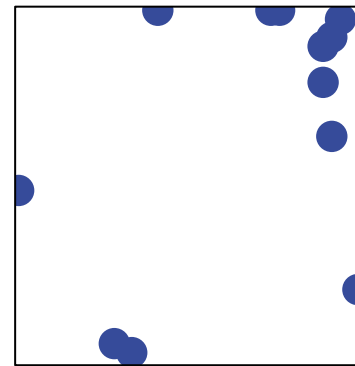
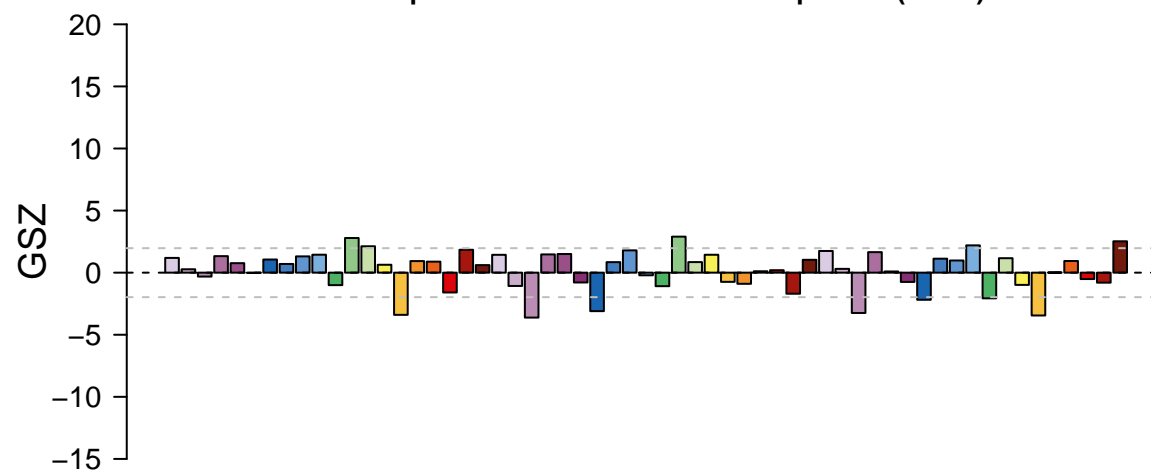
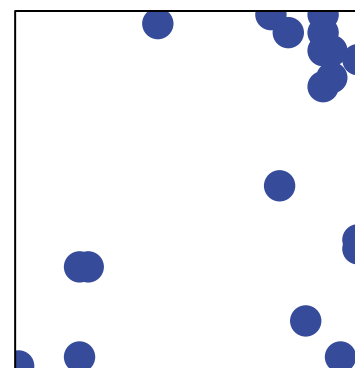
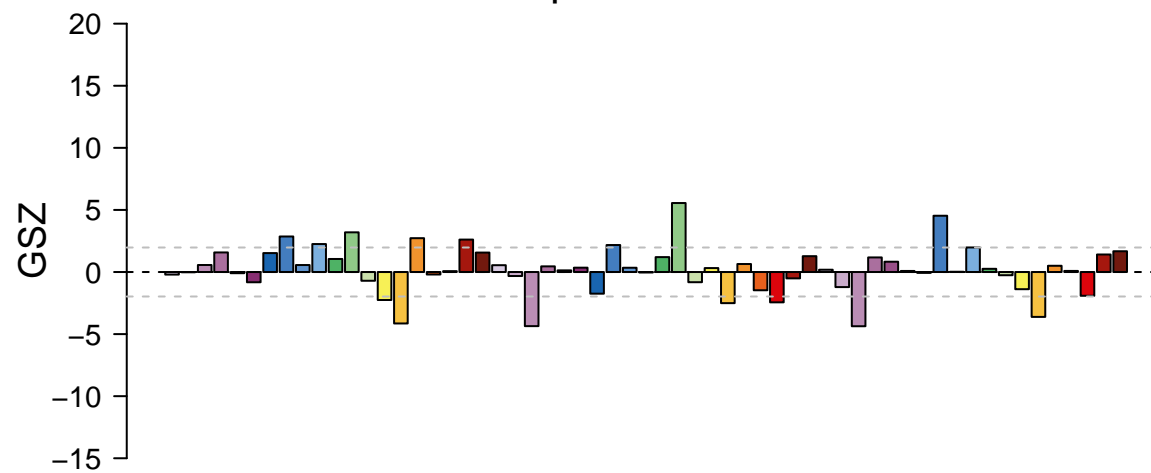


Transcription factors – Basic helix–loop–helix (bHLH)



