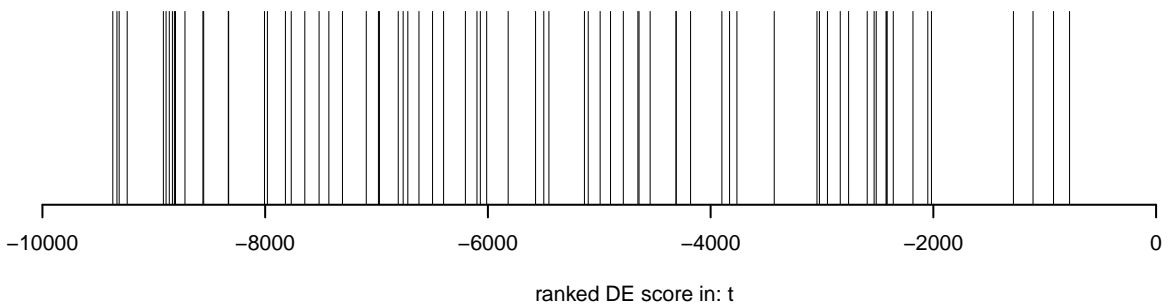
beeswarm plot



histogram

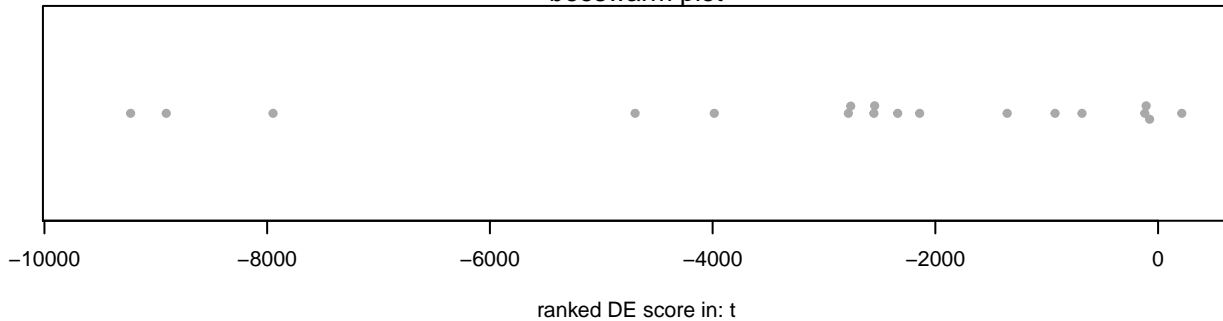


rugplot

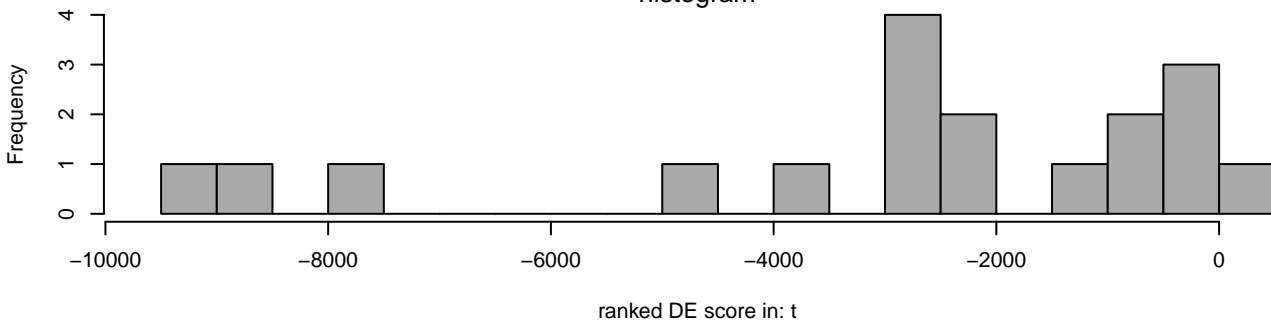


Vasopressin regulates renal water homeostasis via Aquaporins

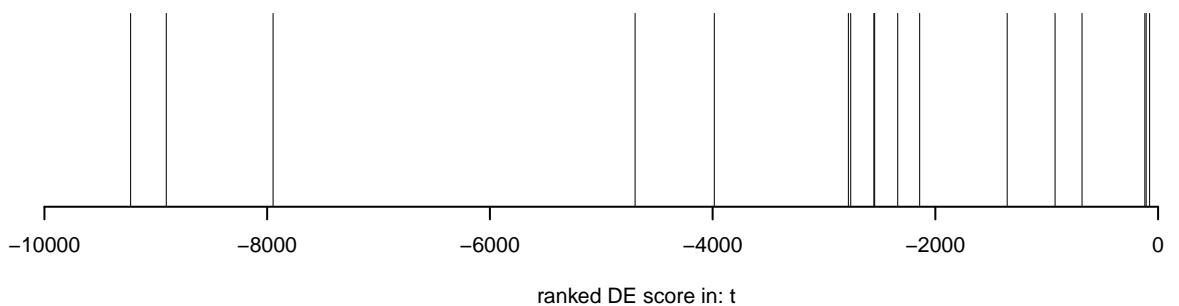
