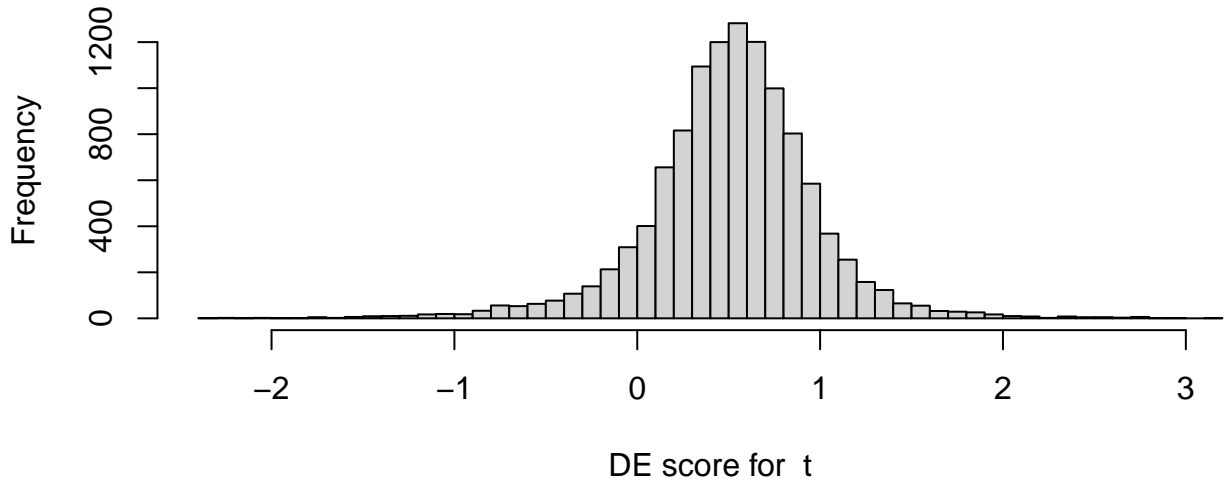
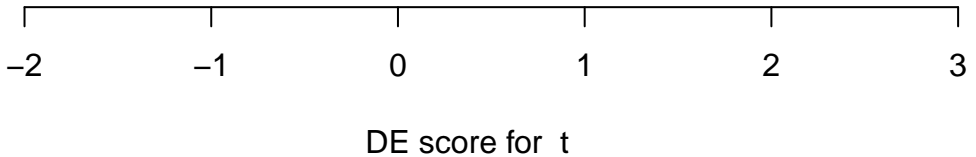


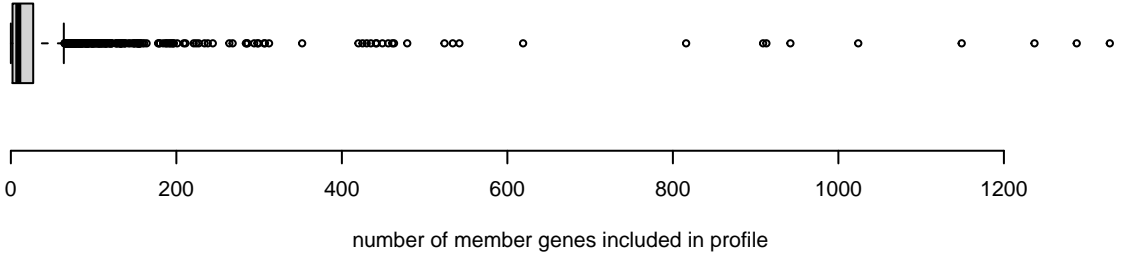
Distribution of DE scores



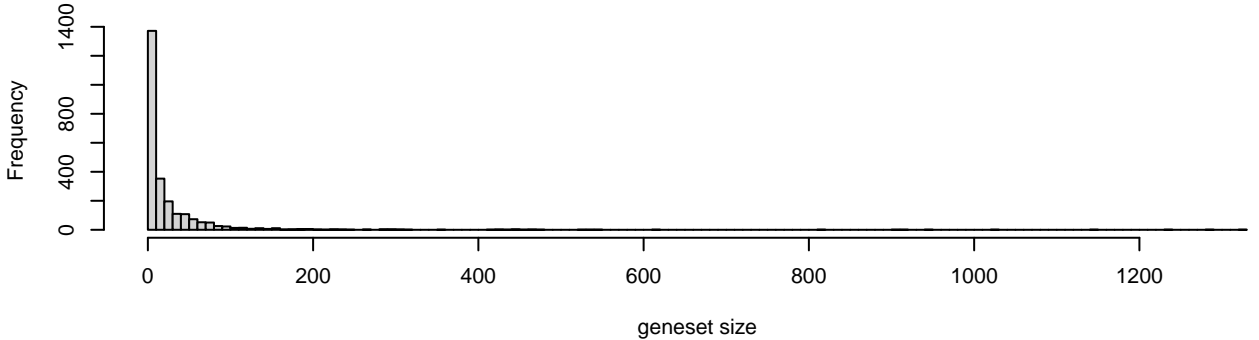
11370 genes in total, 10215 trending up-regulated, 1155 trending down-regulated