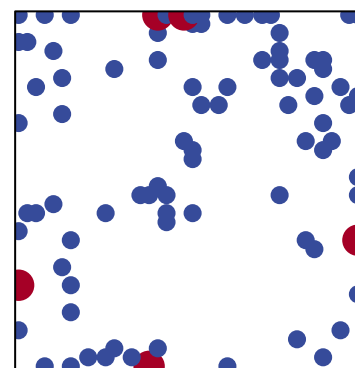
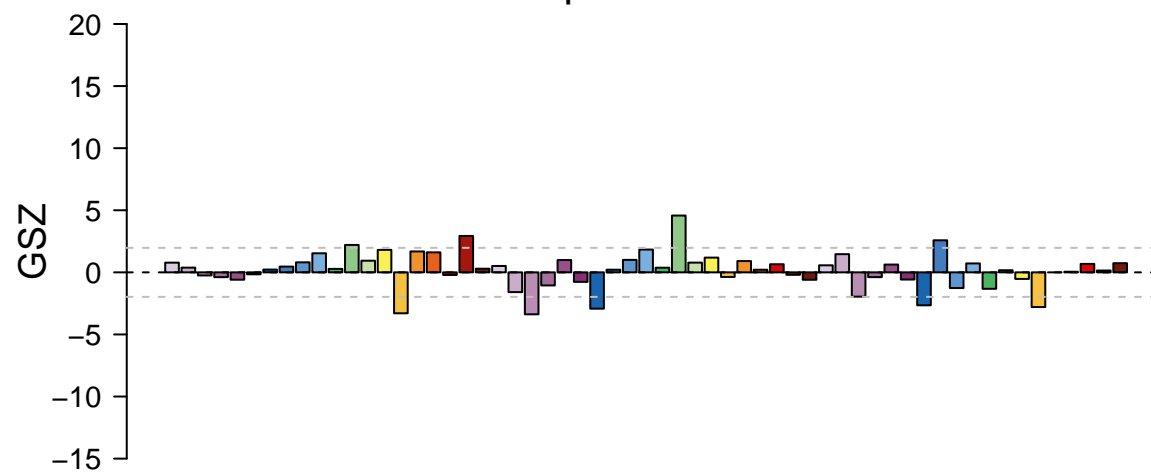
features = 12 , max = 1

Transcription factors – AUXIAA



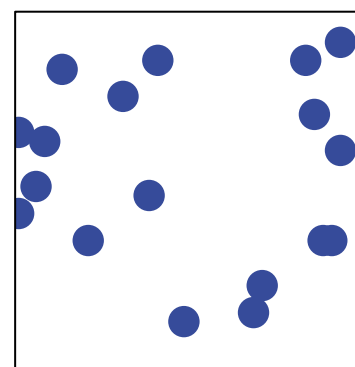
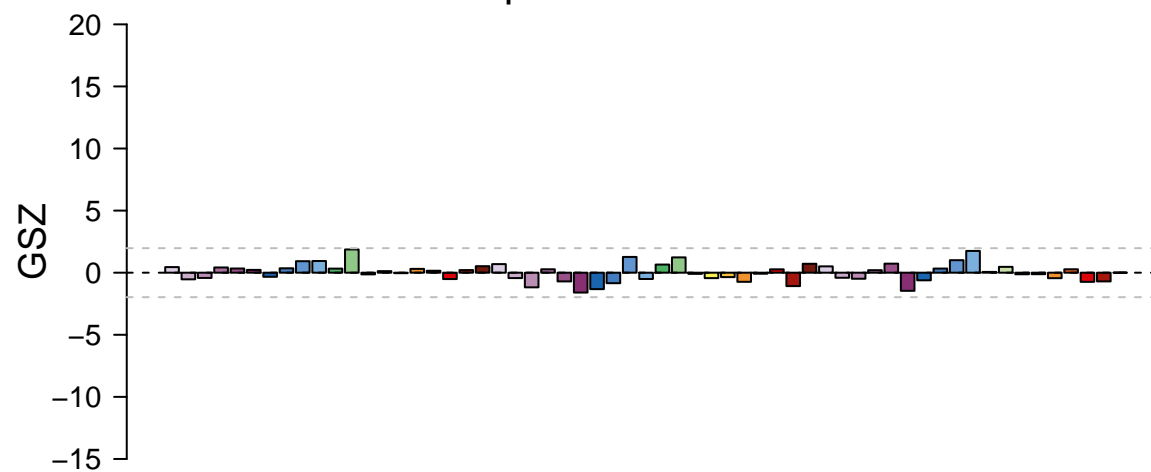
features = 19 , max = 1

