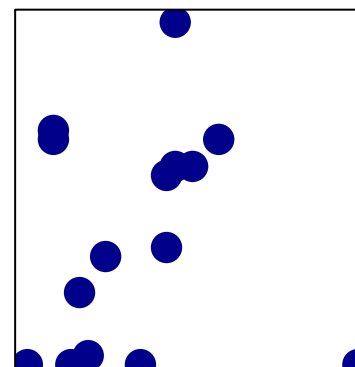
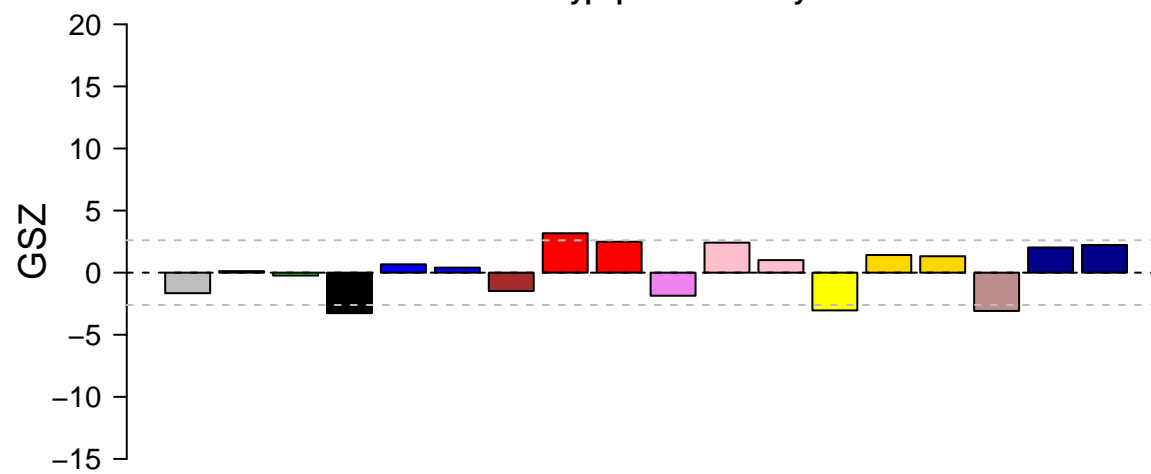
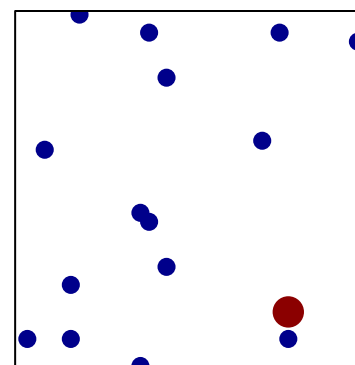
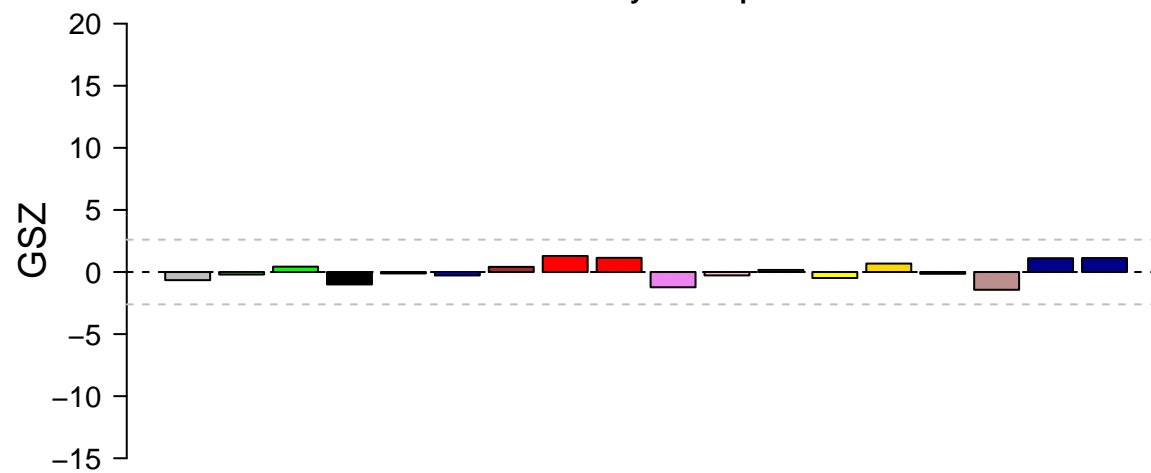


