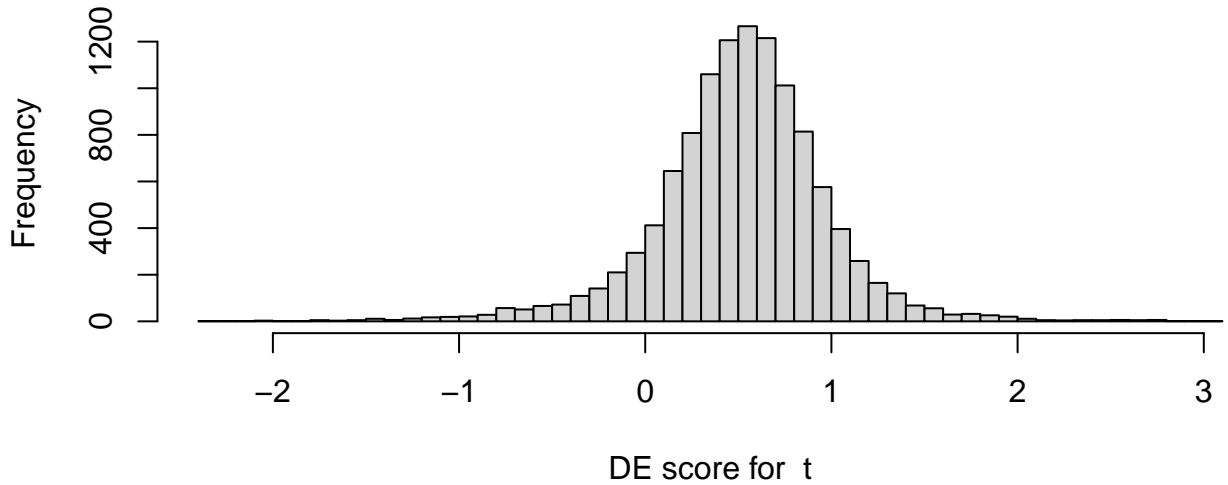
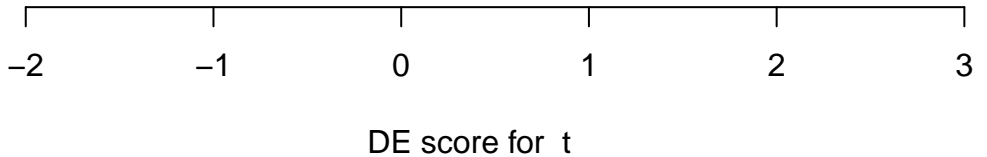


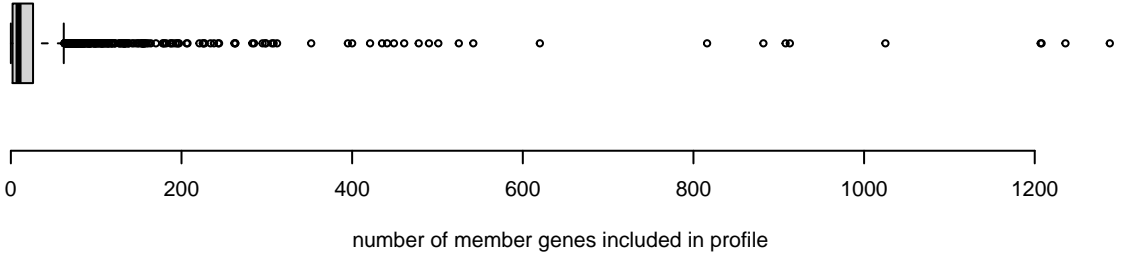
Distribution of DE scores



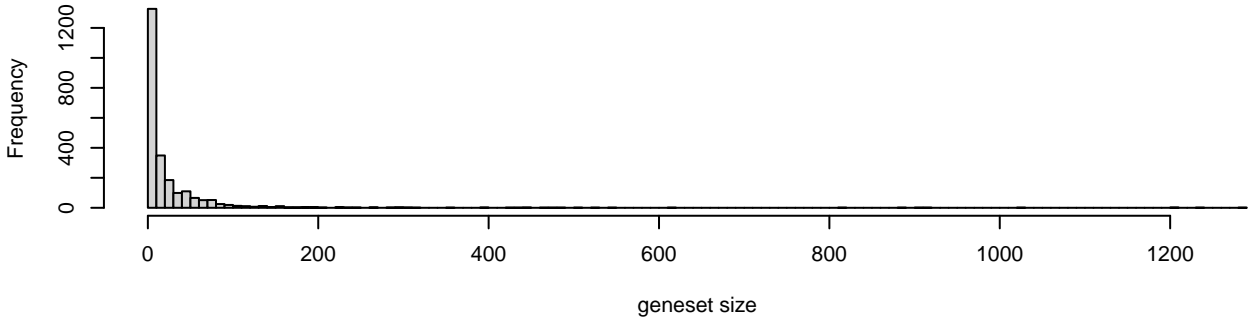
11370 genes in total, 10235 trending up-regulated, 1135 trending down-regulated



