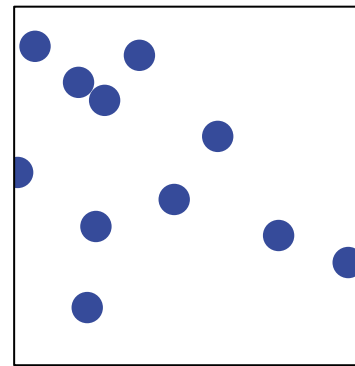
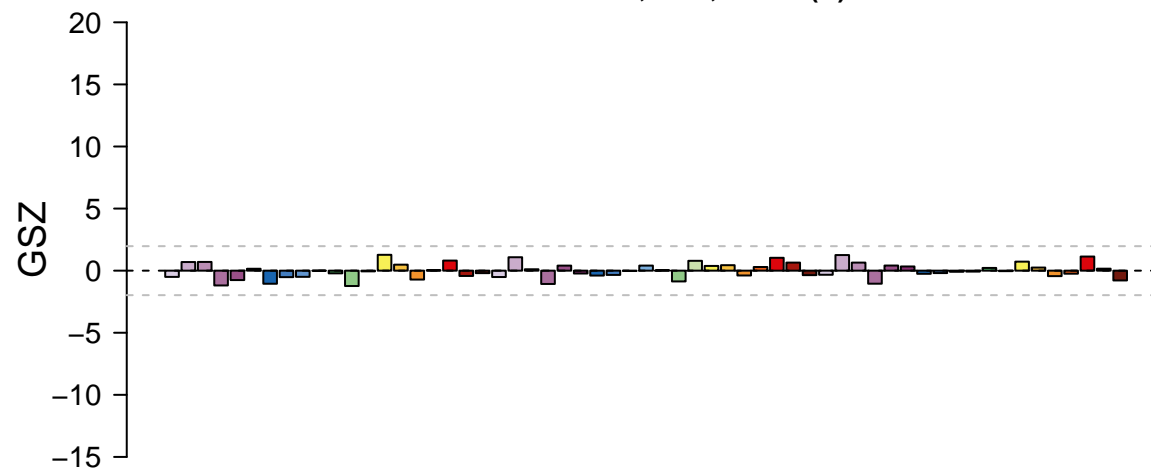
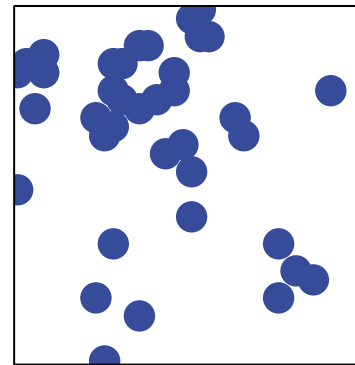
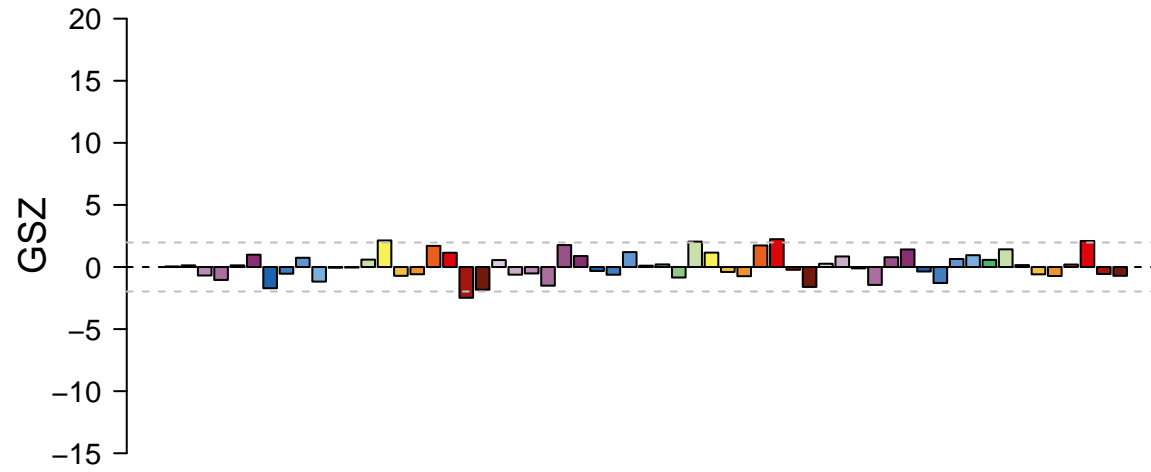


Protein – VAMP, Ykt6, Sec22 (R)



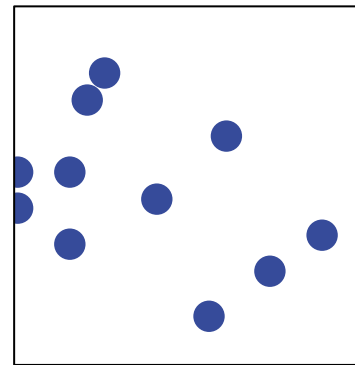
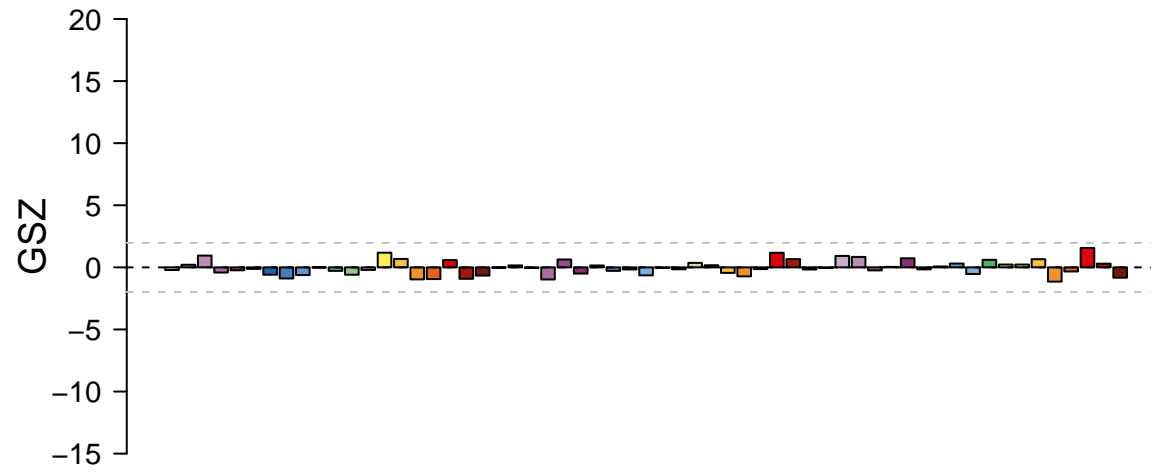
features = 11 , max = 1

