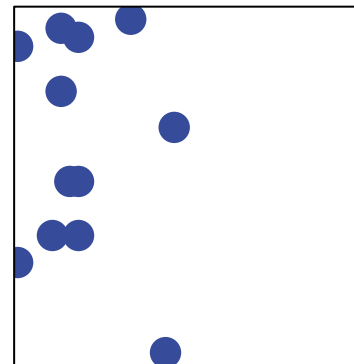
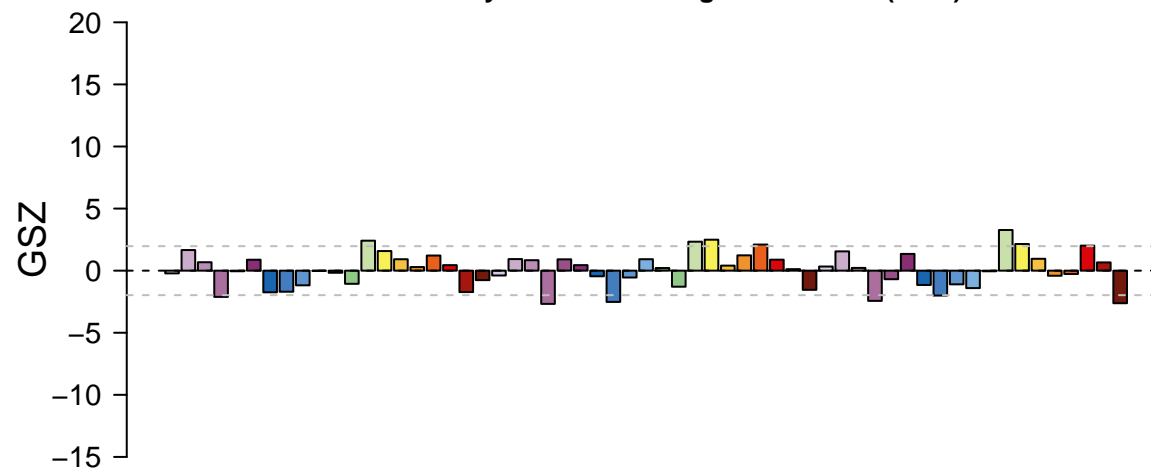
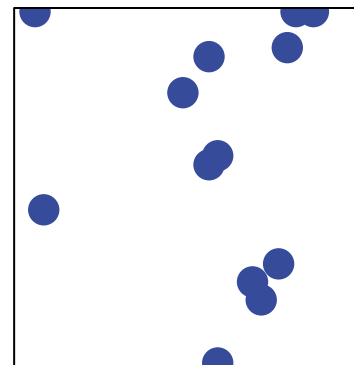
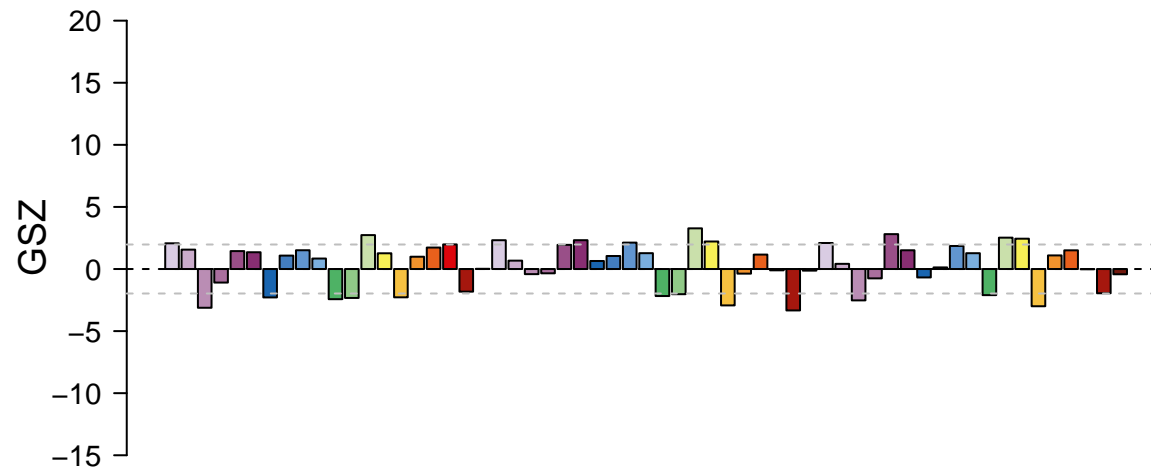


Channel – Cyclic nucleotide–gated channel (CNG)



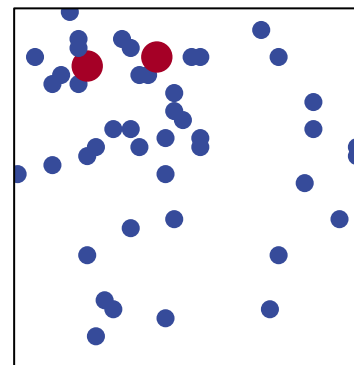
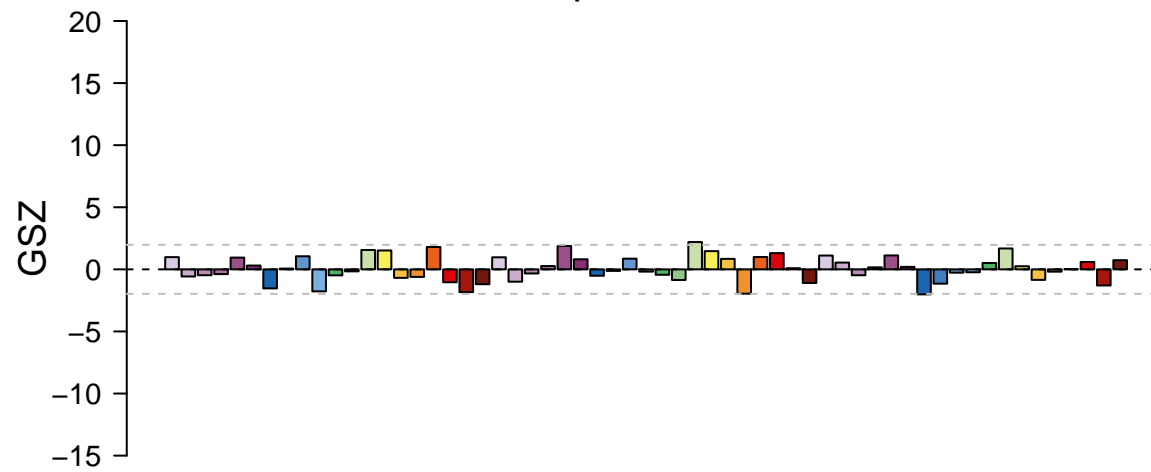
# features = 12 , max = 1