Transcription factors – BHLH



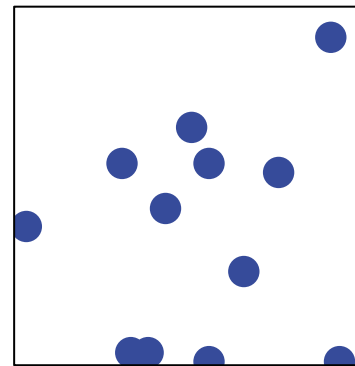
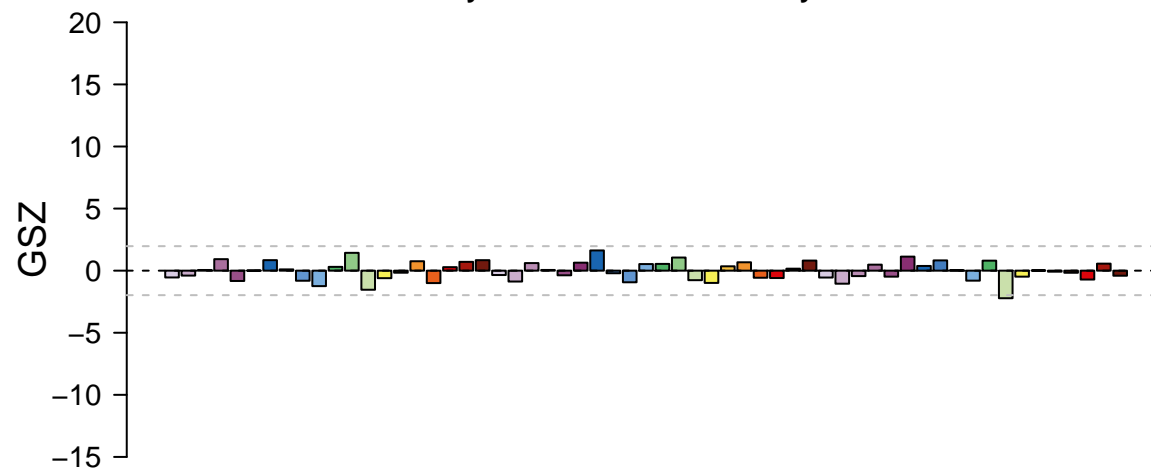
features = 93 , max = 2

Transcription factors – Other zf-DHHC



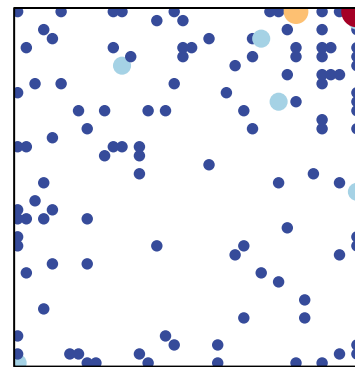
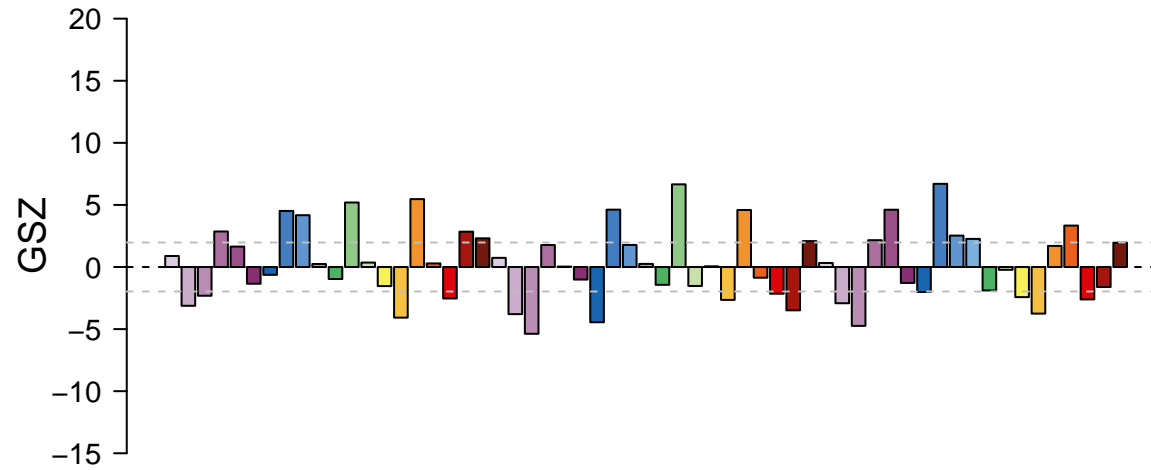
features = 18 , max = 1

