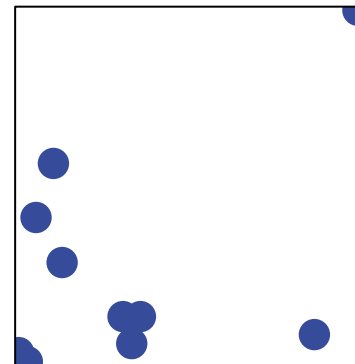
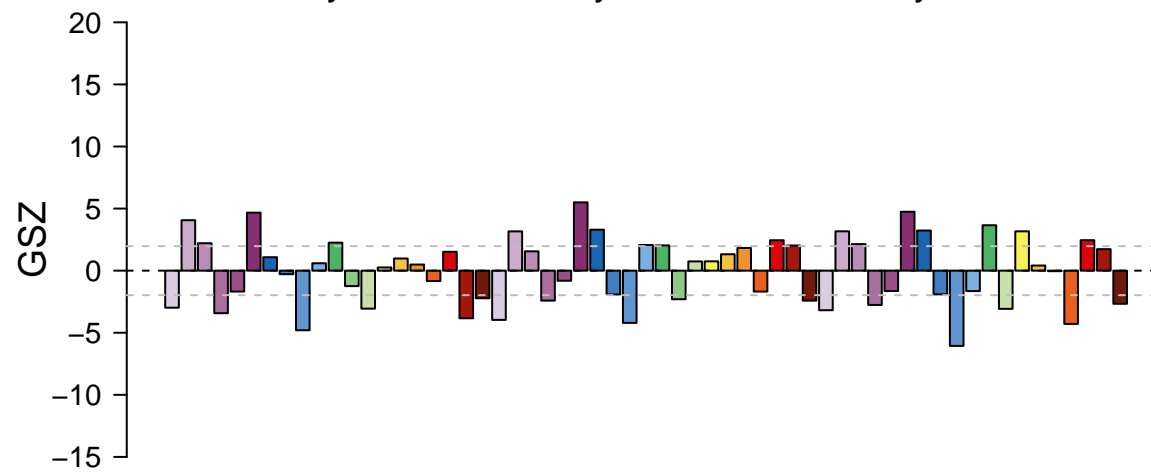
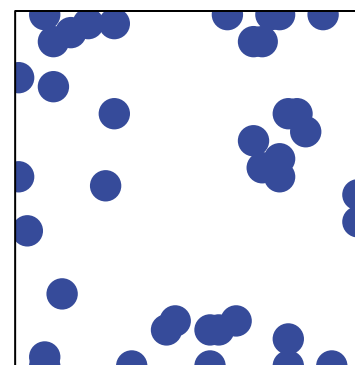
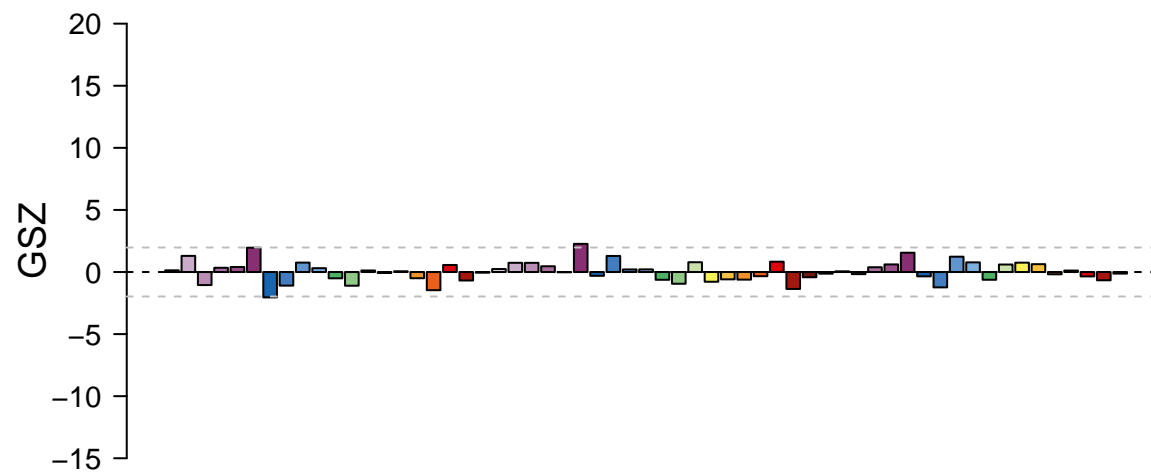


### Biosynthesis of secondary metabolism – ABA biosynthesis



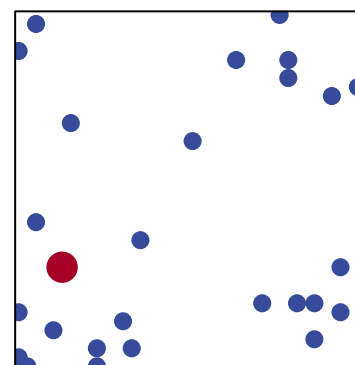
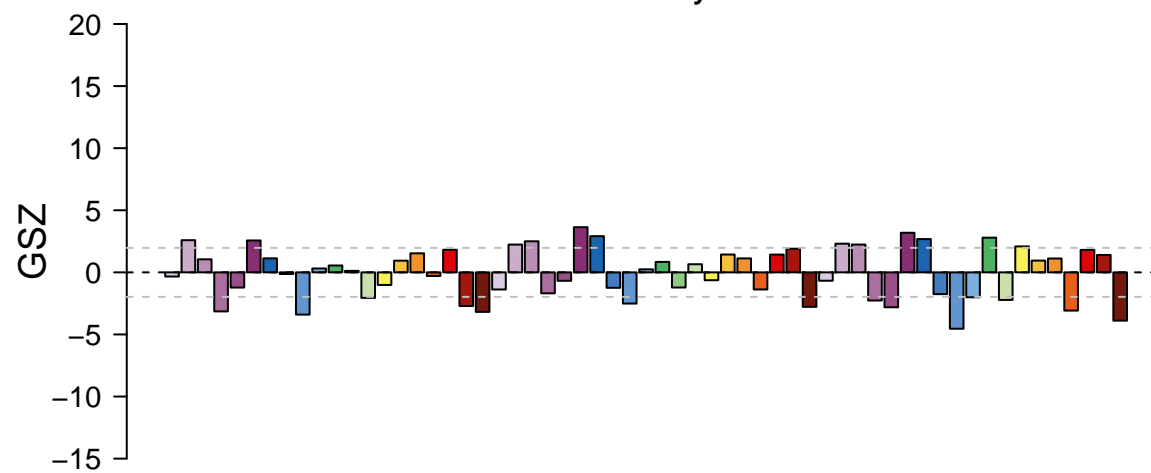
# features = 11 , max = 1