Cofactors and vitamin metabolism – Thiamine metabolism



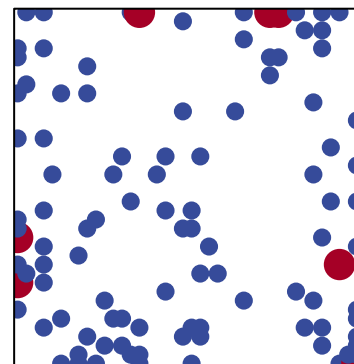
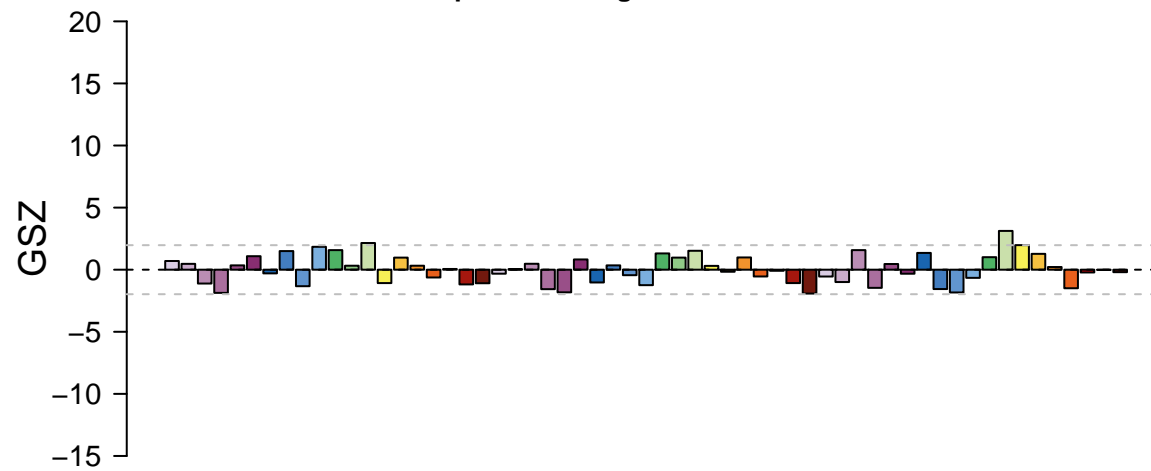
# features = 13 , max = 1

Transcription factors – PHD



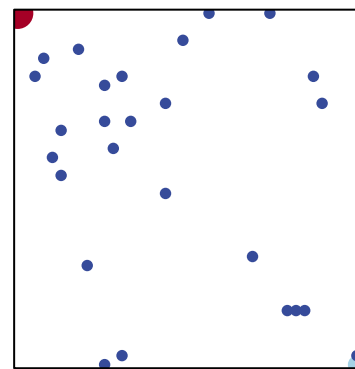
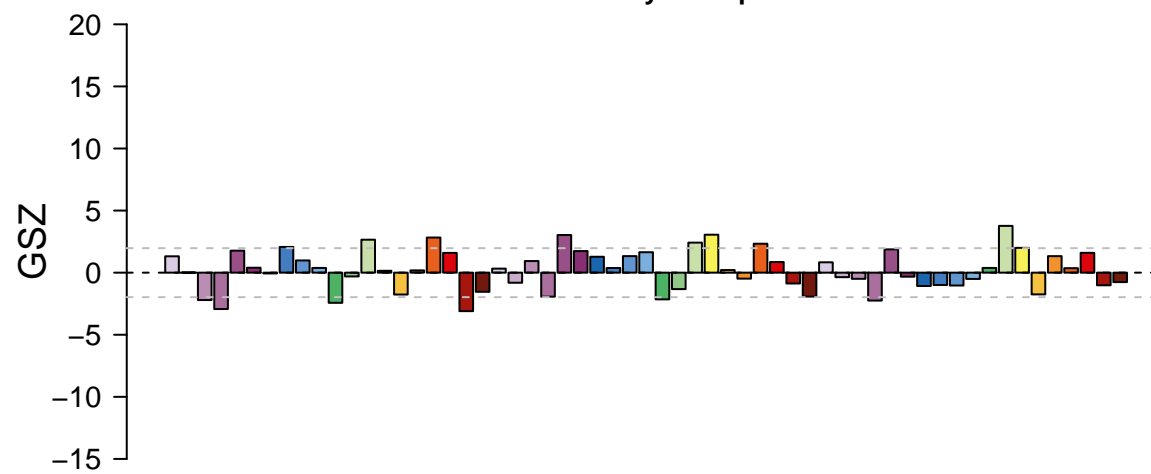
# features = 48 , max = 2