Enzyme – 4.4 Carbon–sulfur lyases



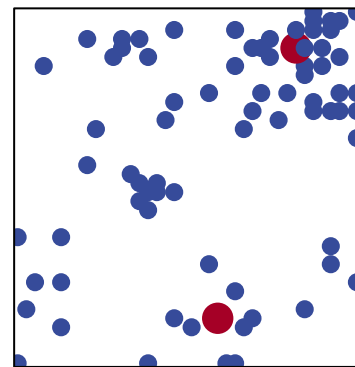
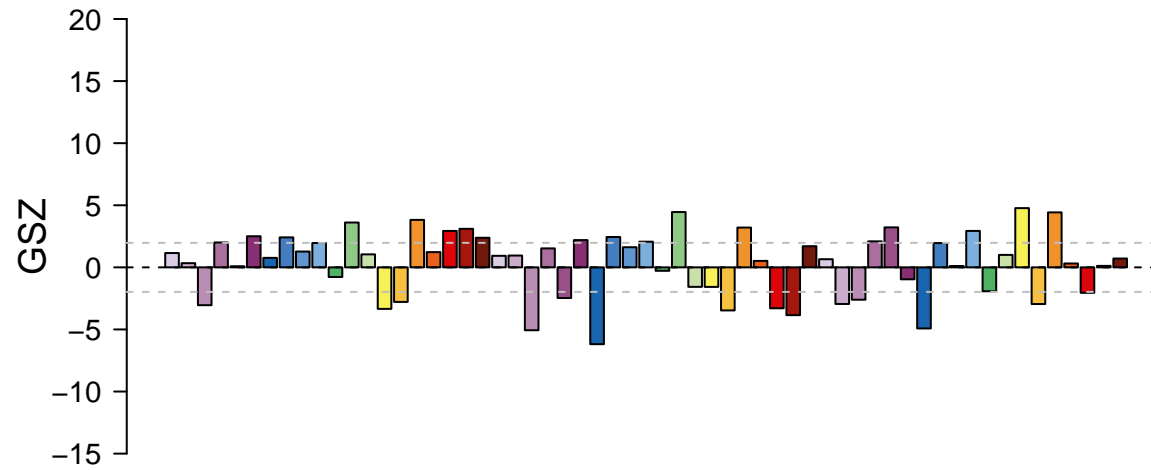
features = 12 , max = 1

Hormone signaling – Auxin signaling



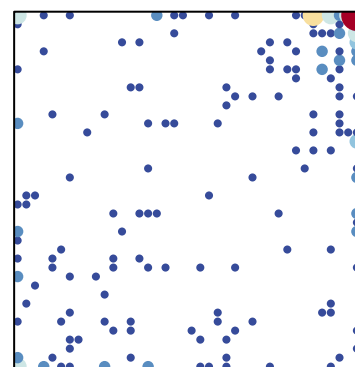
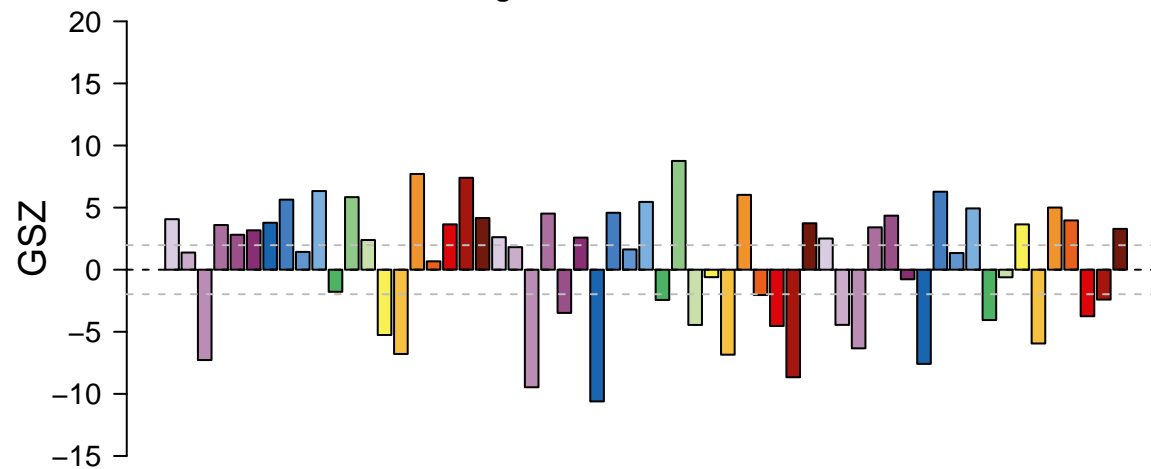
features = 134 , max = 4

