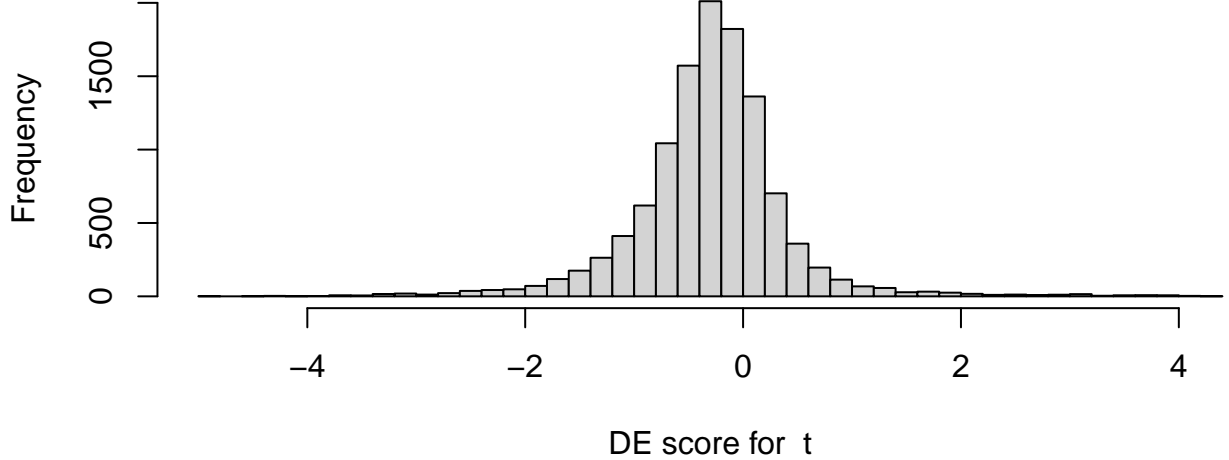


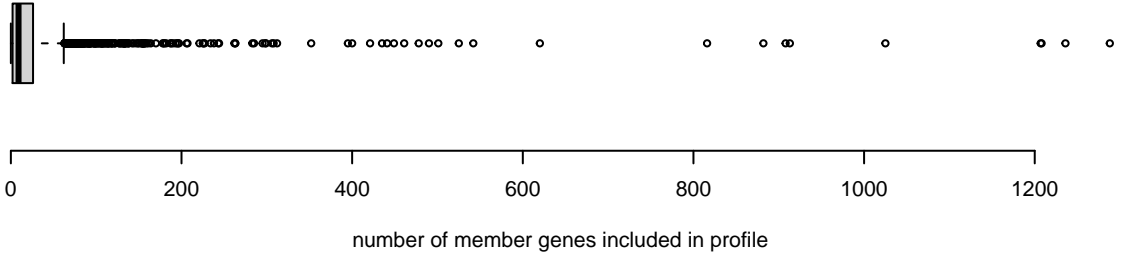
Distribution of DE scores



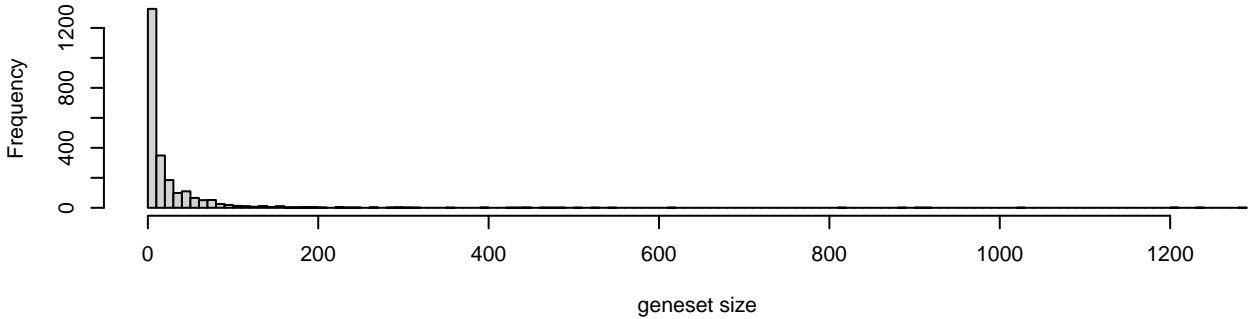
11370 genes in total, 3047 trending up-regulated, 8323 trending down-regulated



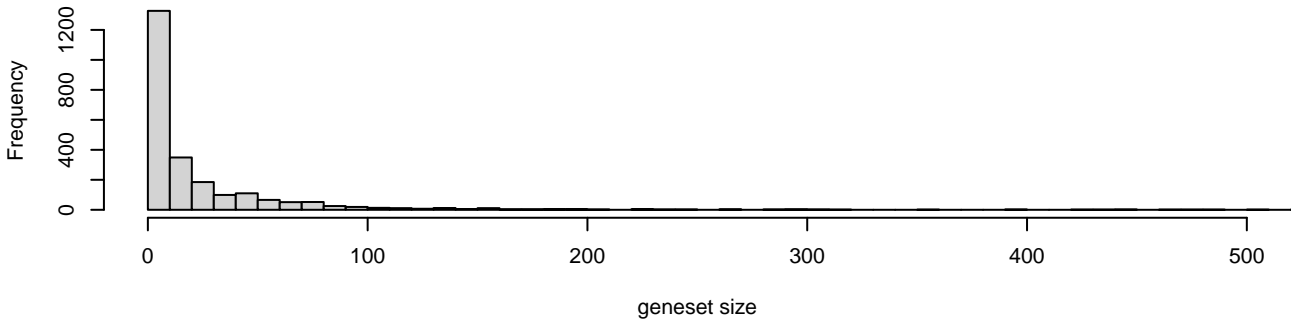
Gene set size



Histogram of geneset size

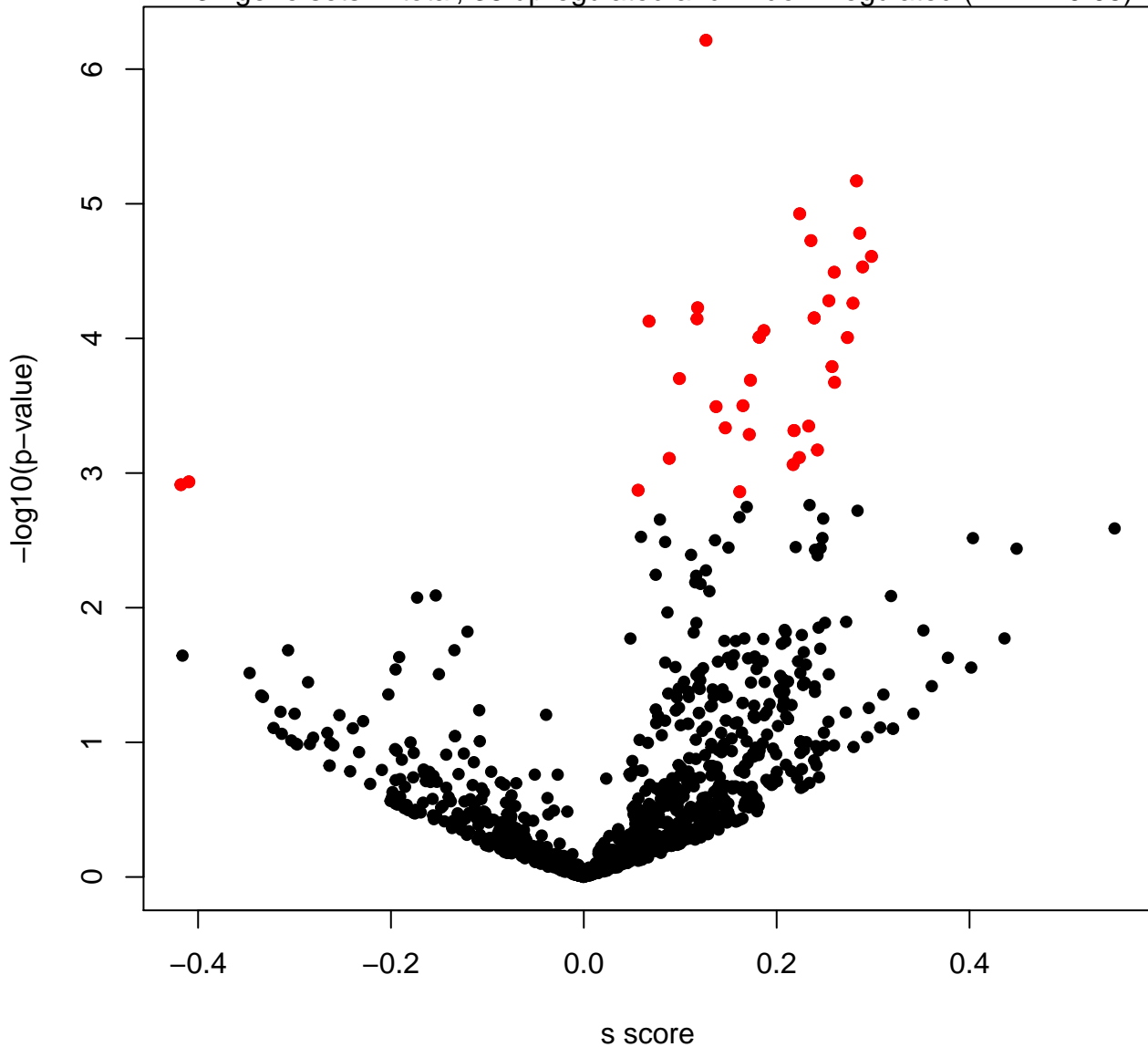


Trimmed histogram of geneset size



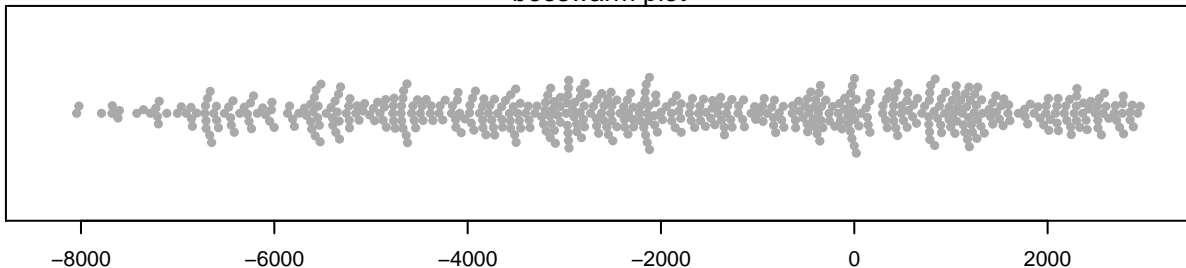
volcano plot of gene set enrichments

1134 gene sets in total, 36 upregulated and 2 downregulated (FDR \leq 0.05)