### Other amino acids metabolism – Selenoamino acid metabolism



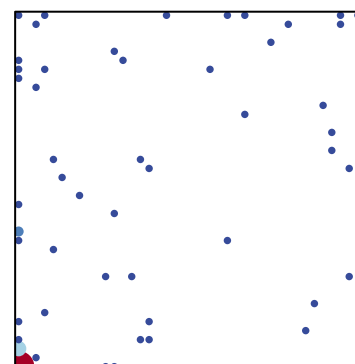
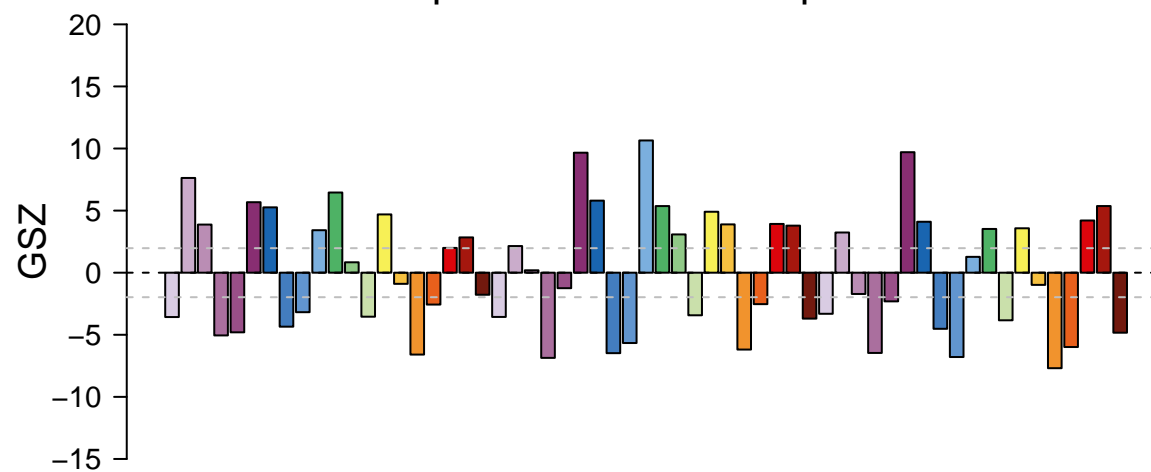
# features = 39 , max = 1

### Carotenoid biosynthesis



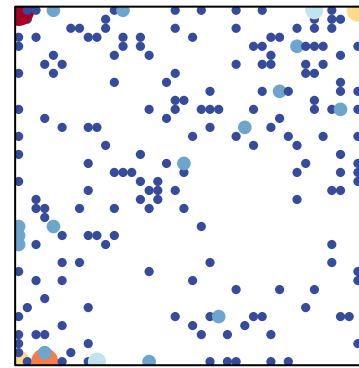
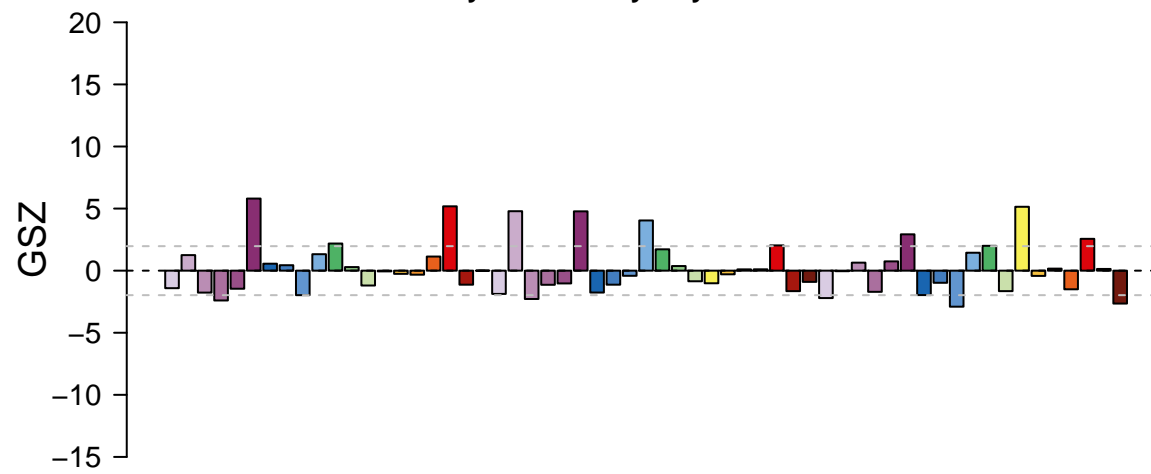
# features = 29 , max = 2