**carboxypeptidase activity**



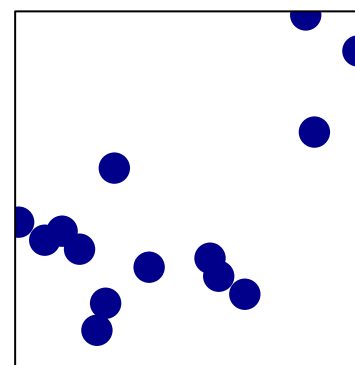
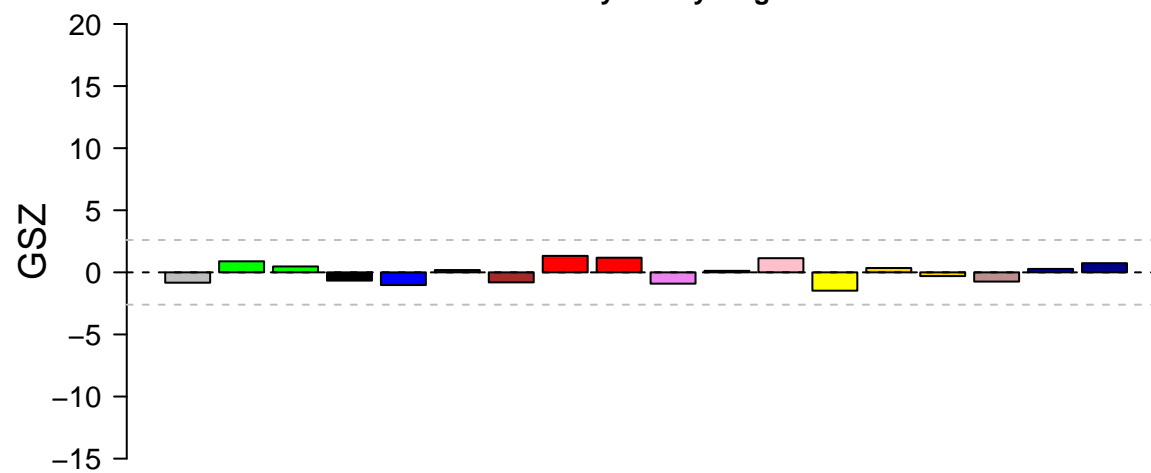
# features = 15 , max = 1

**ceramide biosynthetic process**



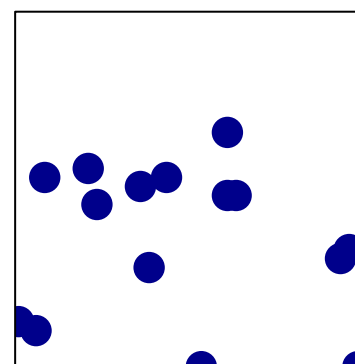
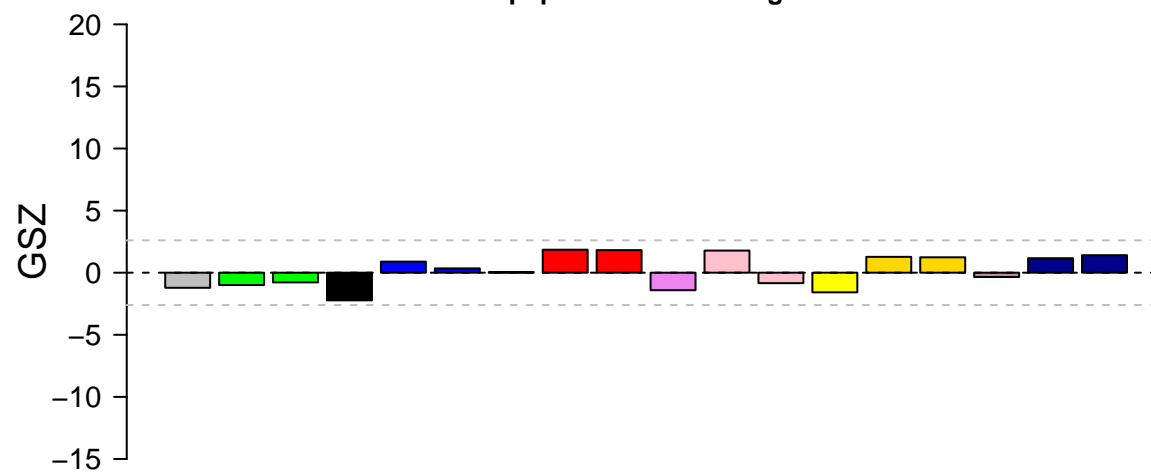
# features = 17 , max = 2