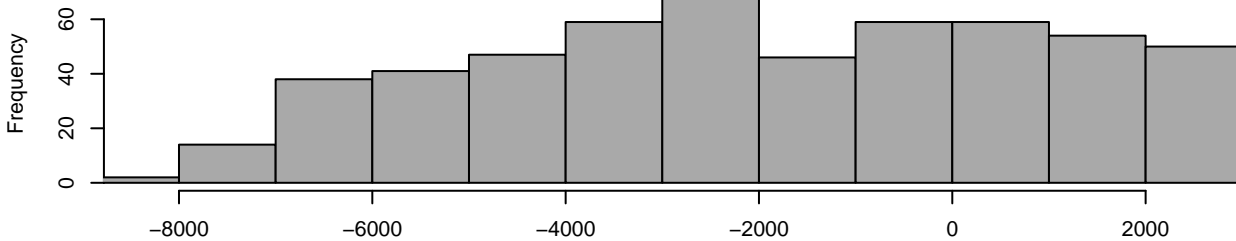
Metabolism of RNA

beeswarm plot



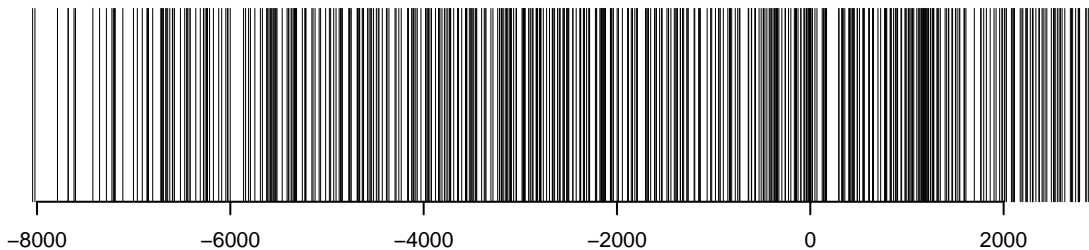
ranked DE score in: t

histogram



ranked DE score in: t

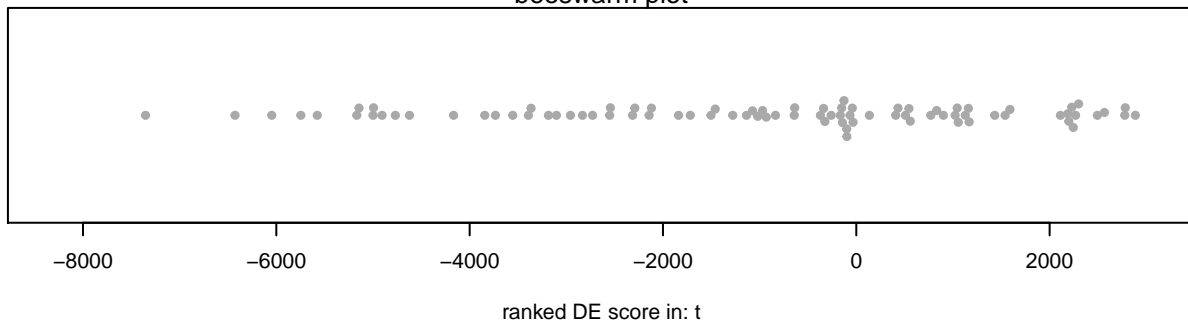
rugplot



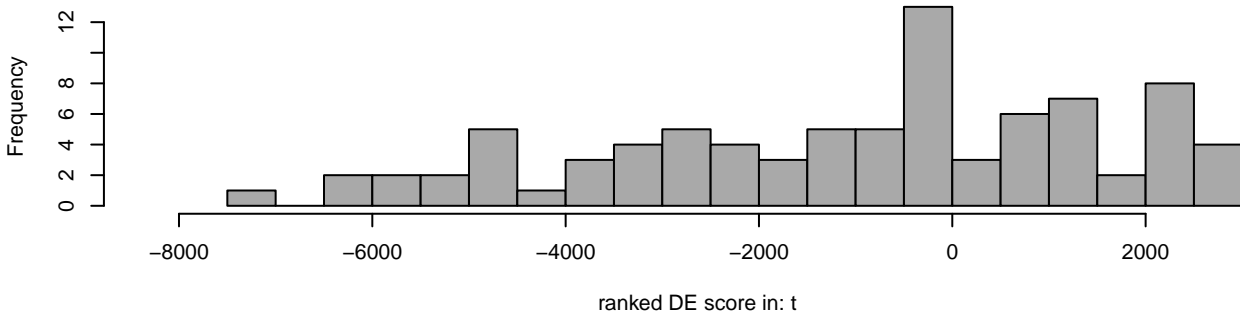
ranked DE score in: t

SRP-dependent cotranslational protein targeting to membrane

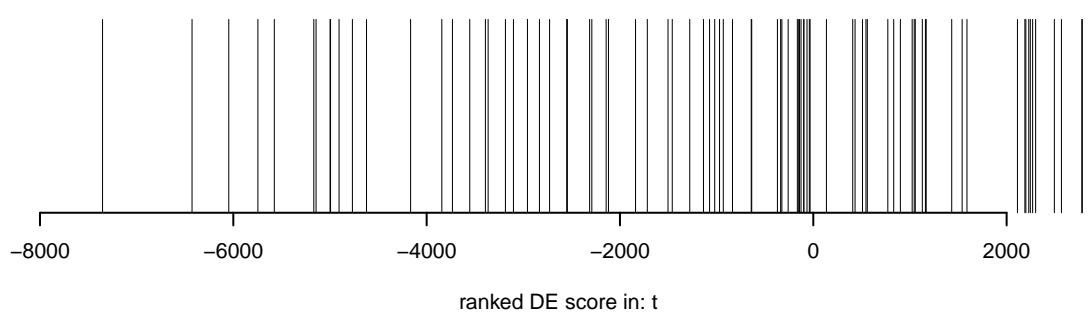
beeswarm plot



histogram

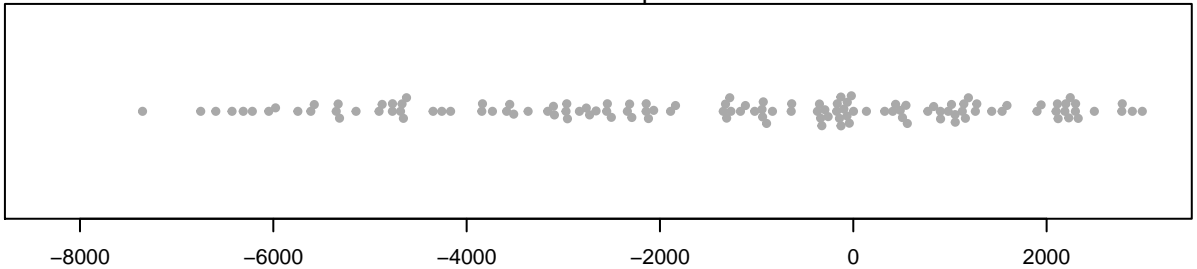


rugplot



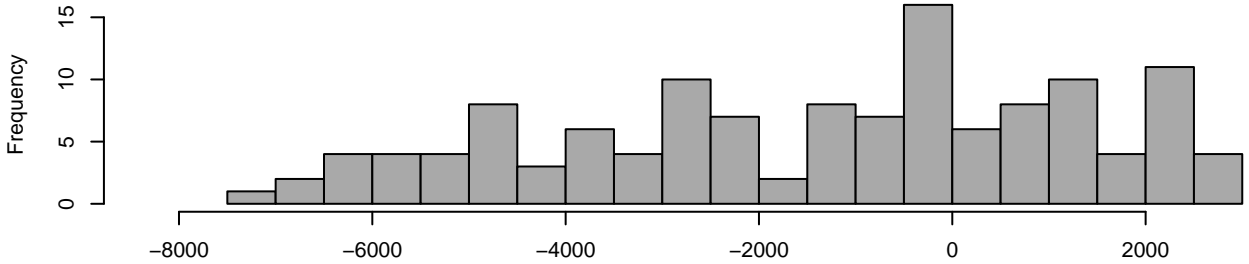