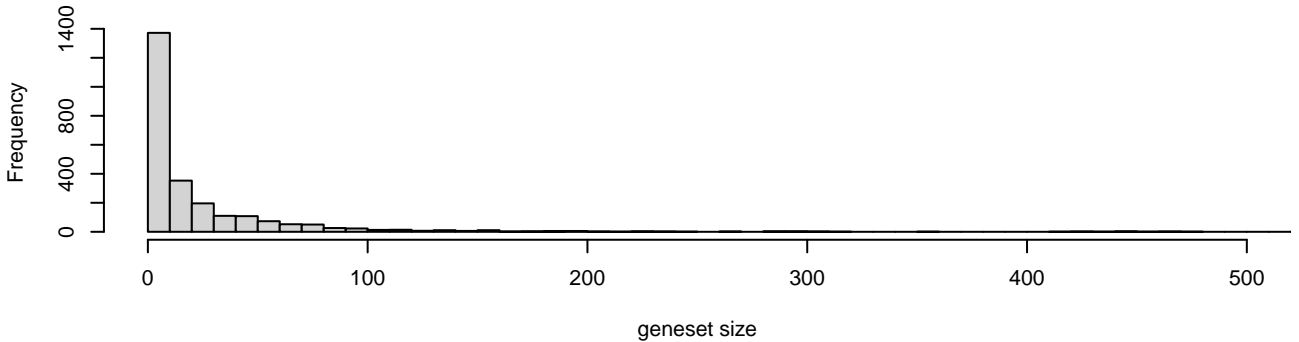
Gene set size



Histogram of geneset size

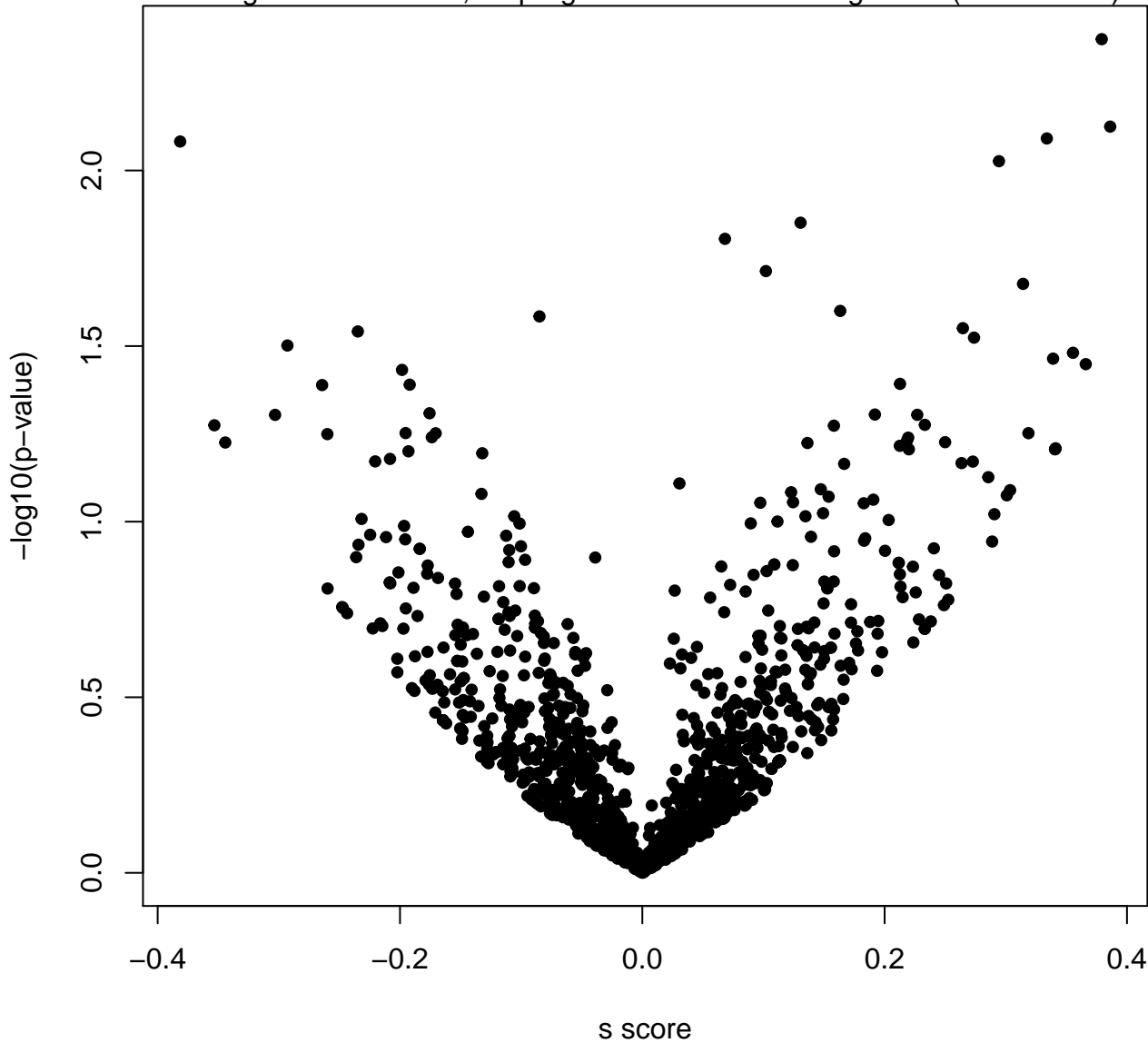


Trimmed histogram of geneset size



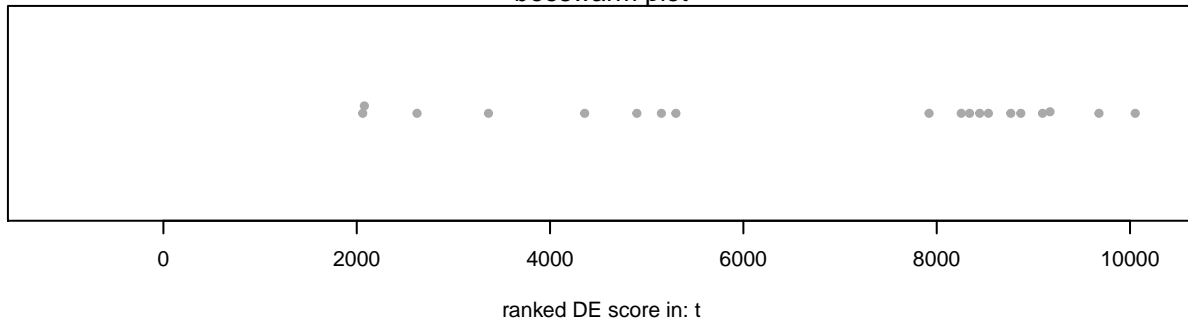
volcano plot of gene set enrichments

1176 gene sets in total, 0 upregulated and 0 downregulated (FDR<=0.05)

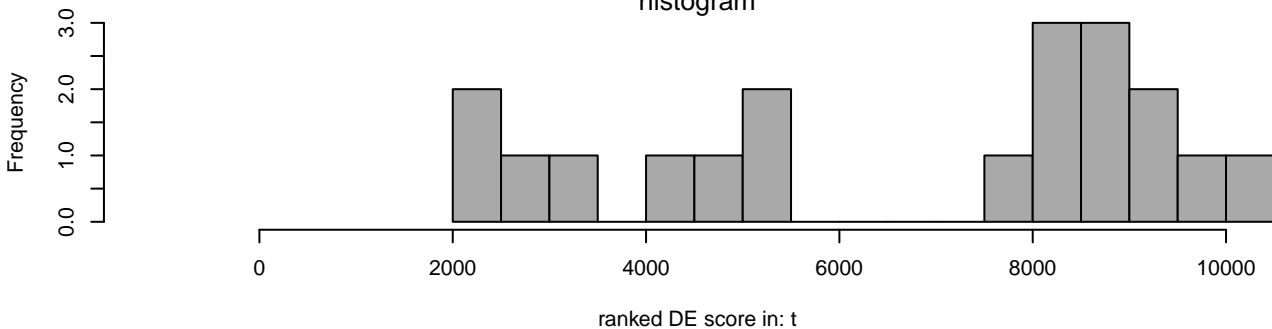


SUMOylation of intracellular receptors

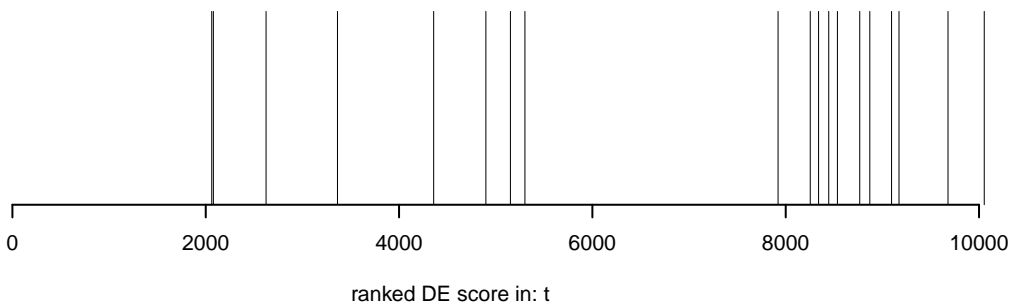
beeswarm plot



histogram

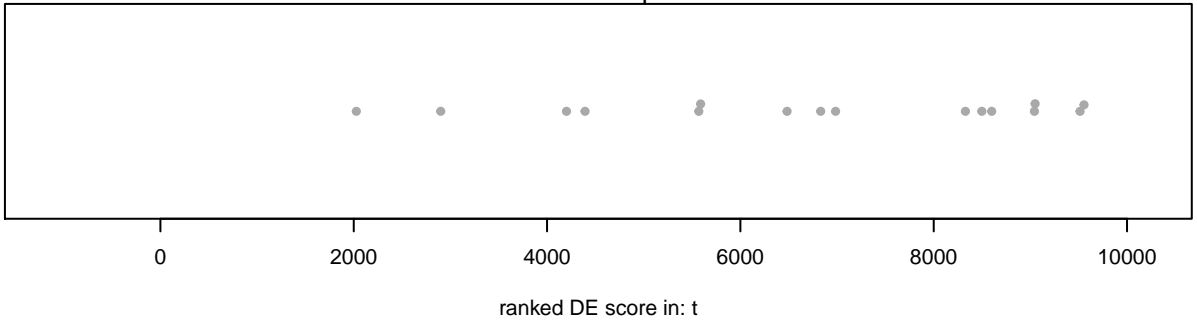


rugplot

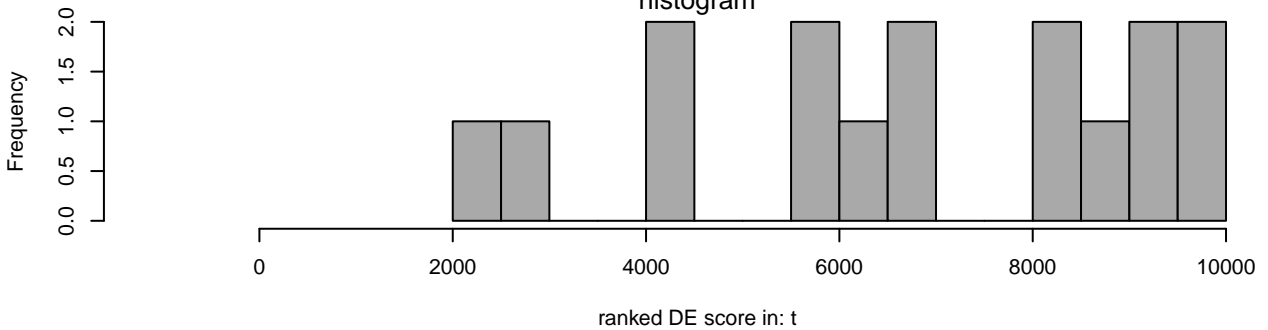


Diseases associated with glycosaminoglycan metabolism

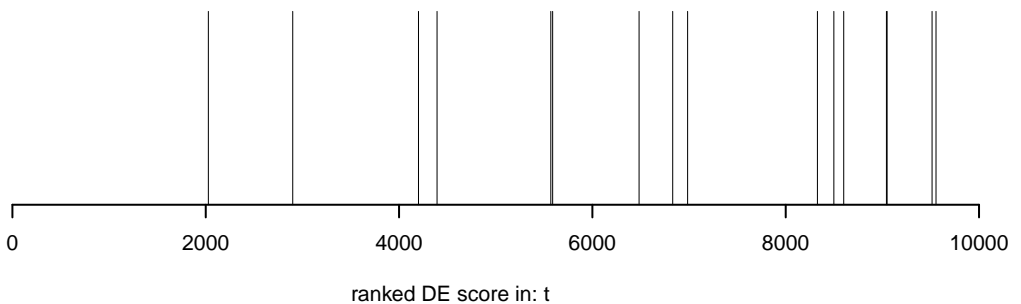
beeswarm plot



histogram

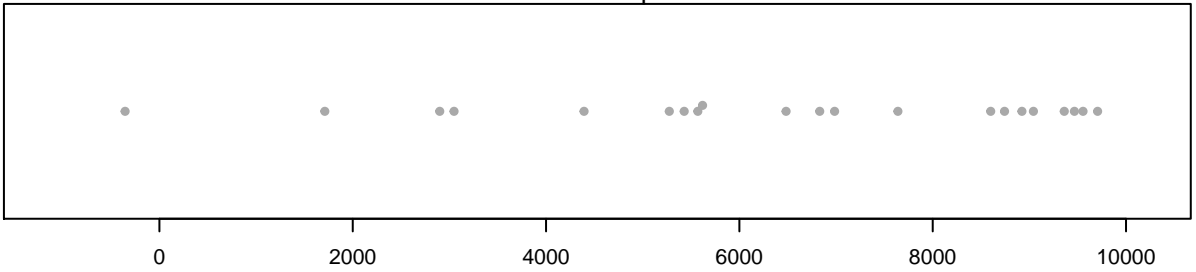


rugplot



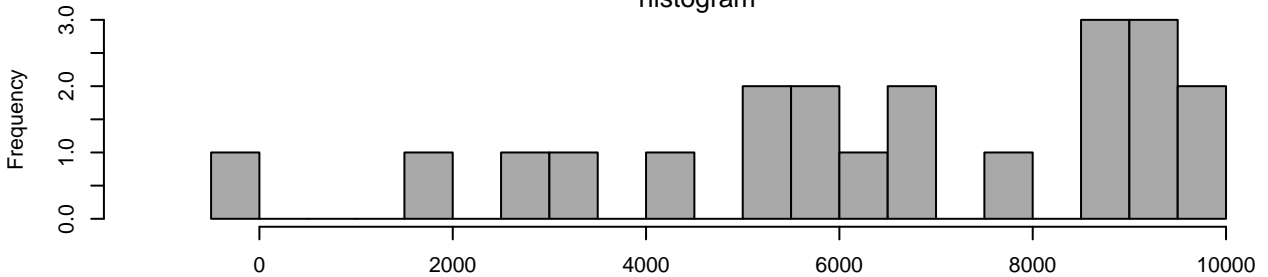
Heparan sulfate/heparin (HS-GAG) metabolism