beeswarm plot



histogram

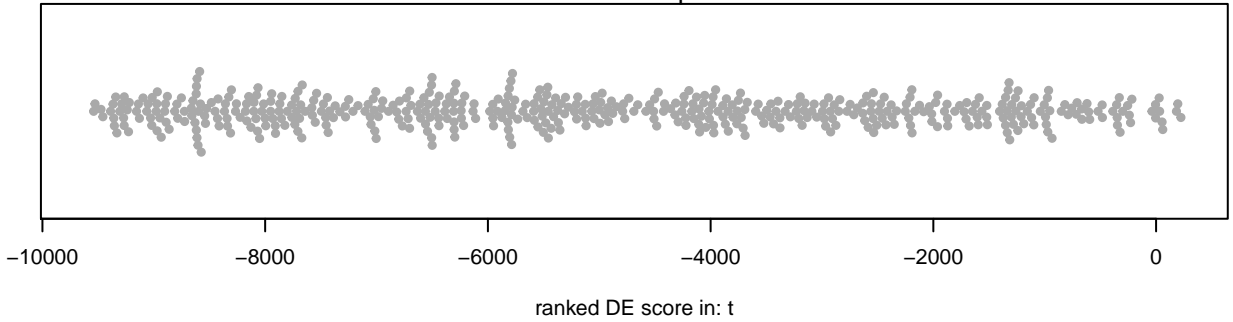


rugplot

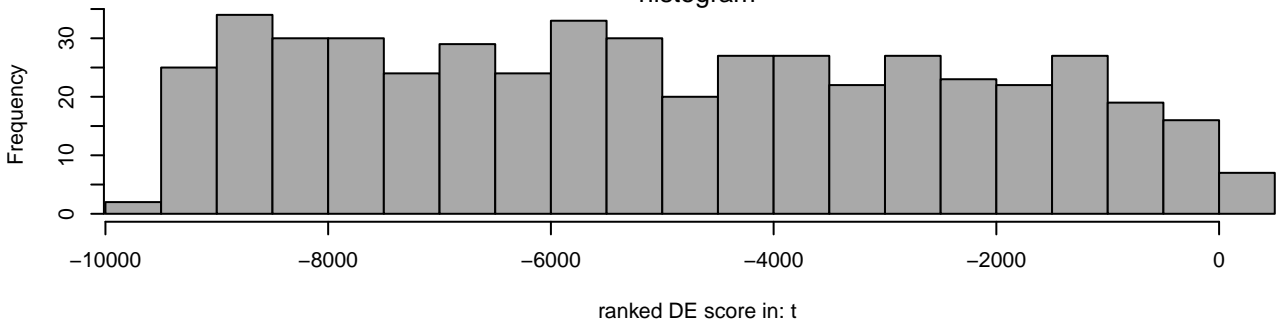


Metabolism of RNA

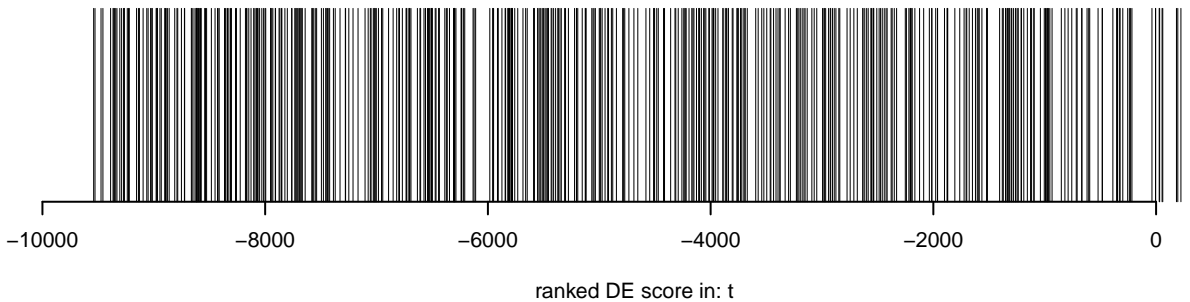
beeswarm plot



histogram

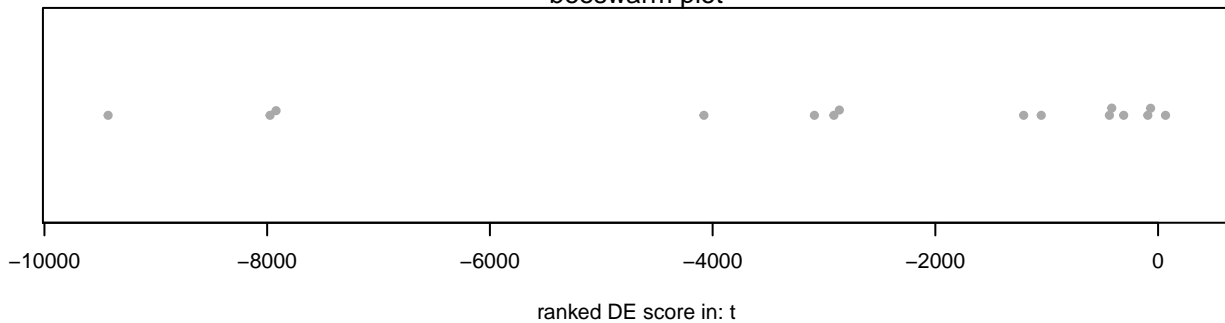


rugplot