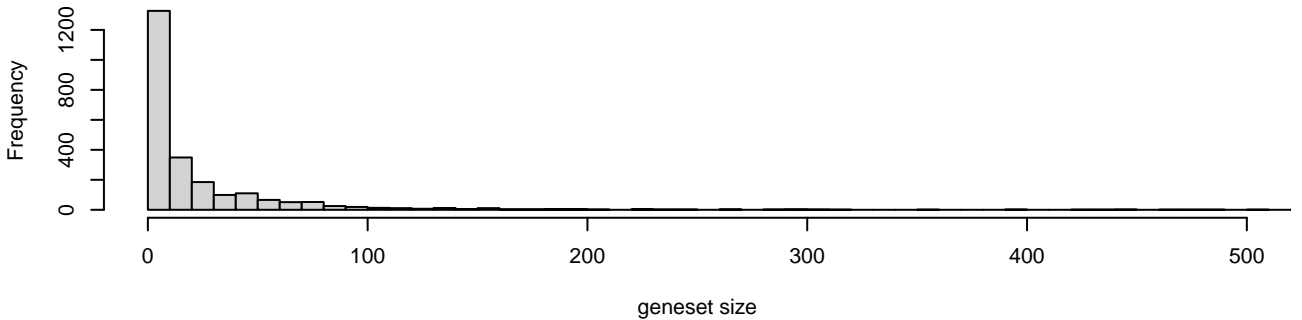
Gene set size



Histogram of geneset size

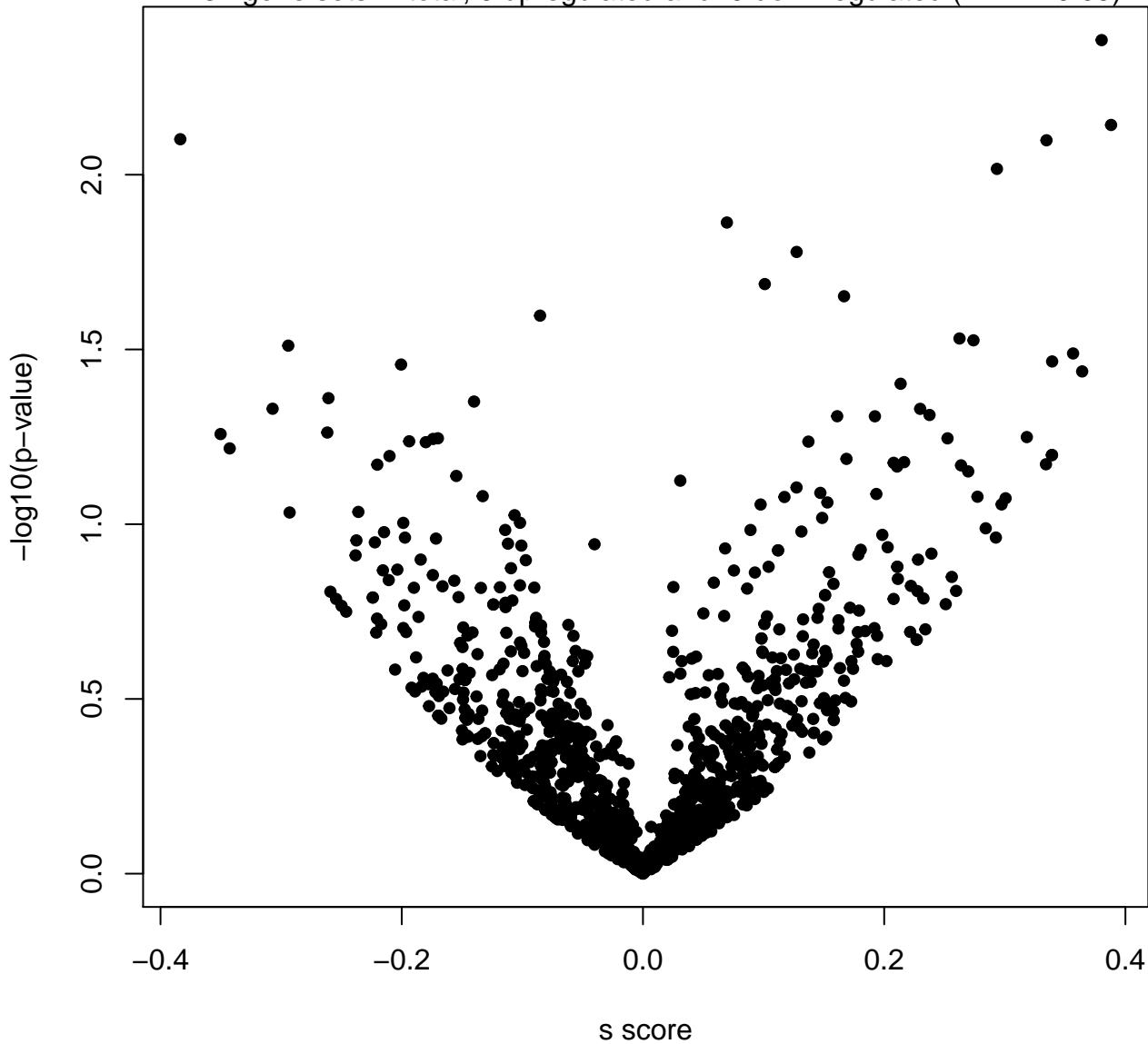


Trimmed histogram of geneset size



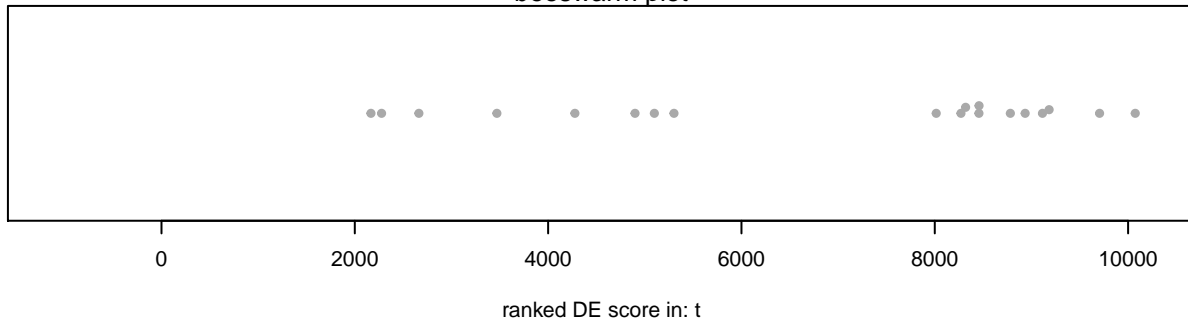
volcano plot of gene set enrichments

1134 gene sets in total, 0 upregulated and 0 downregulated (FDR \leq 0.05)

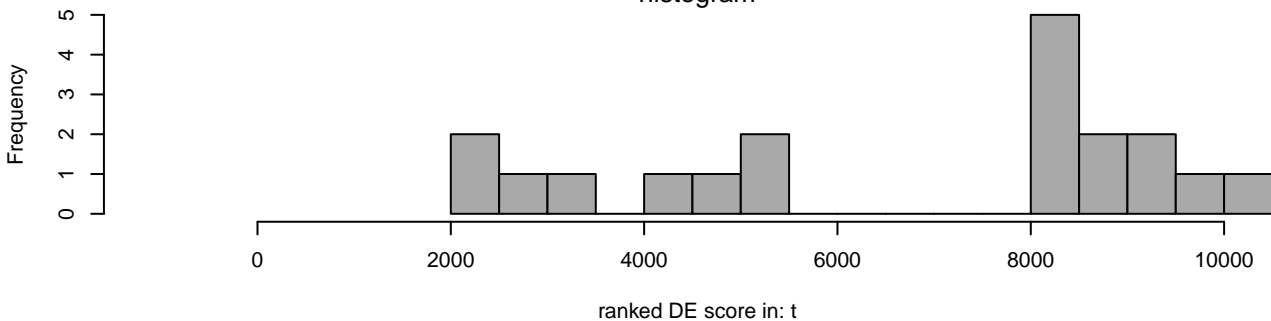


SUMOylation of intracellular receptors

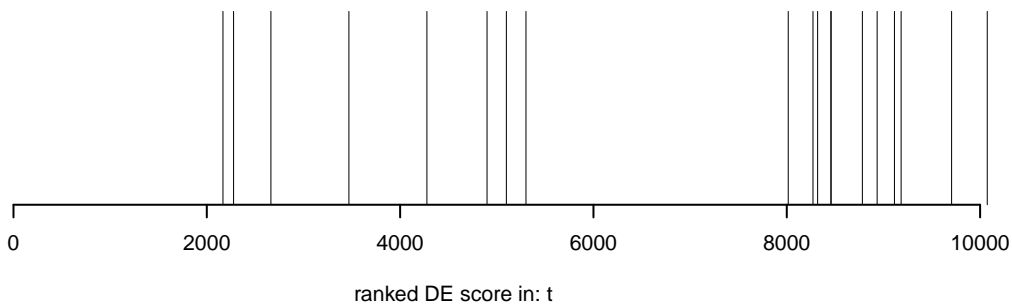
beeswarm plot



histogram

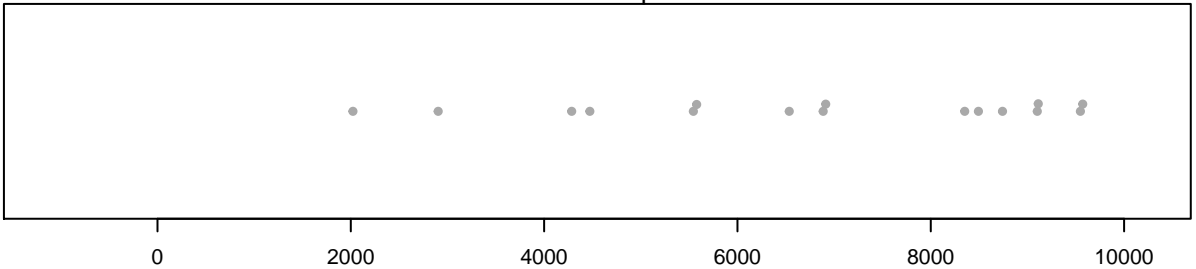


rugplot



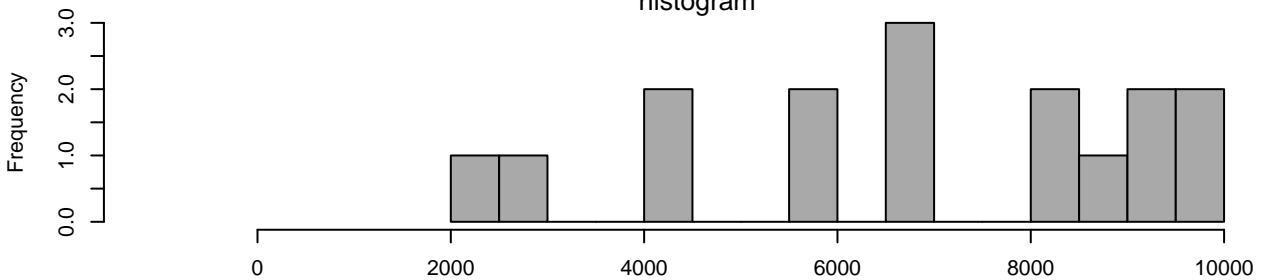
Diseases associated with glycosaminoglycan metabolism

beeswarm plot



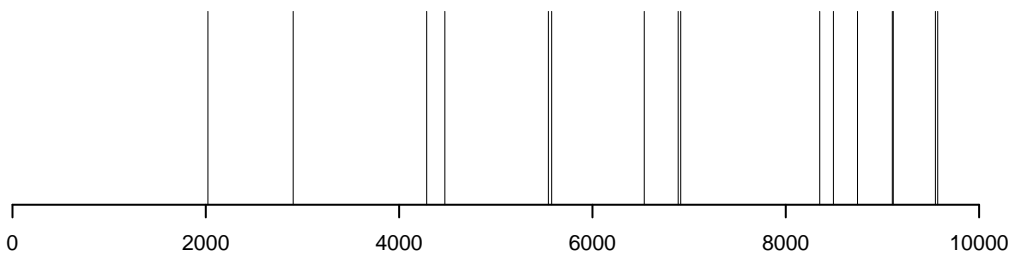
ranked DE score in: t

histogram



ranked DE score in: t

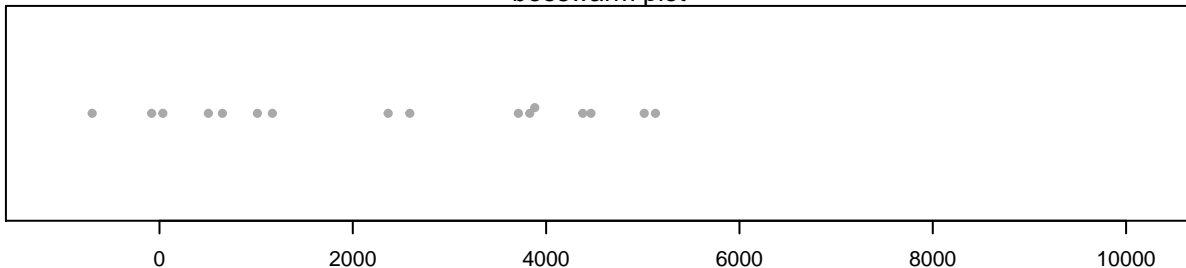
rugplot



ranked DE score in: t

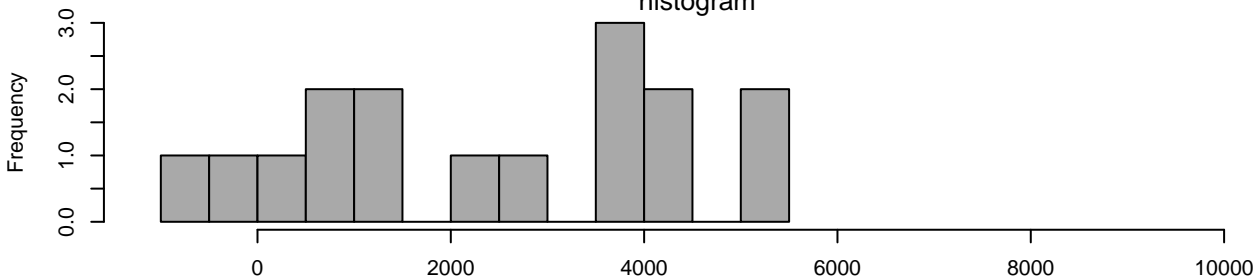
TP53 Regulates Transcription of Genes Involved in G2 Cell Cycle Arrest