Influenza Infection

beeswarm plot



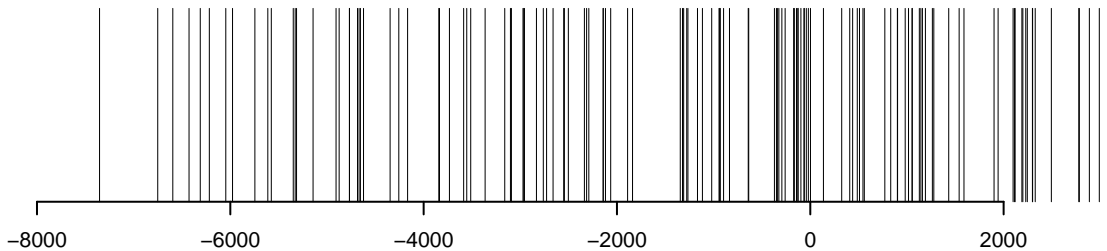
ranked DE score in: t

histogram



ranked DE score in: t

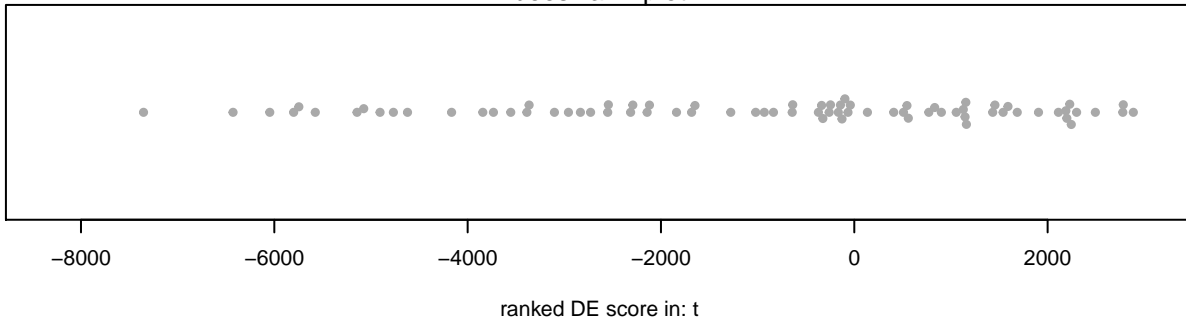
rugplot



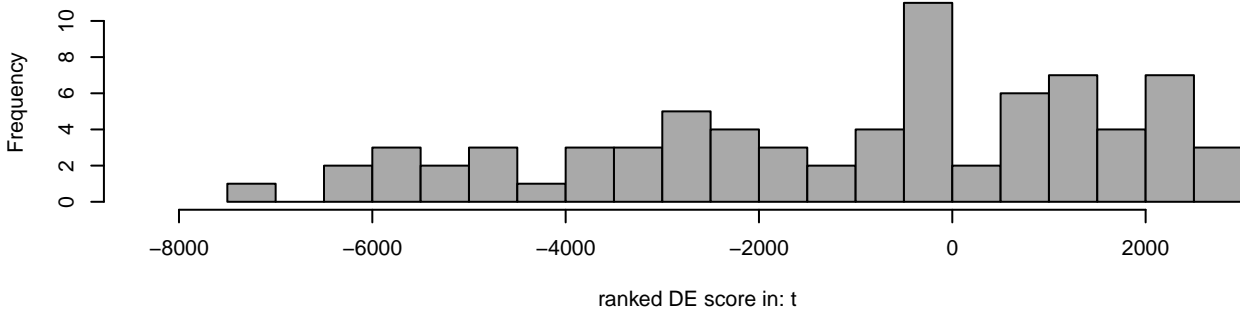
ranked DE score in: t

Formation of a pool of free 40S subunits

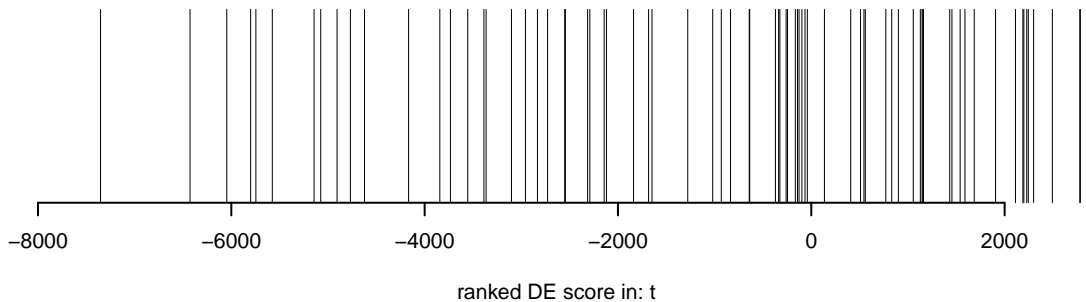
beeswarm plot



histogram

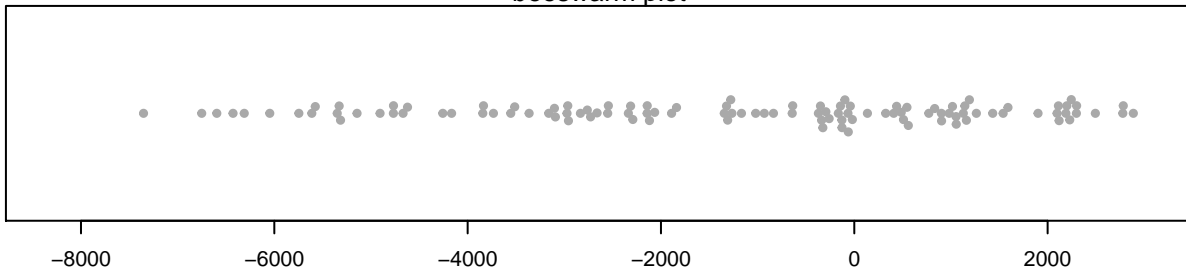


rugplot



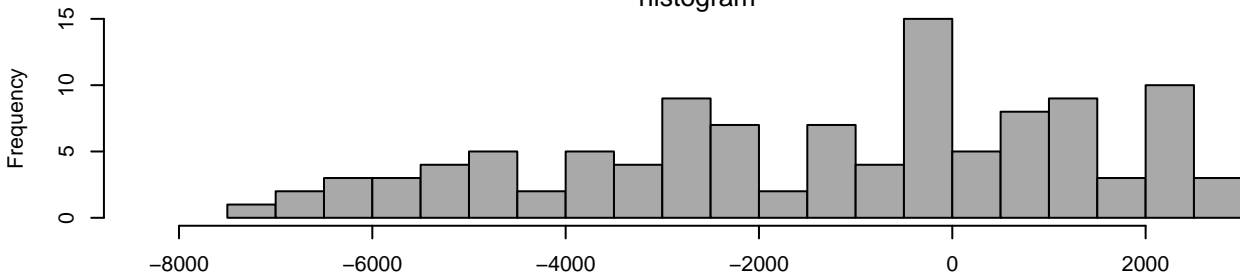
Influenza Viral RNA Transcription and Replication