**endocytic recycling**



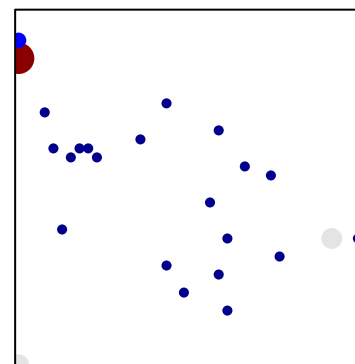
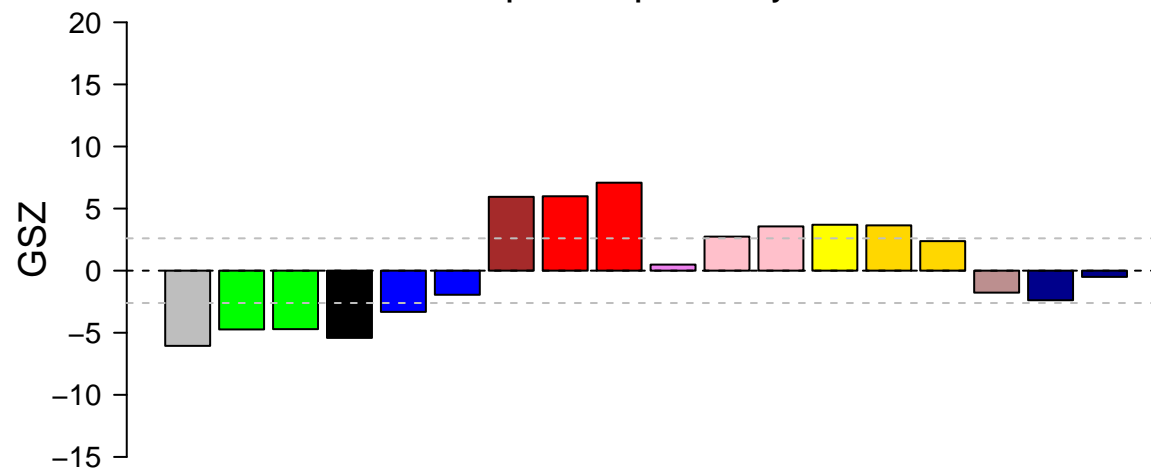
# features = 14 , max = 1

**peptide cross-linking**



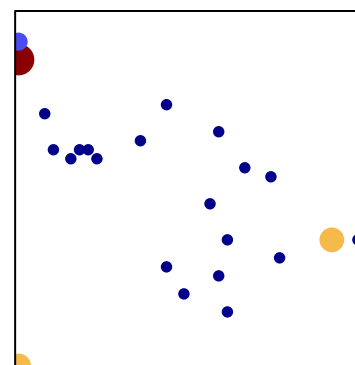
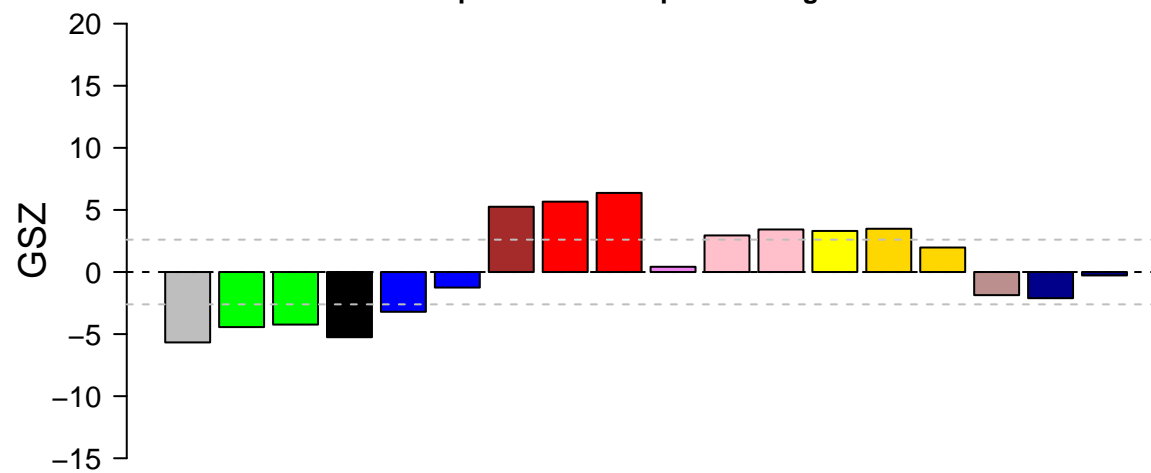
# features = 15 , max = 1