beeswarm plot



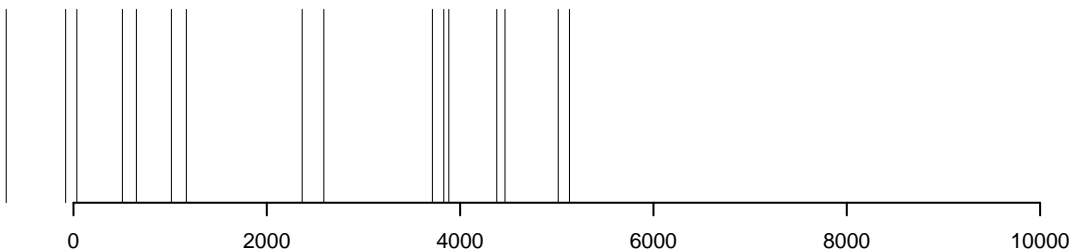
ranked DE score in: t

histogram



ranked DE score in: t

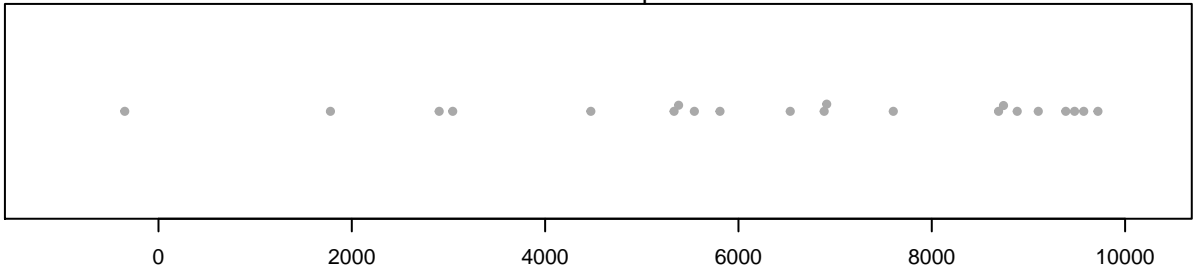
rugplot



ranked DE score in: t

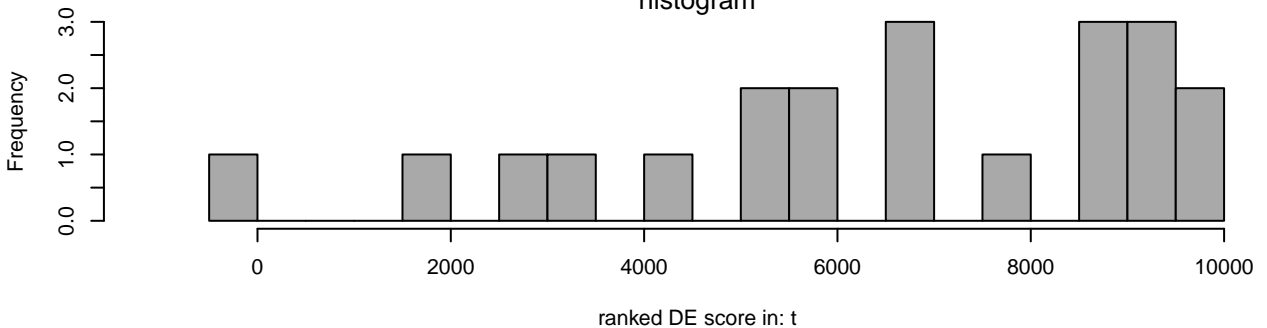
Heparan sulfate/heparin (HS-GAG) metabolism

beeswarm plot



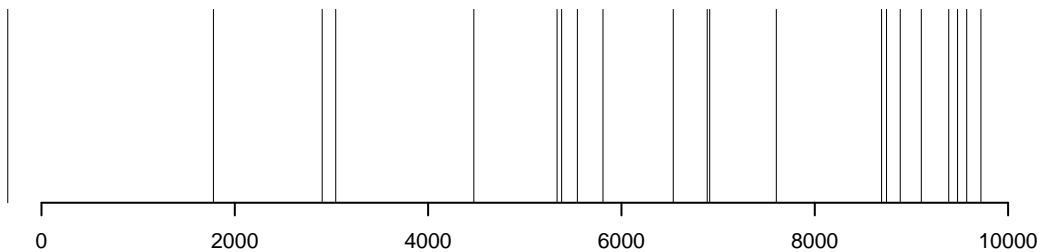
ranked DE score in: t

histogram



ranked DE score in: t

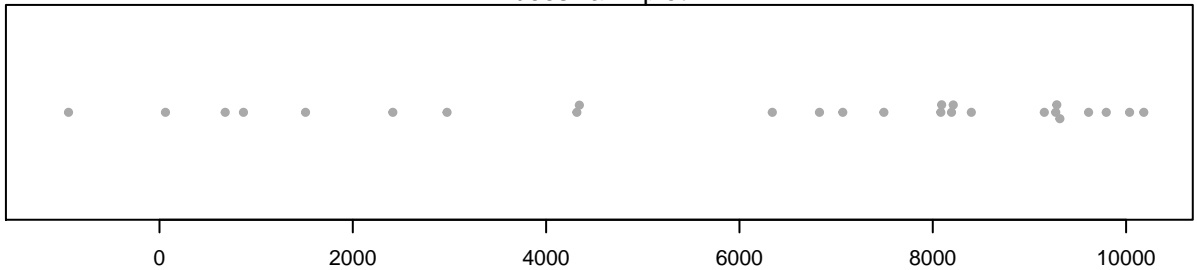
rugplot



ranked DE score in: t

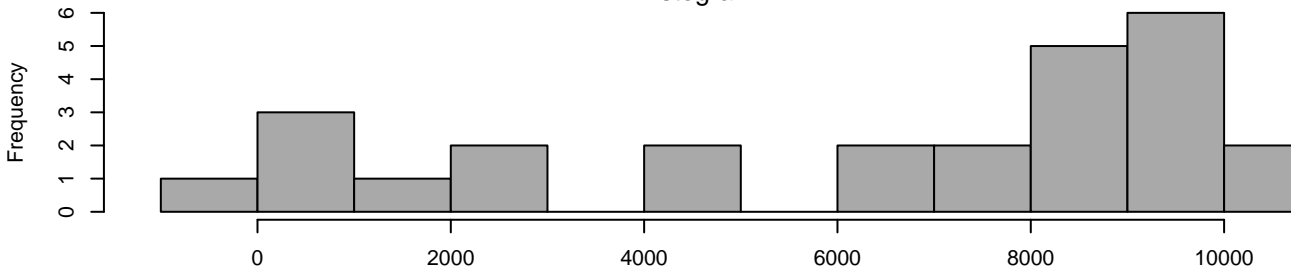
Disassembly of the destruction complex and recruitment of AXIN to the membrane