### Transcription factors – Other transcription factors



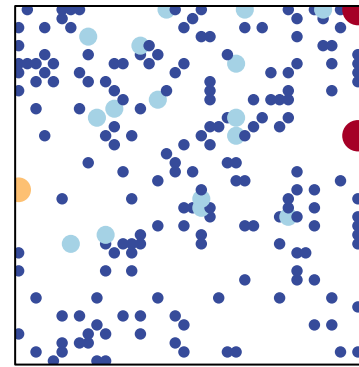
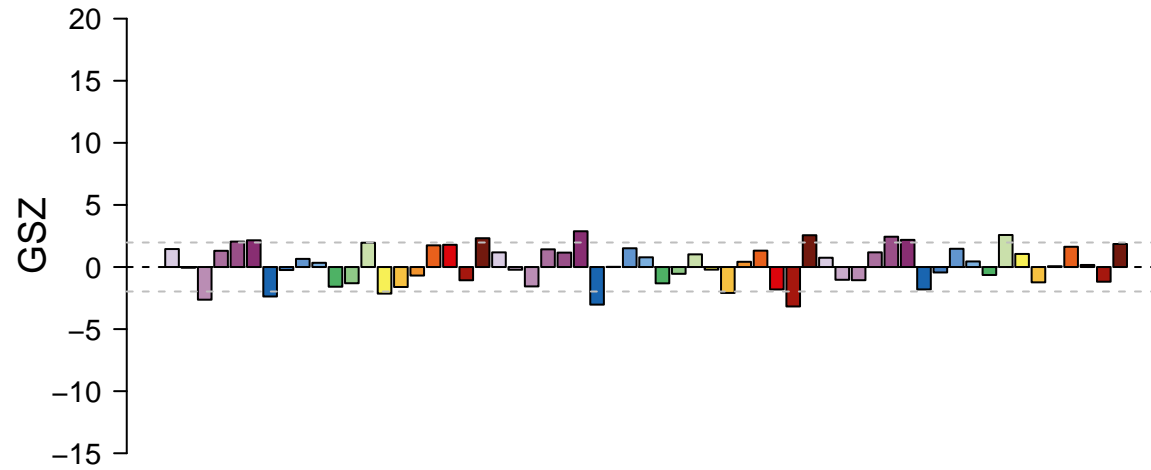
# features = 64 , max = 10

### Enzyme – 2.4 Glycosyltransferases



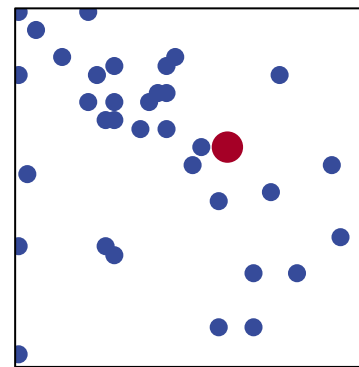
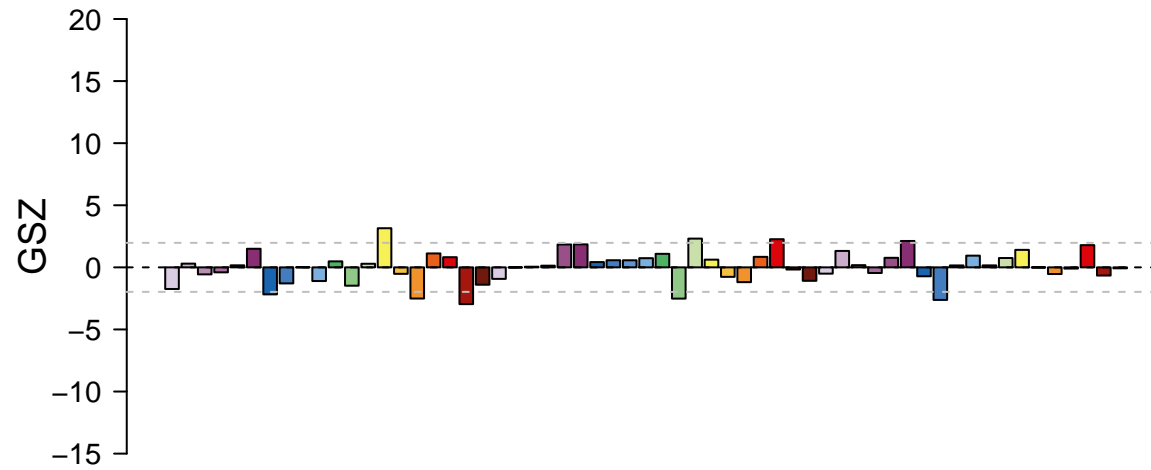
# features = 238 , max = 6

### Enzyme – 3.4 Acting on peptide bonds (peptidases)



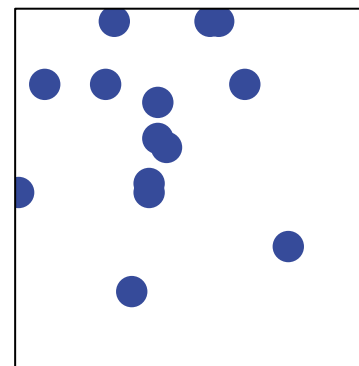
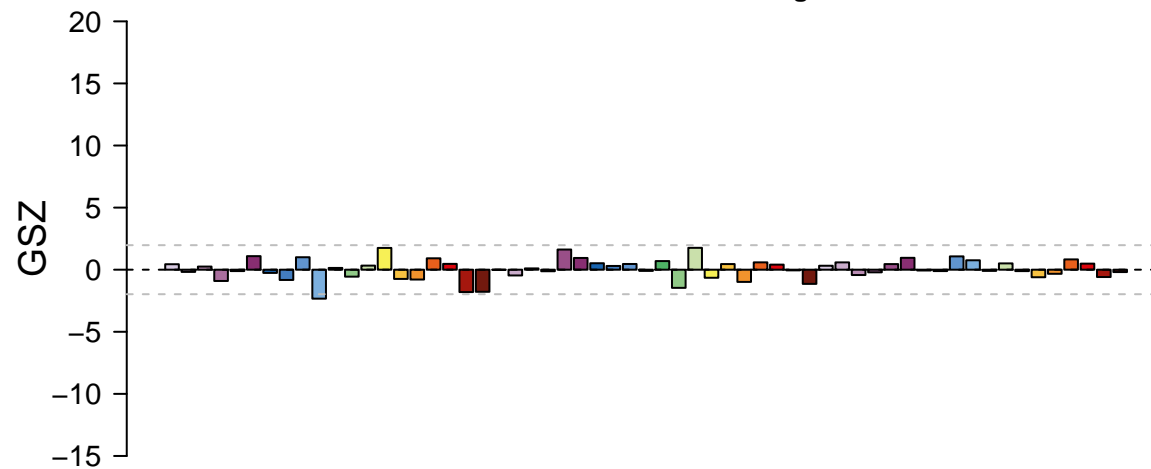
# features = 223 , max = 4