Transporter catalog – Porters cat 66 to 94



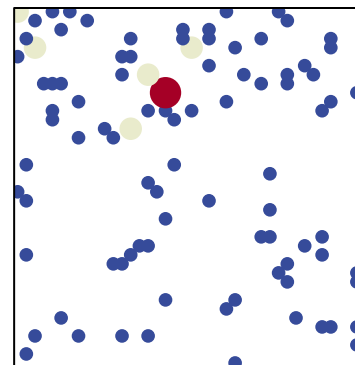
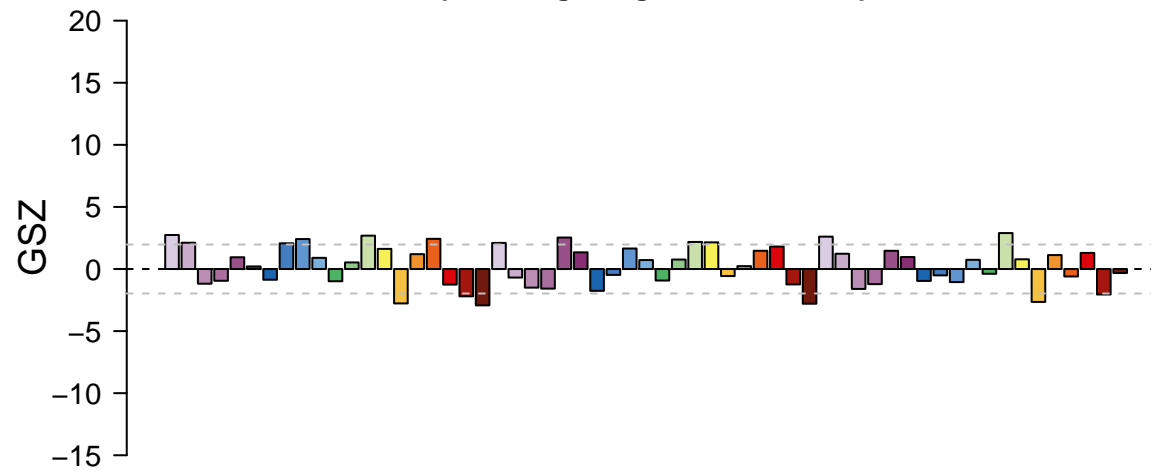
# features = 111 , max = 2

Circadian rhythm – plant



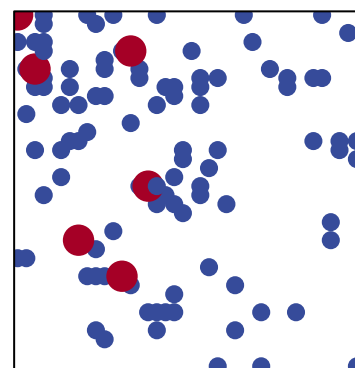
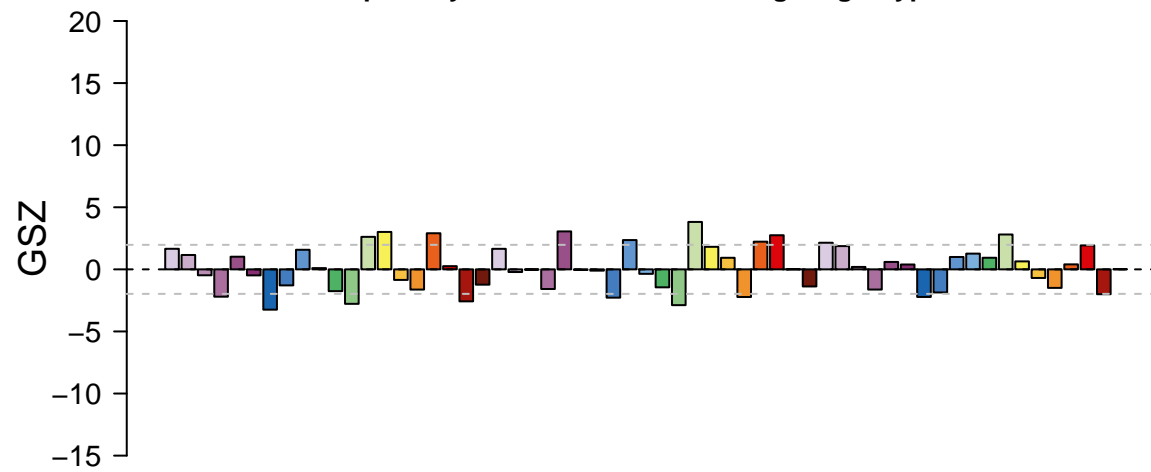
# features = 32 , max = 4

Plant specific signaling – Flower development



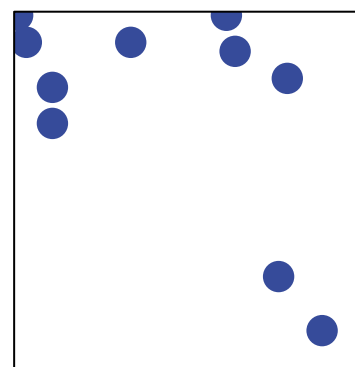
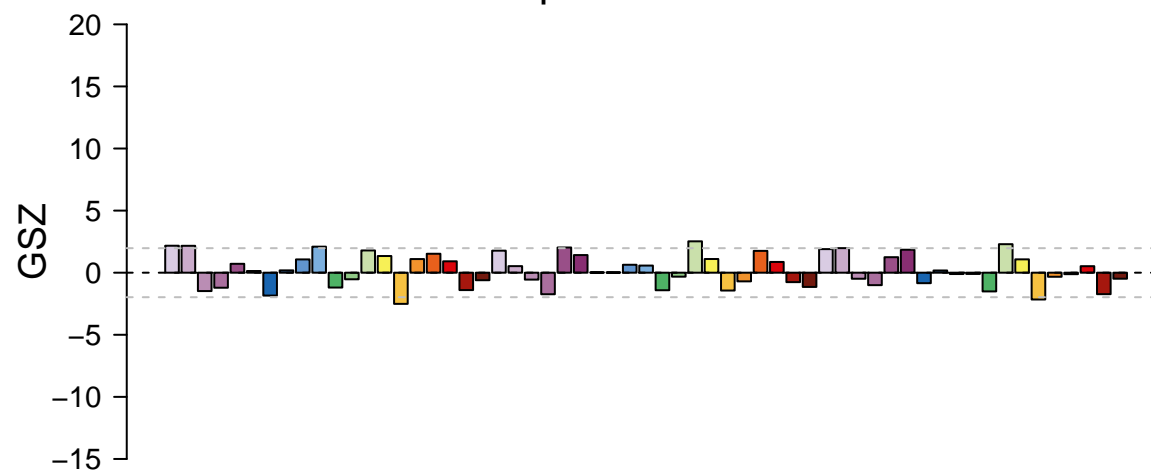
# features = 100 , max = 3