beeswarm plot



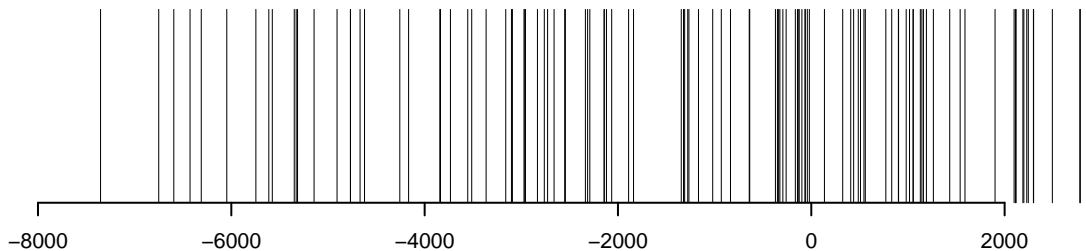
ranked DE score in: t

histogram



ranked DE score in: t

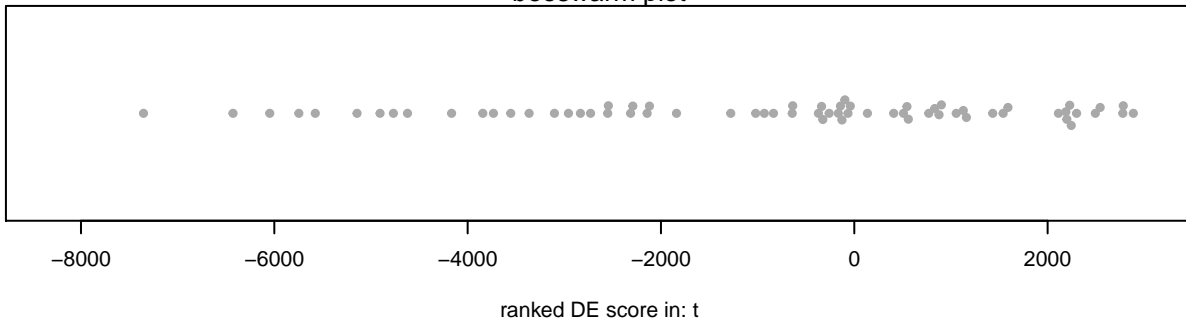
rugplot



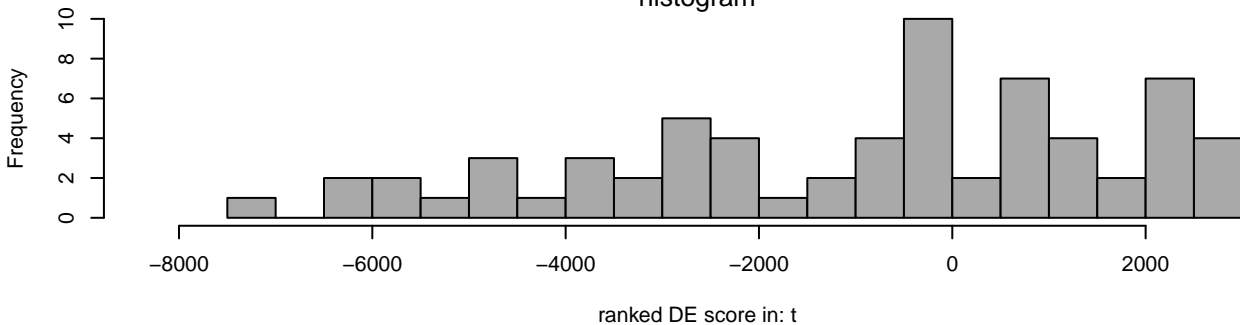
ranked DE score in: t

Peptide chain elongation

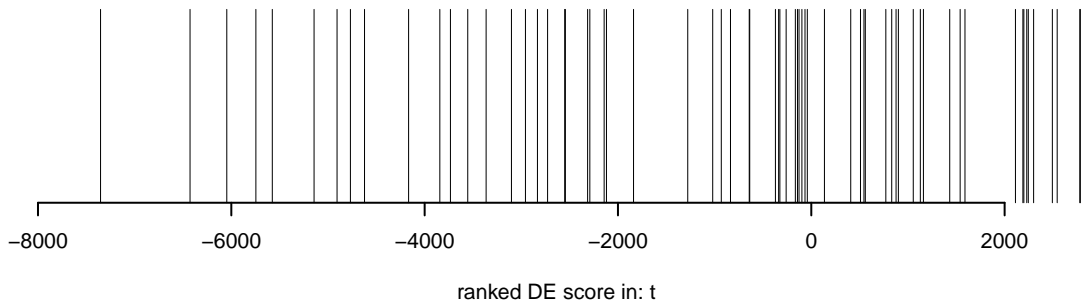
beeswarm plot



histogram

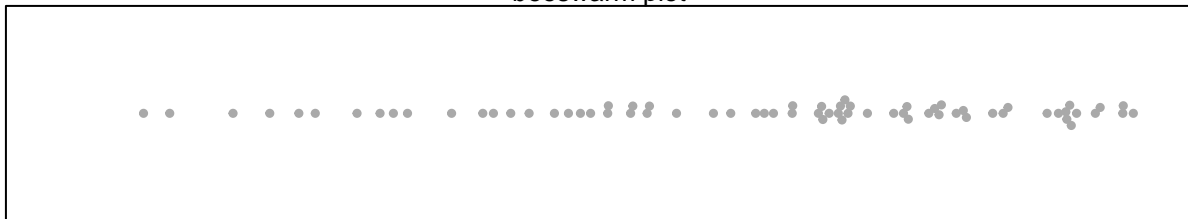


rugplot



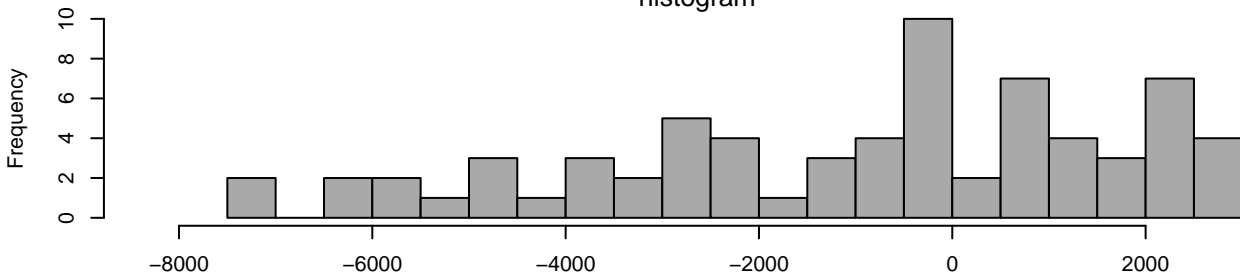
Eukaryotic Translation Elongation

beeswarm plot



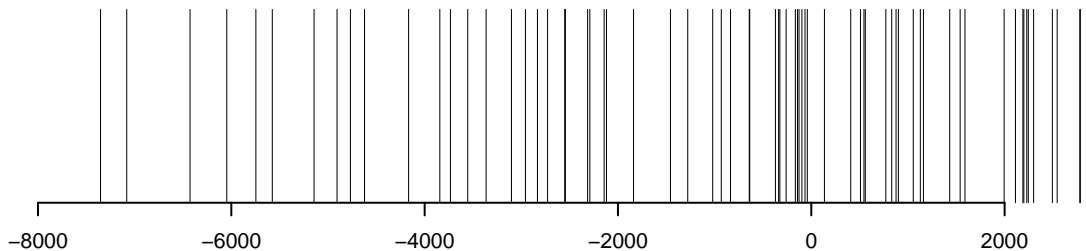
ranked DE score in: t

histogram



ranked DE score in: t

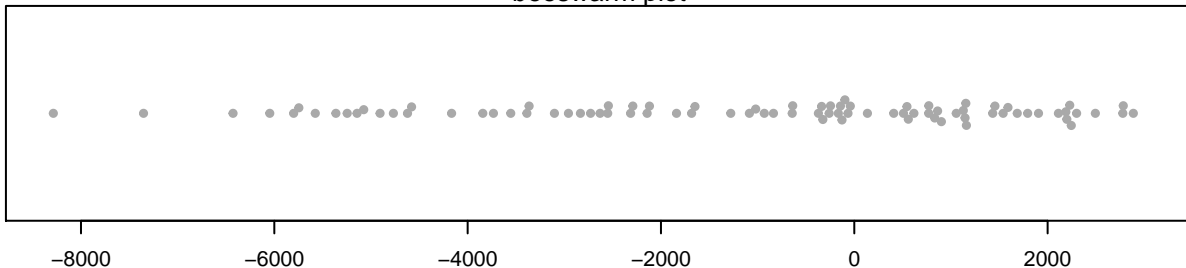
rugplot



ranked DE score in: t

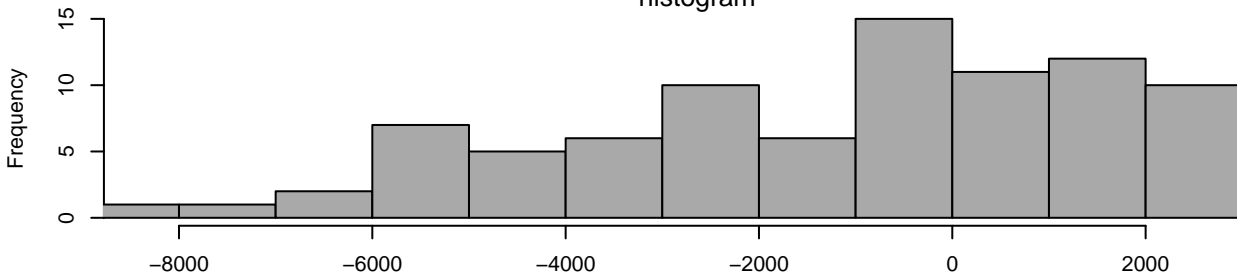
GTP hydrolysis and joining of the 60S ribosomal subunit