### Ribosome biogenesis – Pre-40S particles



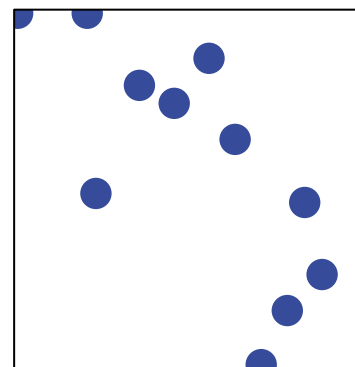
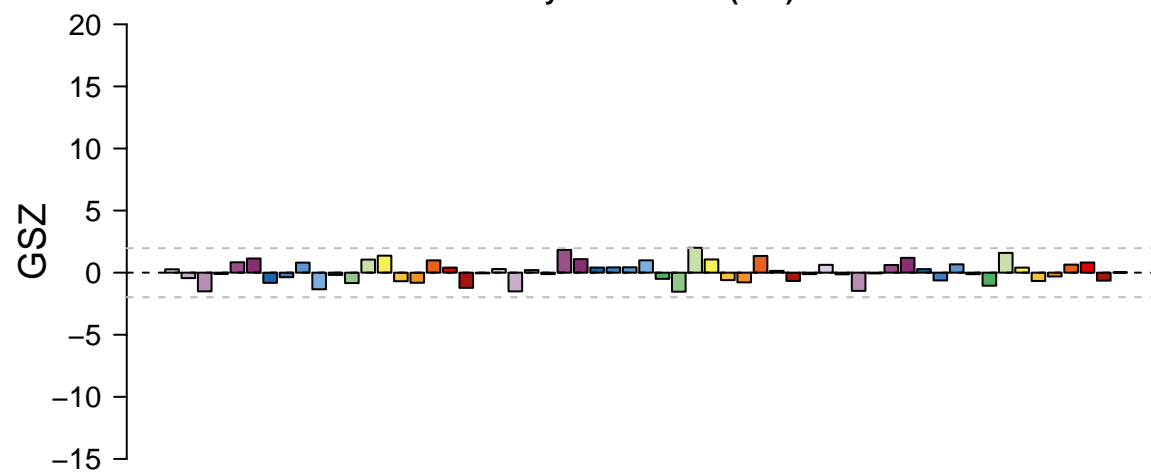
# features = 36 , max = 2

### Proteins involved in snRNP biogenesis



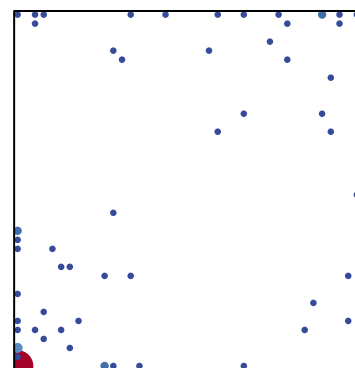
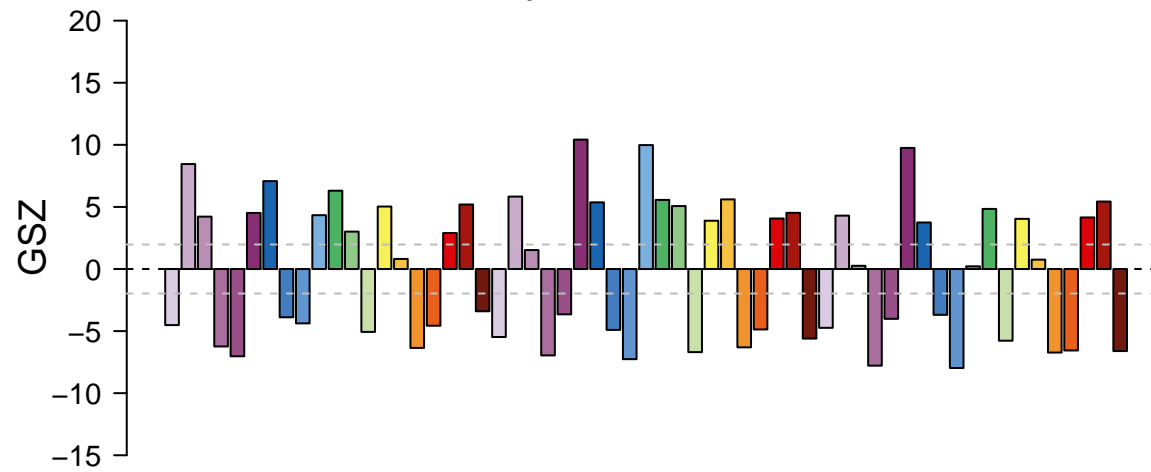
# features = 14 , max = 1

Enzyme – Class II (C/G)