Ubiquitin system – Ubiquitin-specific proteases (UBPs)



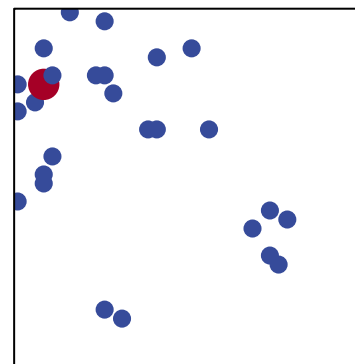
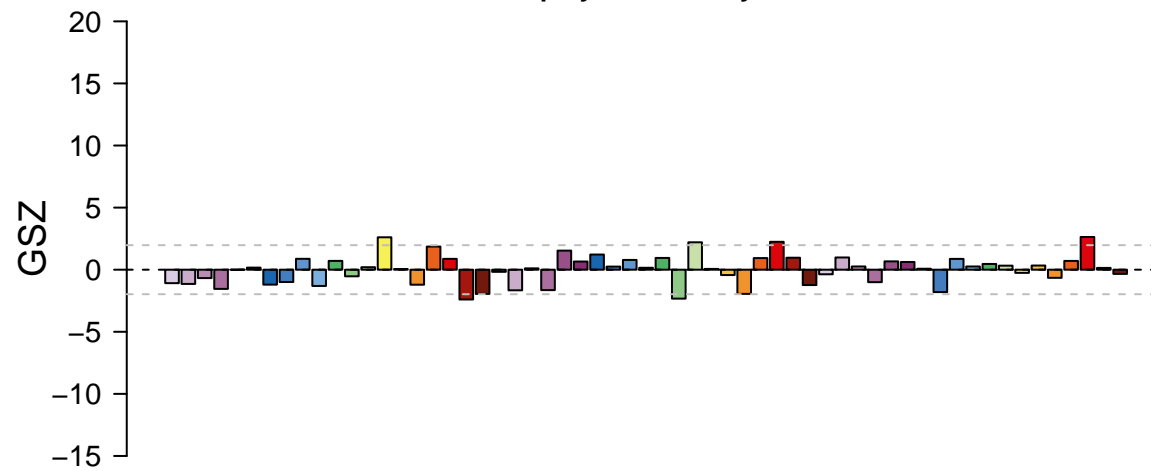
features = 39 , max = 1

Protein – SNAP-25[C] (Qc)



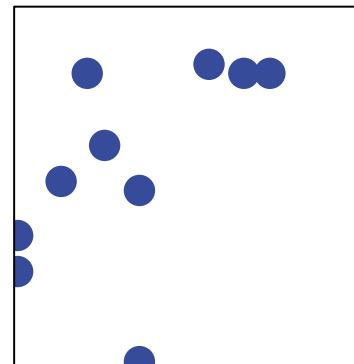
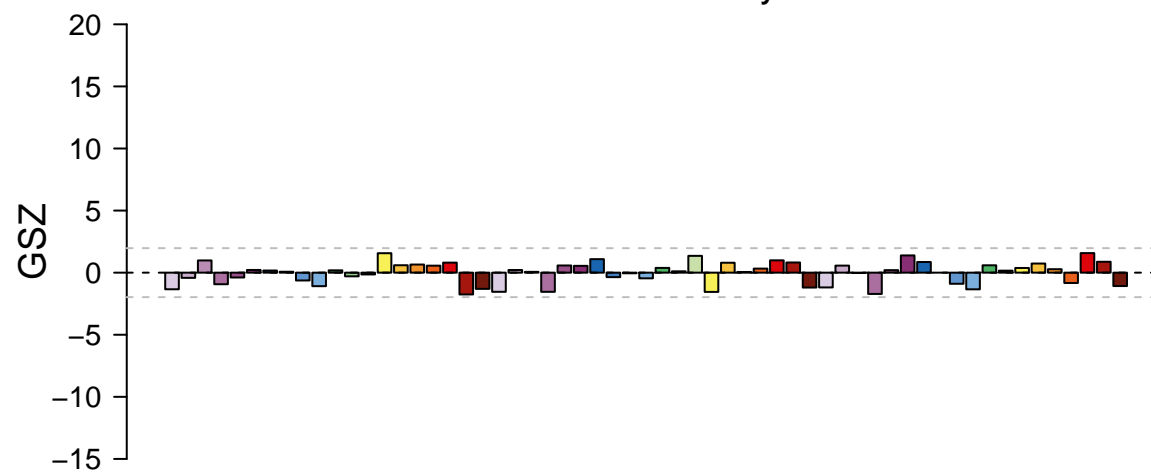
features = 11 , max = 1