beeswarm plot



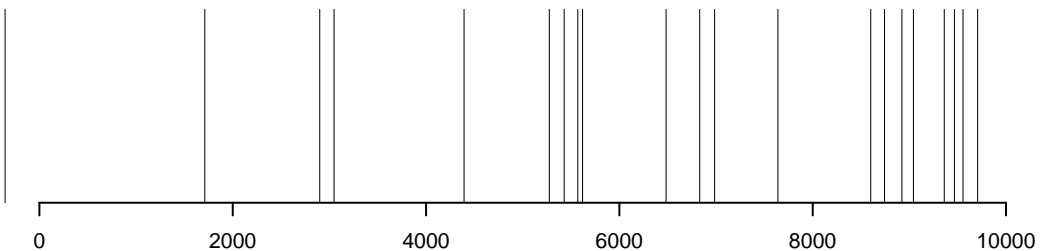
ranked DE score in: t

histogram



ranked DE score in: t

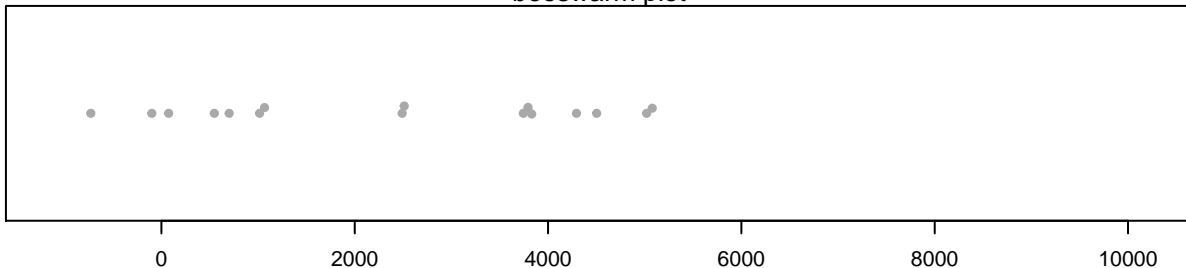
rugplot



ranked DE score in: t

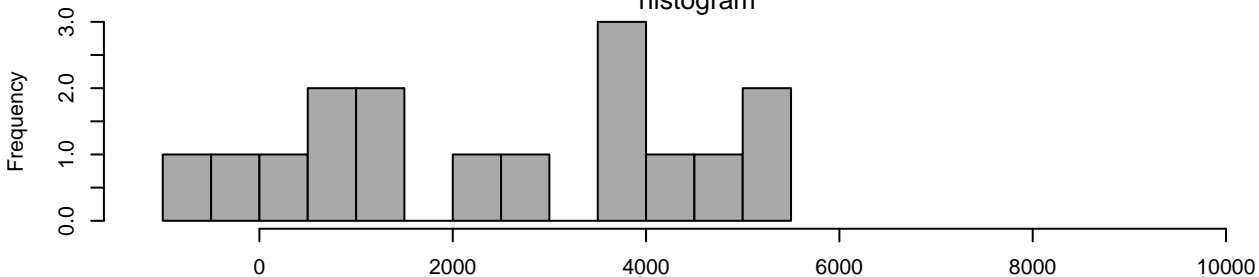
TP53 Regulates Transcription of Genes Involved in G2 Cell Cycle Arrest