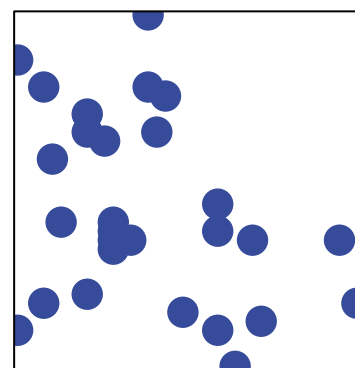
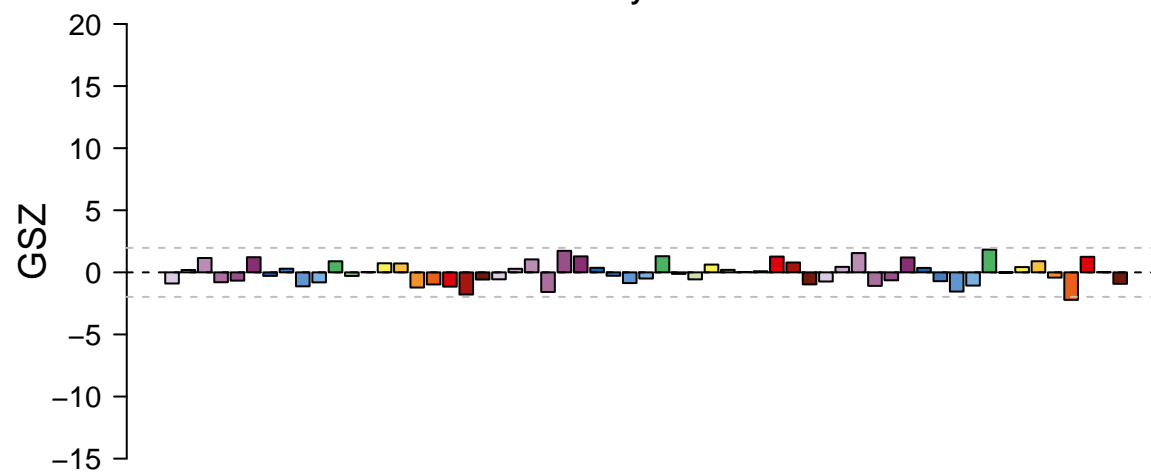
# features = 11 , max = 1

Transcription factors – AP2 EREBP



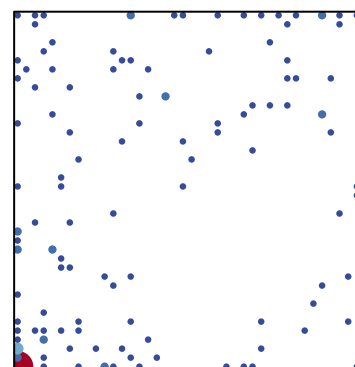
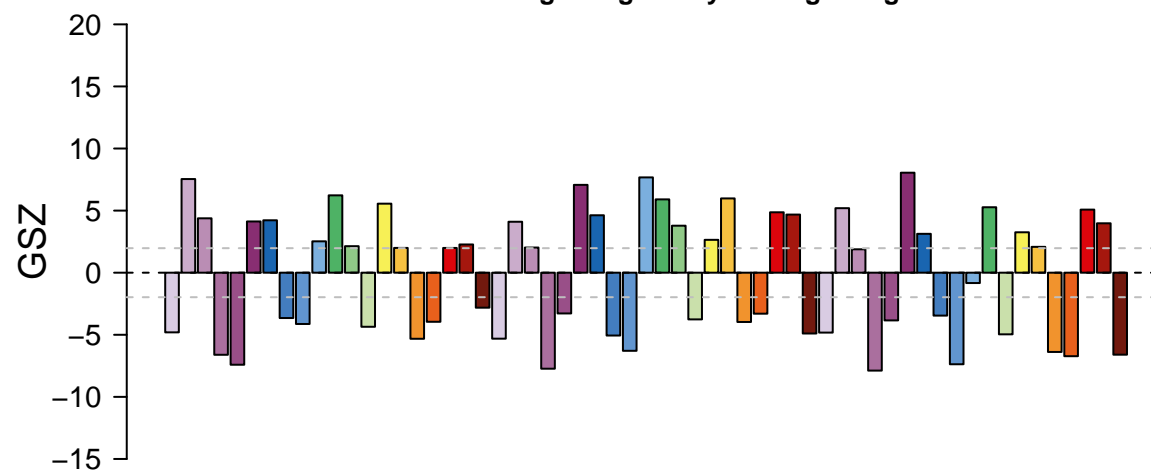
# features = 73 , max = 15

Viral life cycle – HIV-1



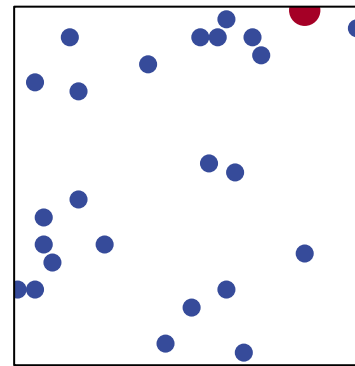
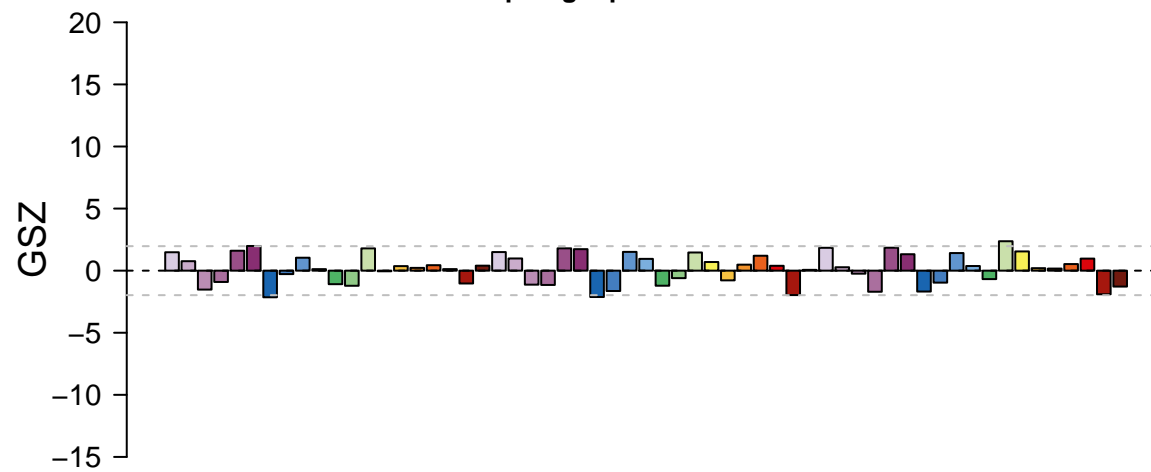
# features = 28 , max = 1