Ubiquitin system – Multi subunit Ring-finger type E3



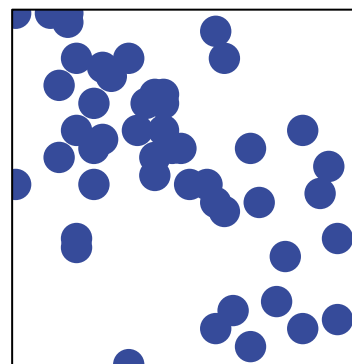
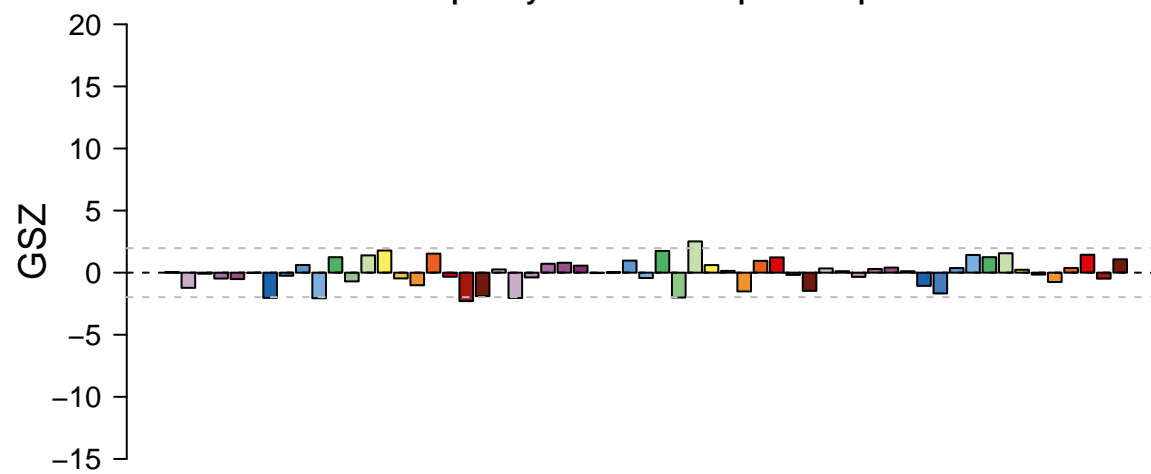
# features = 110 , max = 2

Transcription factors – C2C2-CO



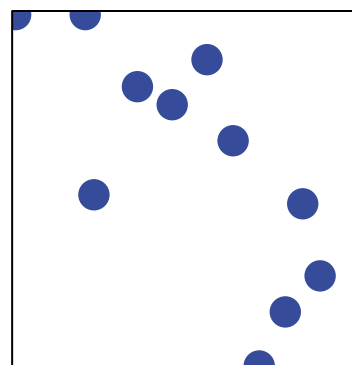
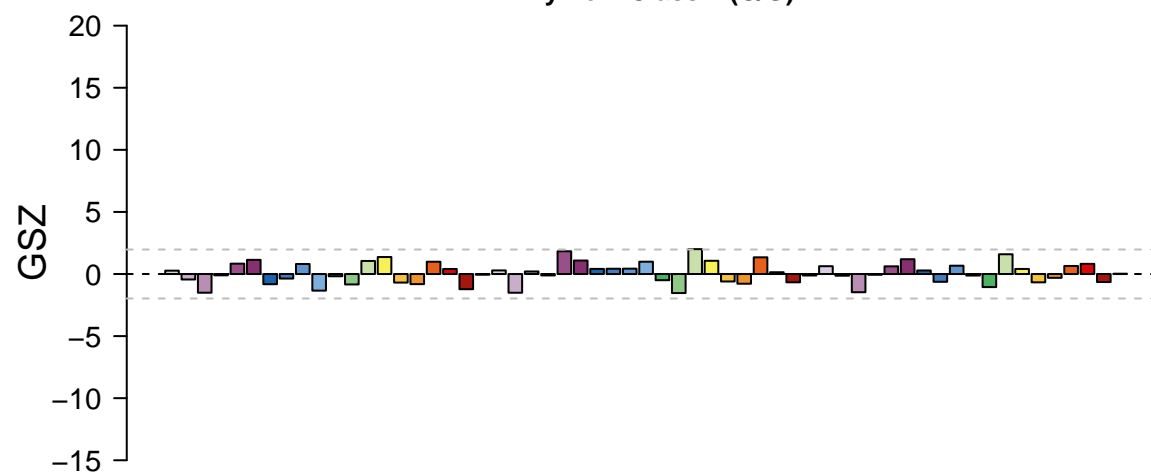
# features = 10 , max = 1