beeswarm plot



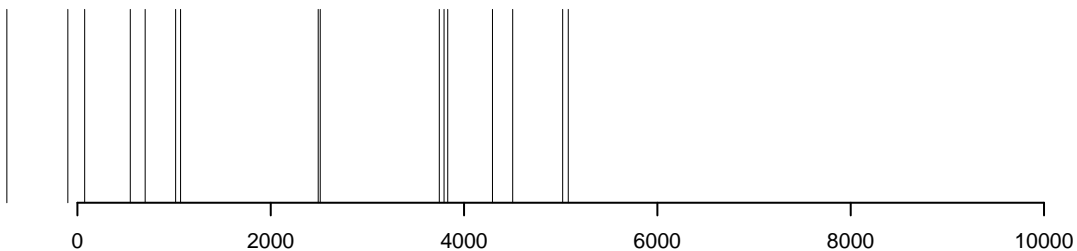
ranked DE score in: t

histogram



ranked DE score in: t

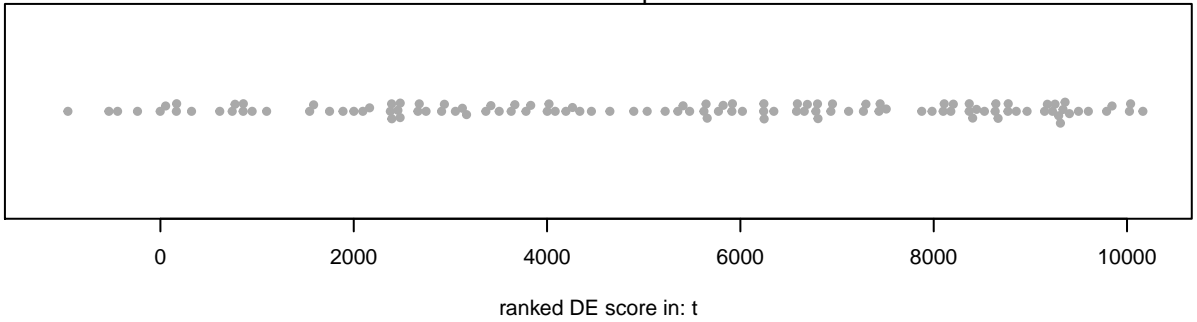
rugplot



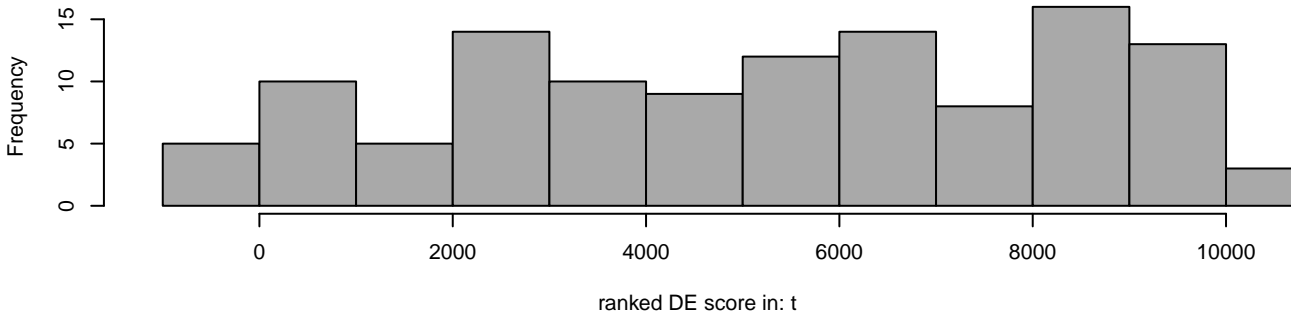
ranked DE score in: t

TCF dependent signaling in response to WNT

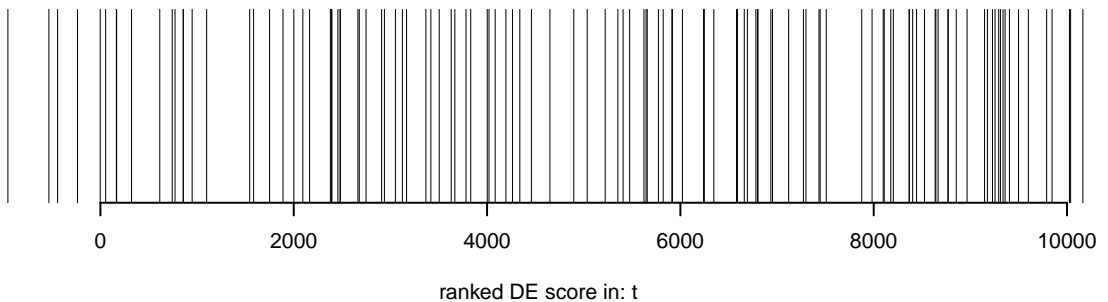
beeswarm plot



histogram

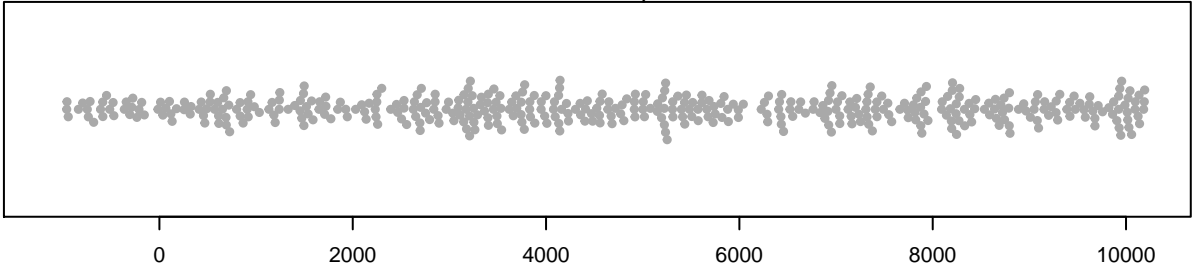


rugplot



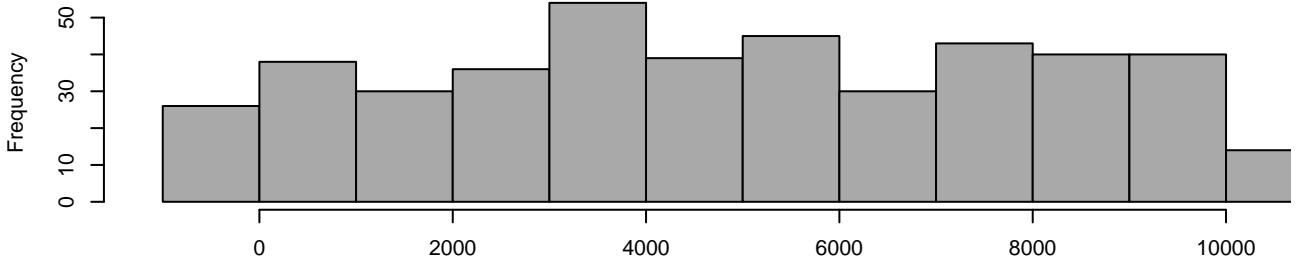
Metabolism of lipids

beeswarm plot



ranked DE score in: t

histogram



ranked DE score in: t

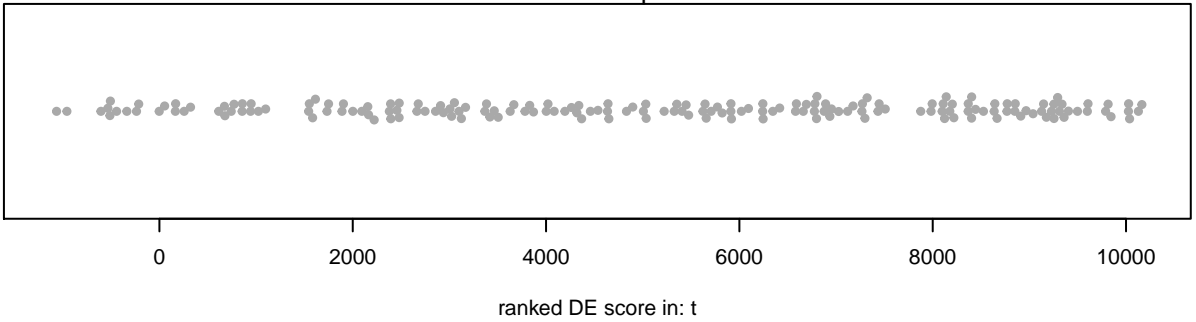
rugplot



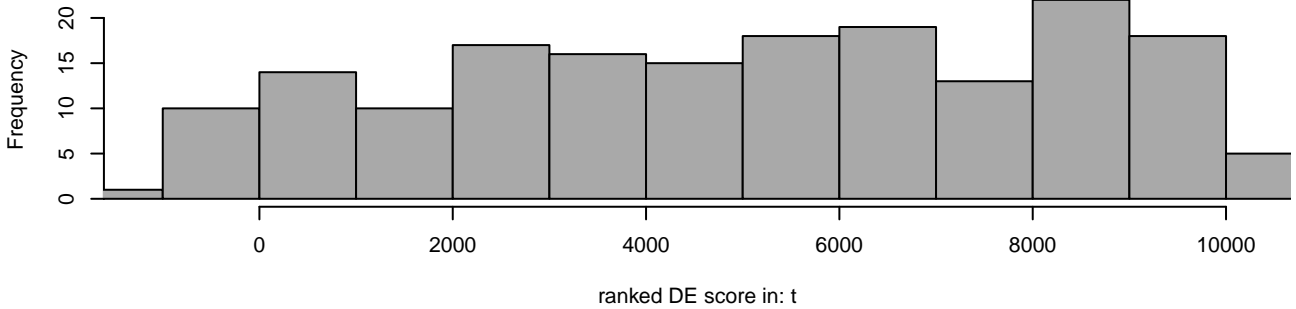
ranked DE score in: t

Signaling by WNT