Hormone signaling – Ethylene signaling



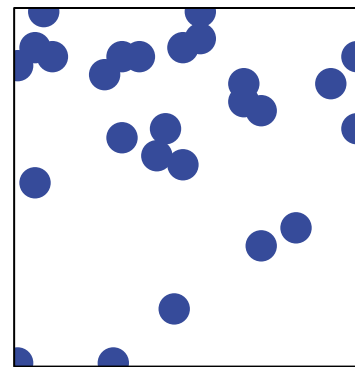
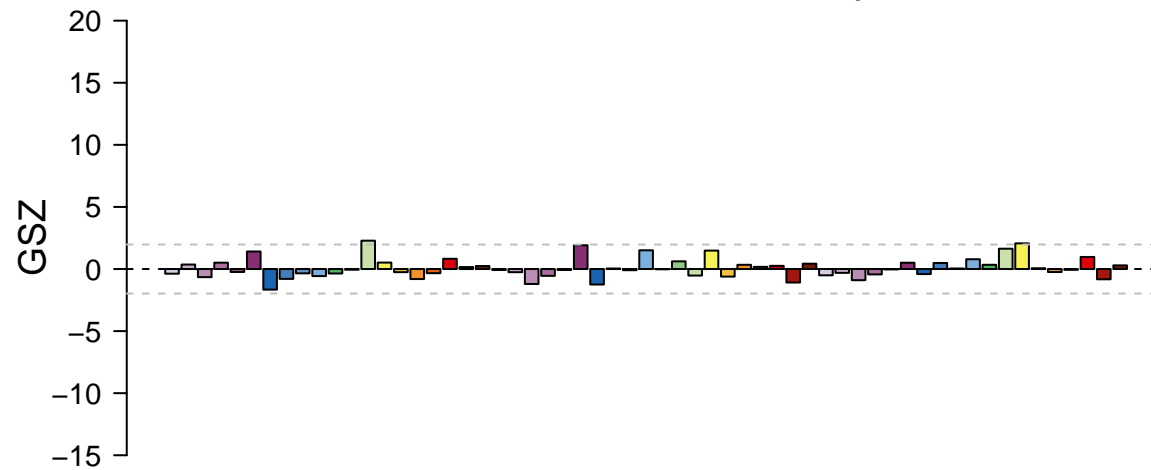
# features = 140 , max = 15

## Sphingolipid metabolism



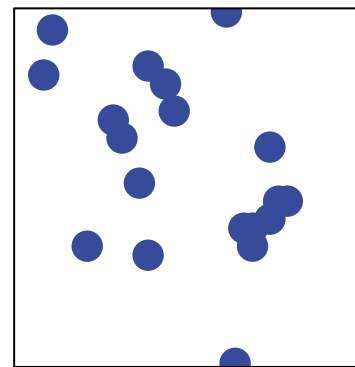
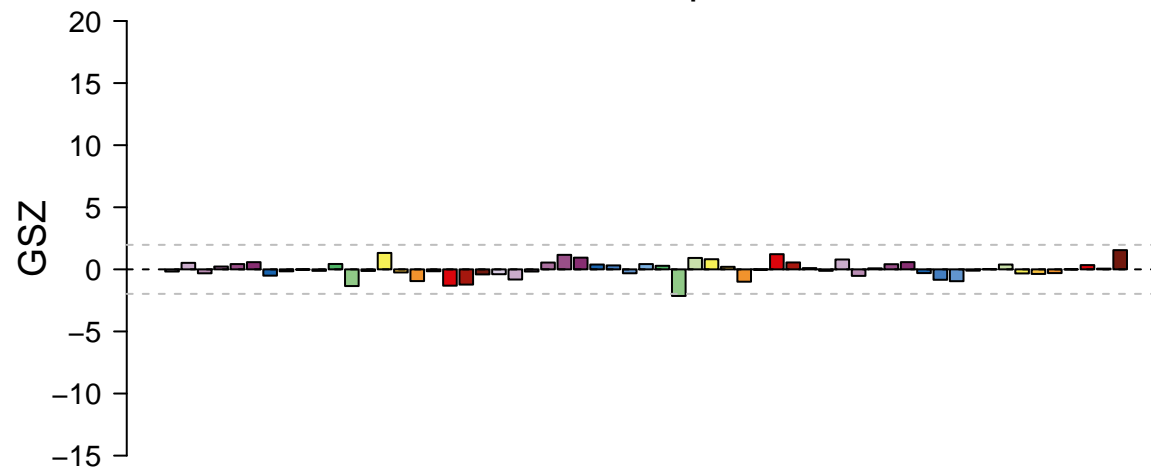
# features = 26 , max = 2