beeswarm plot



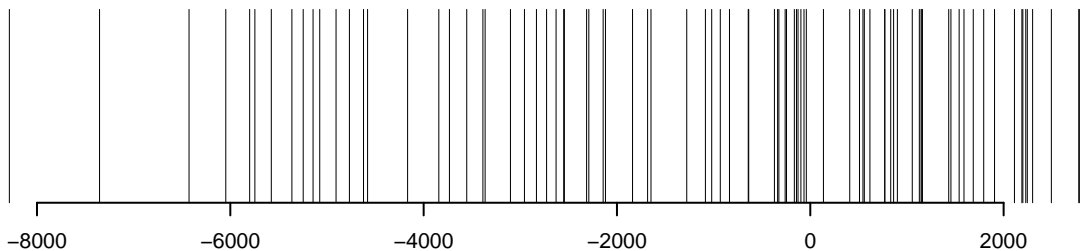
ranked DE score in: t

histogram



ranked DE score in: t

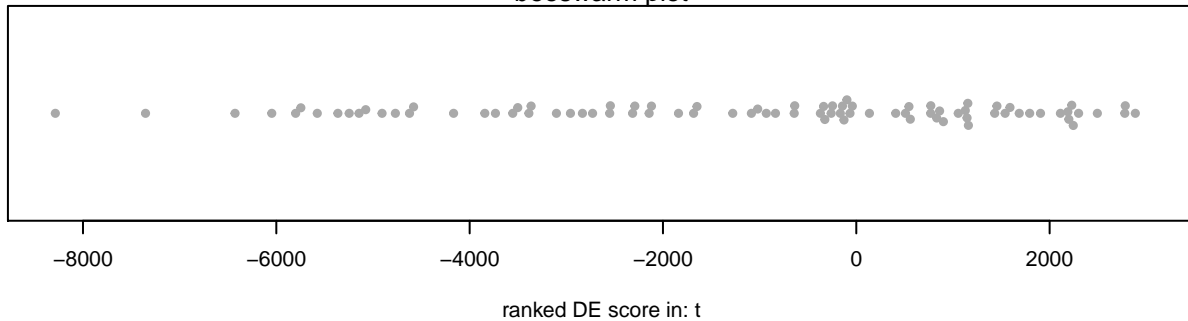
rugplot



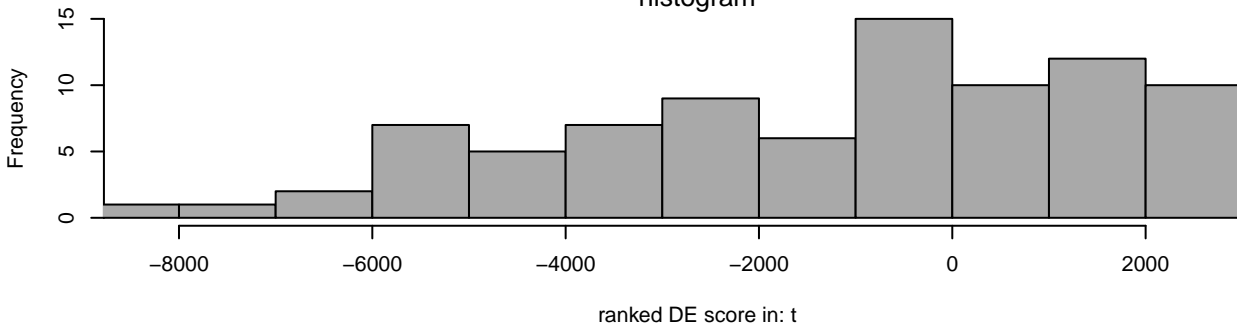
ranked DE score in: t

L13a-mediated translational silencing of Ceruloplasmin expression

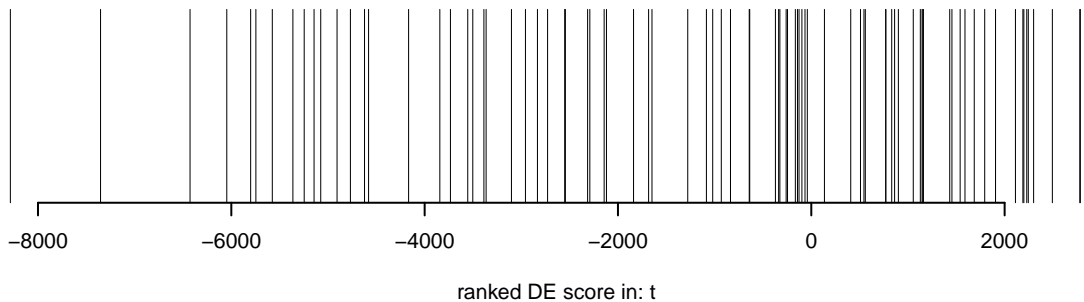
beeswarm plot



histogram

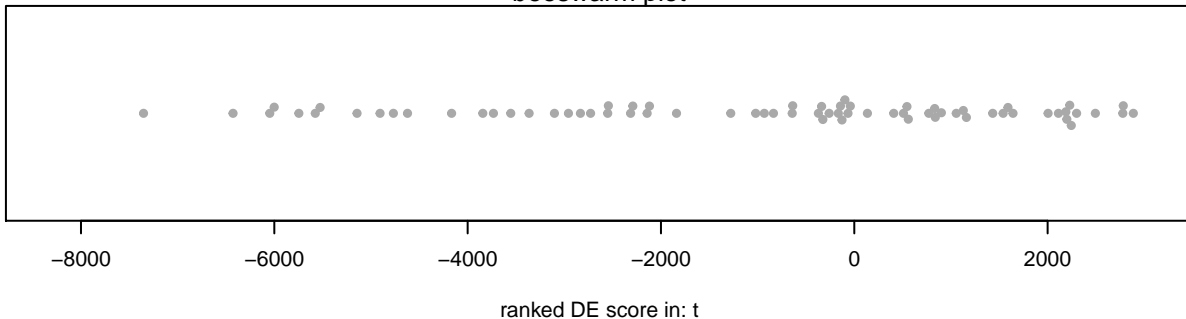


rugplot

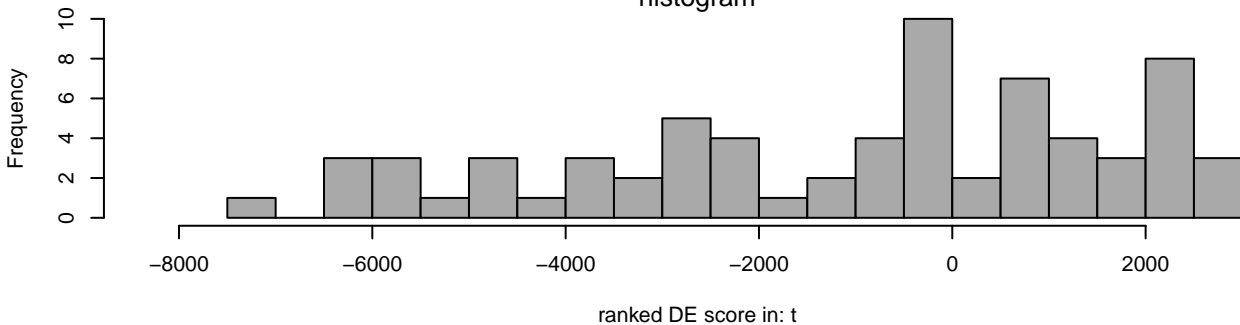


Eukaryotic Translation Termination

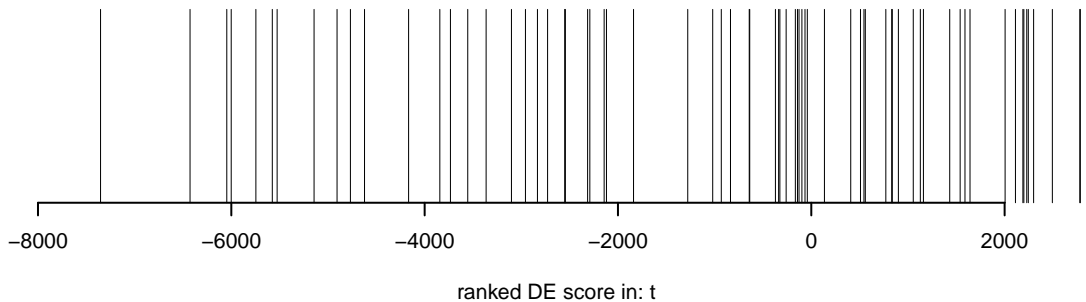
beeswarm plot



histogram

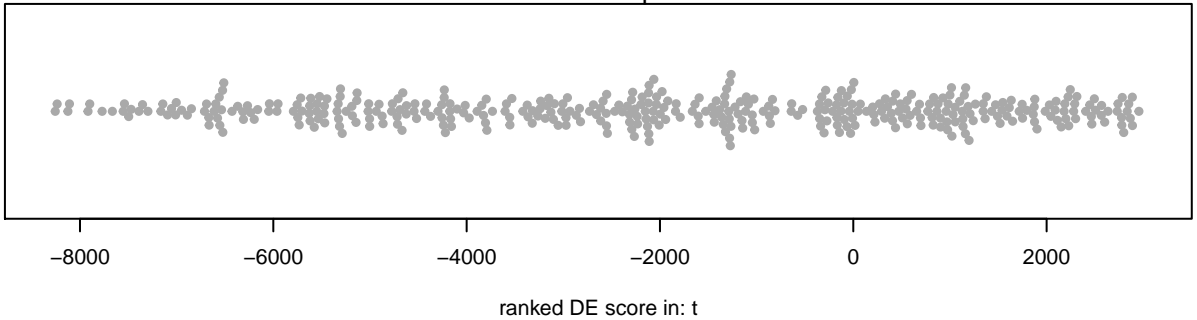


rugplot

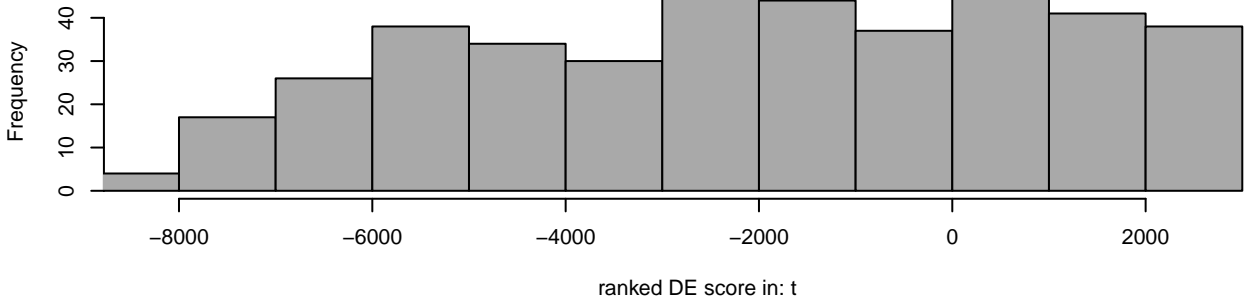


Cellular responses to external stimuli