beeswarm plot



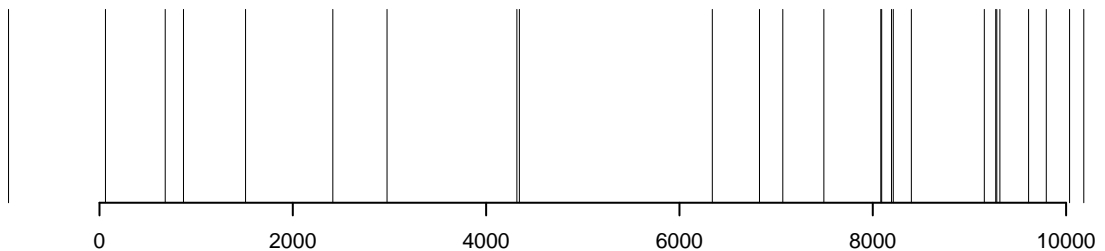
ranked DE score in: t

histogram



ranked DE score in: t

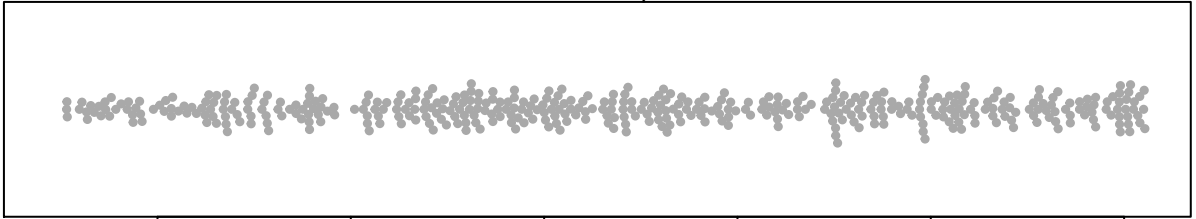
rugplot



ranked DE score in: t

Metabolism of lipids

beeswarm plot



ranked DE score in: t

histogram

Frequency

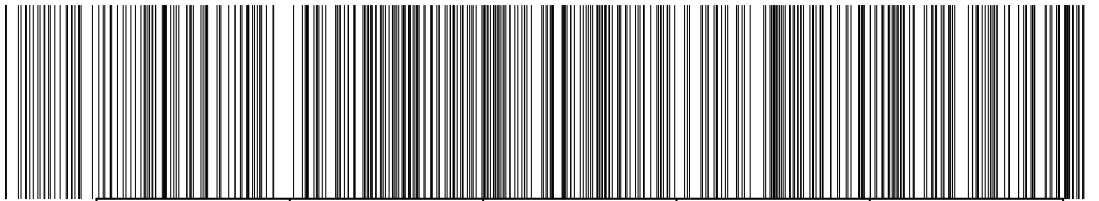
30
10
0



0 2000 4000 6000 8000 10000

ranked DE score in: t

rugplot

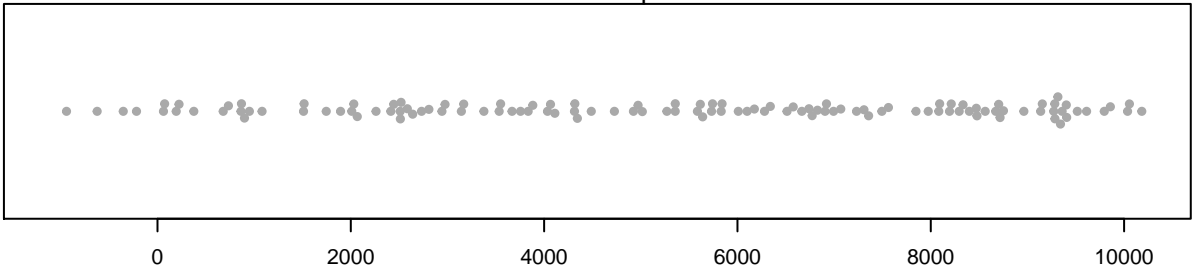


0 2000 4000 6000 8000 10000

ranked DE score in: t

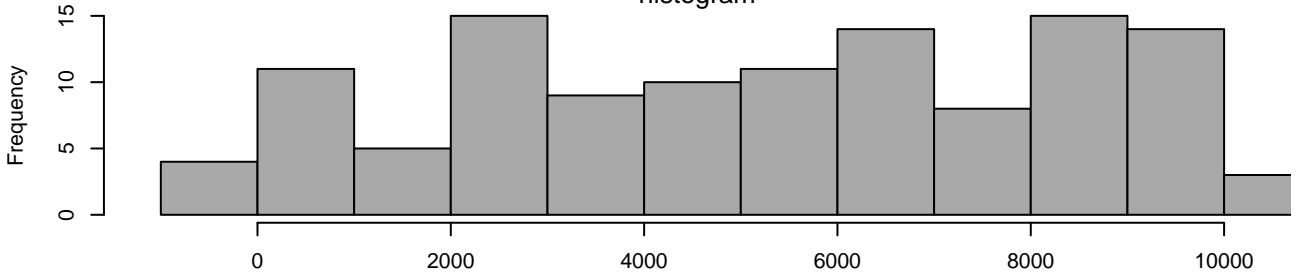
TCF dependent signaling in response to WNT