### Protein – Small GTPases and associated proteins



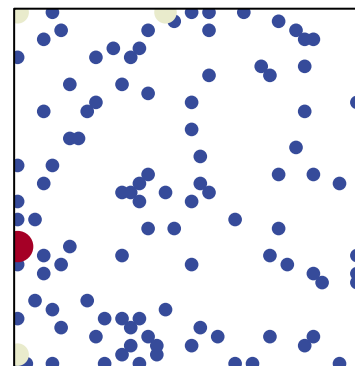
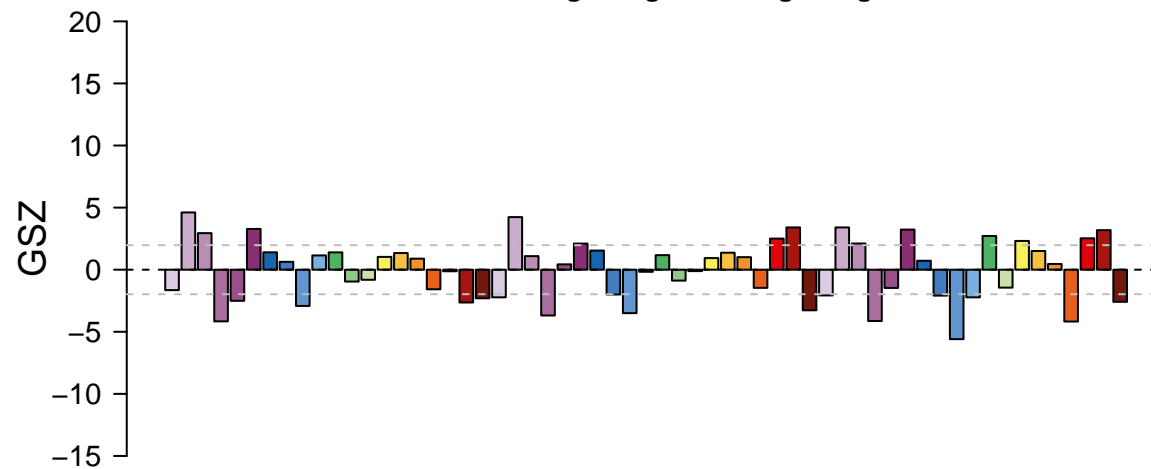
# features = 26 , max = 1

## U1 snRNP components



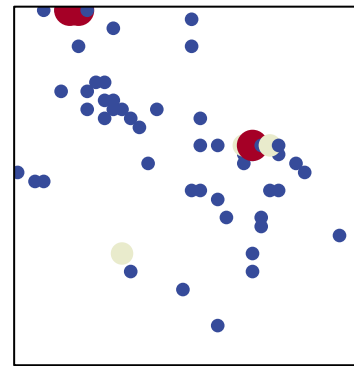
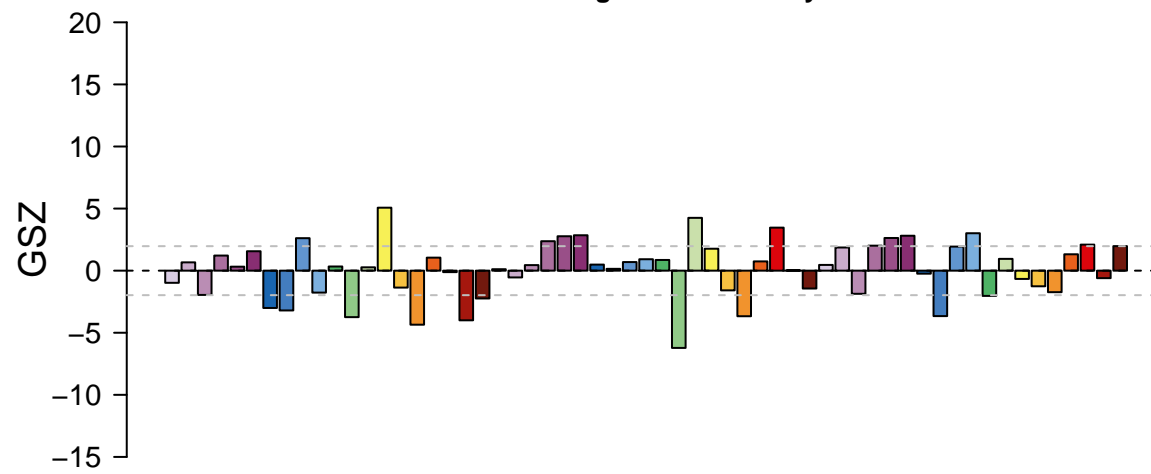
```
# features = 19 , max = 1
```