photoreceptor activity



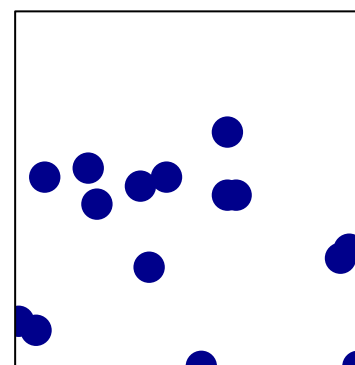
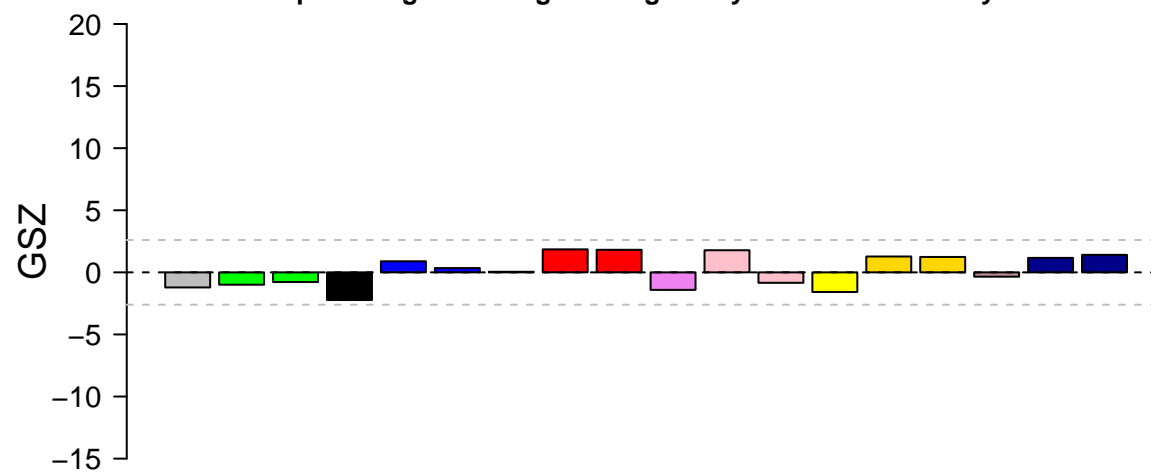
# features = 33 , max = 5

protein-chromophore linkage



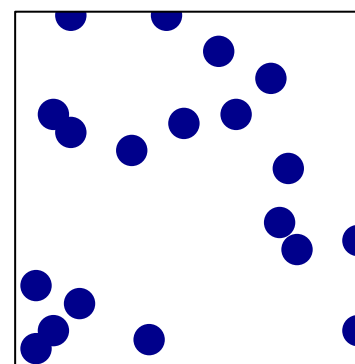
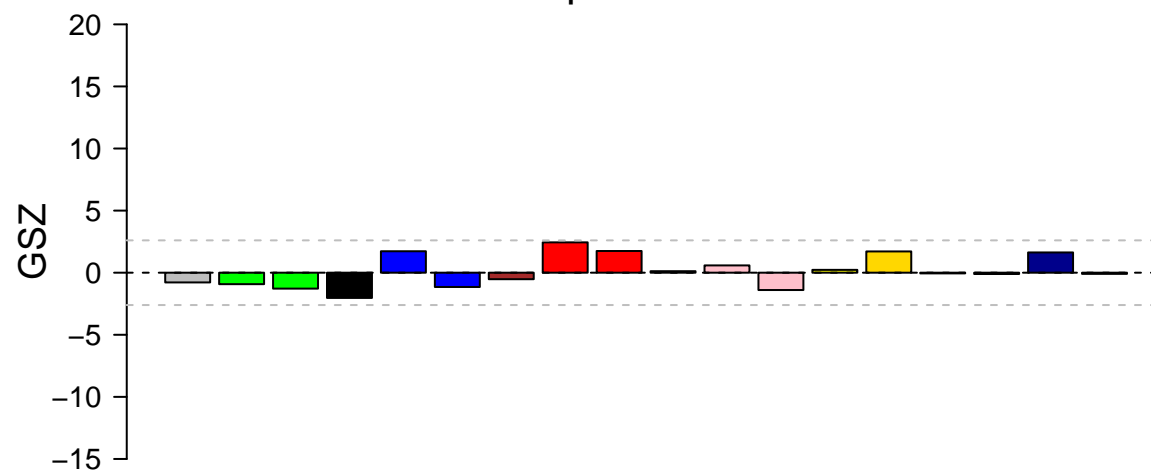
# features = 31 , max = 4