Transport system – Nuclear pore complex



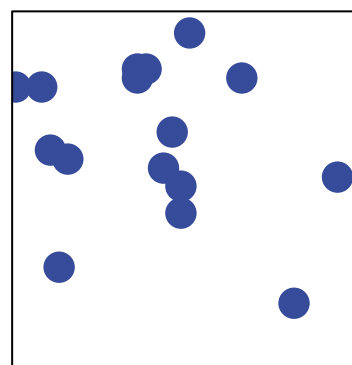
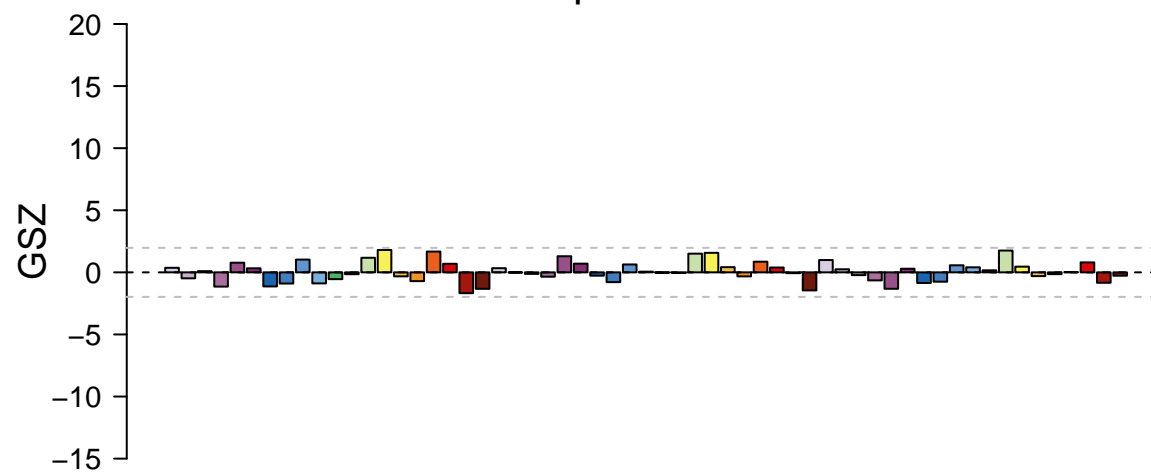
# features = 48 , max = 1

Enzyme – Class II (C/G)



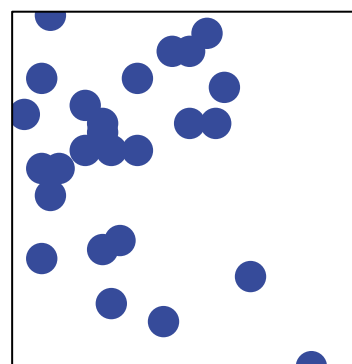
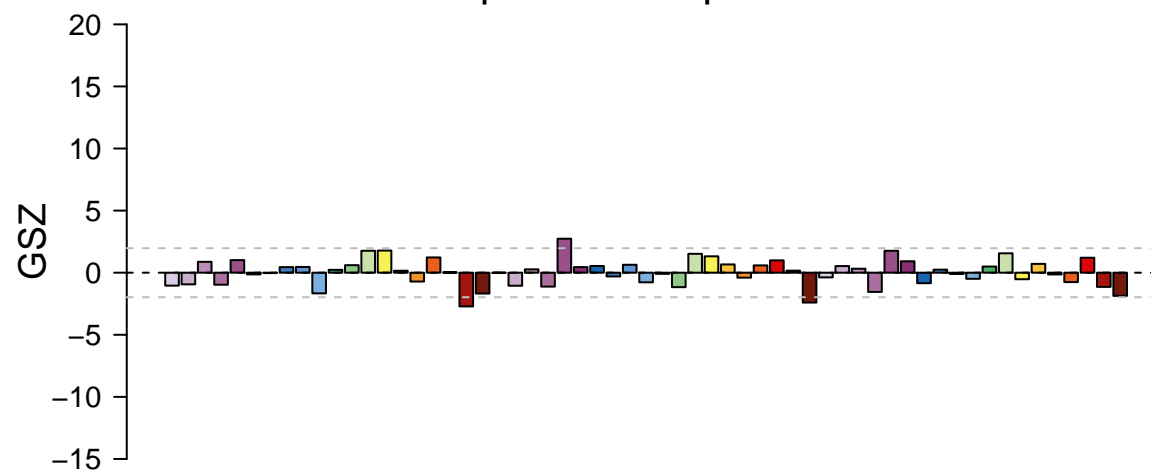
# features = 11 , max = 1

Transcription factors – FHA



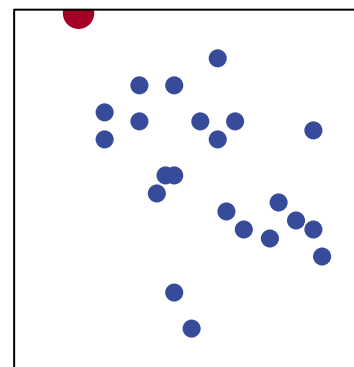
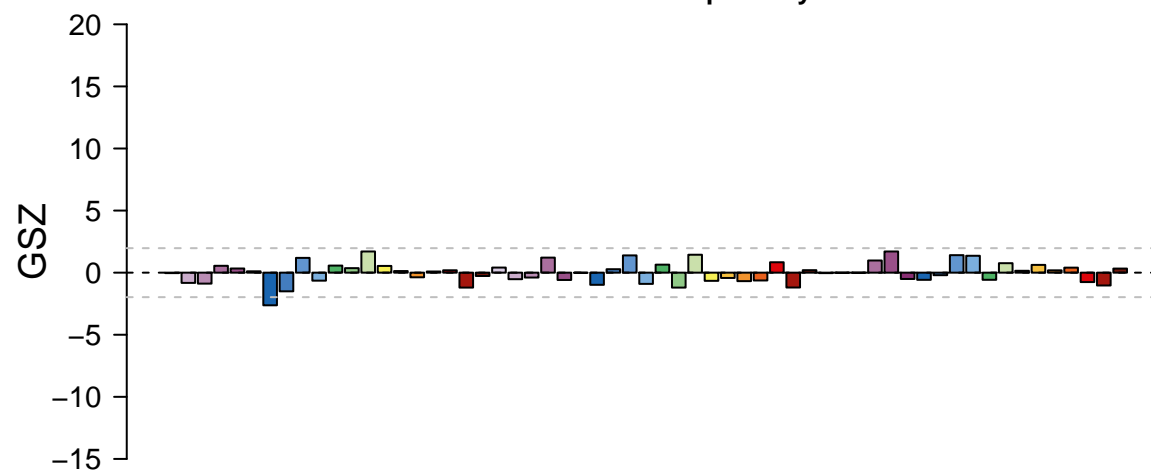
# features = 16 , max = 1