Glycosyltransferase – Structural polysaccharide



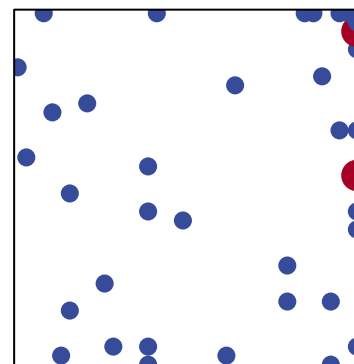
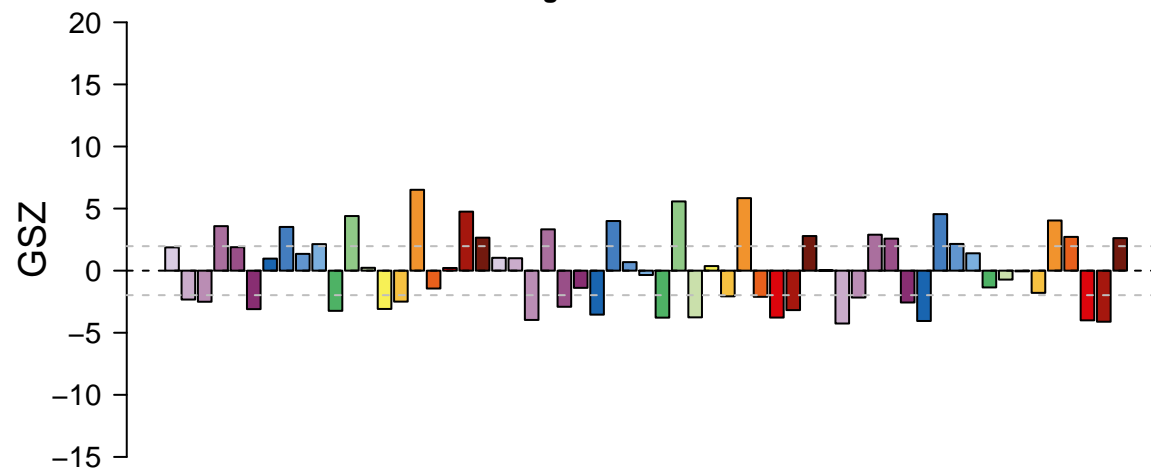
features = 78 , max = 2

Cell growth and death – Cell wall



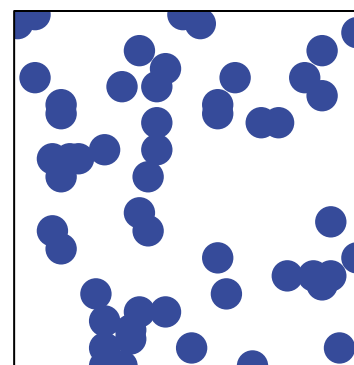
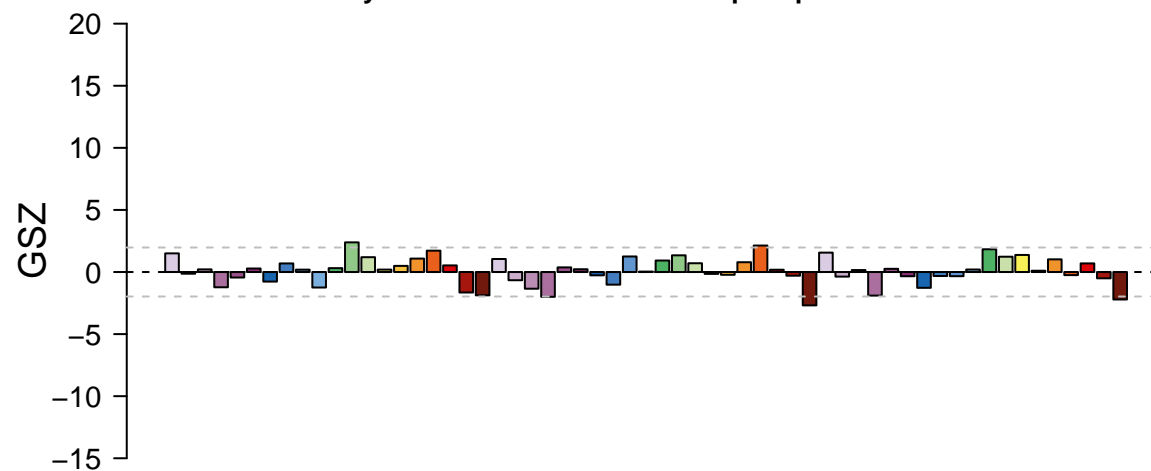
features = 206 , max = 8

Pentose and glucuronate interconversions



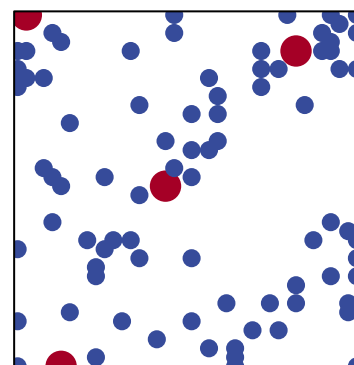
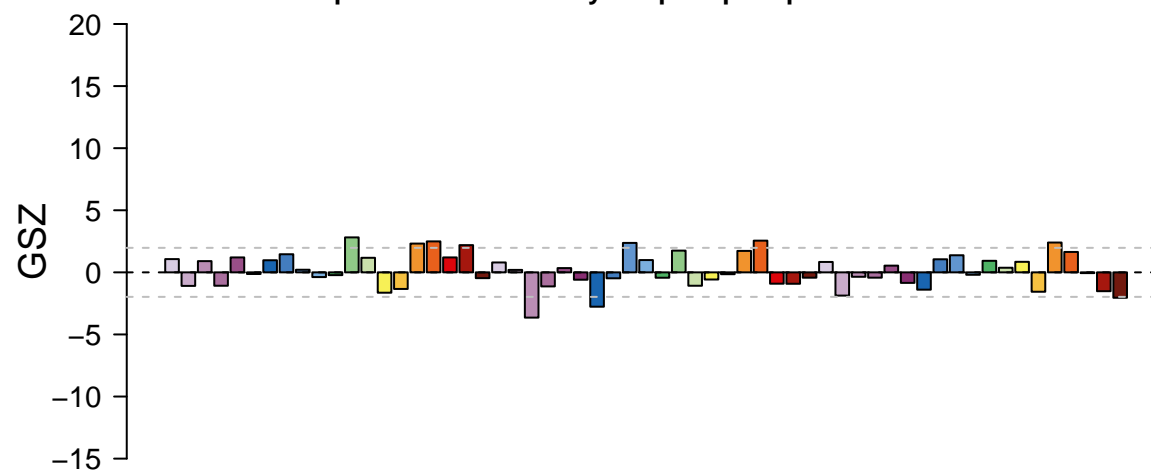
features = 39 , max = 2