beeswarm plot



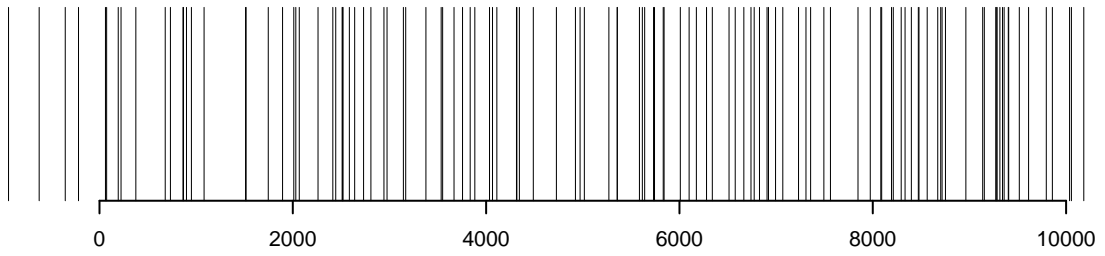
ranked DE score in: t

histogram



ranked DE score in: t

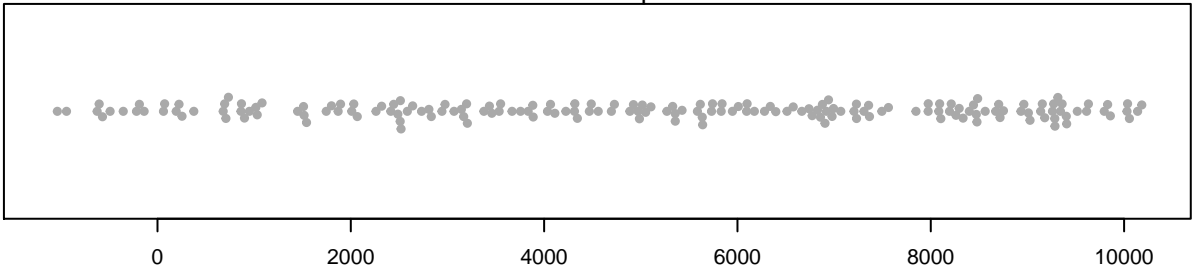
rugplot



ranked DE score in: t

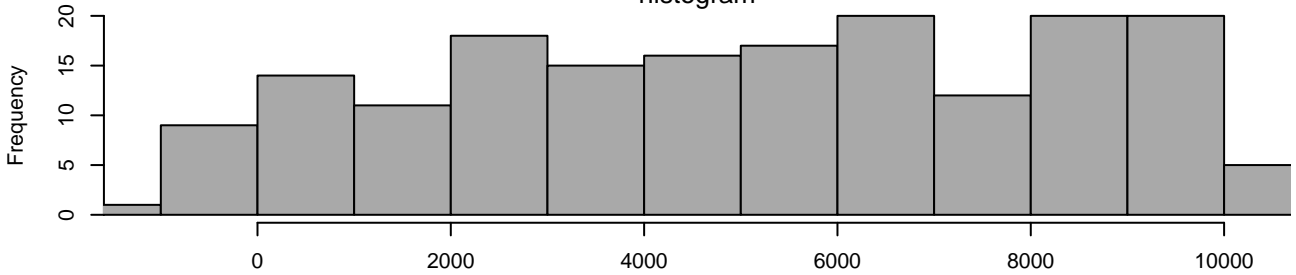
Signaling by WNT

beeswarm plot



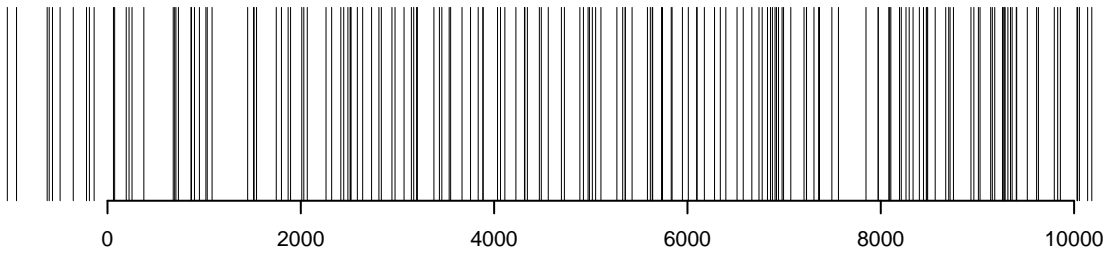
ranked DE score in: t

histogram



ranked DE score in: t

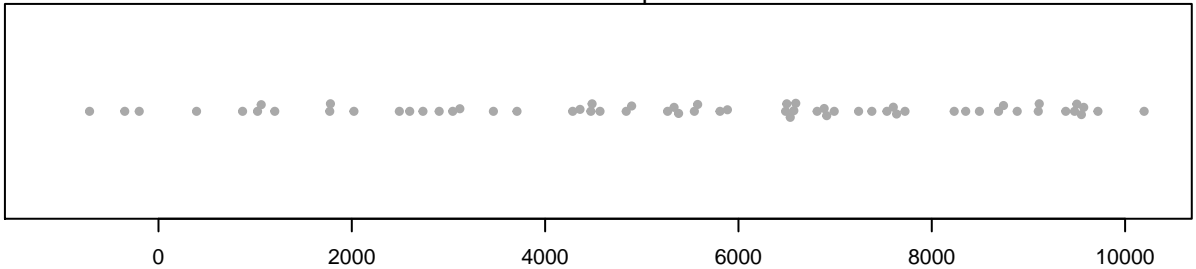
rugplot



ranked DE score in: t

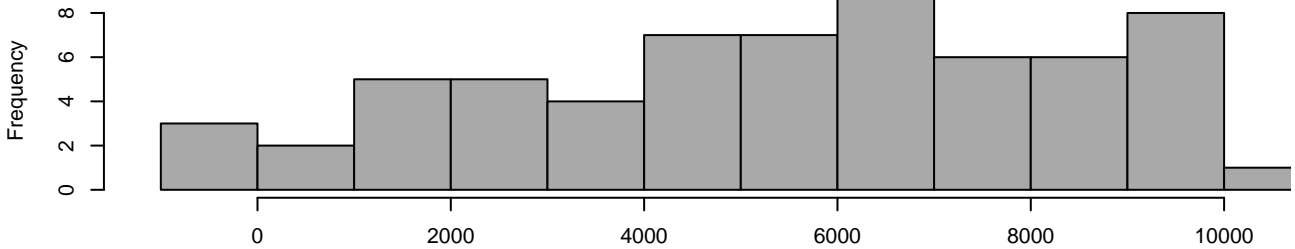
Glycosaminoglycan metabolism

beeswarm plot



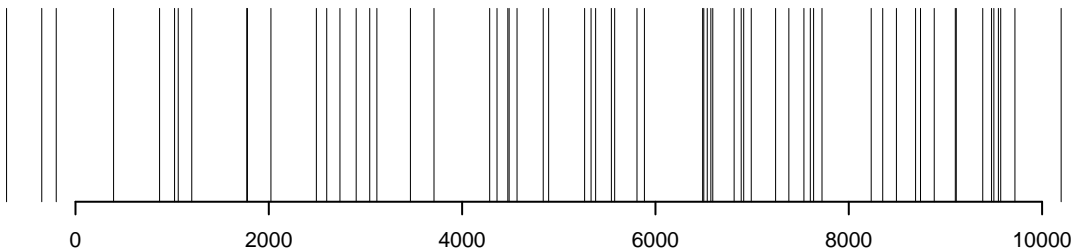
ranked DE score in: t

histogram



ranked DE score in: t

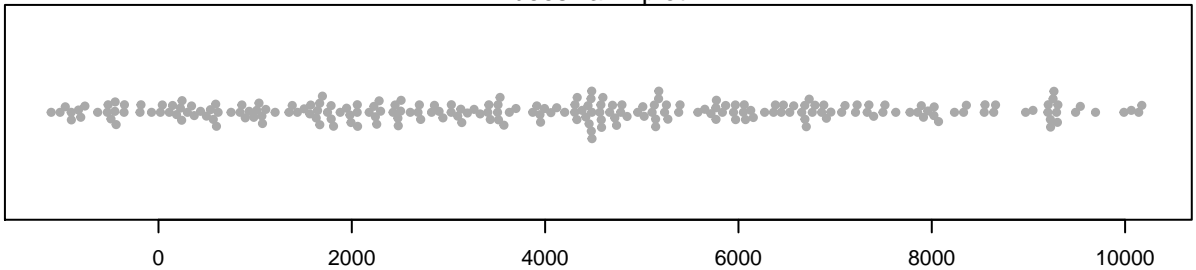
rugplot



ranked DE score in: t

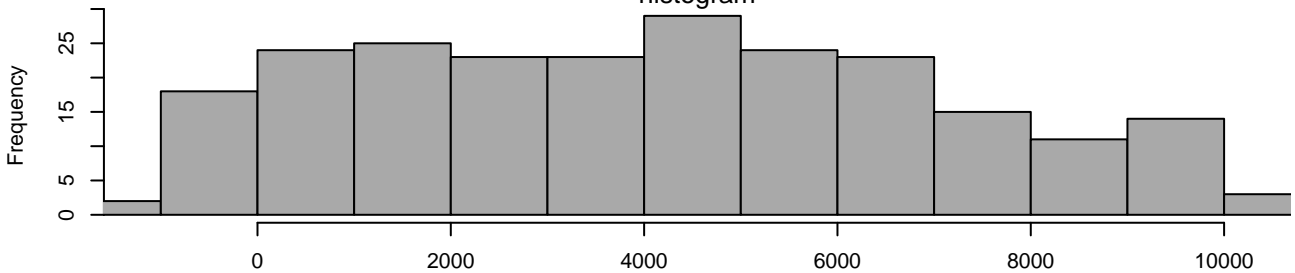
Translation

beeswarm plot



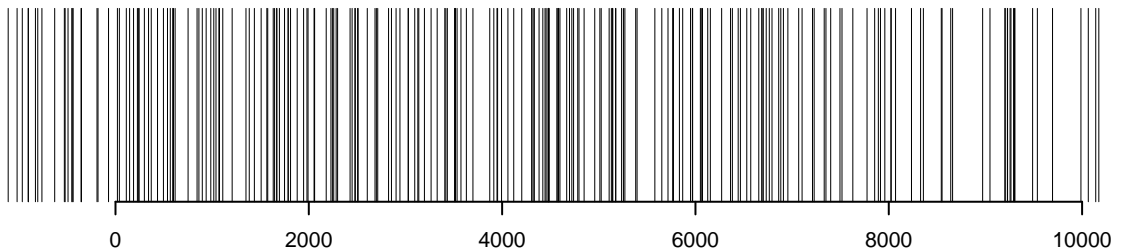
ranked DE score in: t

histogram



ranked DE score in: t

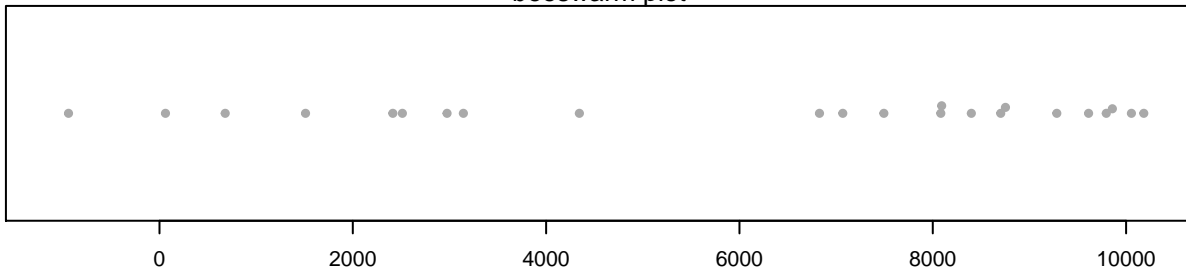
rugplot



ranked DE score in: t

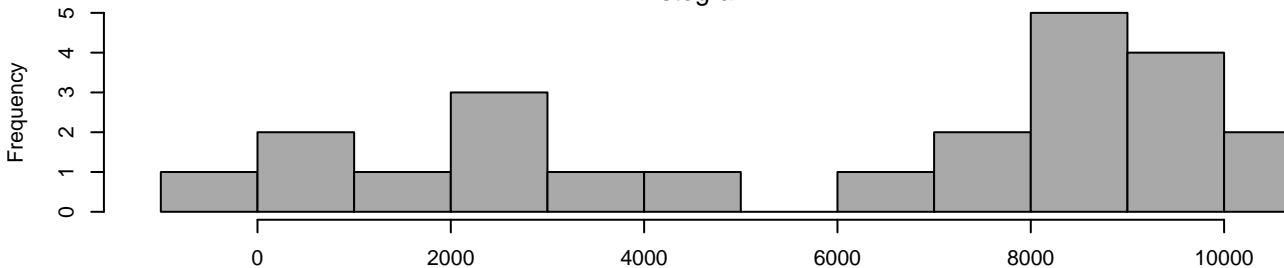
Signaling by WNT in cancer

beeswarm plot



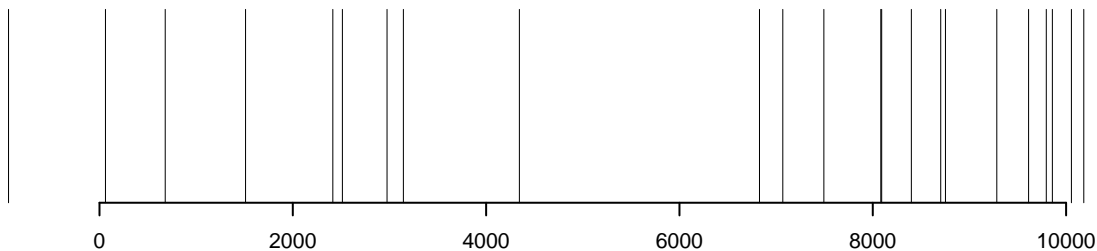
ranked DE score in: t

histogram



ranked DE score in: t

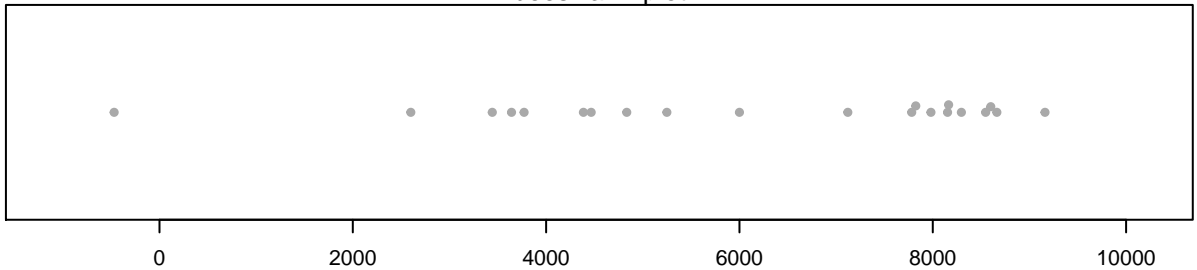
rugplot



ranked DE score in: t

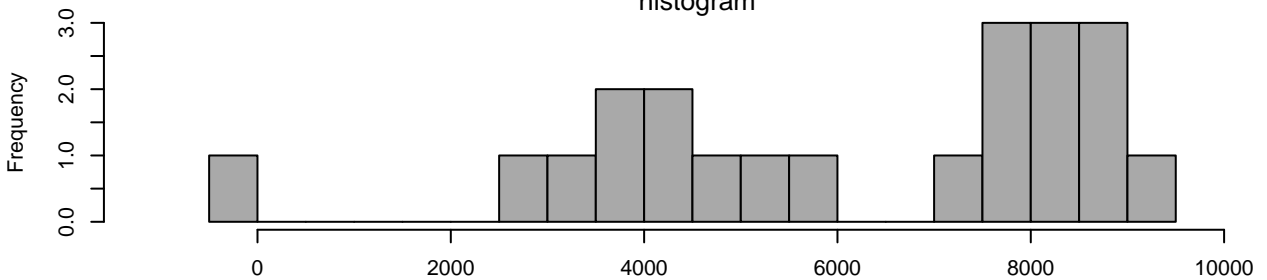
Processing of Capped Intronless Pre-mRNA