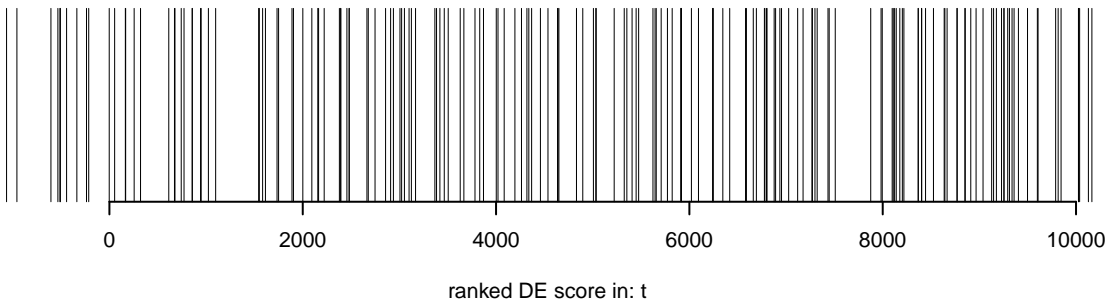
beeswarm plot



histogram

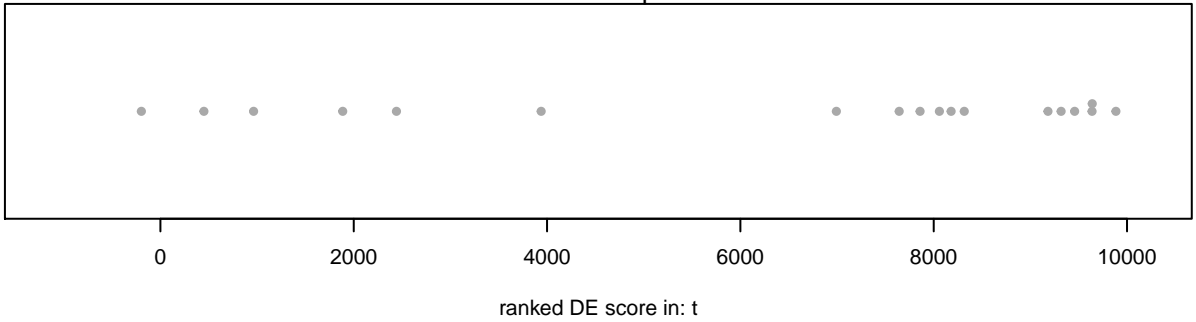


rugplot

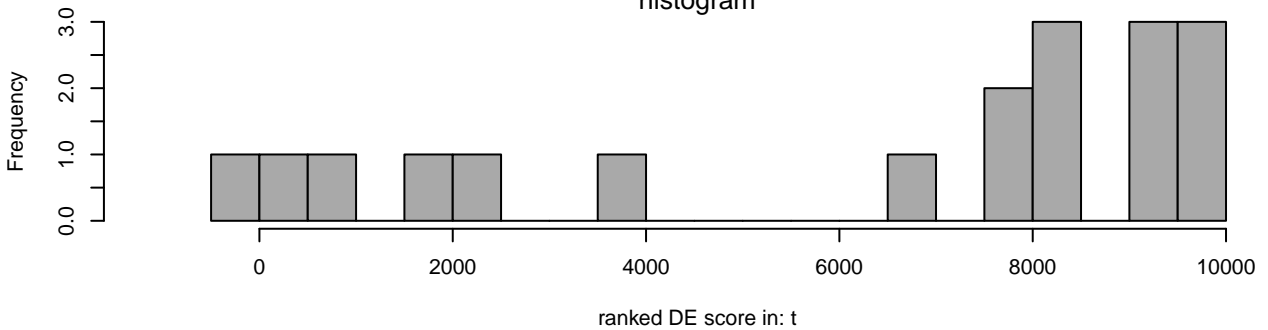


Sensory processing of sound by outer hair cells of the cochlea

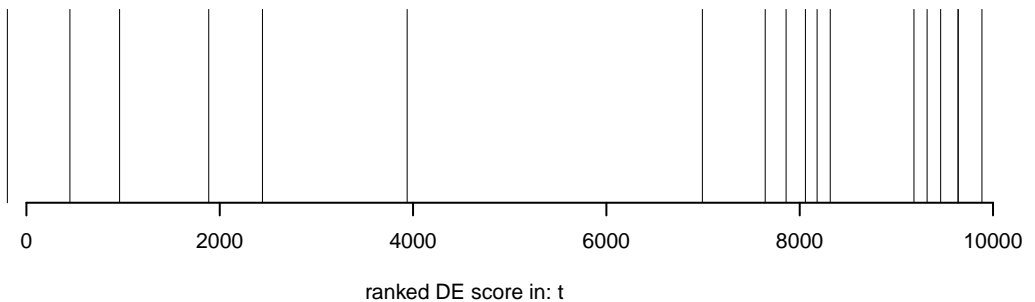
beeswarm plot



histogram

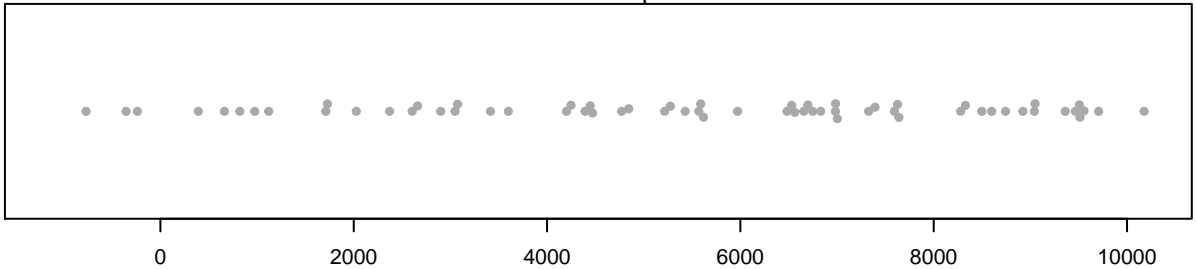


rugplot



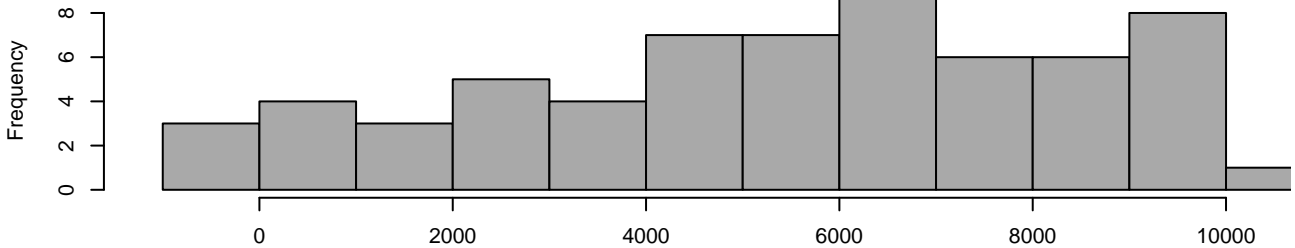
Glycosaminoglycan metabolism

beeswarm plot



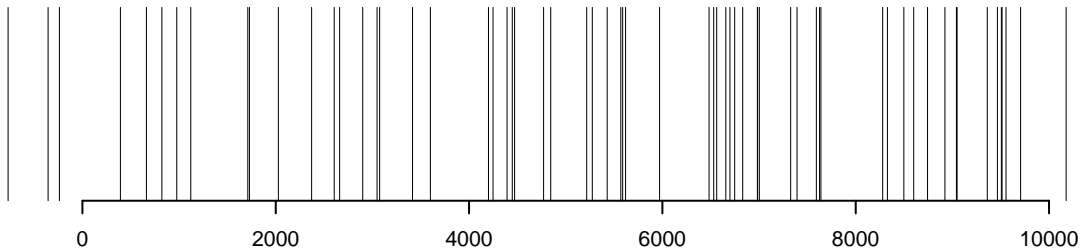
ranked DE score in: t

histogram



ranked DE score in: t

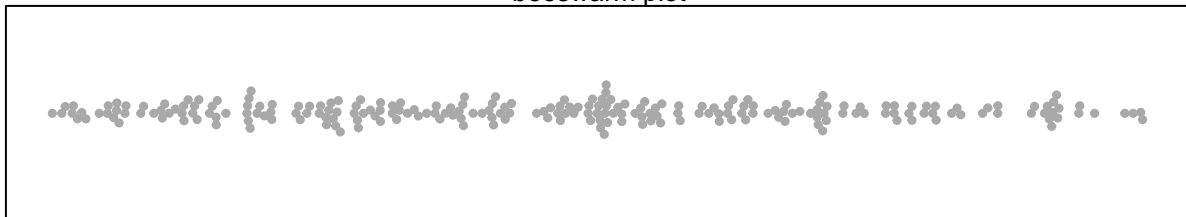
rugplot



ranked DE score in: t

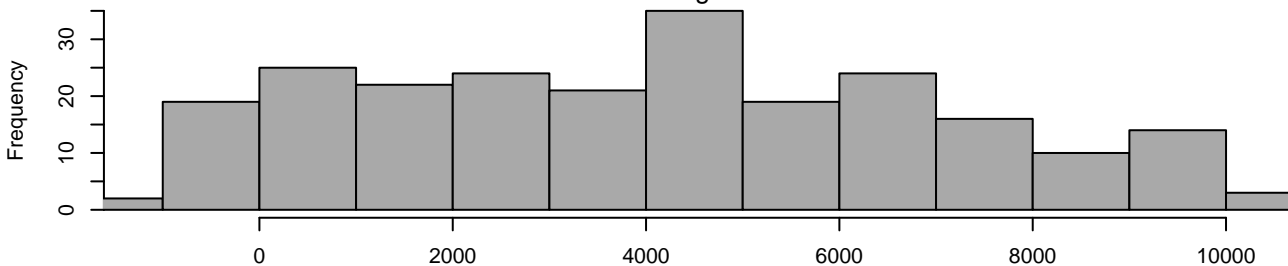
Translation

beeswarm plot