Carbohydrate metabolism – Inositol phosphate metabolism



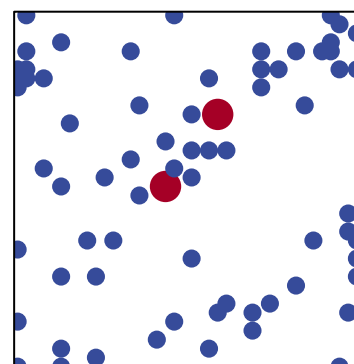
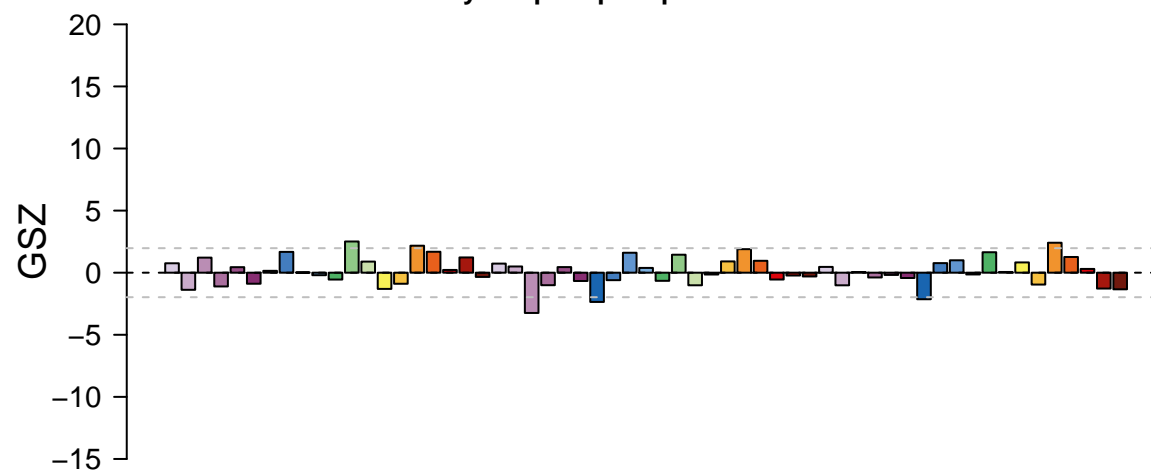
features = 54 , max = 1

Lipid metabolism – Glycerophospholipid metabolism



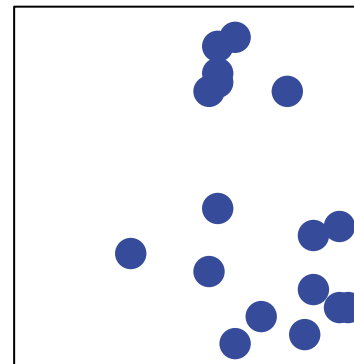
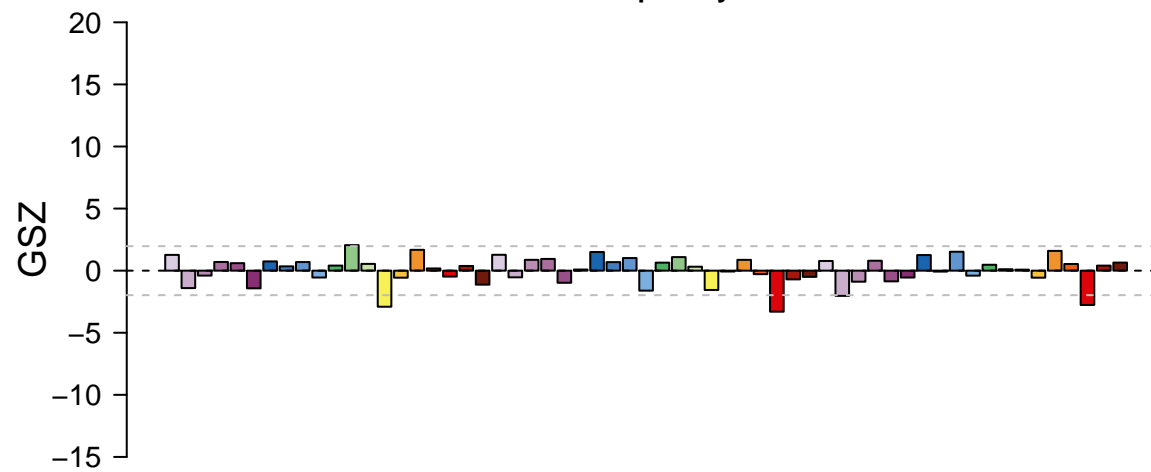
features = 90 , max = 2