beeswarm plot



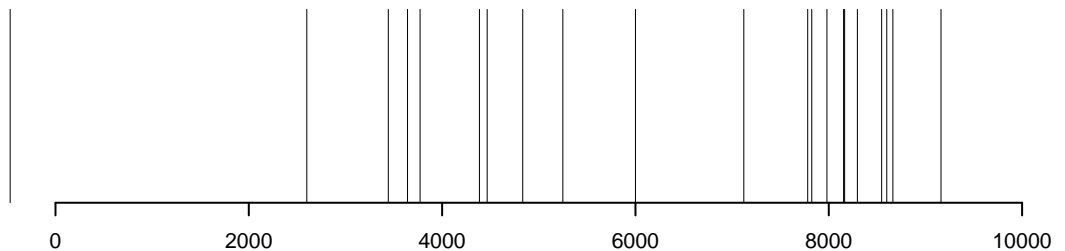
ranked DE score in: t

histogram



ranked DE score in: t

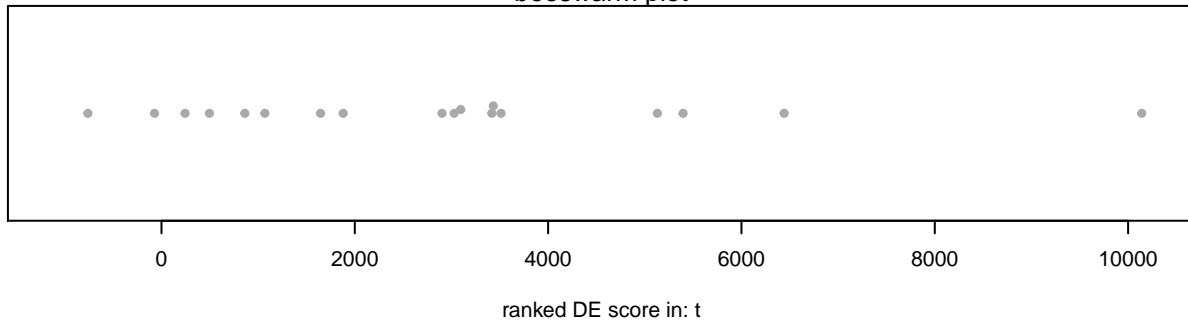
rugplot



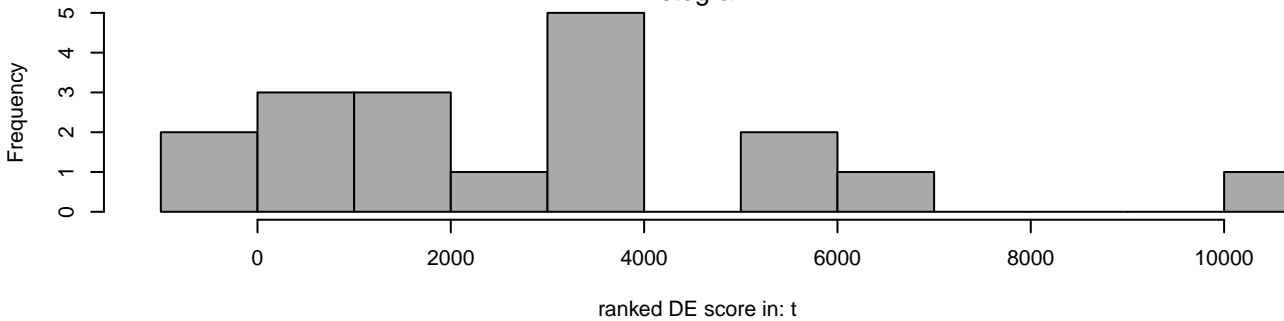
ranked DE score in: t

Mitochondrial tRNA aminoacylation

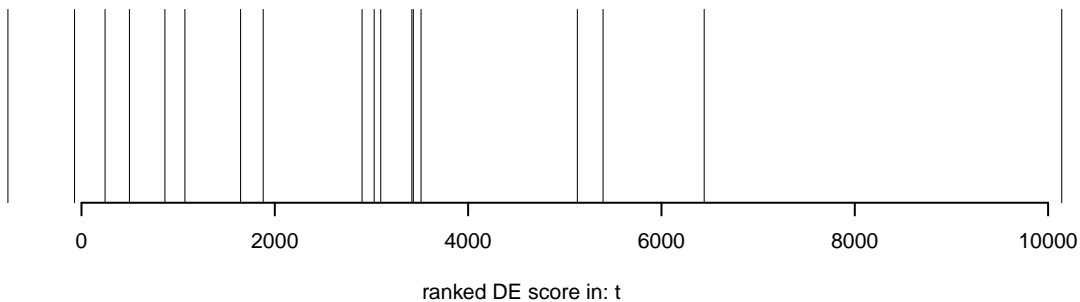
beeswarm plot



histogram

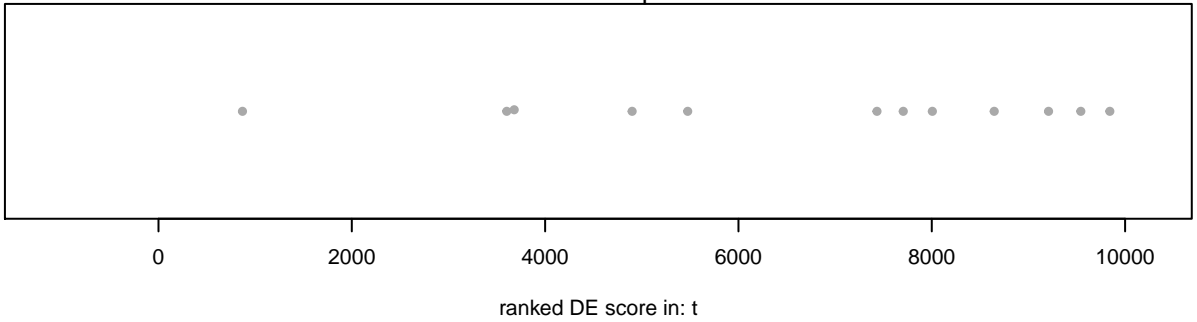


rugplot

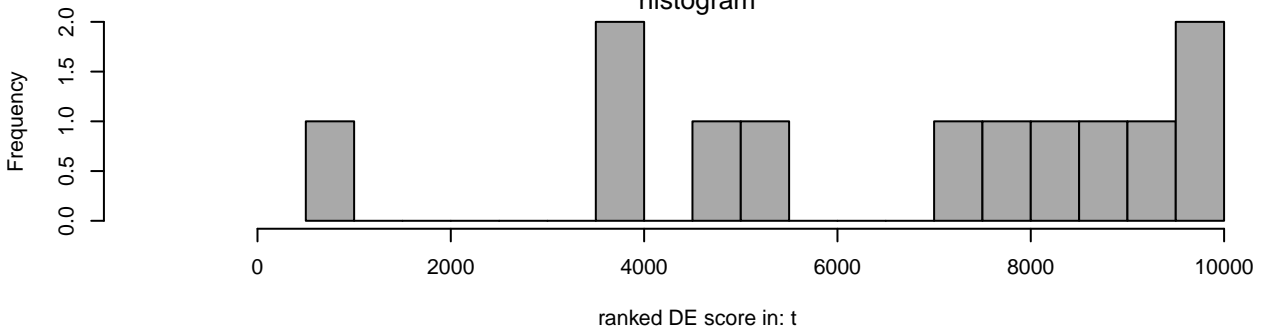


Signaling by Hippo

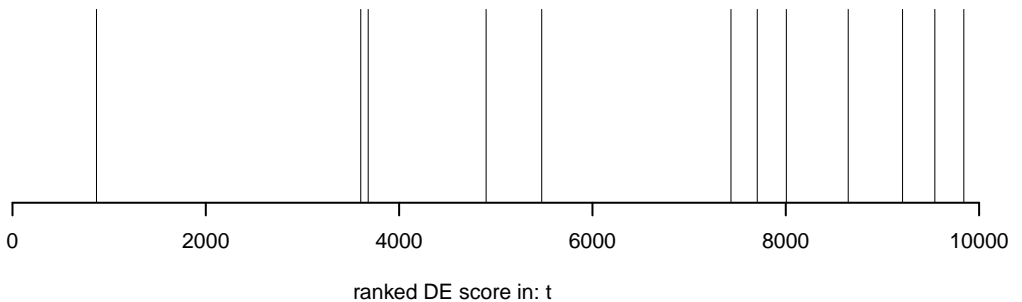
beeswarm plot



histogram

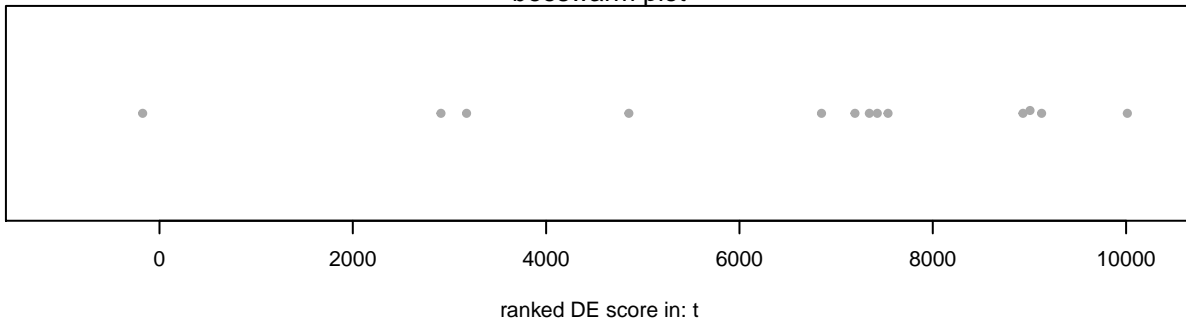


rugplot

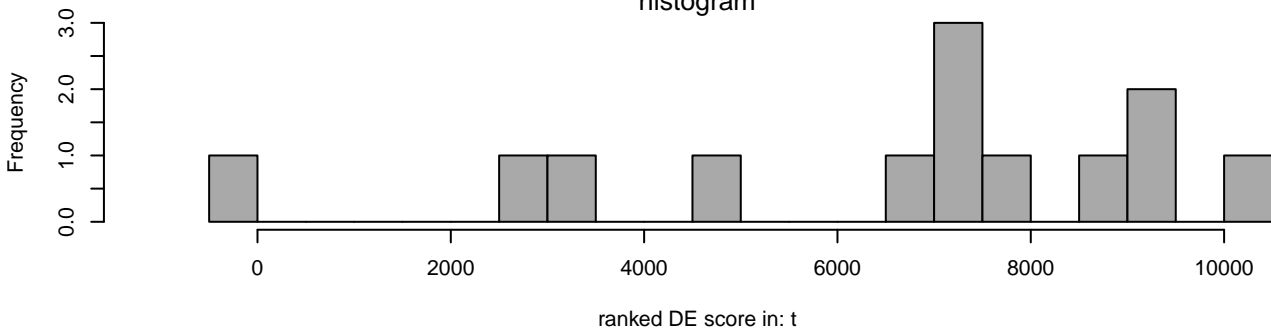


MET promotes cell motility