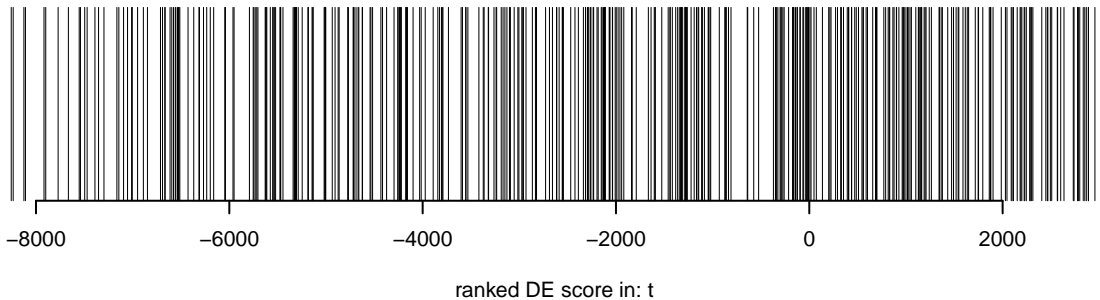
beeswarm plot



histogram

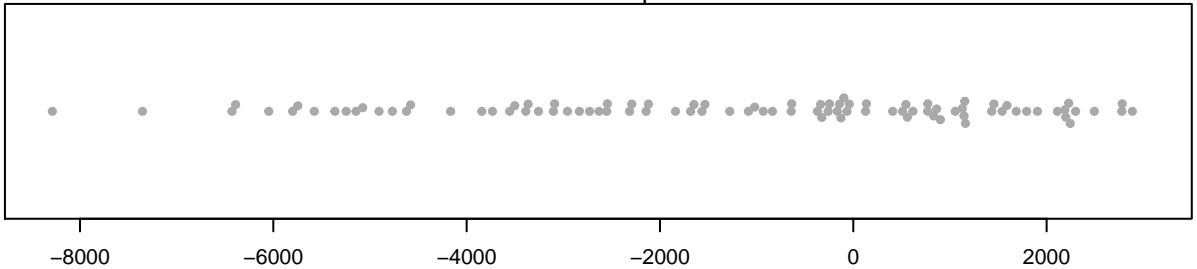


rugplot



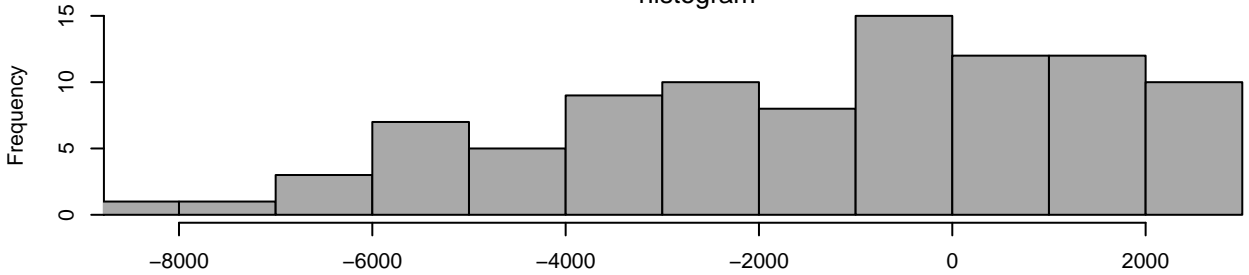
Cap-dependent Translation Initiation

beeswarm plot



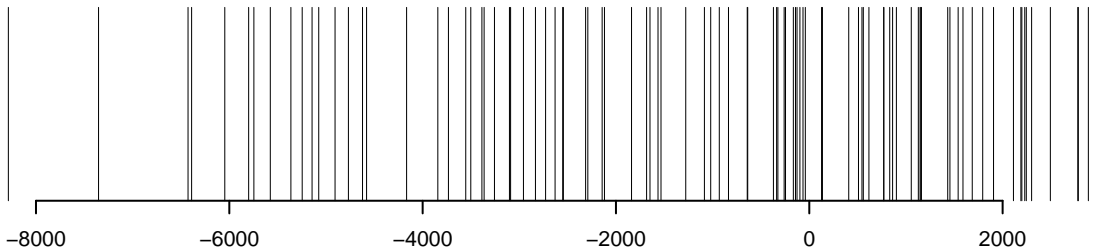
ranked DE score in: t

histogram



ranked DE score in: t

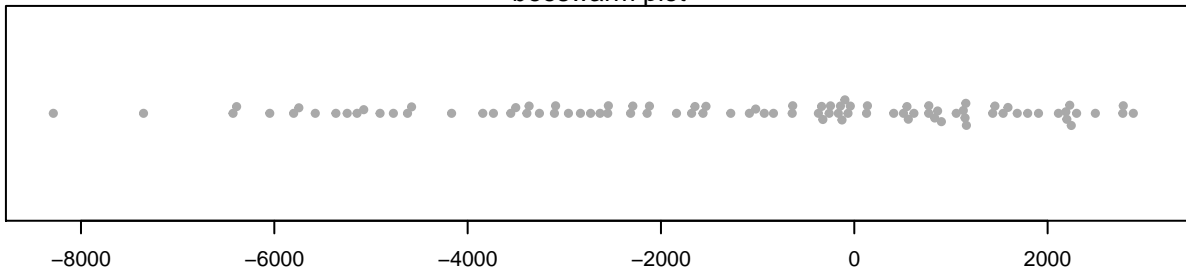
rugplot



ranked DE score in: t

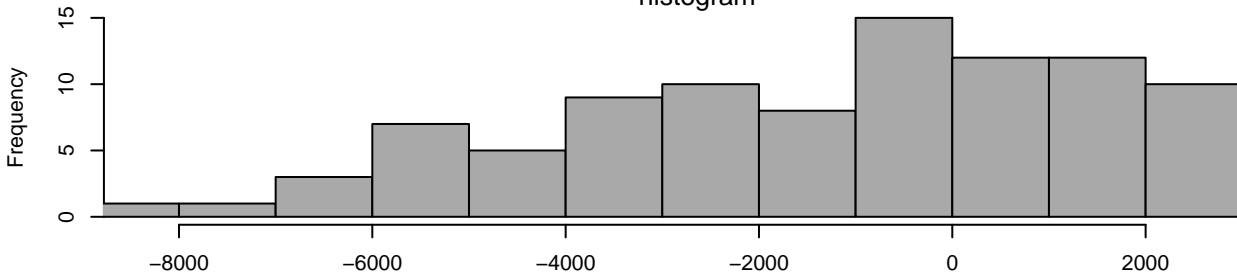
Eukaryotic Translation Initiation

beeswarm plot



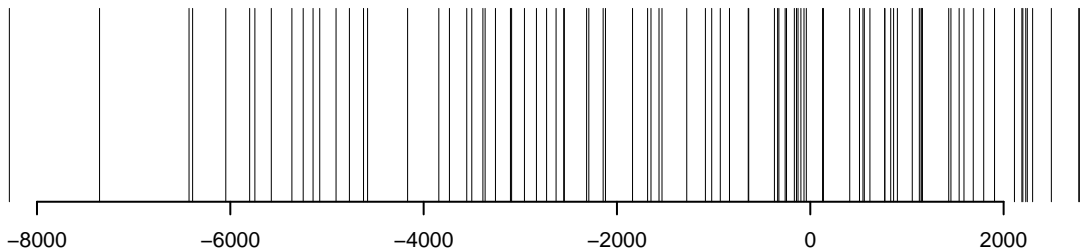
ranked DE score in: t

histogram



ranked DE score in: t

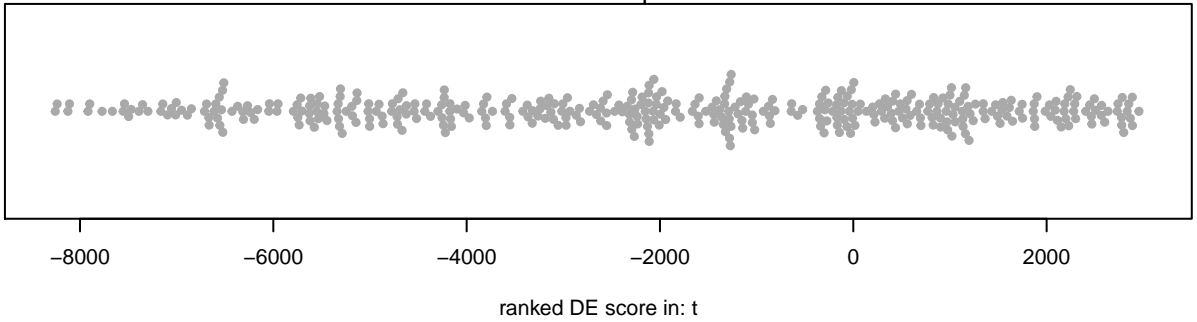
rugplot



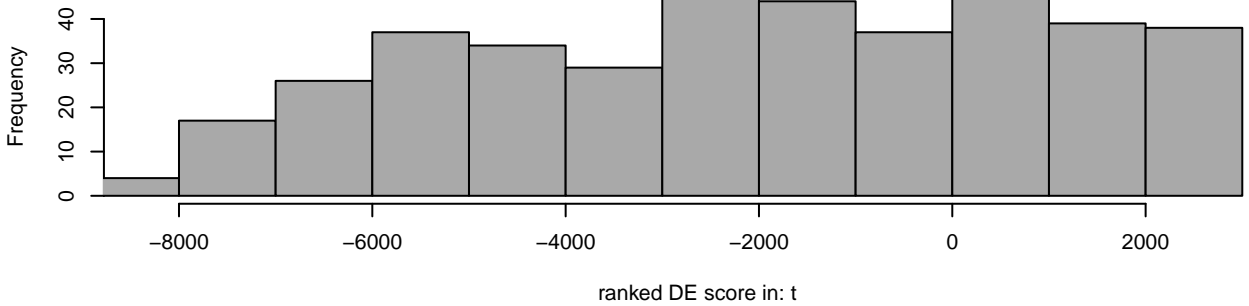
ranked DE score in: t