Transcription factors – Orphans FAR-RED



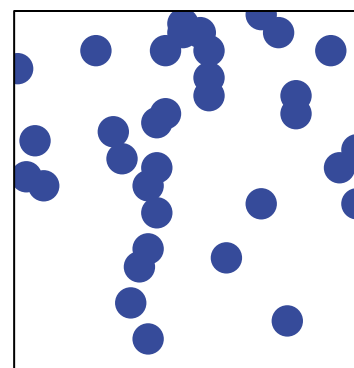
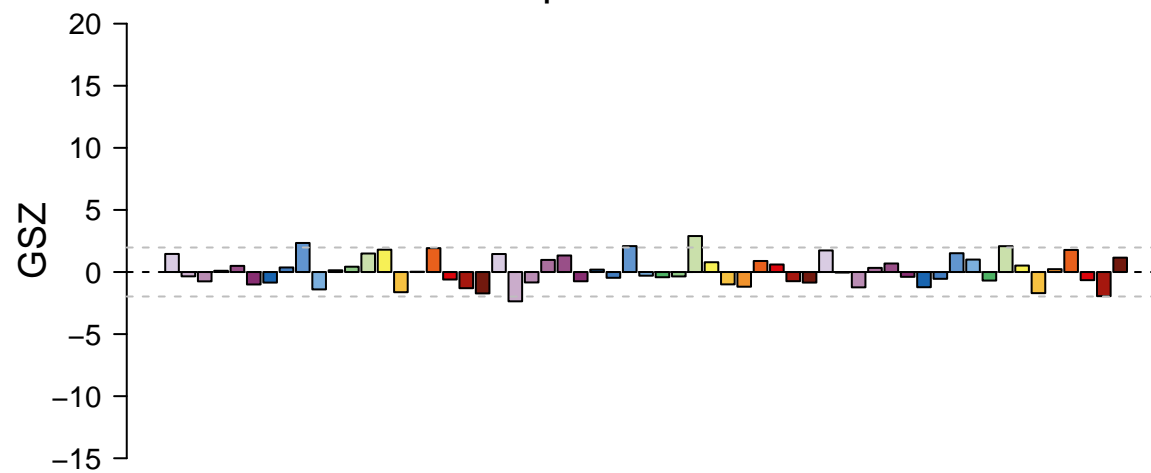
# features = 26 , max = 1

Protein – Retrieval pathways



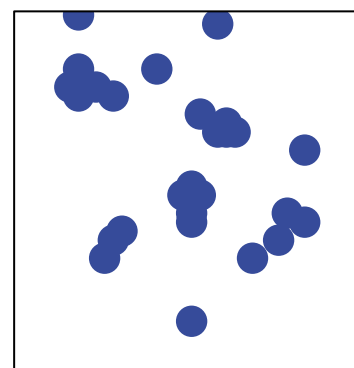
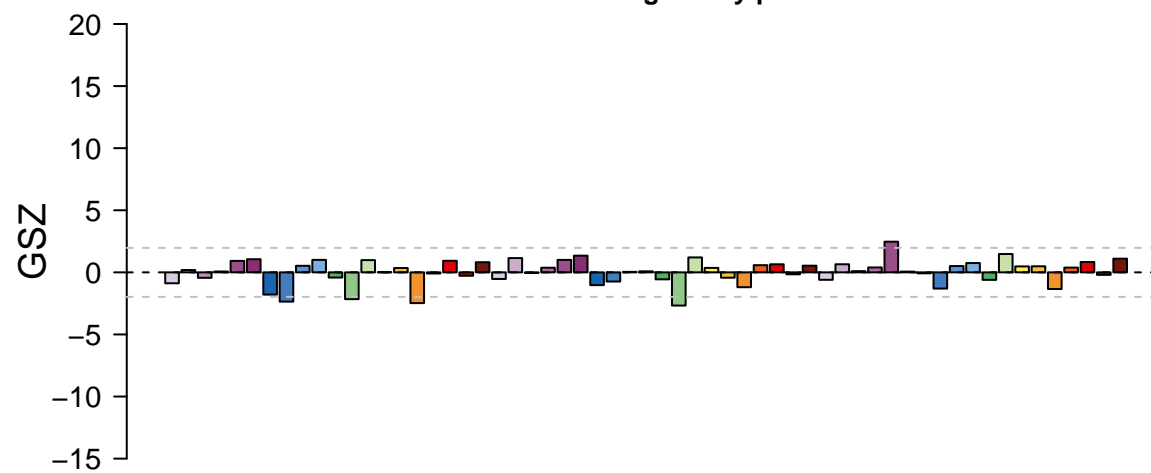
# features = 24 , max = 2