Glycerophospholipid metabolism



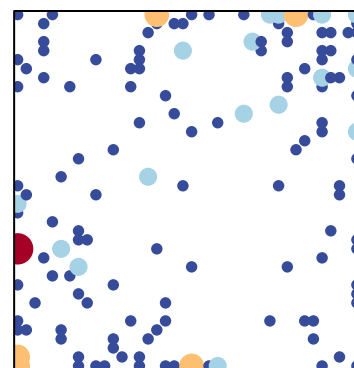
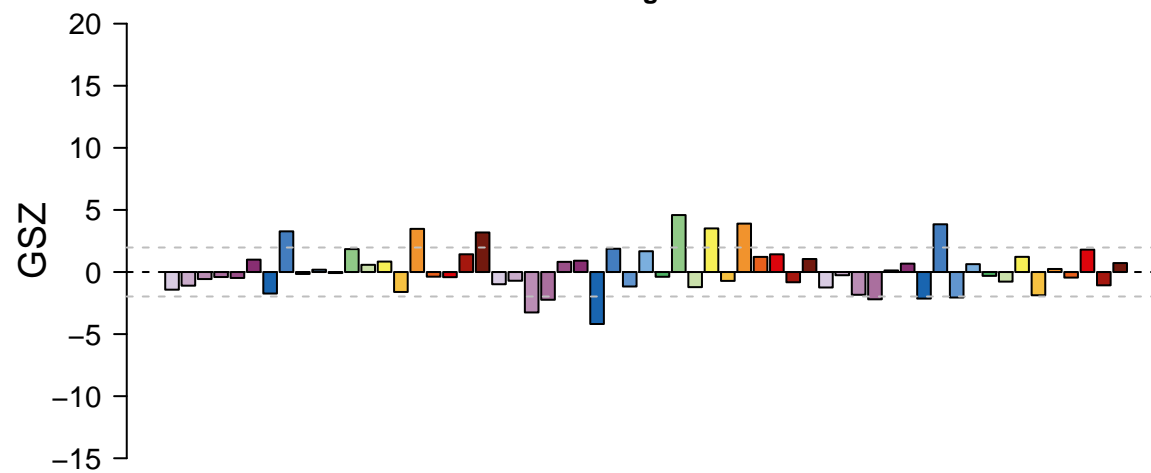
features = 72 , max = 2

One carbon pool by folate



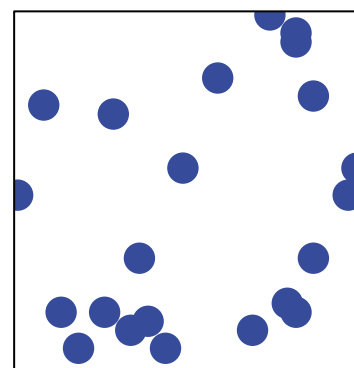
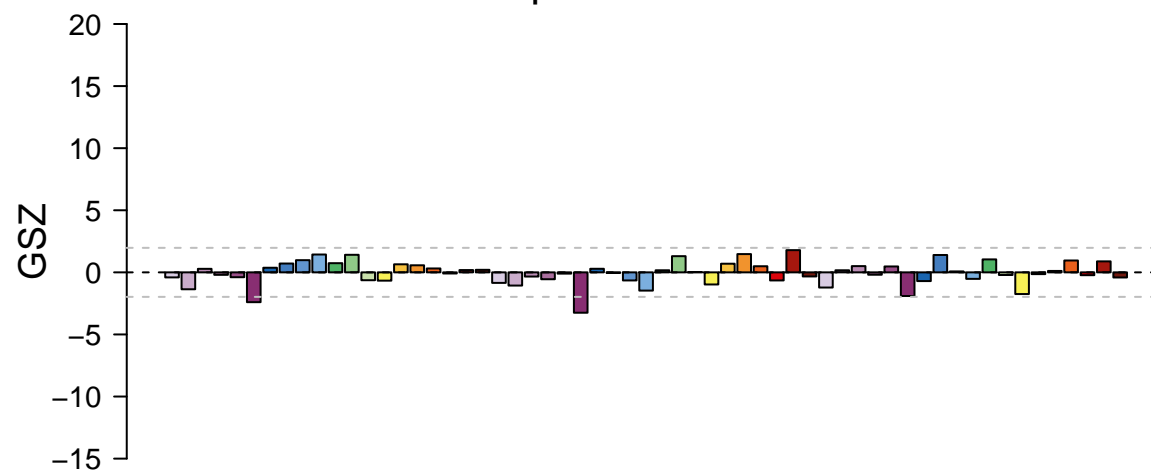
features = 17 , max = 1