Cellular responses to stress

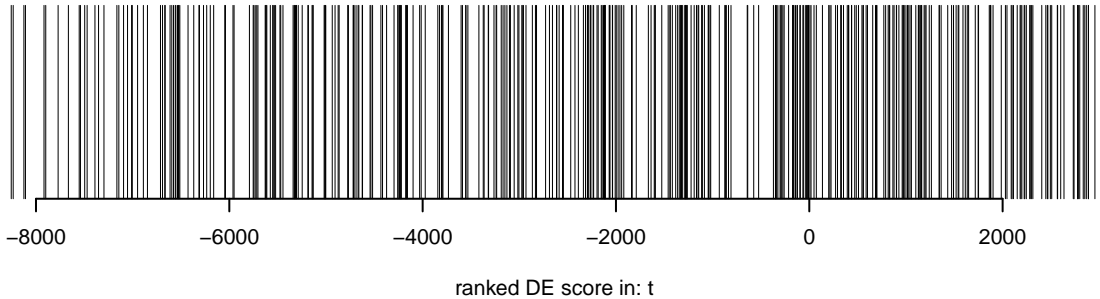
beeswarm plot



histogram

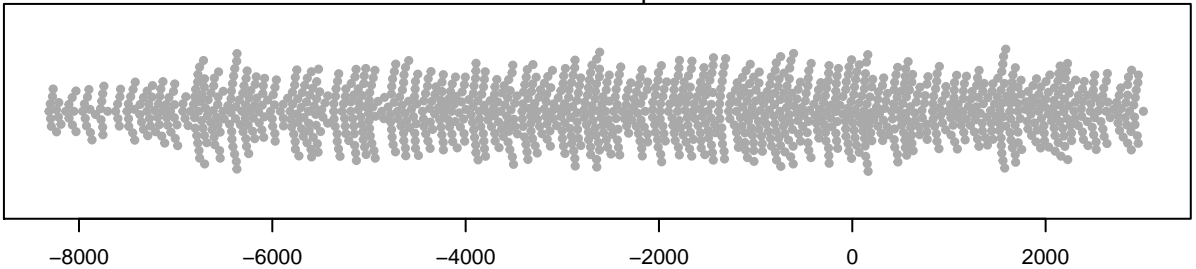


rugplot



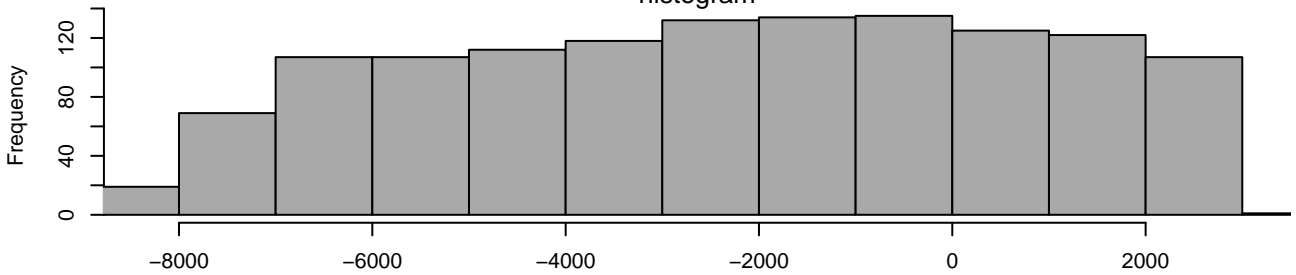
Metabolism of proteins

beeswarm plot



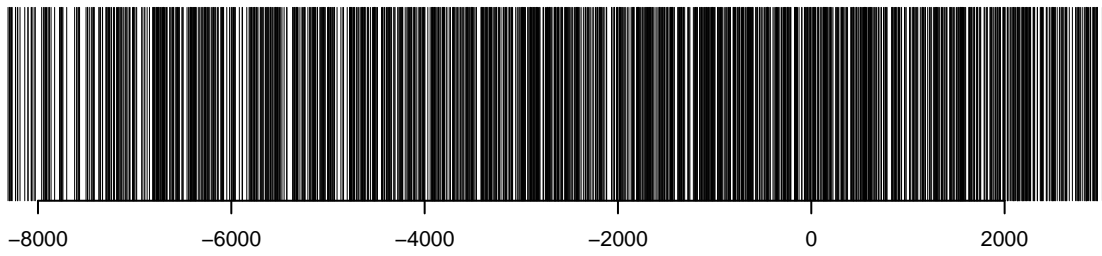
ranked DE score in: t

histogram



ranked DE score in: t

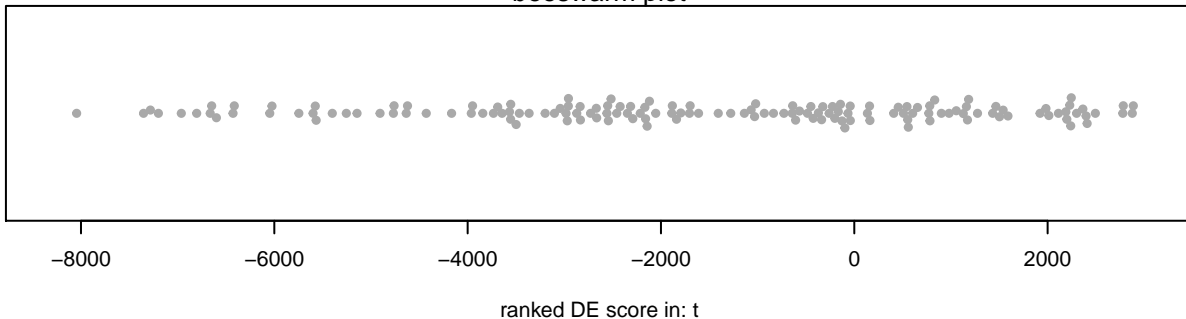
rugplot



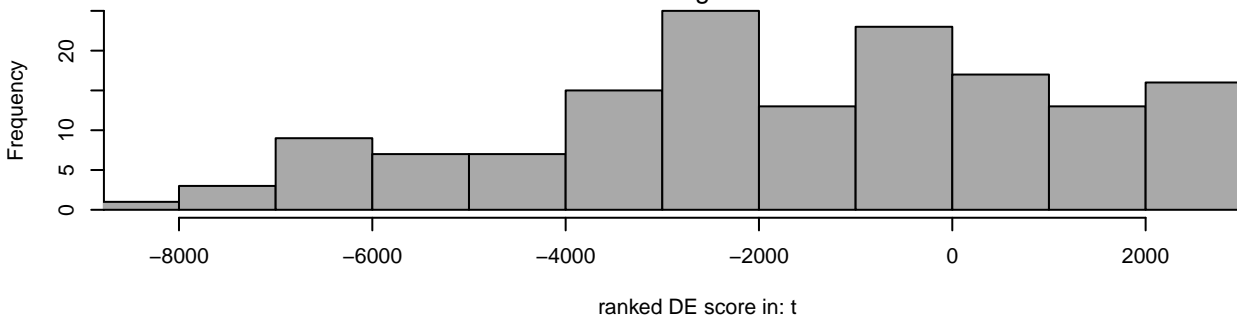
ranked DE score in: t

Major pathway of rRNA processing in the nucleolus and cytosol

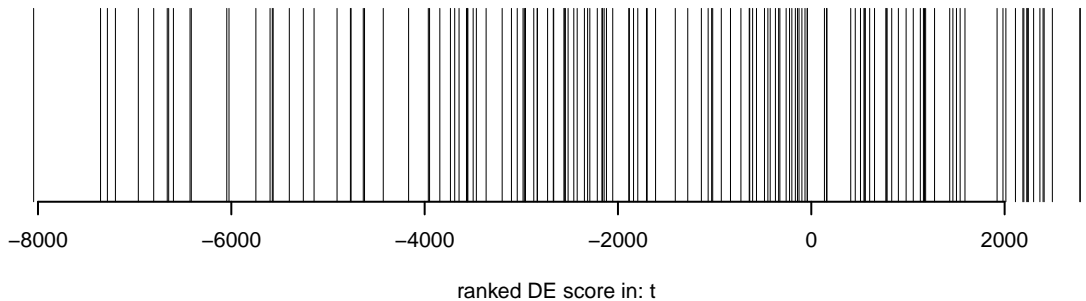
beeswarm plot



histogram

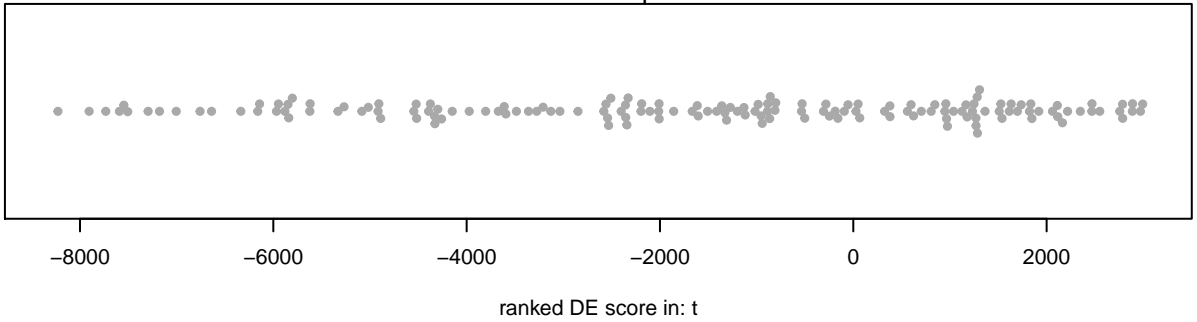


rugplot

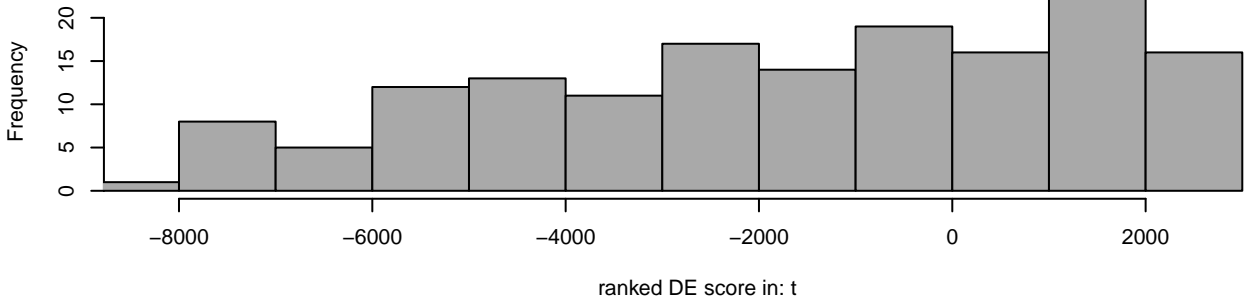


Chromatin modifying enzymes