RNA polymerase III system



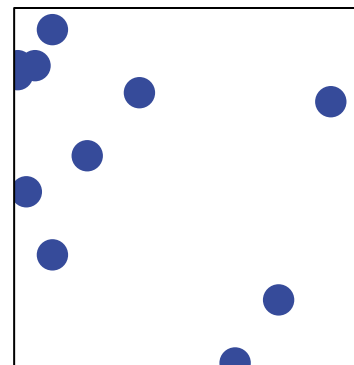
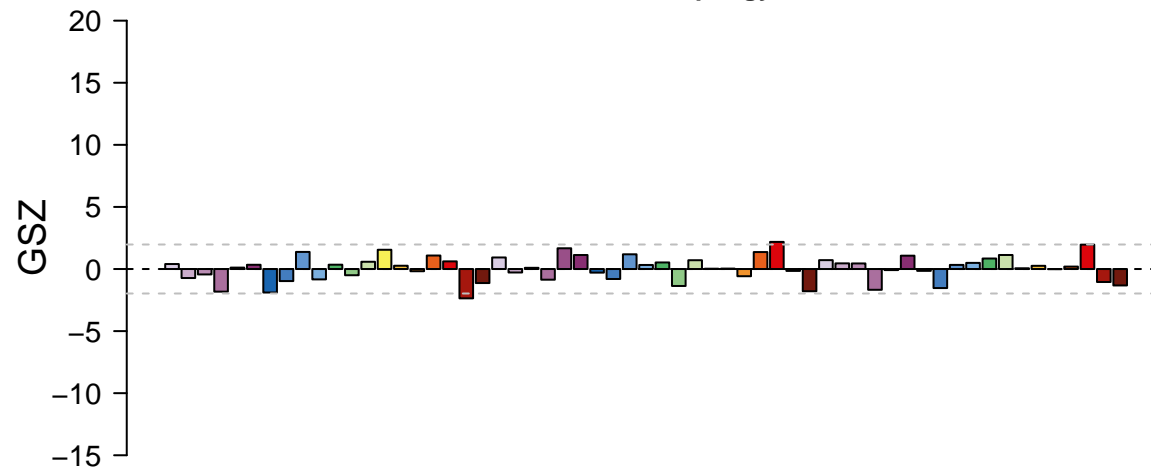
features = 28 , max = 2

Kinase – Wnk family



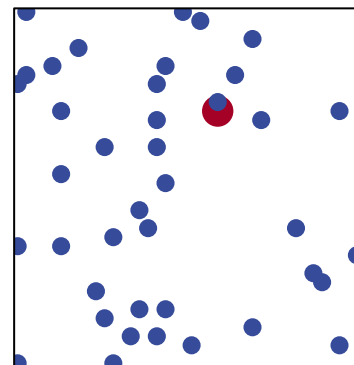
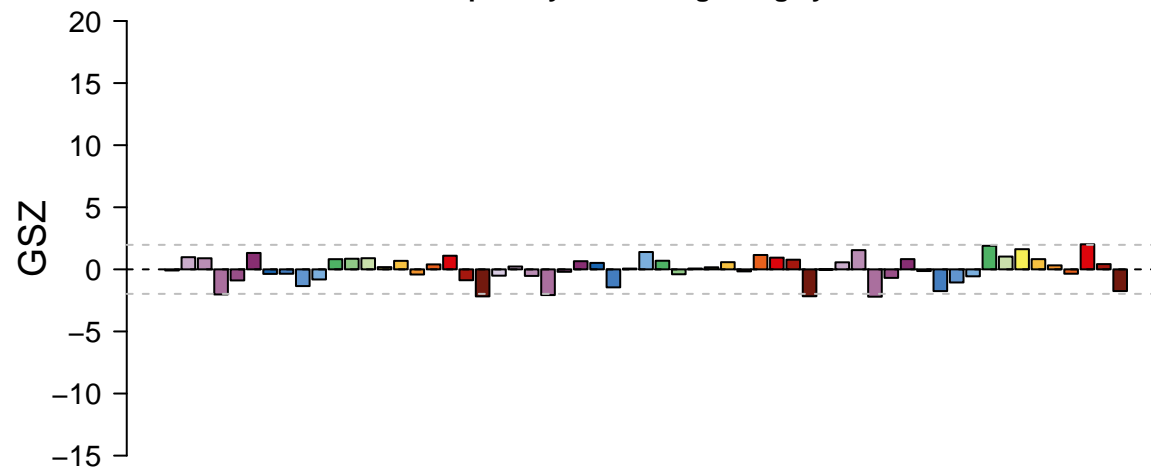
features = 10 , max = 1

Protein – Mitophagy



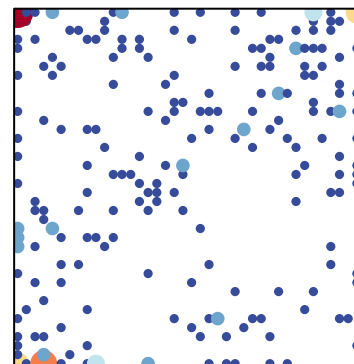
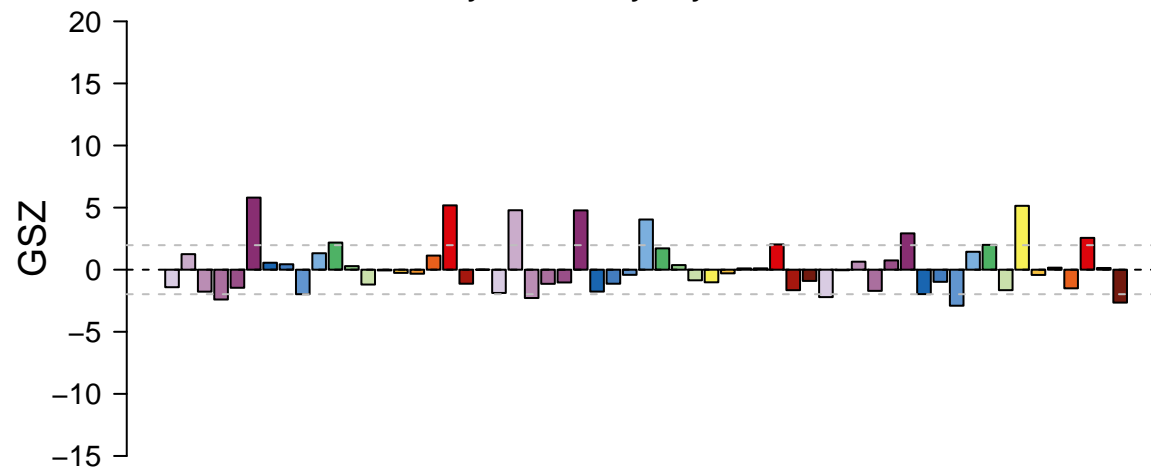
features = 11 , max = 1