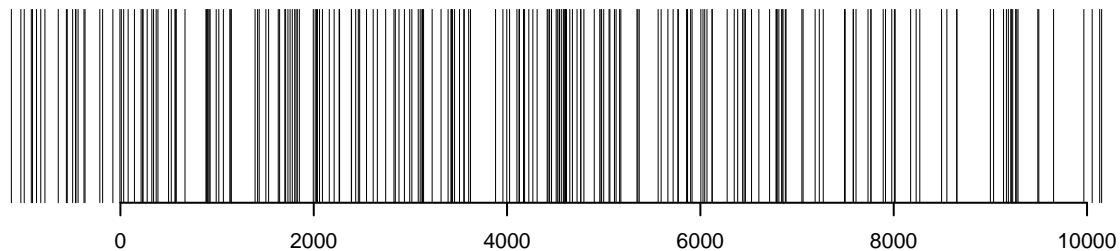
ranked DE score in: t

histogram



ranked DE score in: t

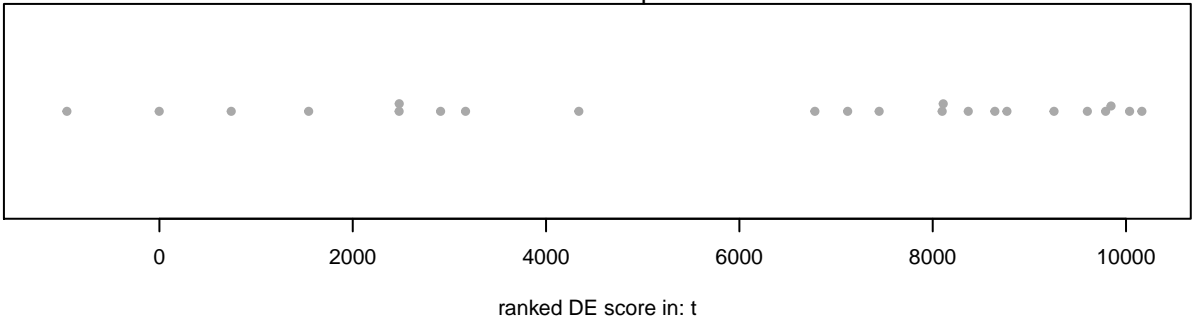
rugplot



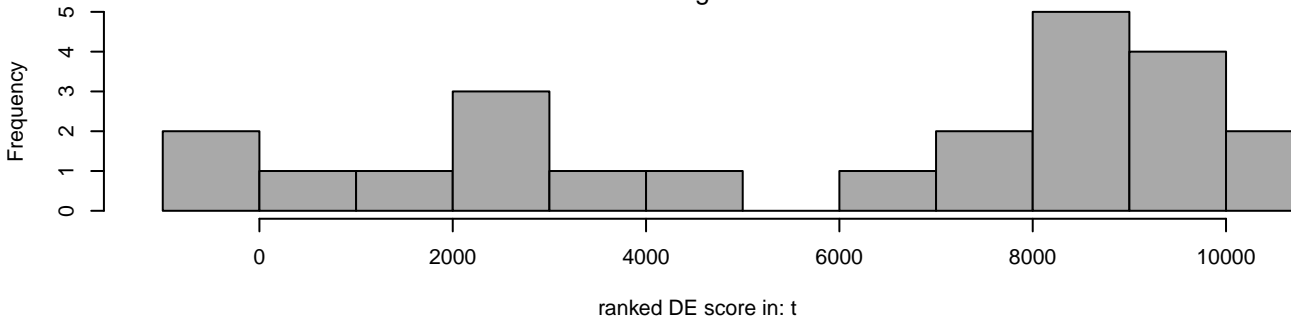
ranked DE score in: t

Signaling by WNT in cancer

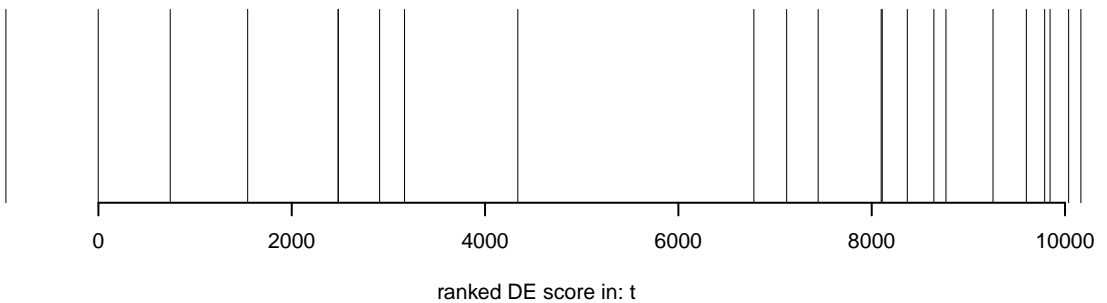
beeswarm plot



histogram

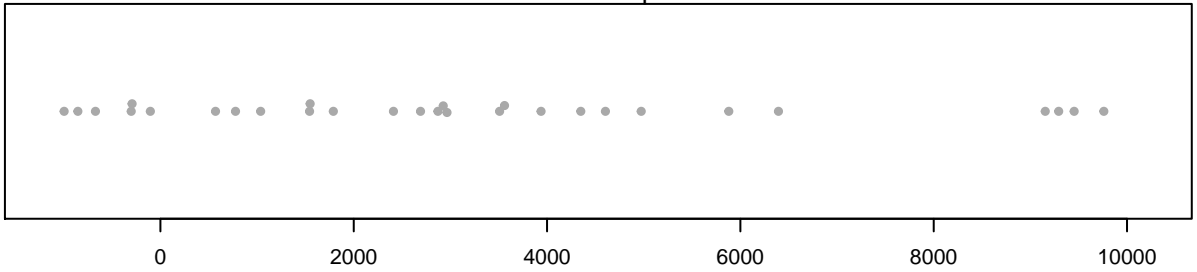


rugplot



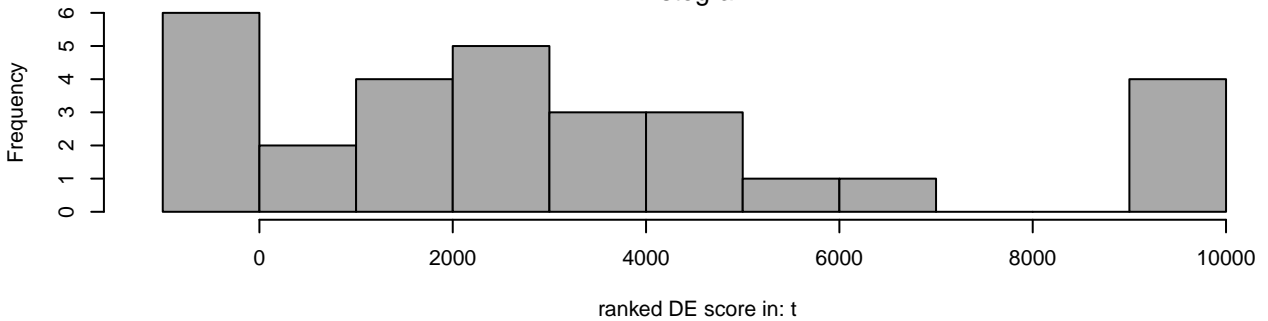
RHOH GTPase cycle

beeswarm plot



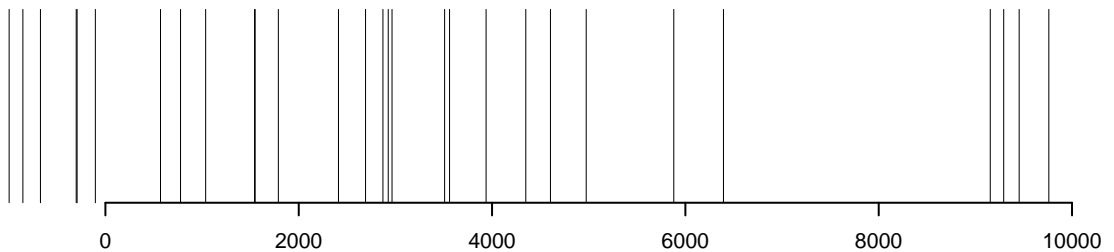
ranked DE score in: t

histogram



ranked DE score in: t

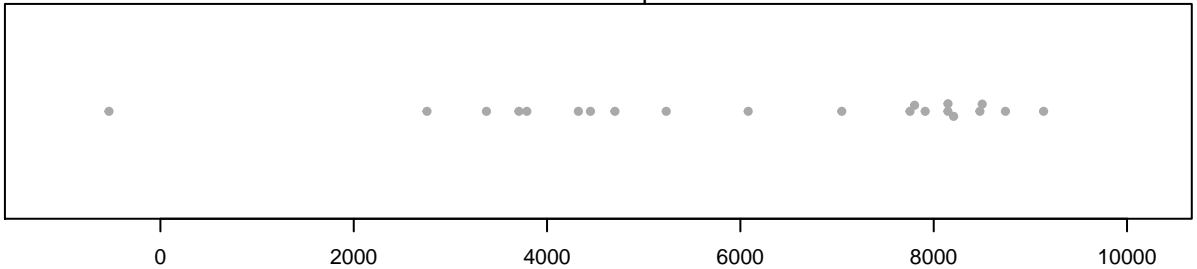
rugplot



ranked DE score in: t

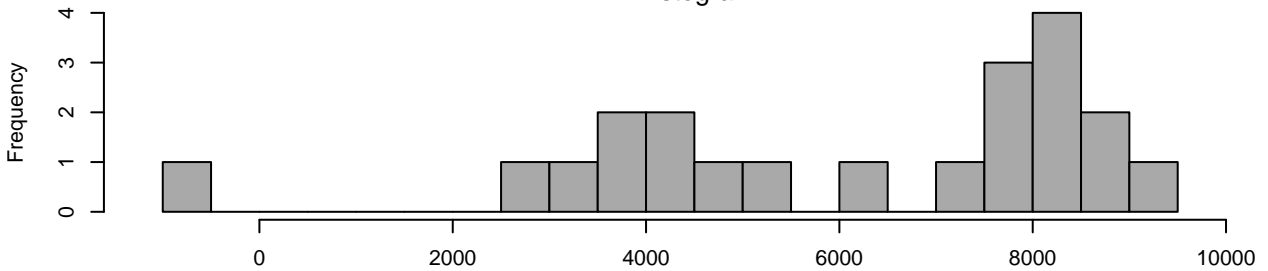
Processing of Capped Intronless Pre-mRNA