protein-glutamine gamma-glutamyltransferase activity



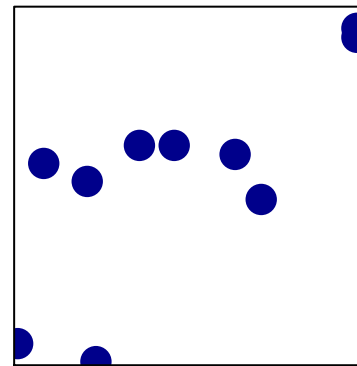
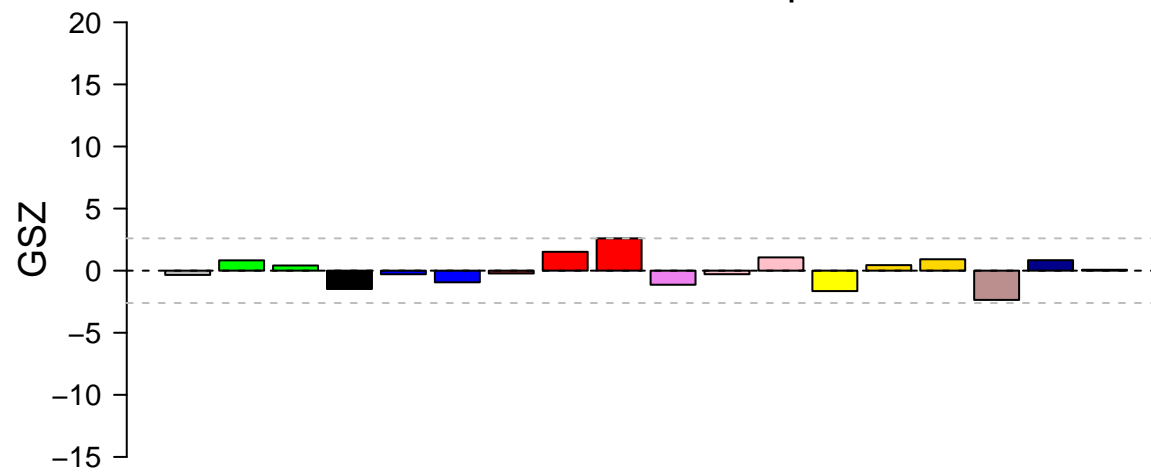
# features = 15 , max = 1