Phosphatidylinositol signaling system



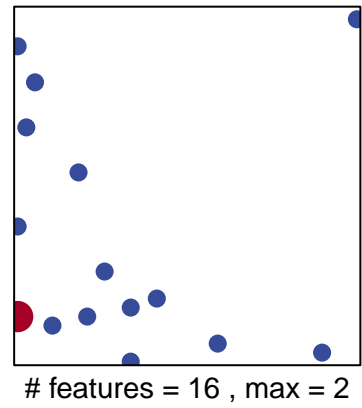
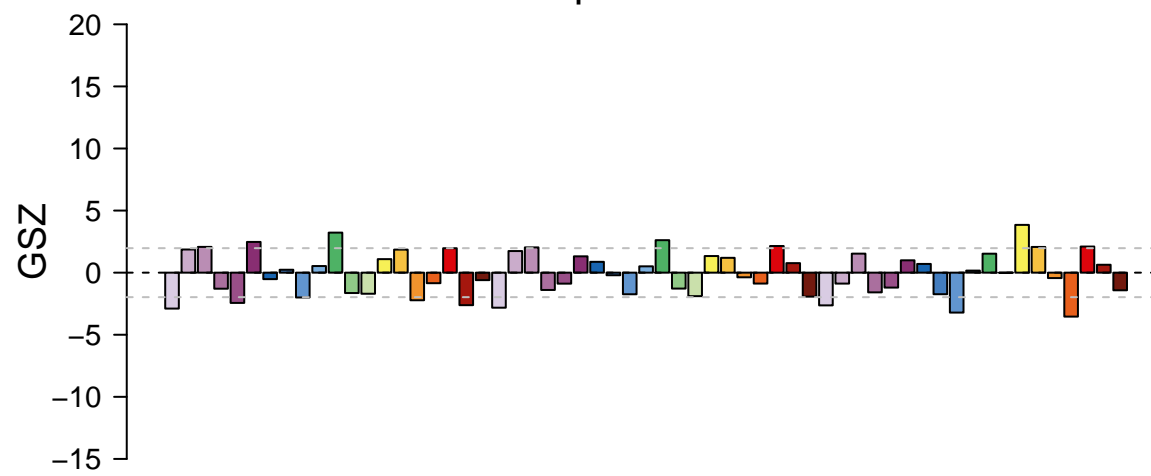
features = 42 , max = 2

Enzyme – 2.4 Glycosyltransferases

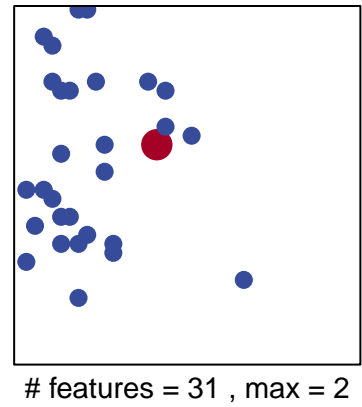
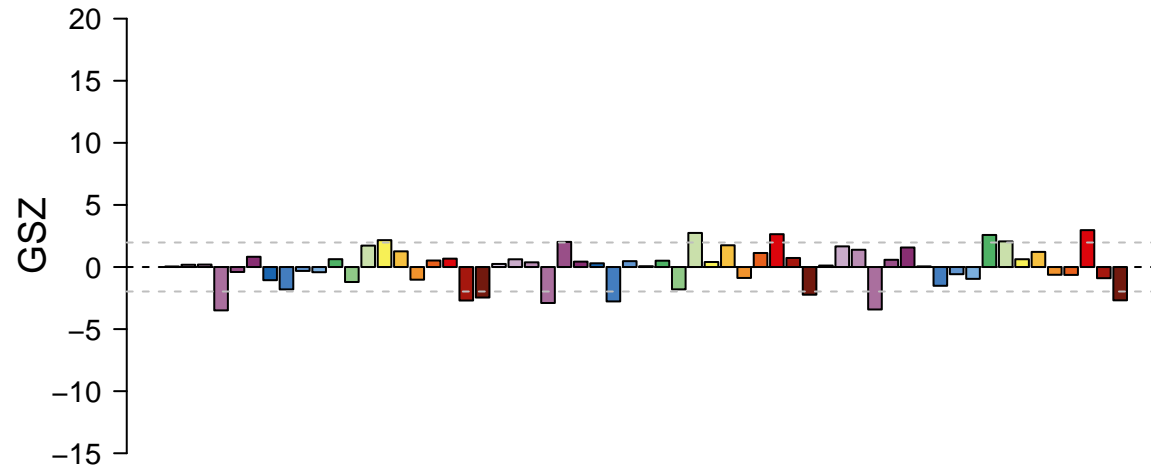