Plant hormone signal transduction



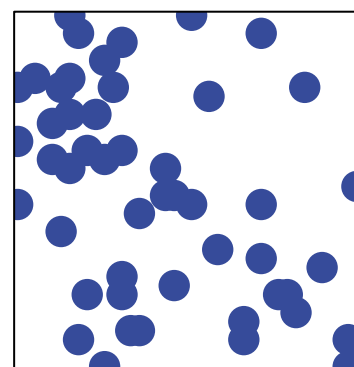
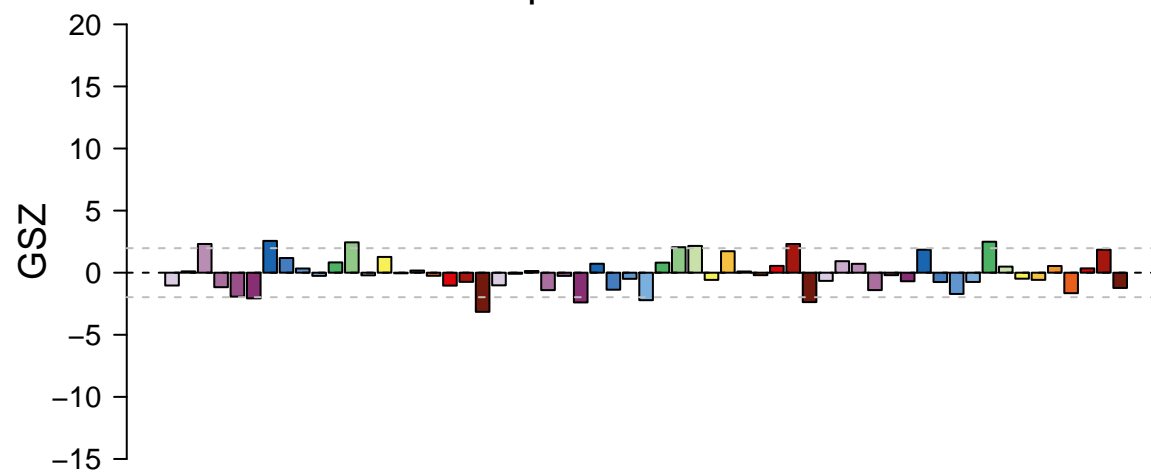
features = 168 , max = 4

Transcription factors – C2C2–DOF



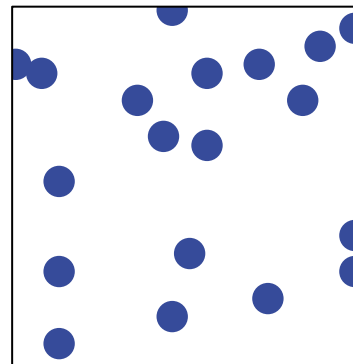
features = 22 , max = 1

Transcription factors – MYBrelated



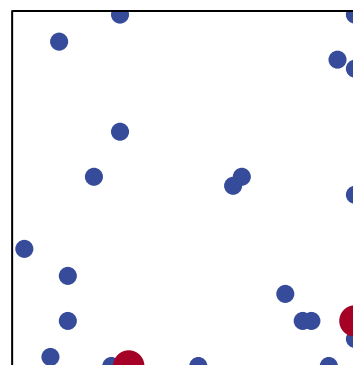
features = 50 , max = 1

Ether lipid metabolism



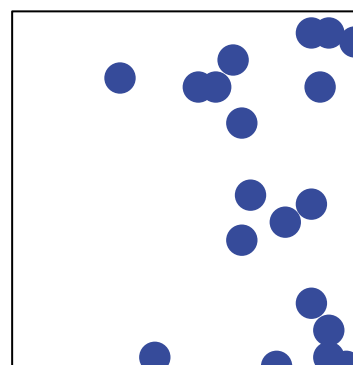
features = 19 , max = 1

Nitrogen metabolism



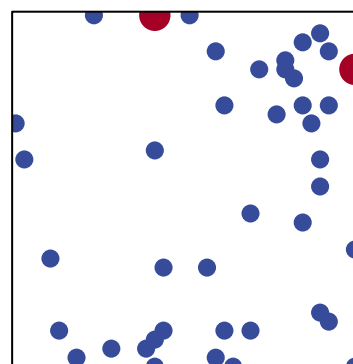
features = 25 , max = 2

Cofactors and vitamin metabolism – Ubiquinone biosynthesis



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

Fructose and mannose metabolism



```
# features = 44 , max = 2
```