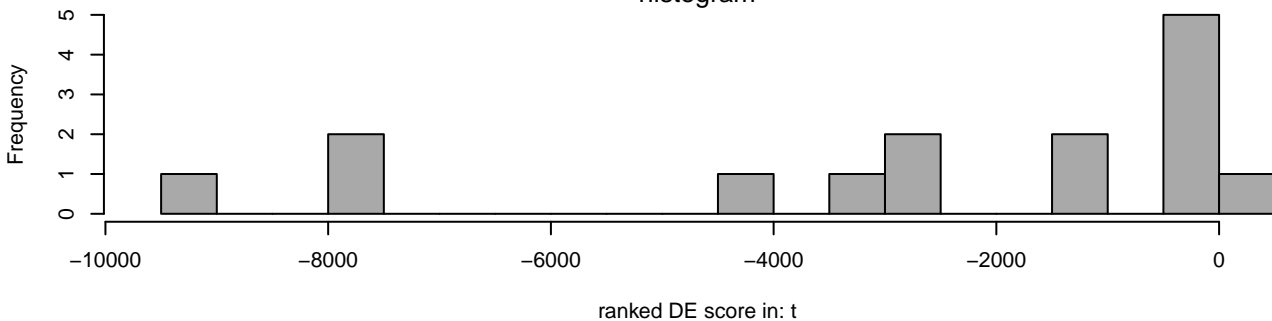


CREB1 phosphorylation through NMDA receptor-mediated activation of RAS signaling

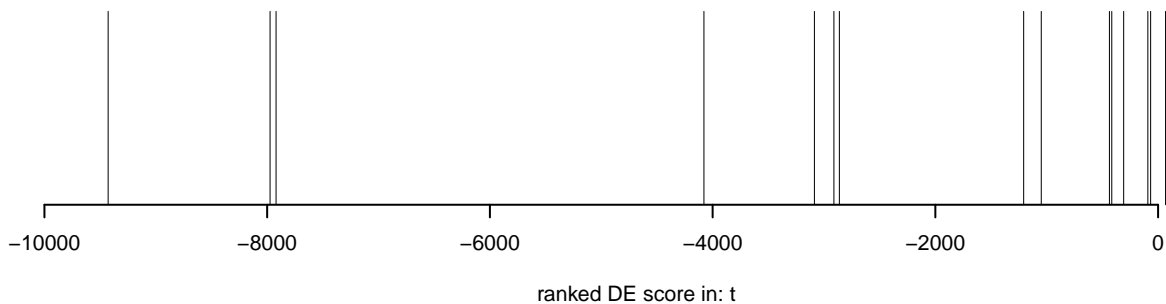
beeswarm plot



histogram

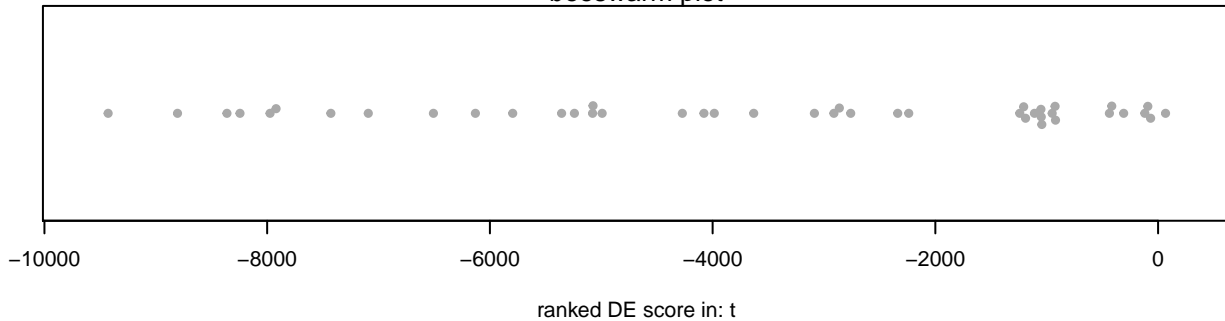


rugplot