## Hormone signaling – ABA signaling



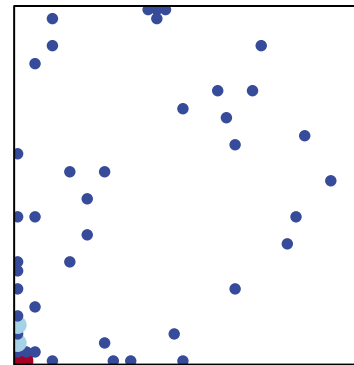
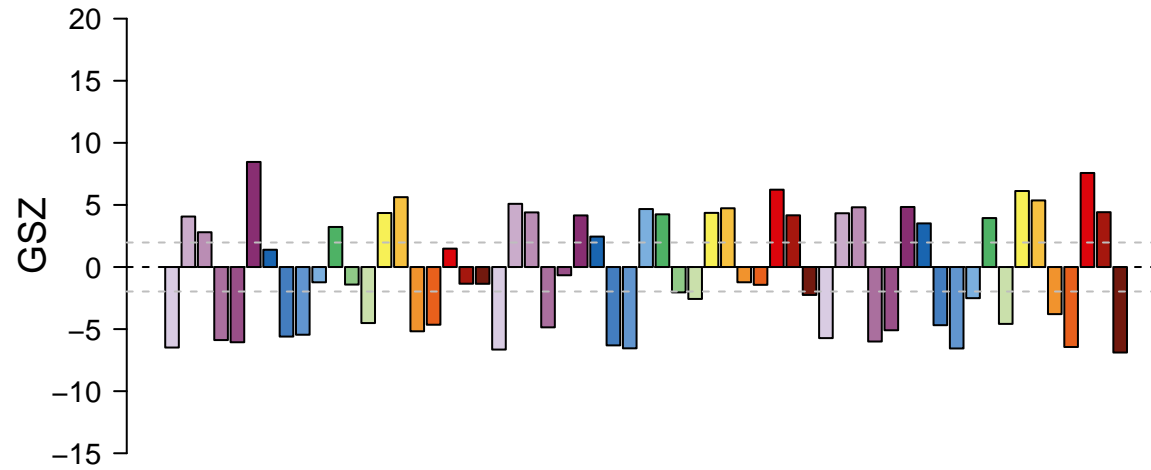
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# features = 111 , max = 3
```

### Ribosome biogenesis in eukaryotes



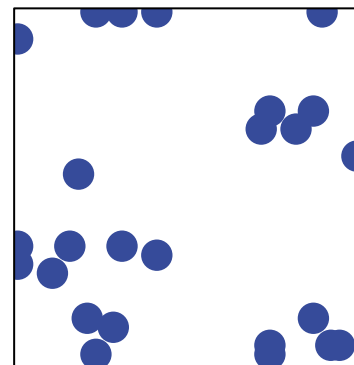
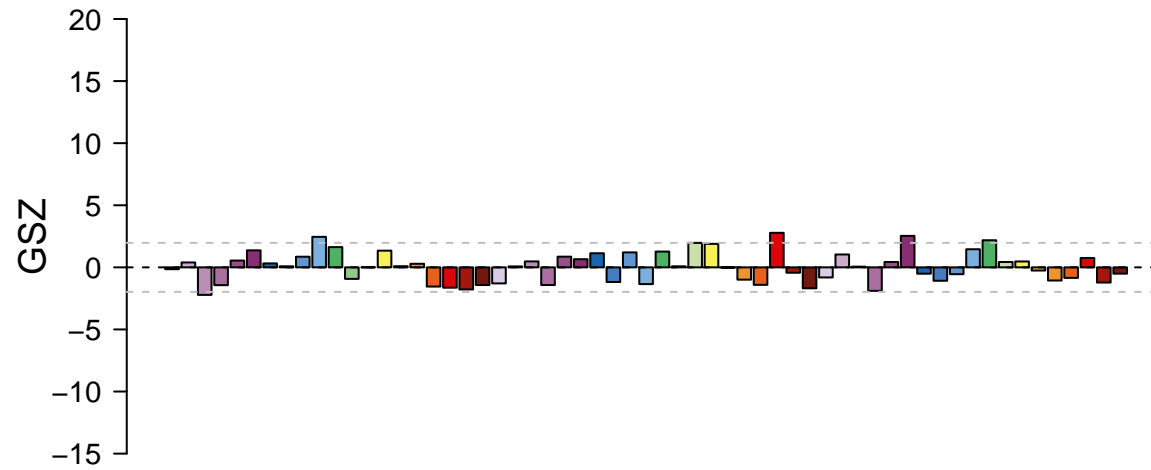
# features = 64 , max = 3

### Transcription factors – NAC



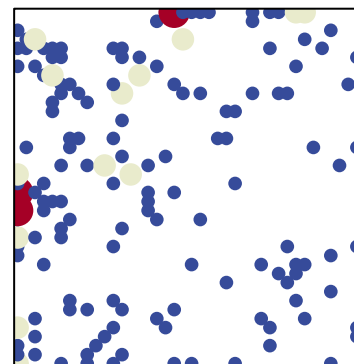
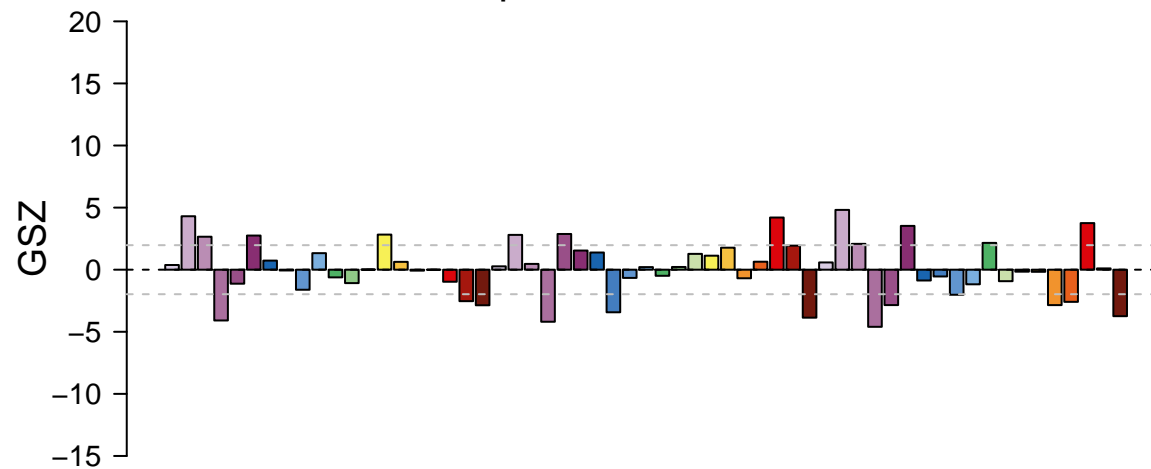
# features = 49 , max = 4

### Biosynthesis of secondary metabolism – Zeatin biosynthesis



# features = 25 , max = 1

### Transcription factors – Other zf-C3HC4



# features = 170 , max = 3