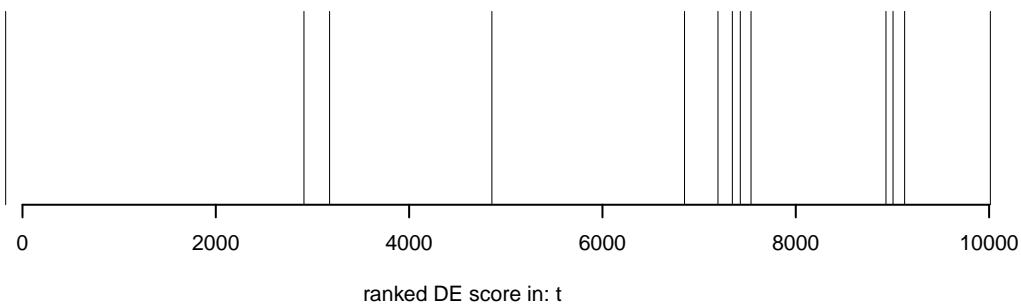
beeswarm plot



histogram

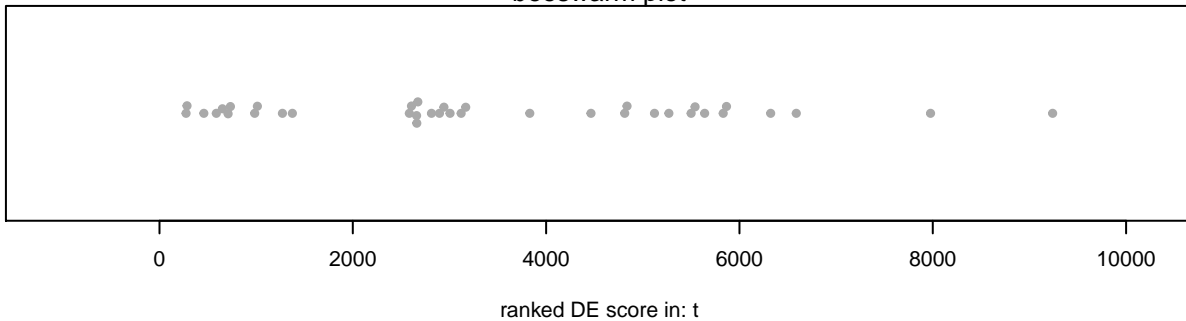


rugplot

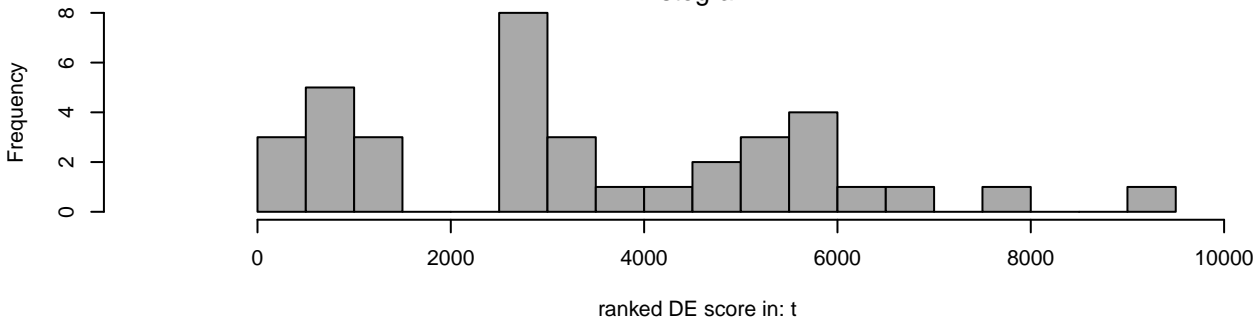


Transcriptional activity of SMAD2/SMAD3:SMAD4 heterotrimer

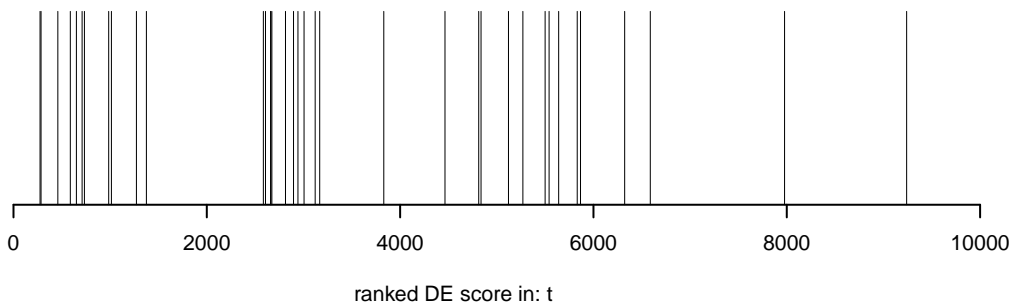
beeswarm plot



histogram

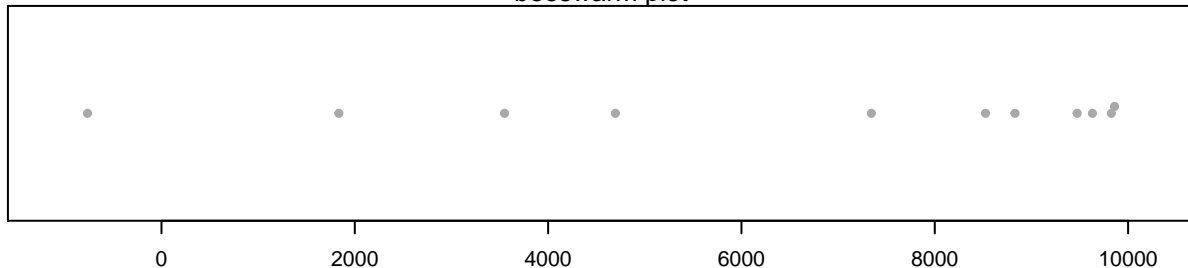


rugplot



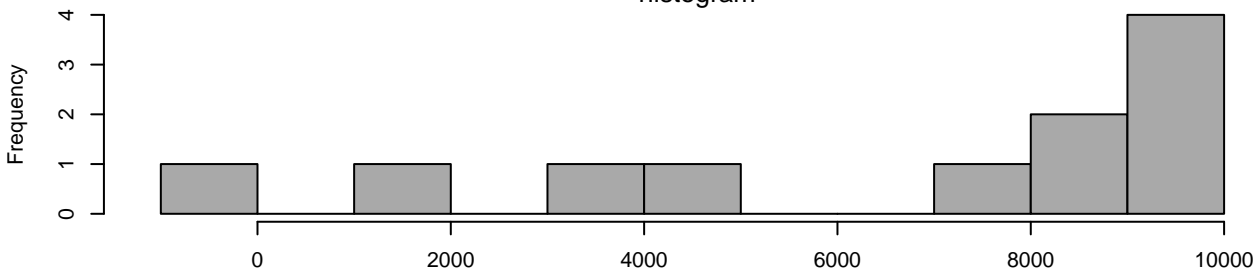
SHC1 events in ERBB2 signaling

beeswarm plot



ranked DE score in: t

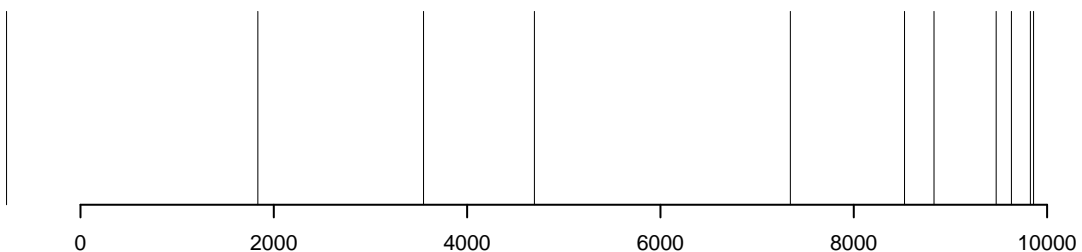
histogram



Frequency

ranked DE score in: t

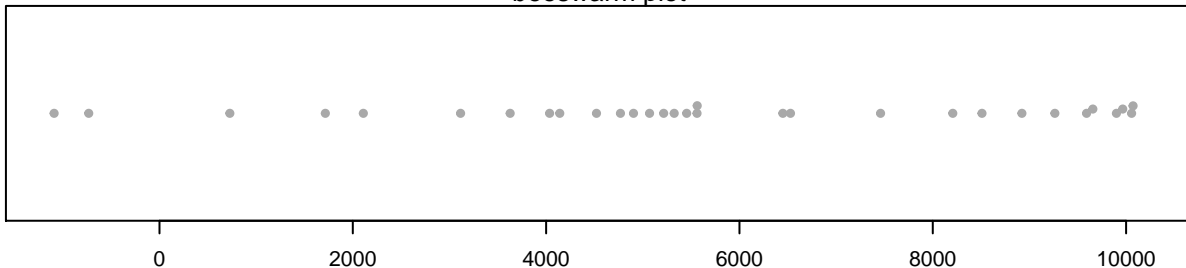
rugplot



ranked DE score in: t

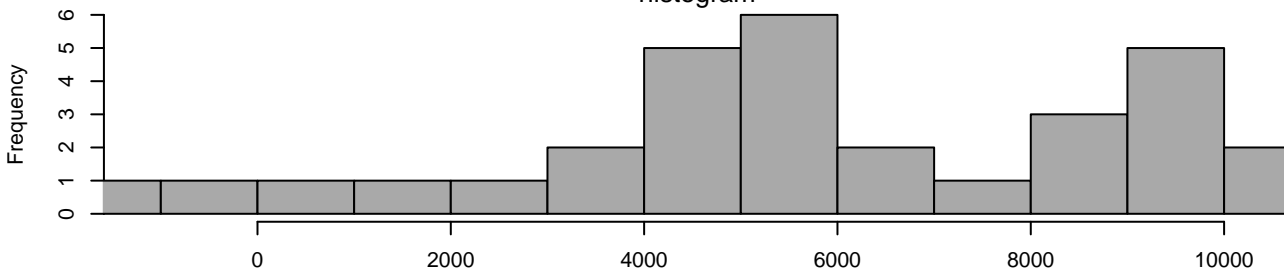