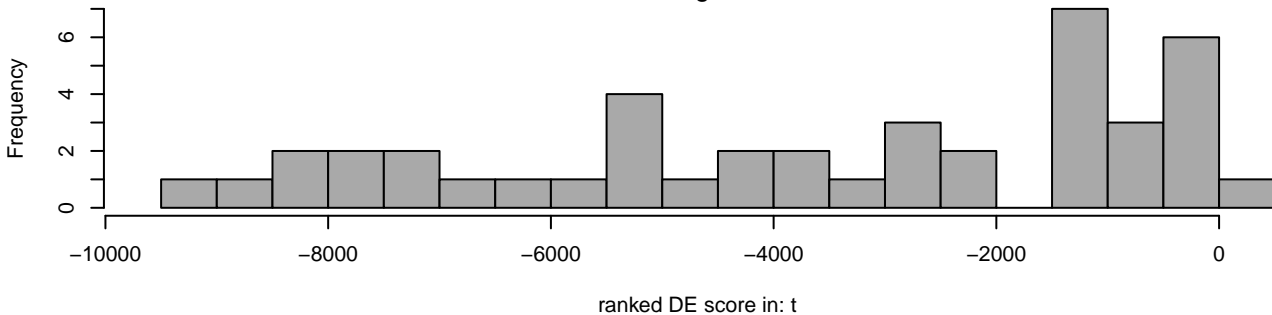


Activation of NMDA receptors and postsynaptic events

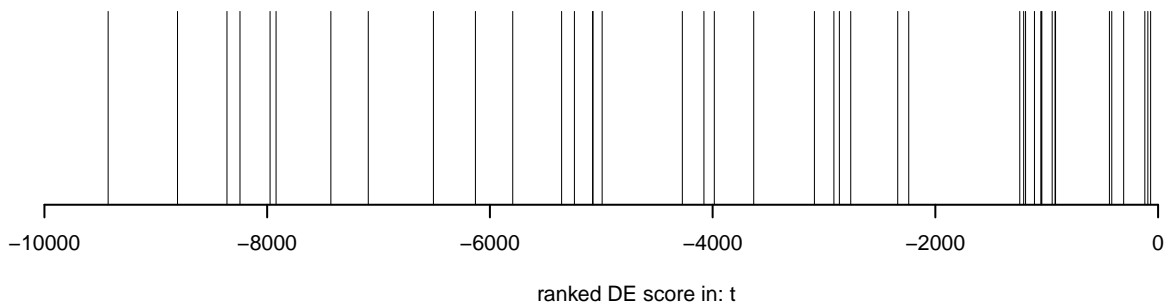
beeswarm plot



histogram

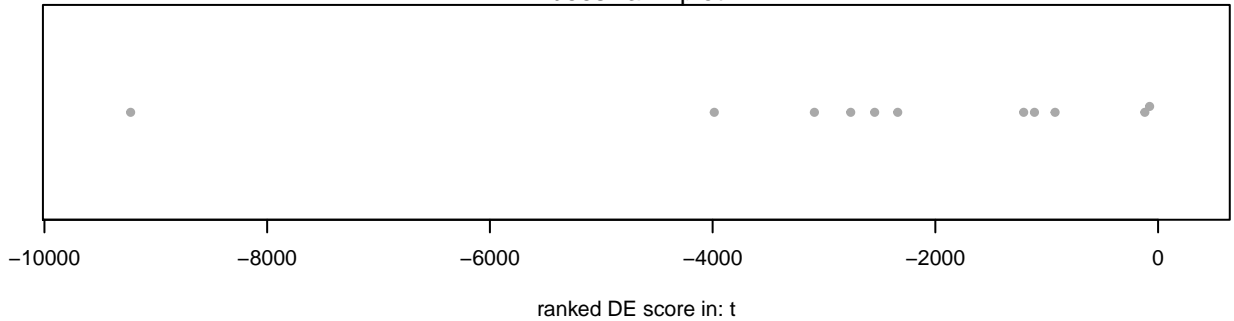


rugplot

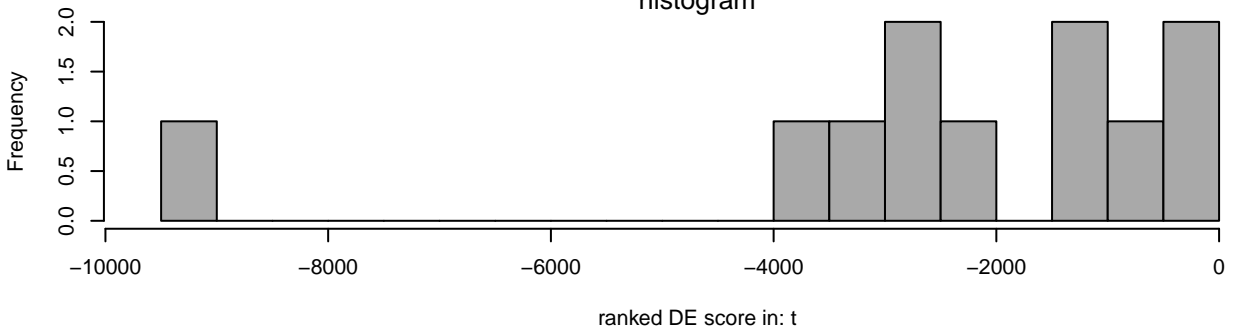