features = 238 , max = 6

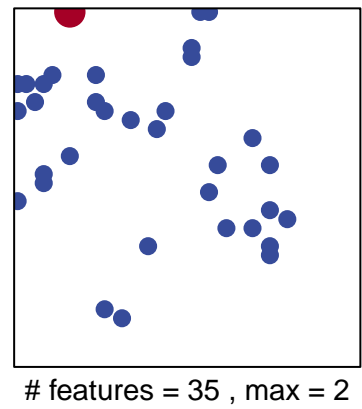
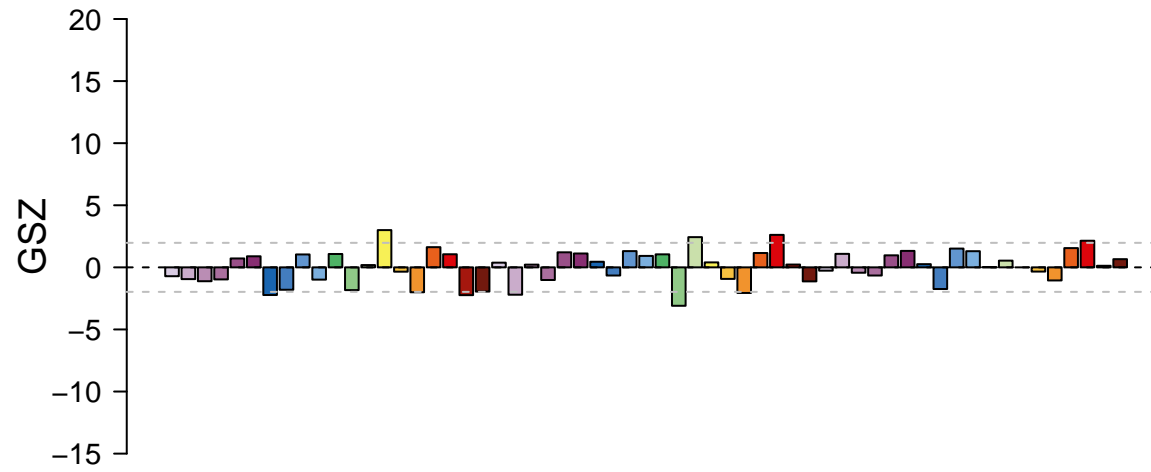
Transcription factors – HSF



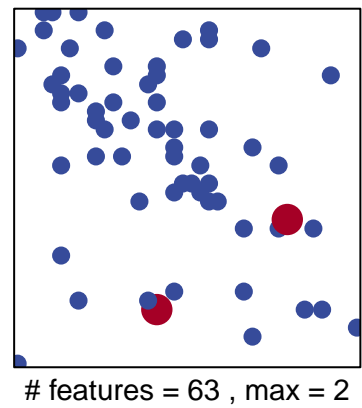
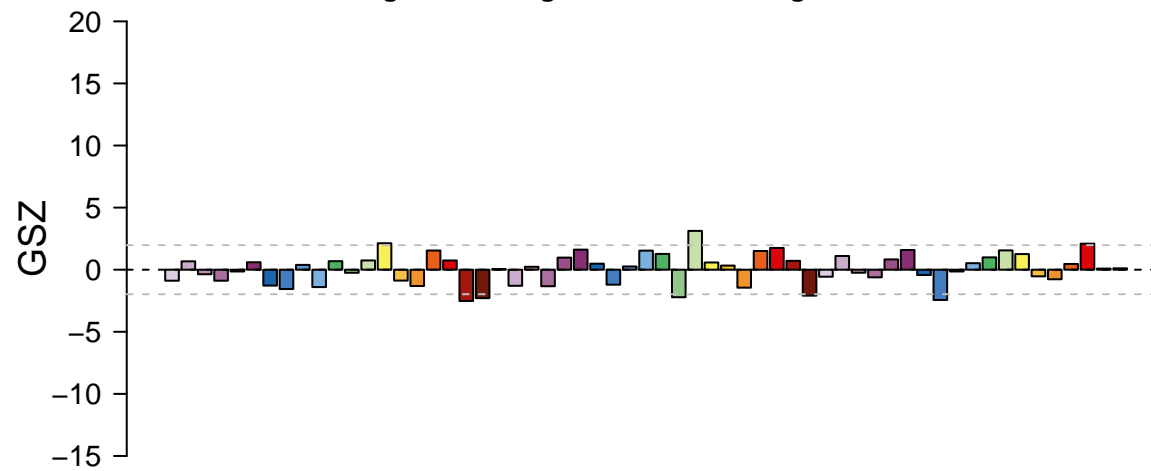
Autophagy – other



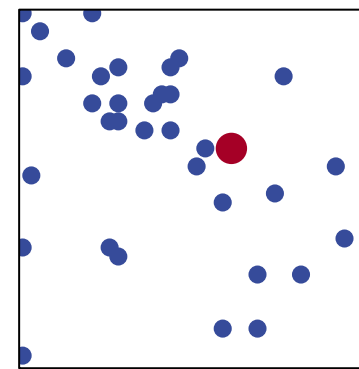
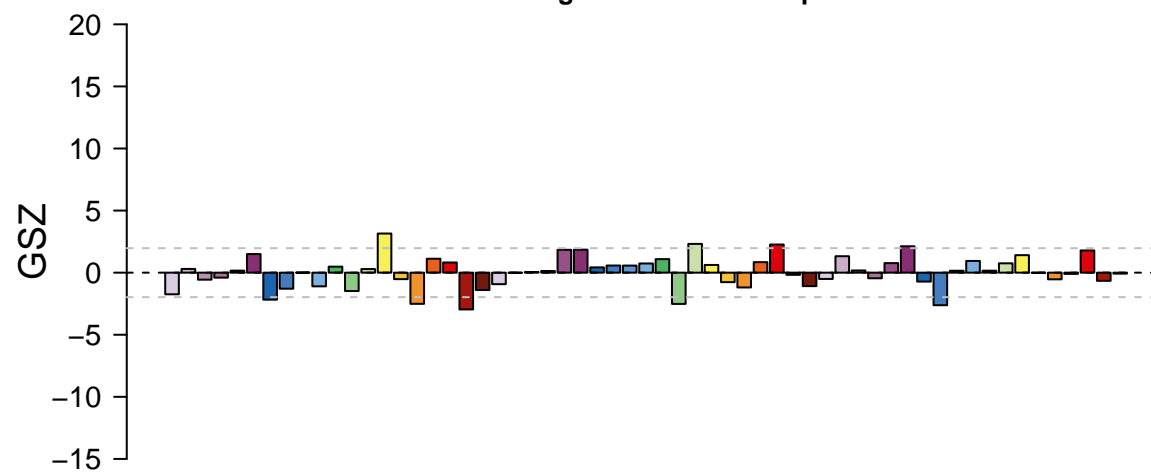
RNA polymerase