response to stress



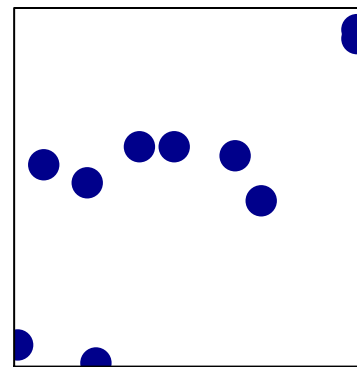
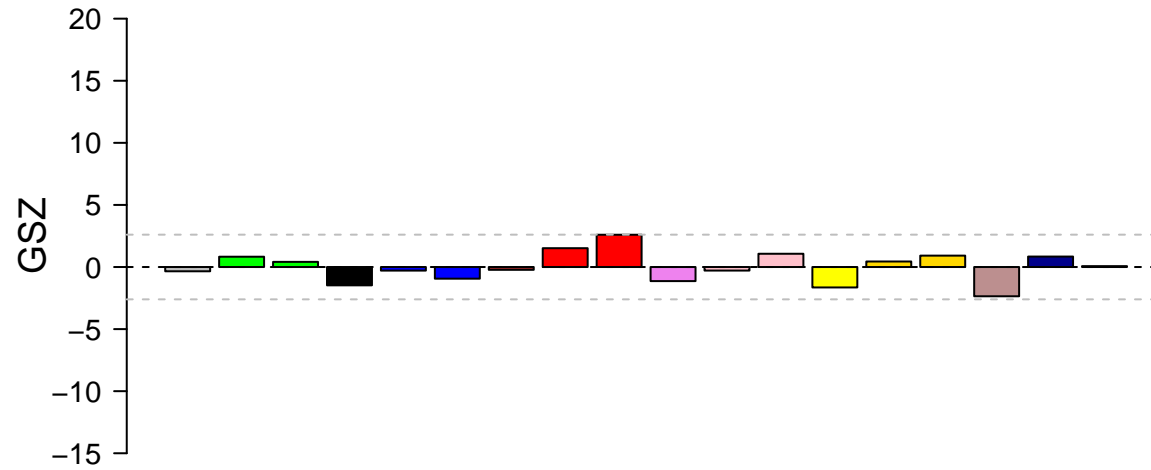
# features = 19 , max = 1

urea transmembrane transport



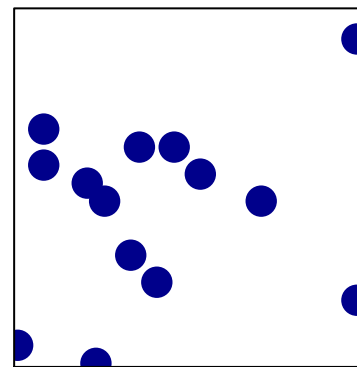
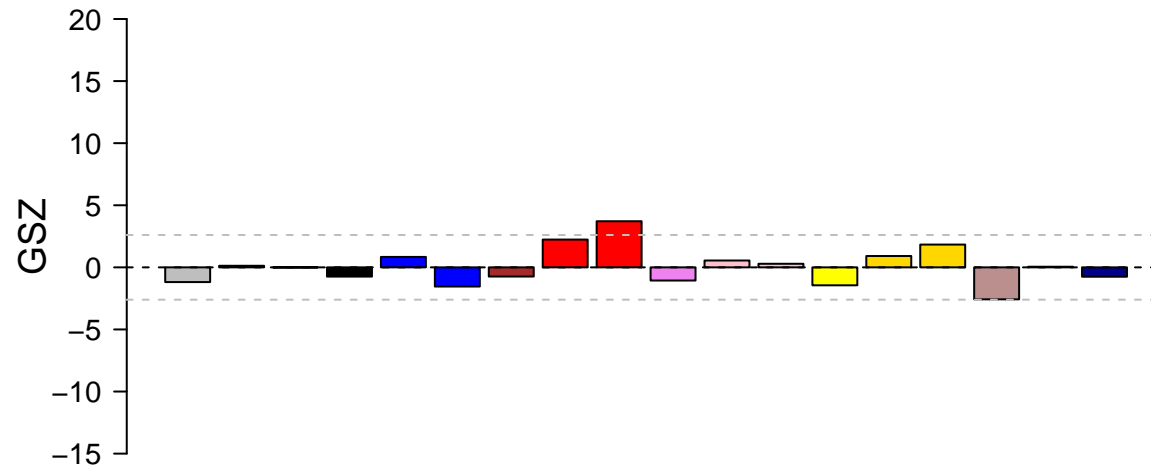
# features = 10 , max = 1