PKA-mediated phosphorylation of CREB

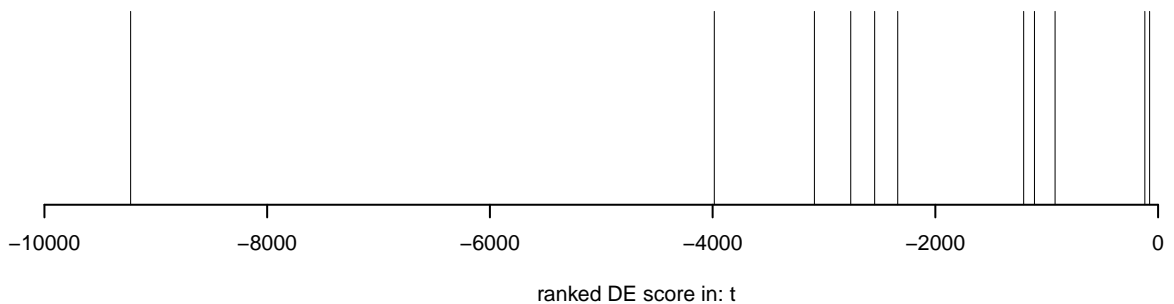
beeswarm plot



histogram

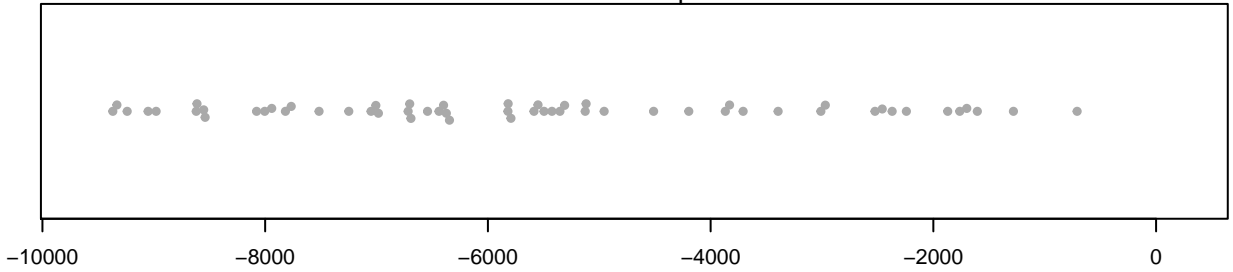


rugplot



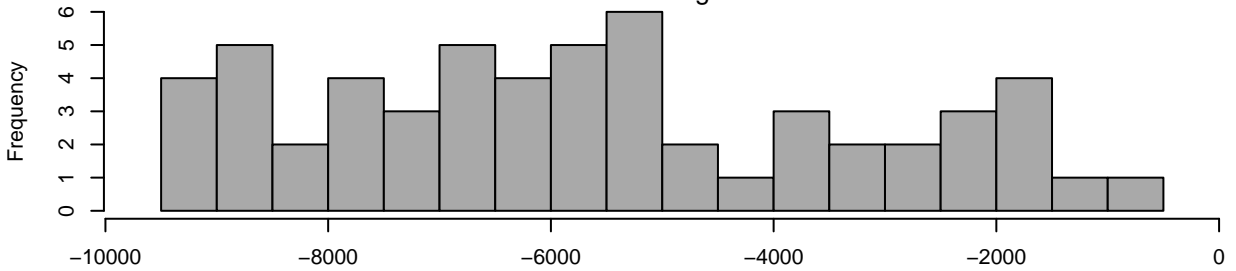
G1/S DNA Damage Checkpoints

beeswarm plot



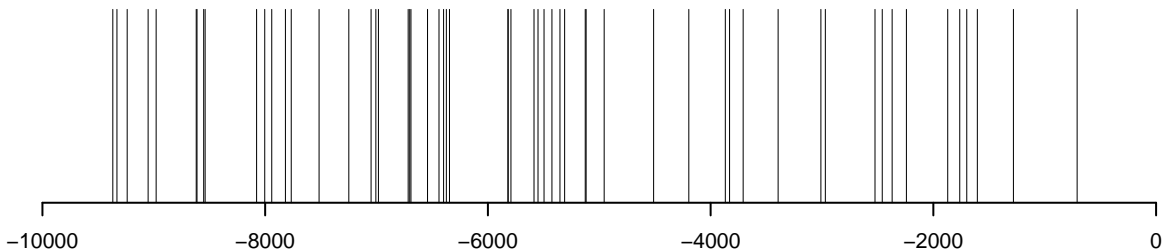