Messenger RNA Biogenesis – mRNA degradation factors

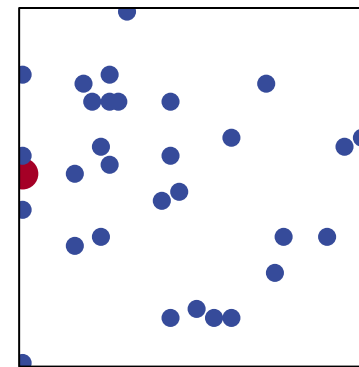
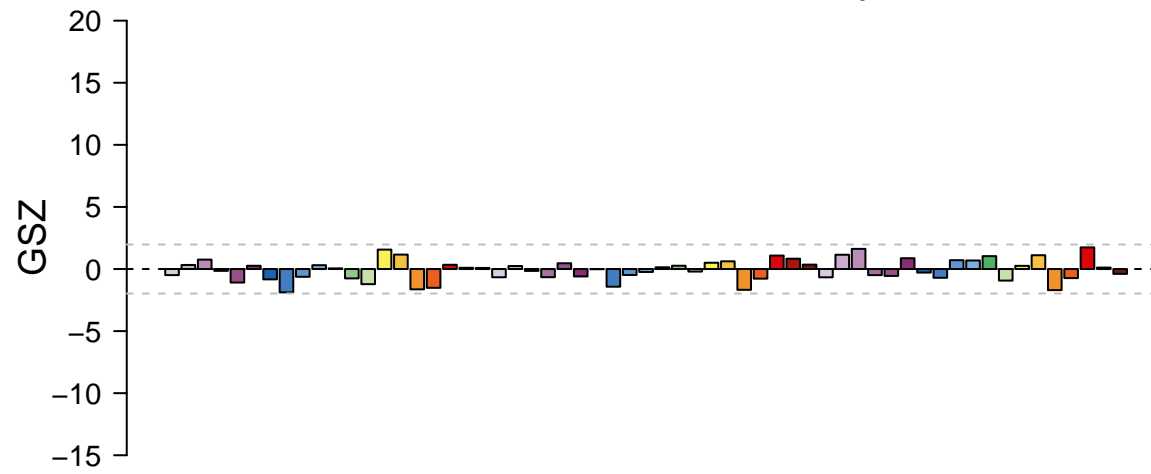


Ribosome biogenesis – Pre-40S particles



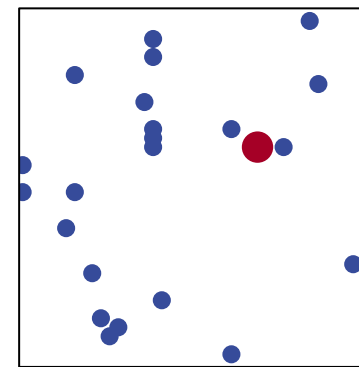
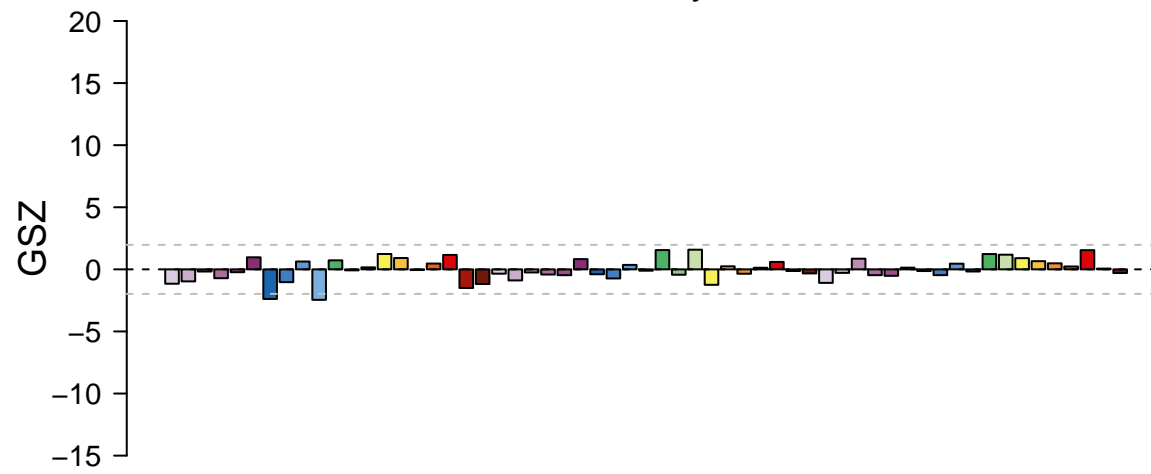
features = 36 , max = 2