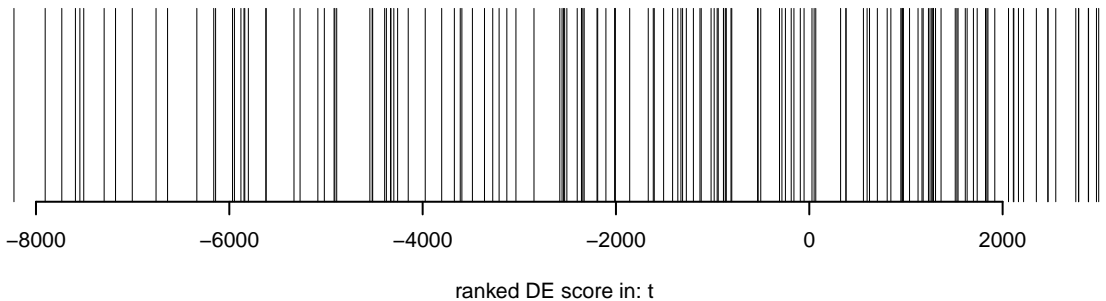
beeswarm plot



histogram

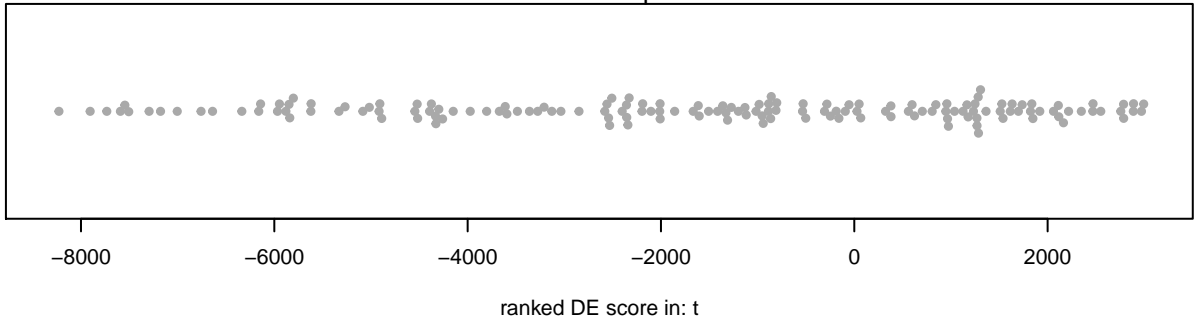


rugplot

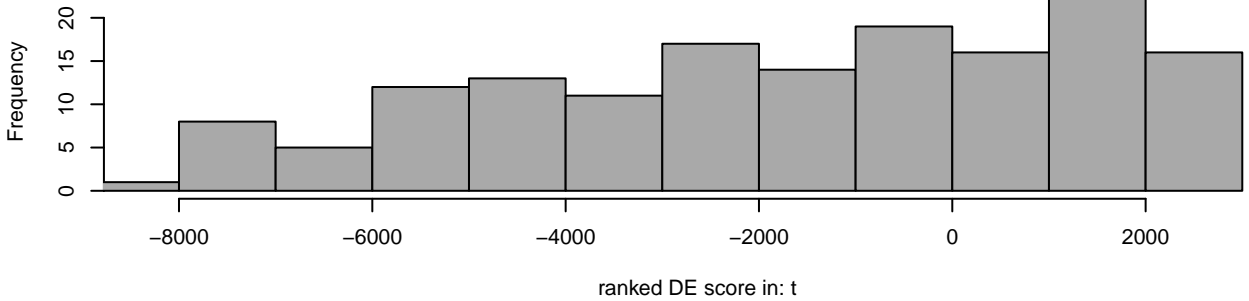


Chromatin organization

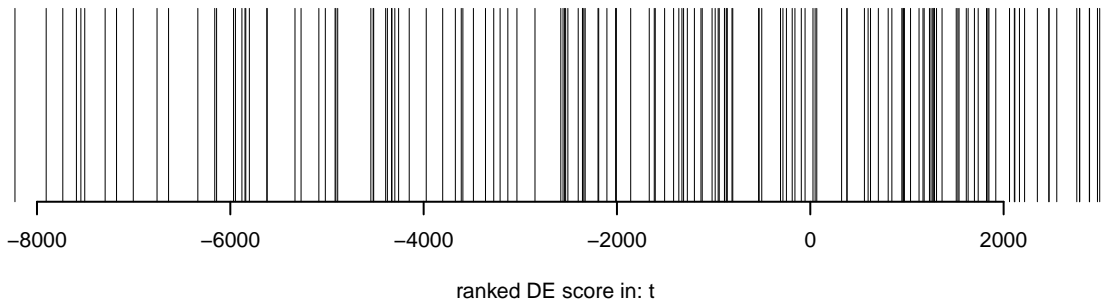
beeswarm plot



histogram

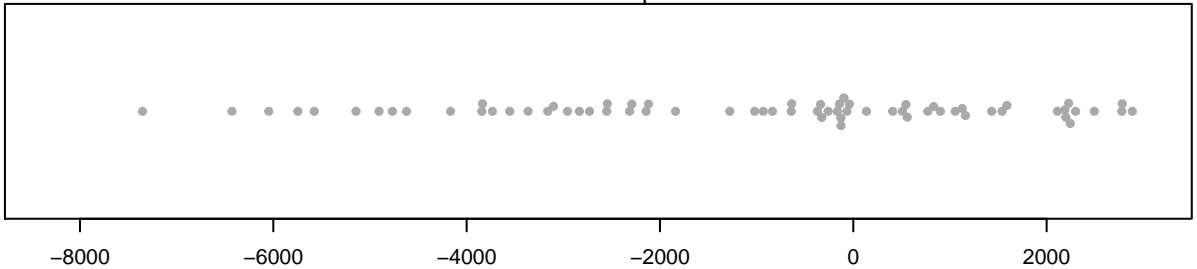


rugplot



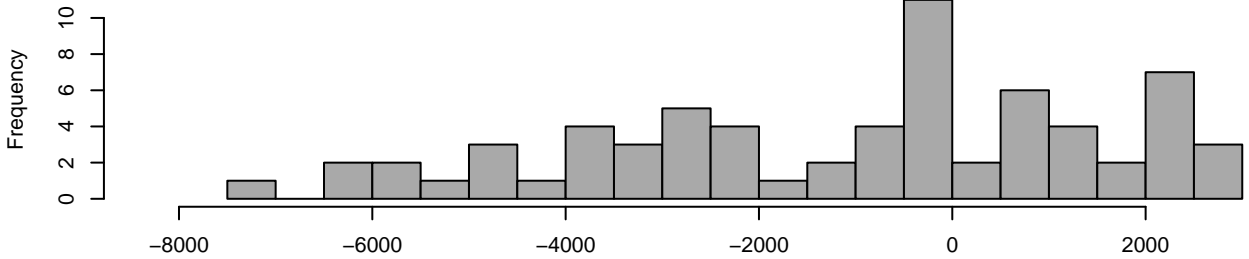
Viral mRNA Translation

beeswarm plot



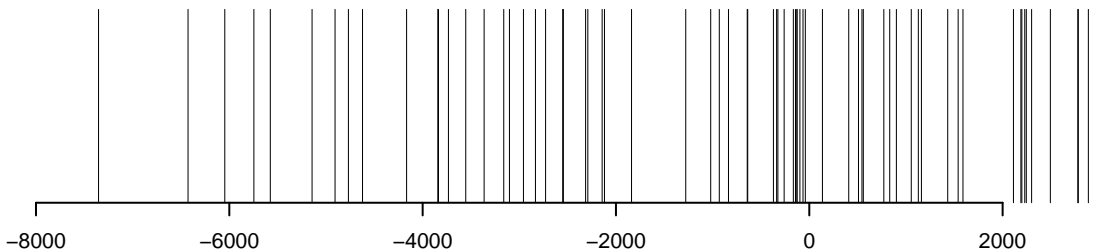
ranked DE score in: t

histogram



ranked DE score in: t

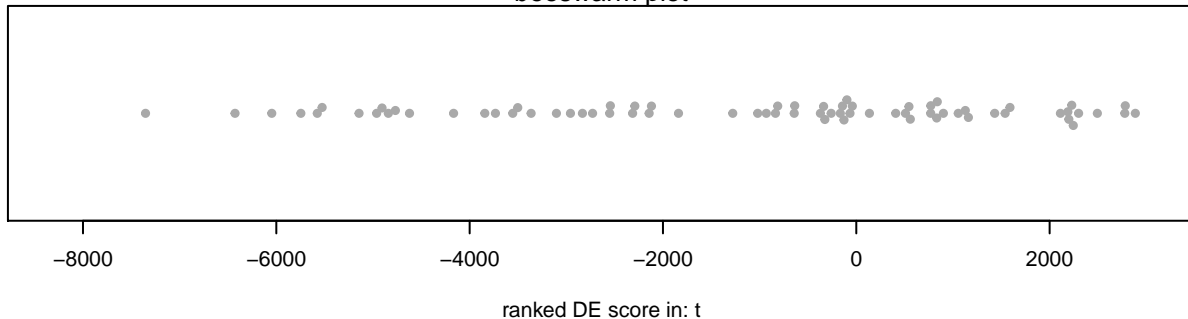
rugplot



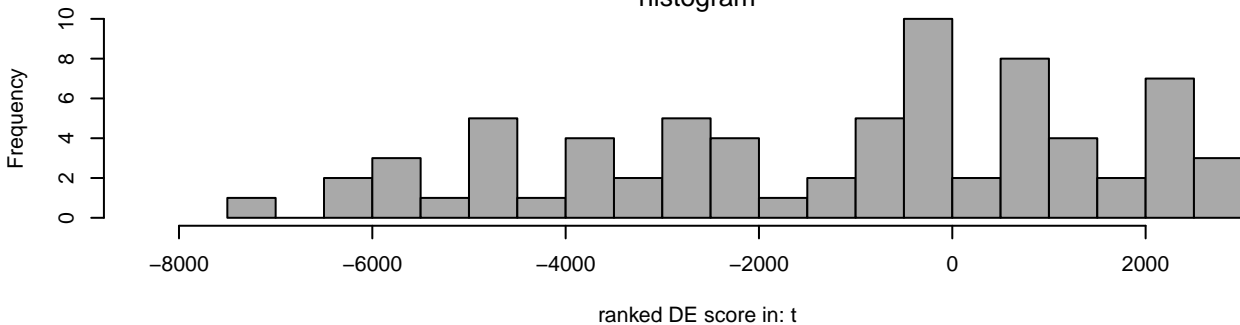
ranked DE score in: t

Nonsense Mediated Decay (NMD) independent of the Exon Junction Complex (EJC)

beeswarm plot



histogram



rugplot