ranked DE score in: t

histogram



ranked DE score in: t

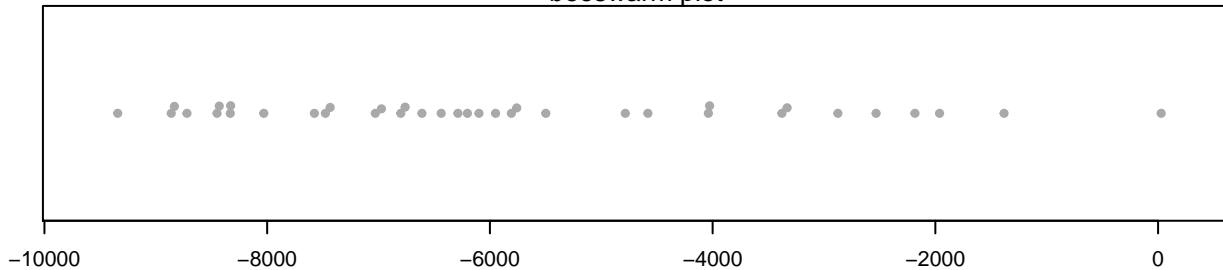
rugplot



ranked DE score in: t

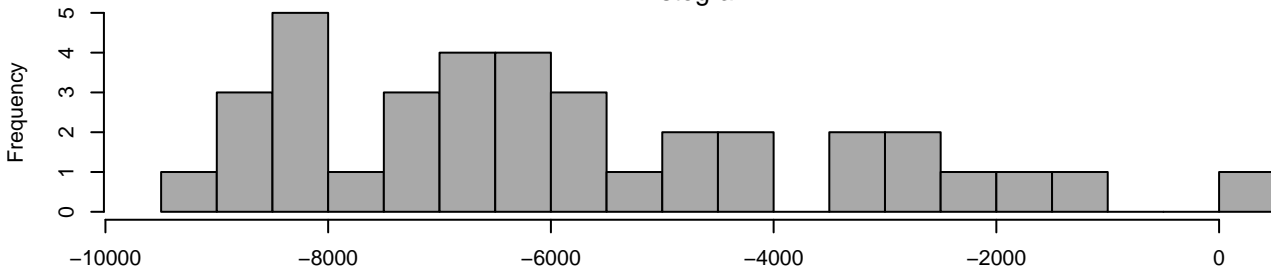
HIV Transcription Initiation

beeswarm plot



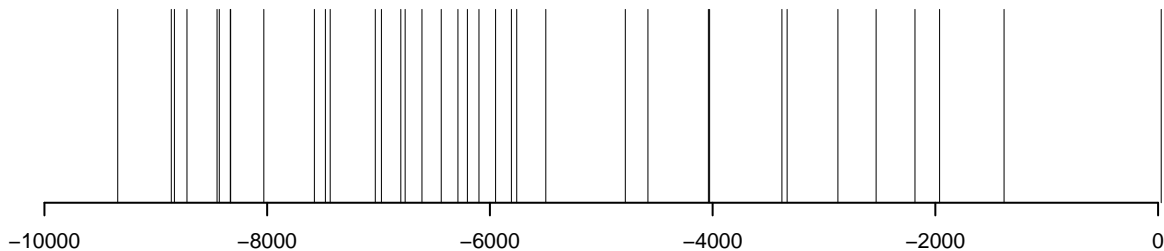