SNARE interactions in vesicular transport



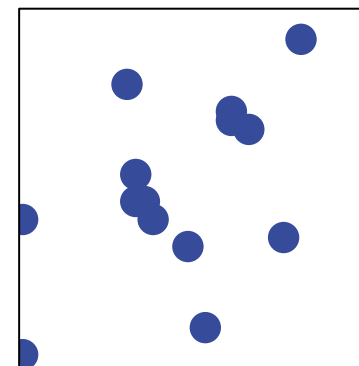
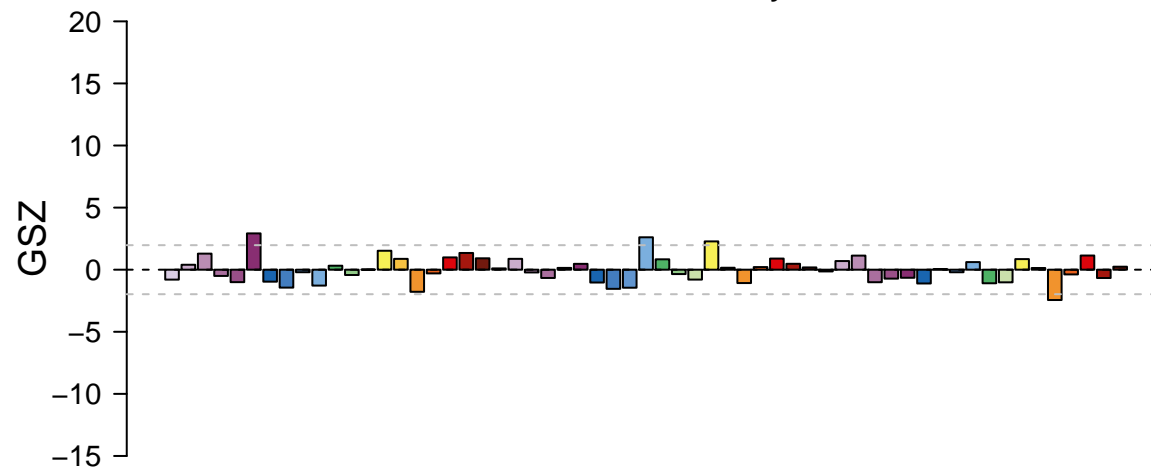
features = 32 , max = 2

Mitochondrial dynamics



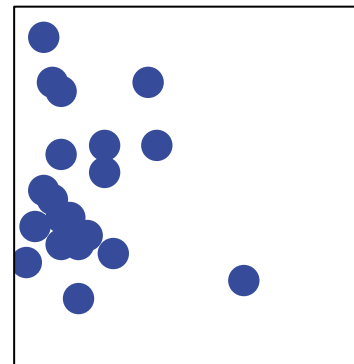
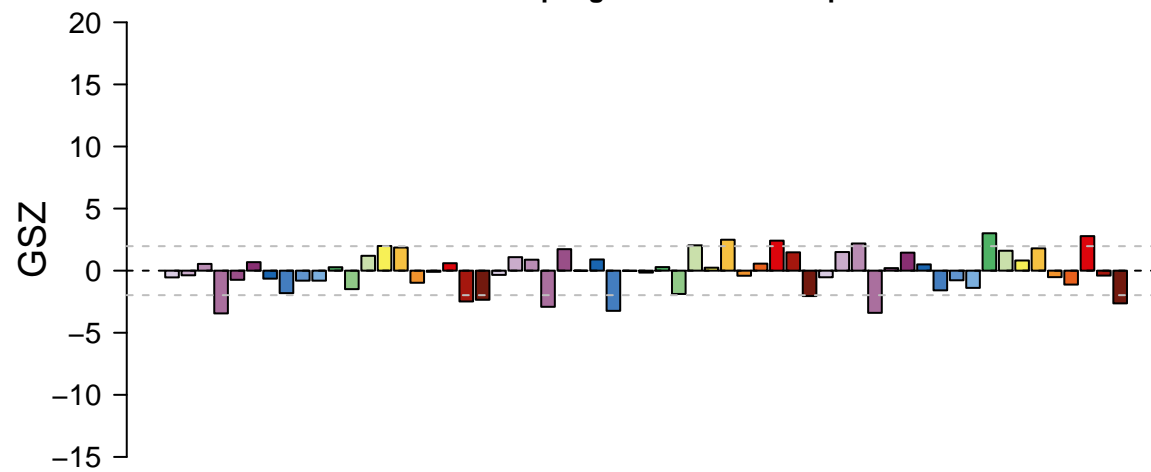
features = 24 , max = 2

Kinase – CDPK family



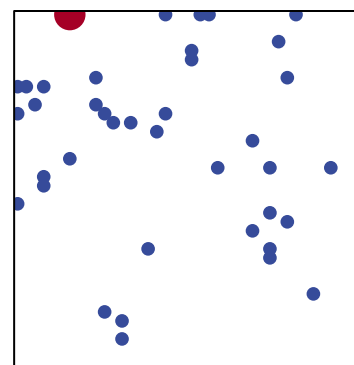
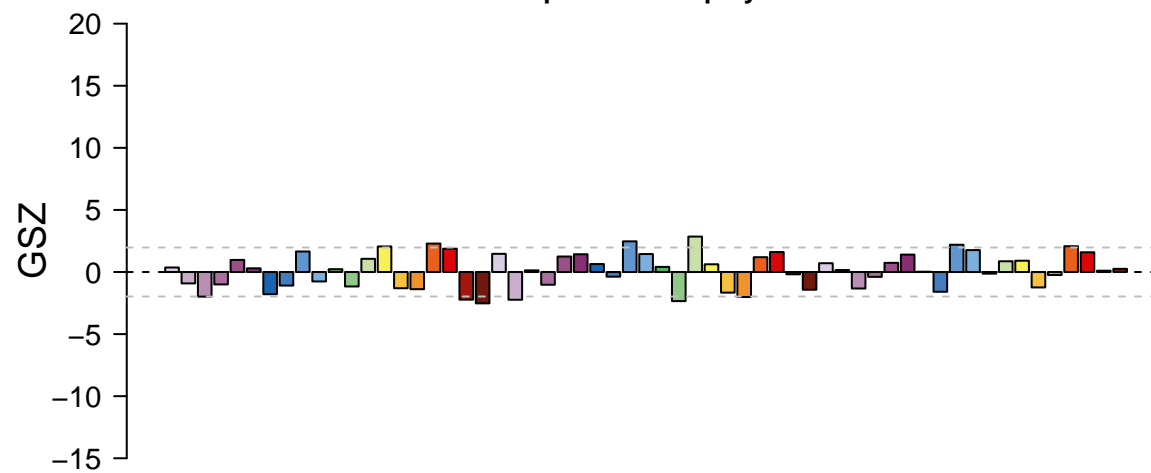
features = 14 , max = 1