beeswarm plot



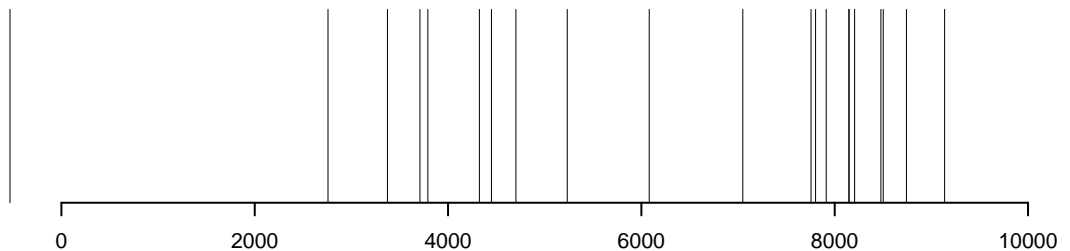
ranked DE score in: t

histogram



ranked DE score in: t

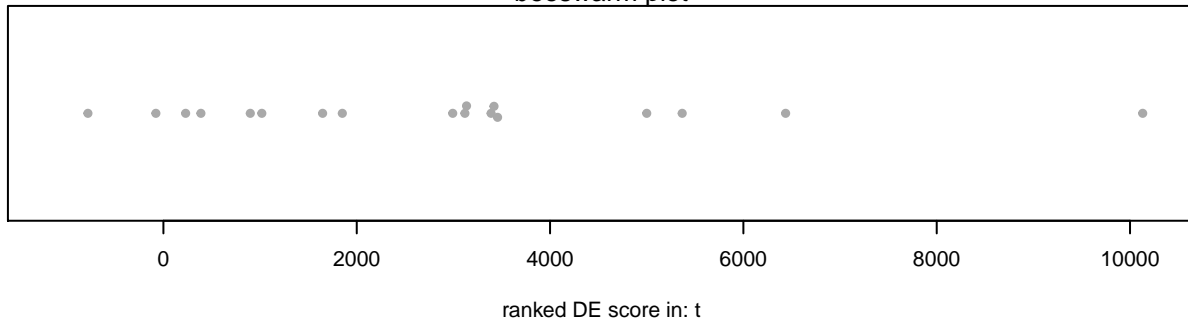
rugplot



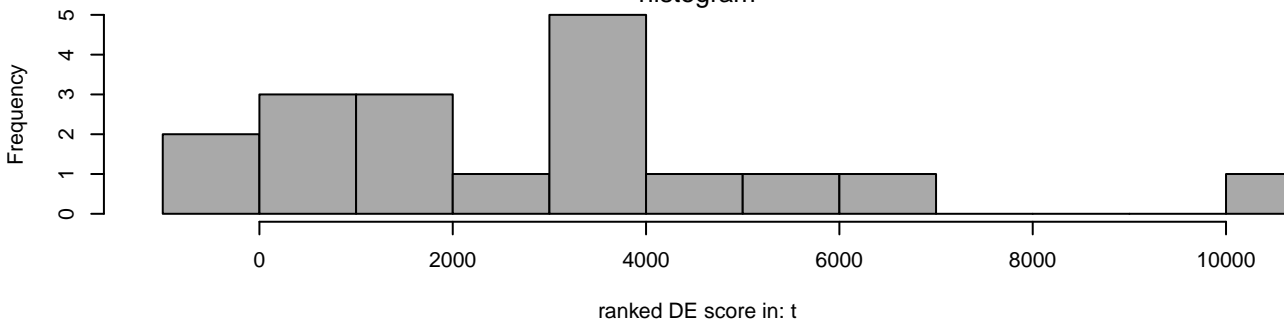
ranked DE score in: t

Mitochondrial tRNA aminoacylation

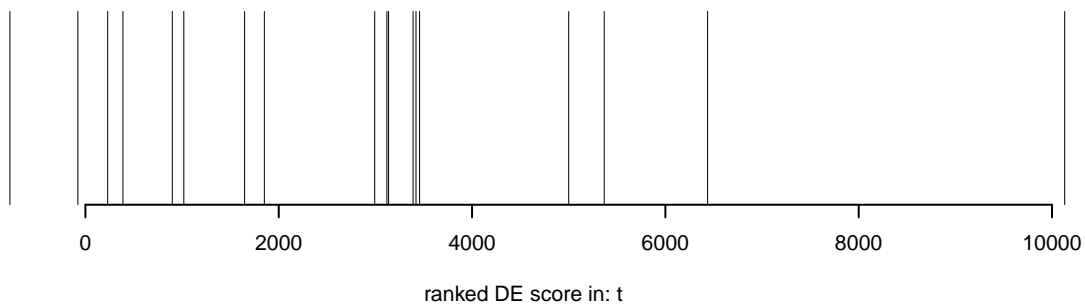
beeswarm plot



histogram

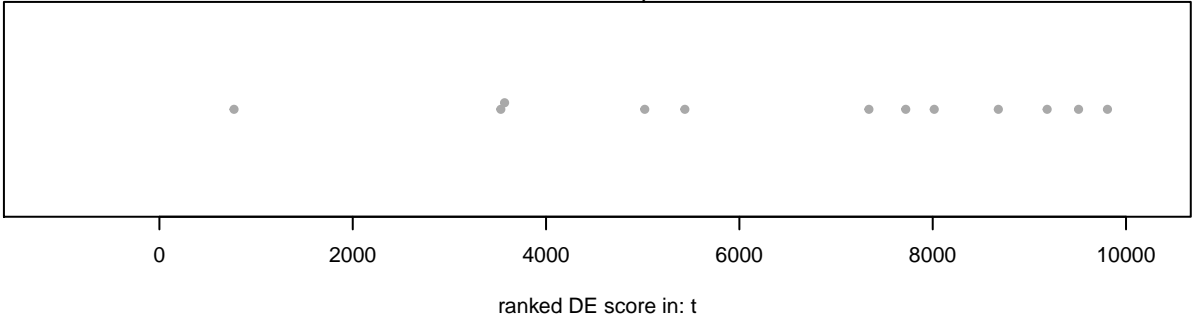


rugplot

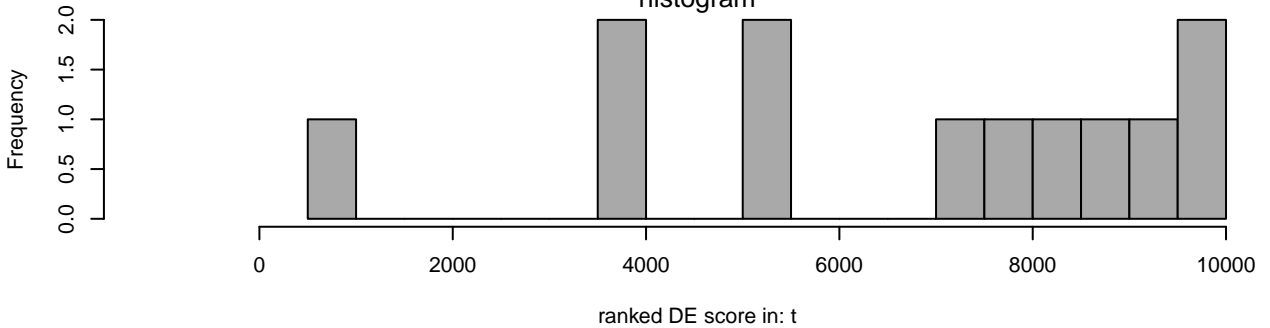


Signaling by Hippo

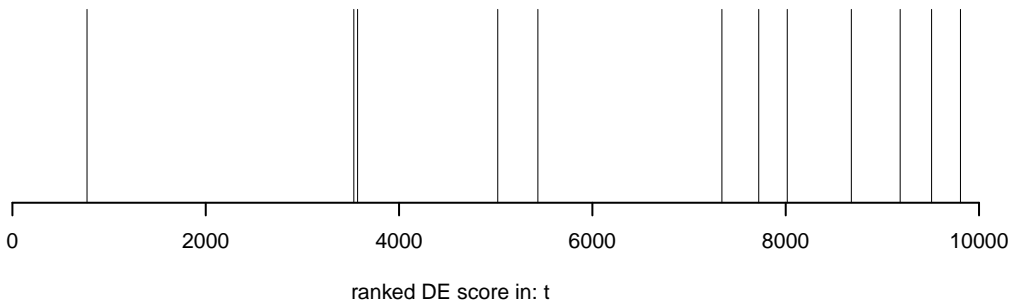
beeswarm plot



histogram

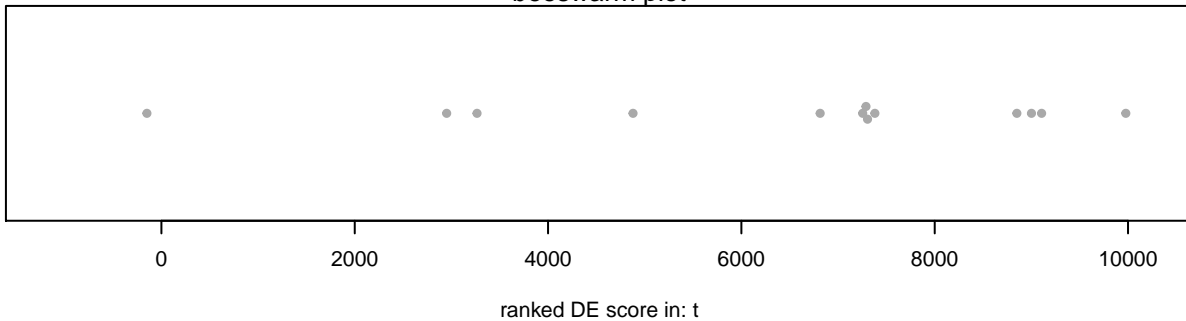


rugplot

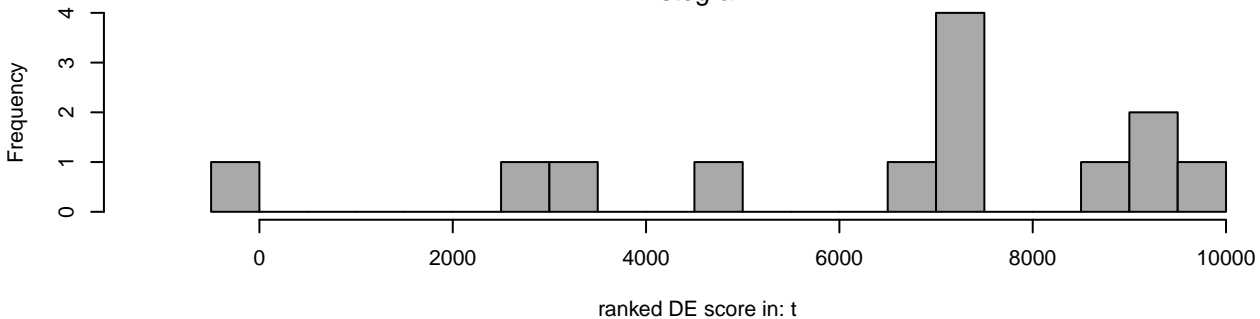


MET promotes cell motility

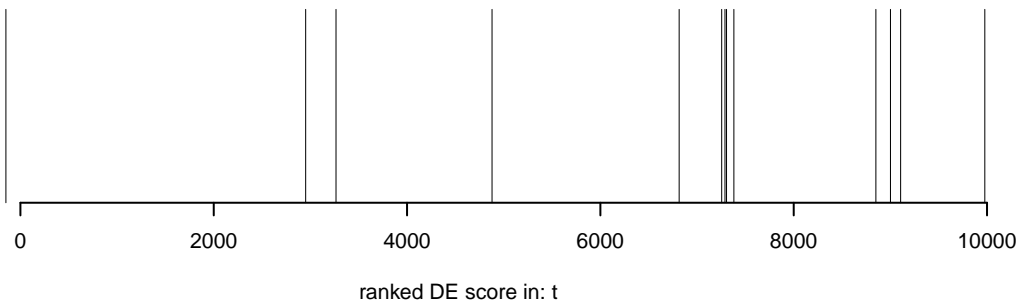
beeswarm plot



histogram

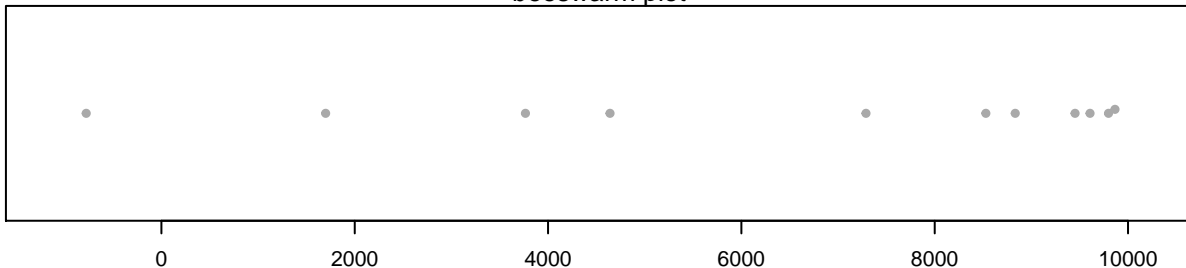


rugplot