Transcription factors – SET PCG



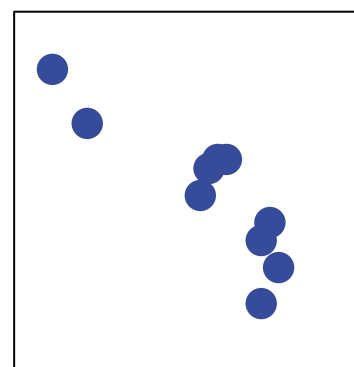
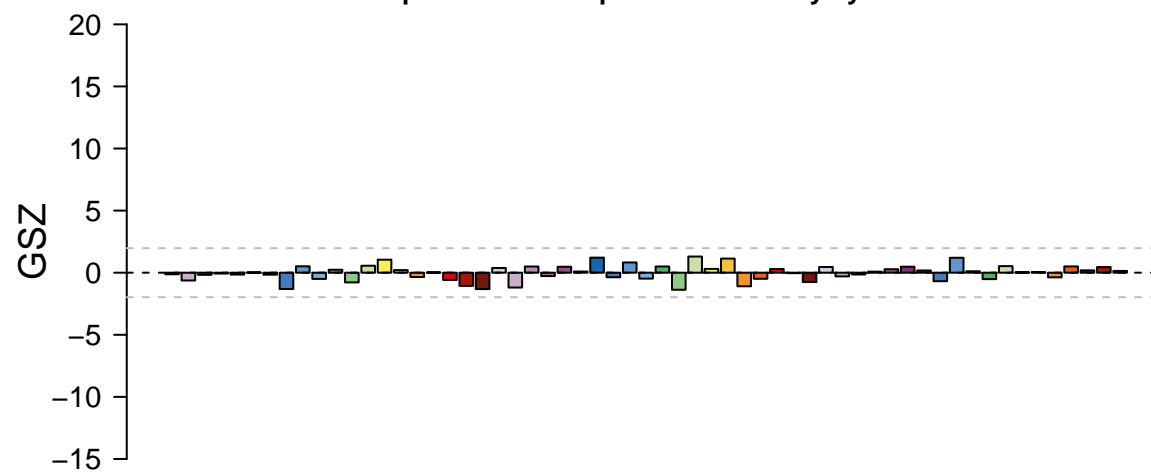
# features = 34 , max = 1

Proteasome – Regulatory particles



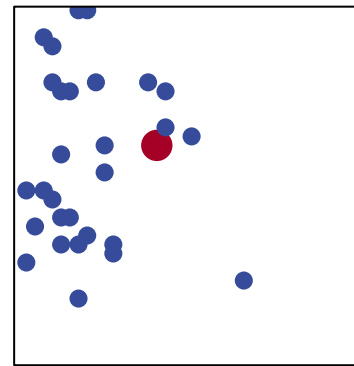
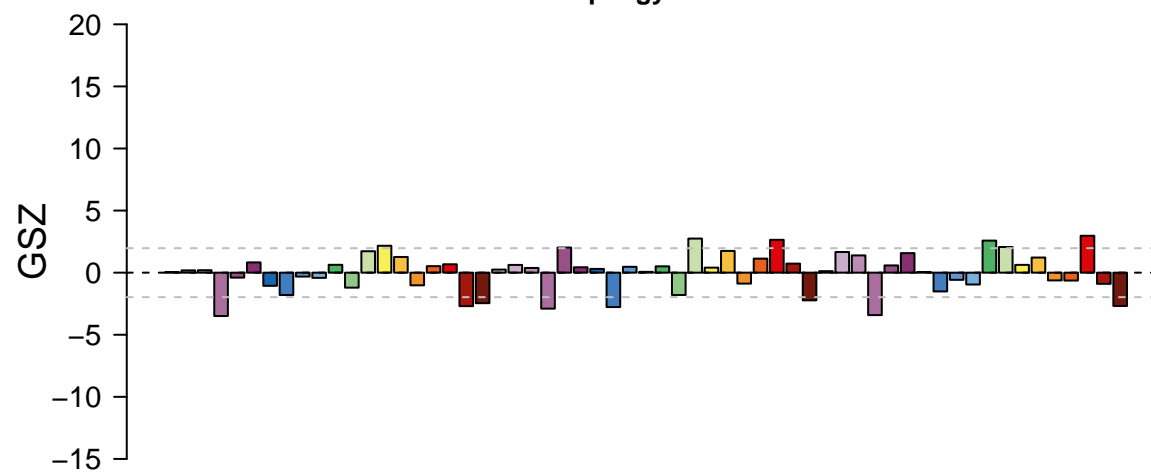
# features = 28 , max = 1

Replication and repair – Sulfur relay system



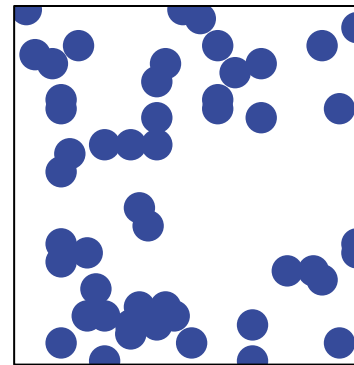
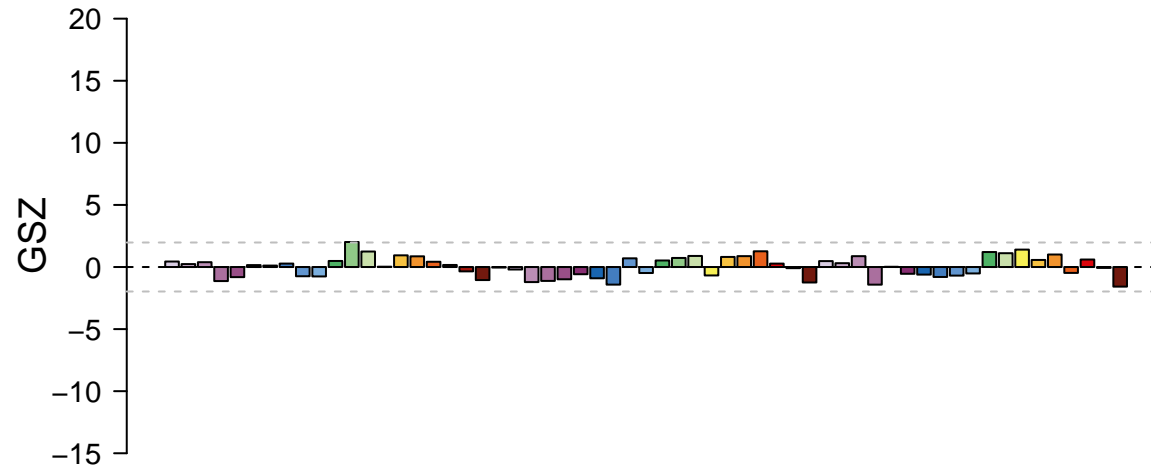
# features = 10 , max = 1