ranked DE score in: t

histogram



ranked DE score in: t

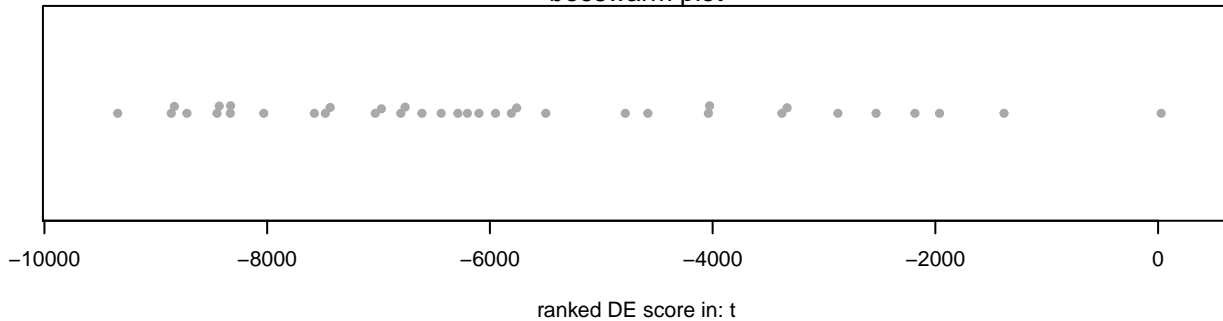
rugplot



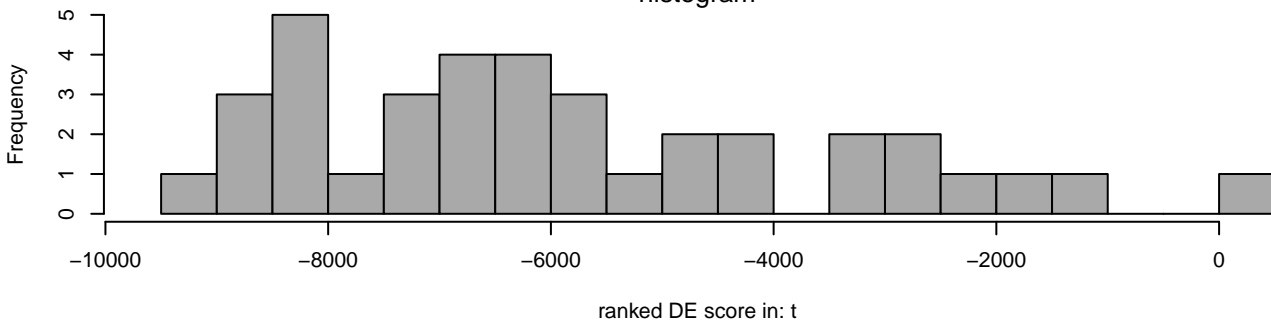
ranked DE score in: t

RNA Polymerase II HIV Promoter Escape

beeswarm plot



histogram



rugplot