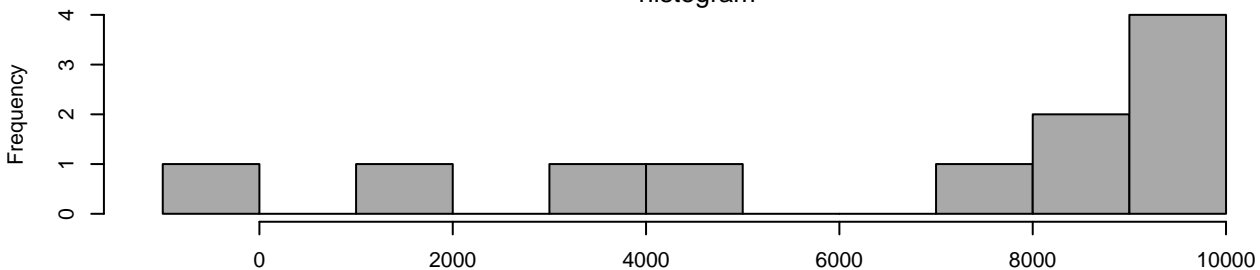
SHC1 events in ERBB2 signaling

beeswarm plot



ranked DE score in: t

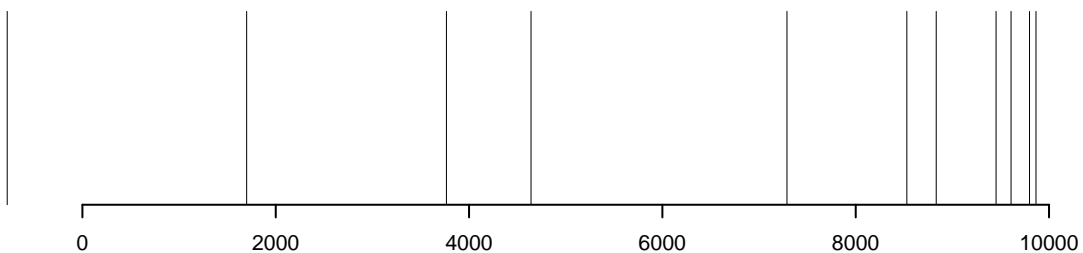
histogram



Frequency

ranked DE score in: t

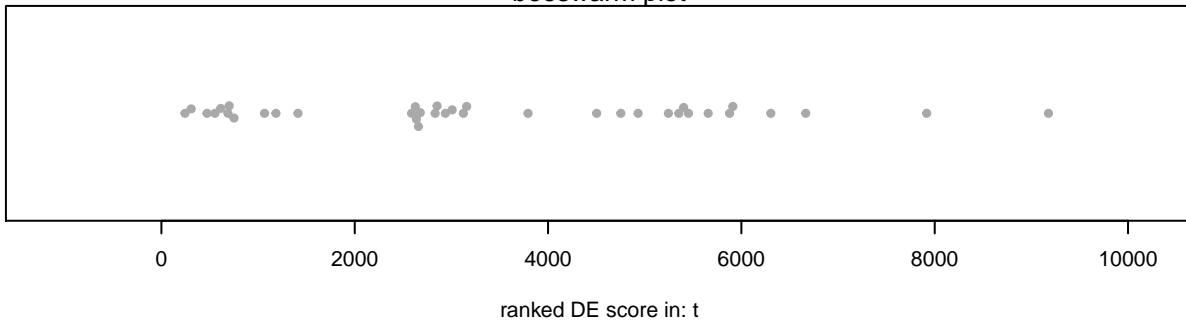
rugplot



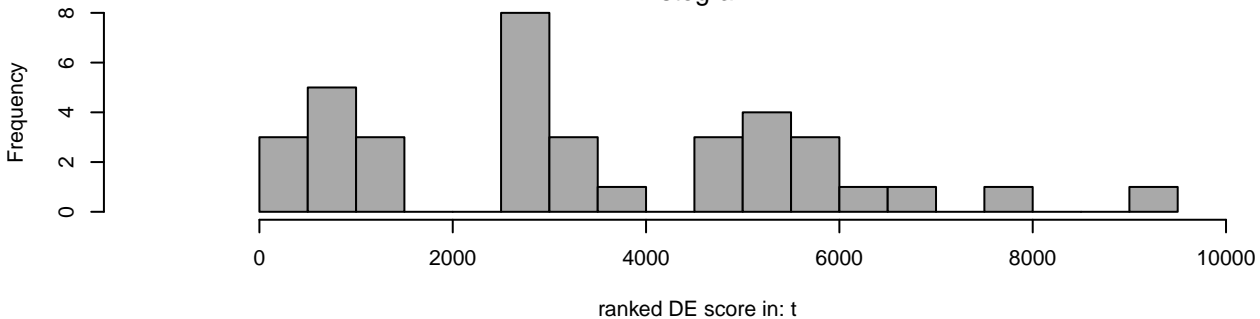
ranked DE score in: t

Transcriptional activity of SMAD2/SMAD3:SMAD4 heterotrimer

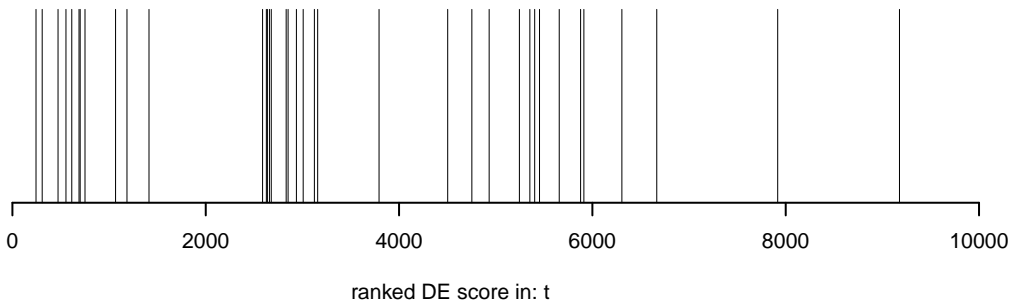
beeswarm plot



histogram

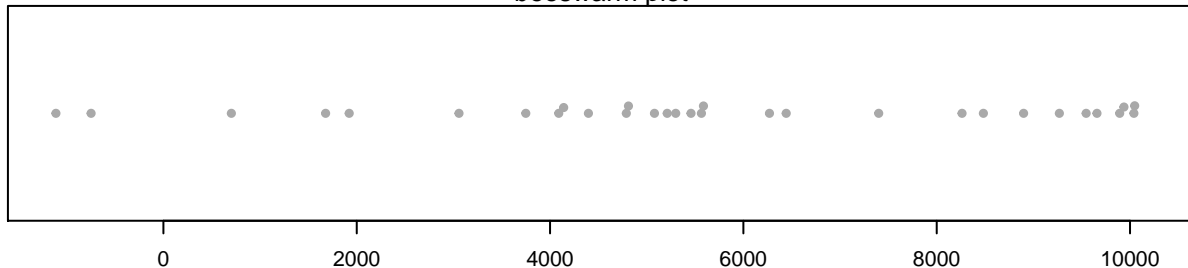


rugplot



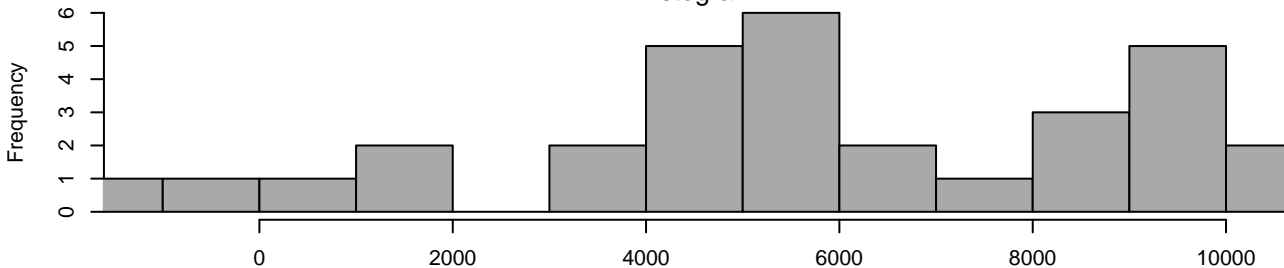
Post-translational modification: synthesis of GPI-anchored proteins

beeswarm plot



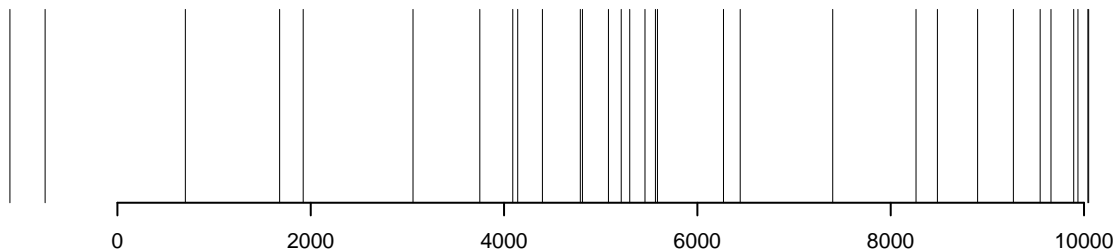
ranked DE score in: t

histogram



ranked DE score in: t

rugplot



ranked DE score in: t