### Autophagy – other



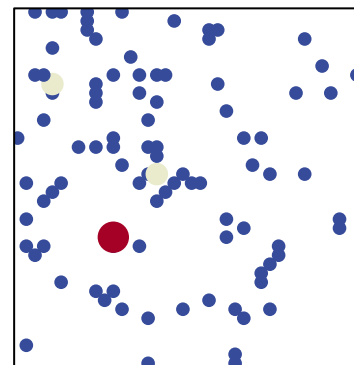
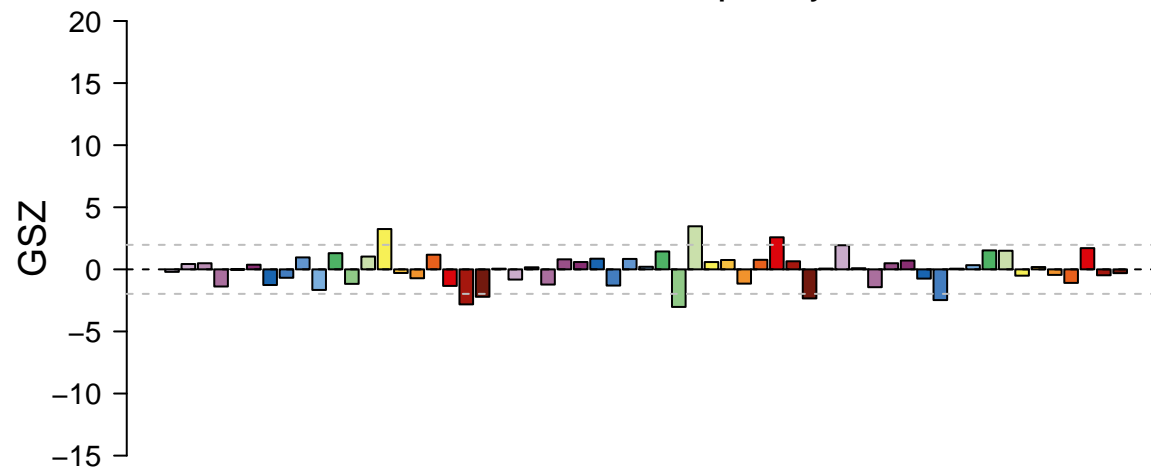
# features = 31 , max = 2

### Inositol phosphate metabolism



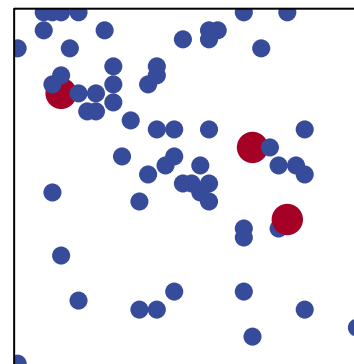
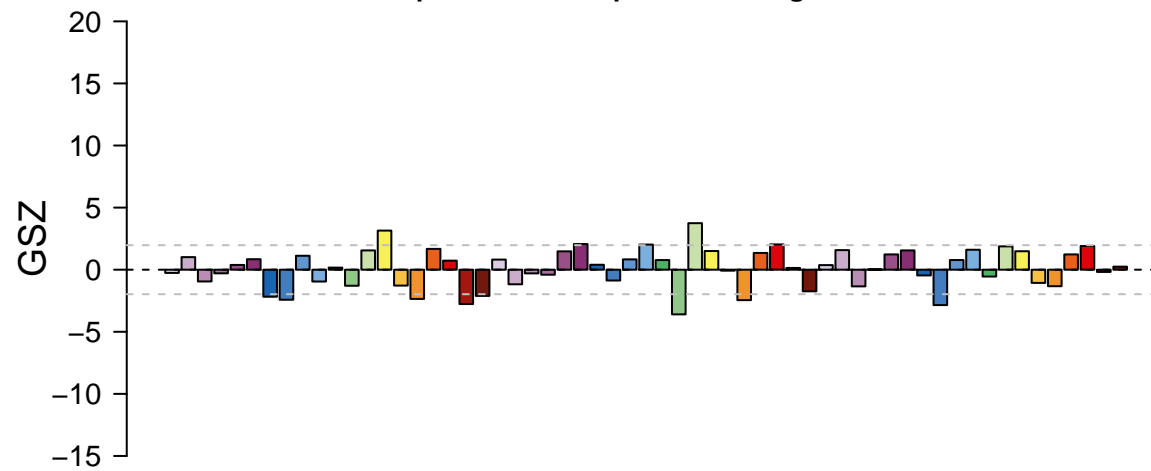
# features = 50 , max = 1

### mRNA surveillance pathway



# features = 99 , max = 3

### Replication and repair – RNA degradation



# features = 67 , max = 2