Post-translational modification: synthesis of GPI-anchored proteins

beeswarm plot



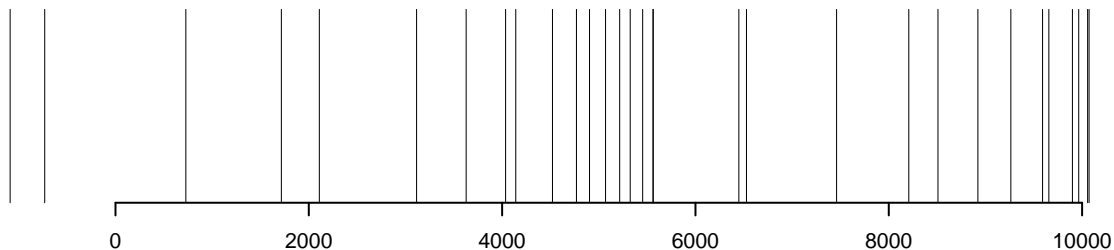
ranked DE score in: t

histogram



ranked DE score in: t

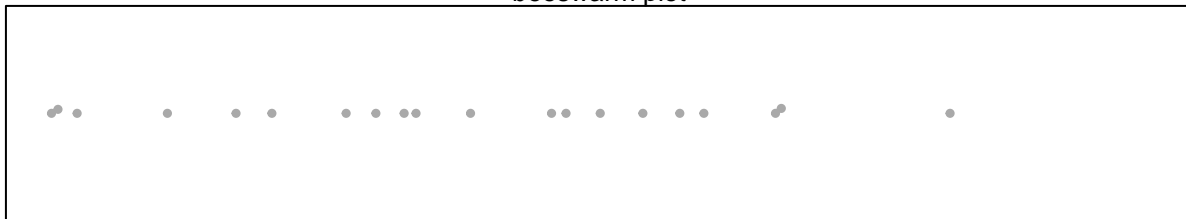
rugplot



ranked DE score in: t

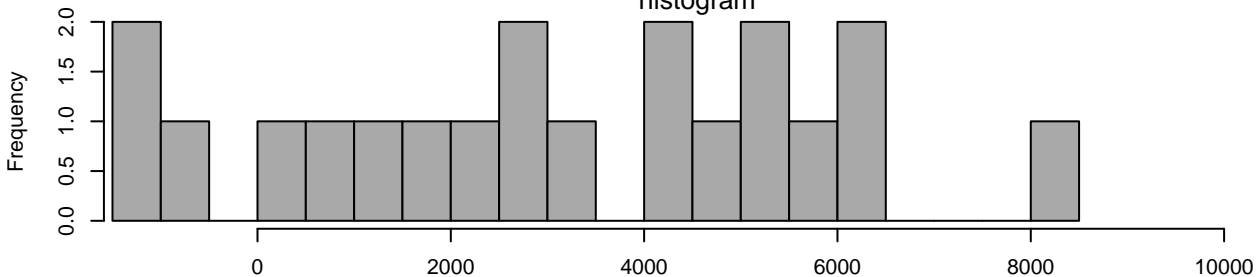
TNFR1-induced NFkappaB signaling pathway

beeswarm plot



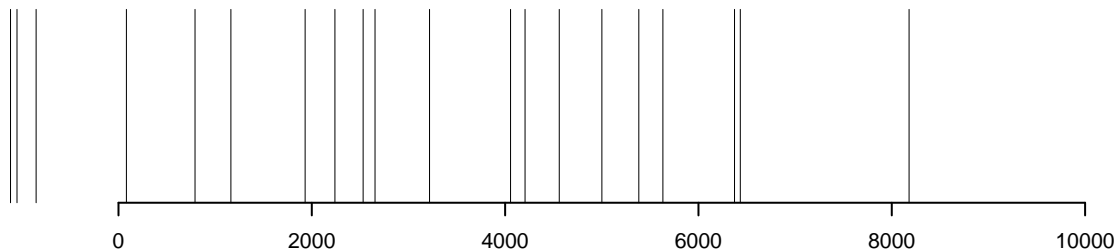
ranked DE score in: t

histogram



ranked DE score in: t

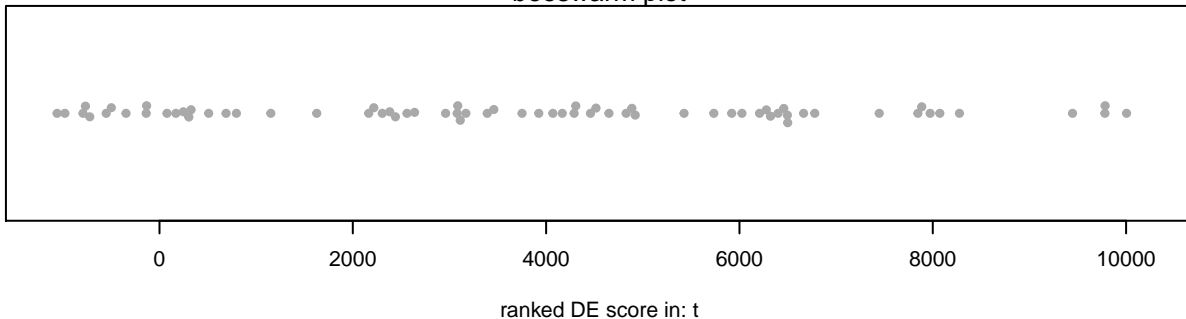
rugplot



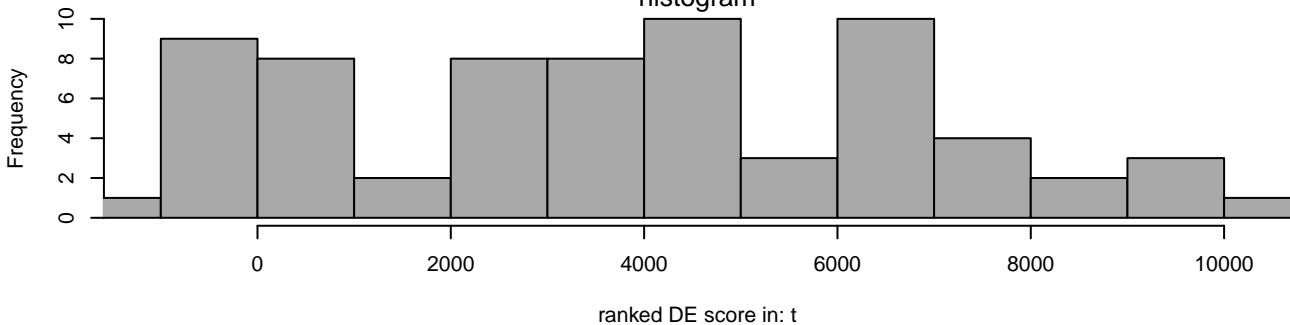
ranked DE score in: t

Muscle contraction

beeswarm plot



histogram



rugplot