Protein – Autophagosome formation proteins



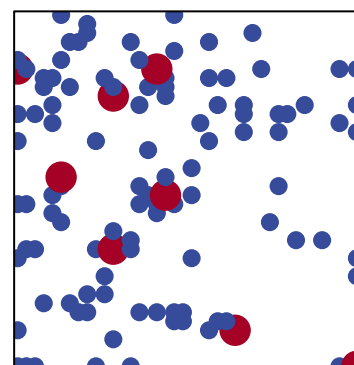
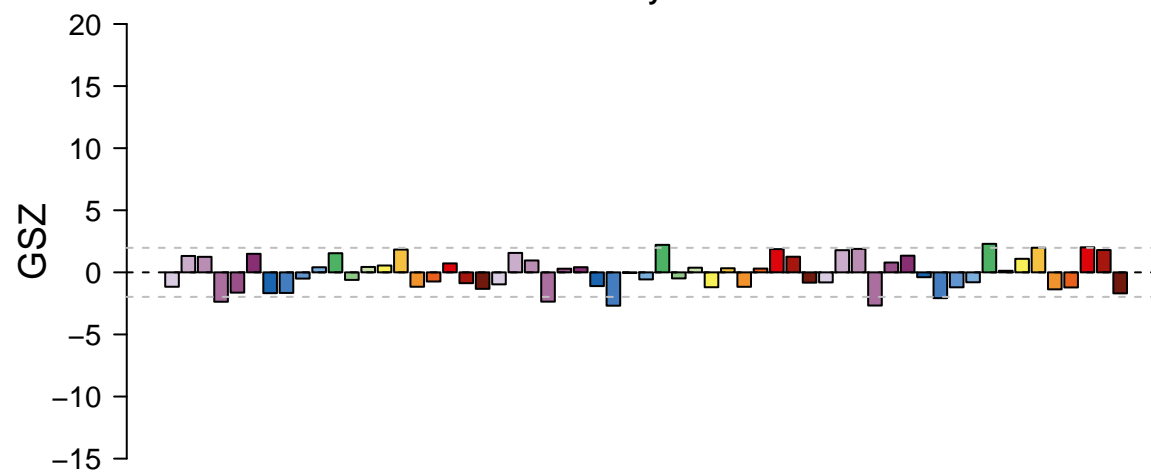
features = 20 , max = 1

Transcription – RNA polymerase



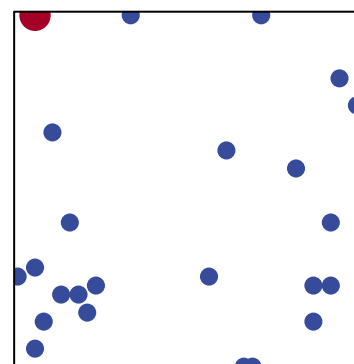
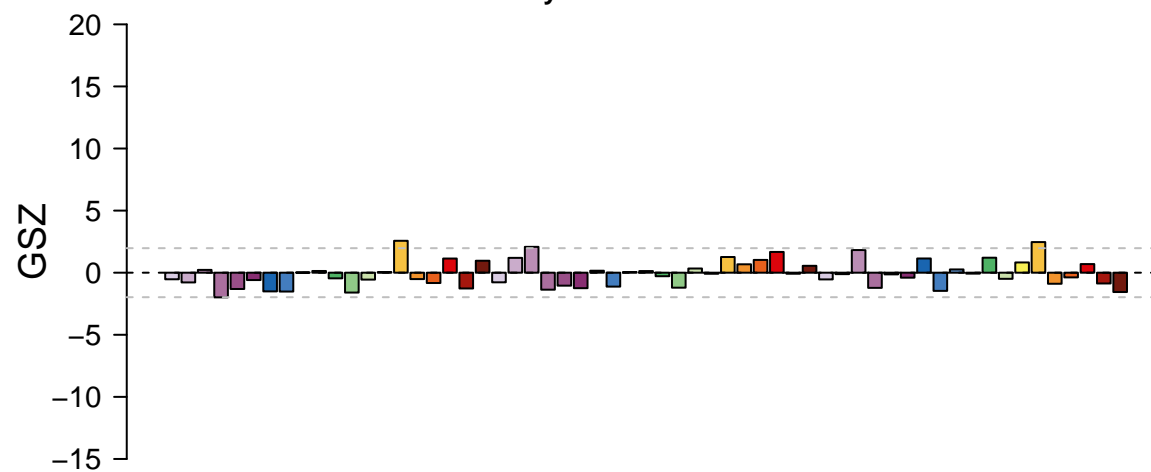
features = 41 , max = 3

Endocytosis



features = 119 , max = 2

Phenylalanine metabolism



features = 25 , max = 2