urea transmembrane transporter activity



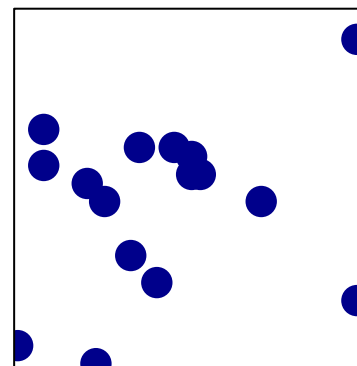
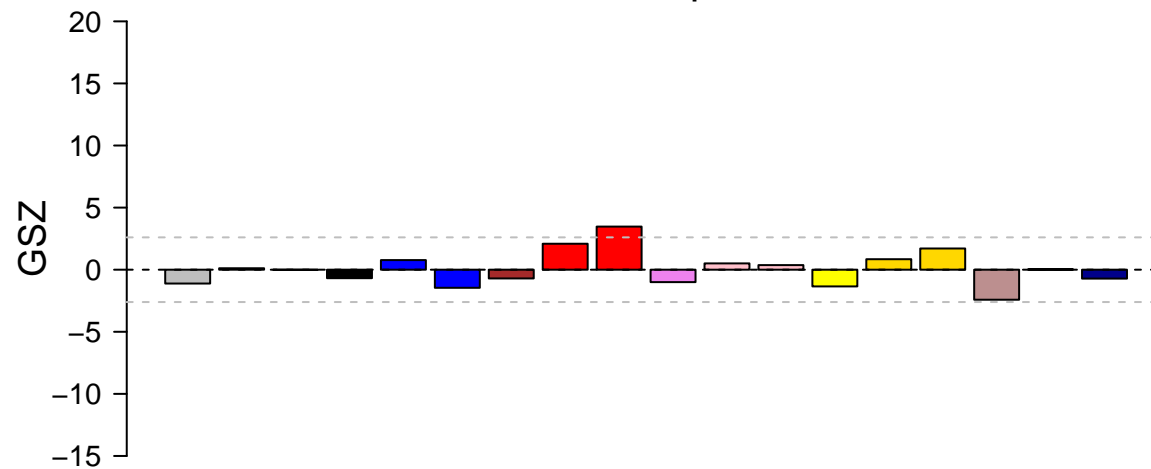
# features = 10 , max = 1

water transmembrane transporter activity



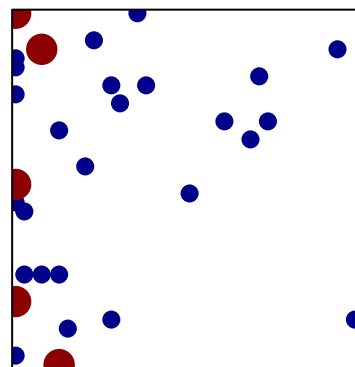
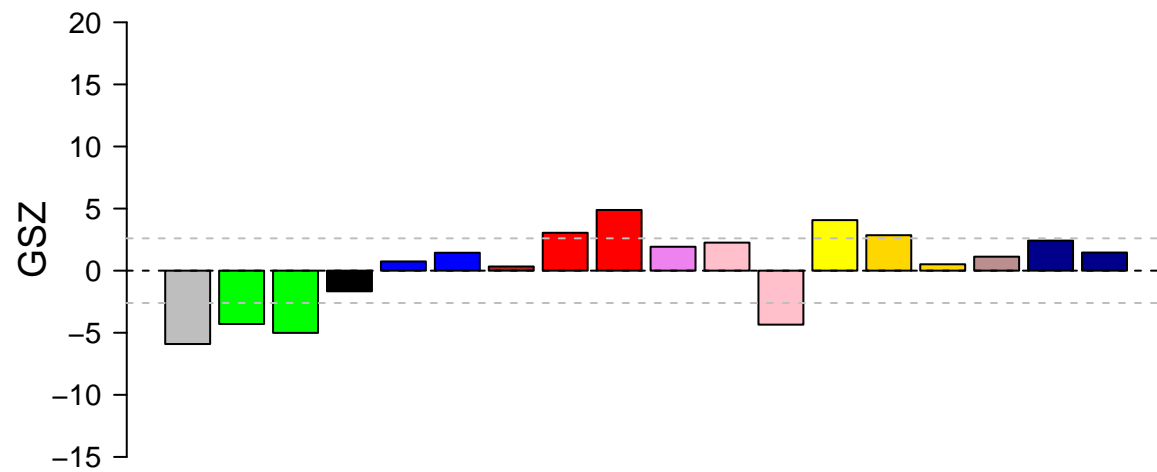
# features = 14 , max = 1

water transport



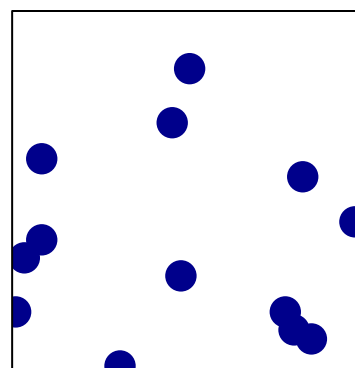
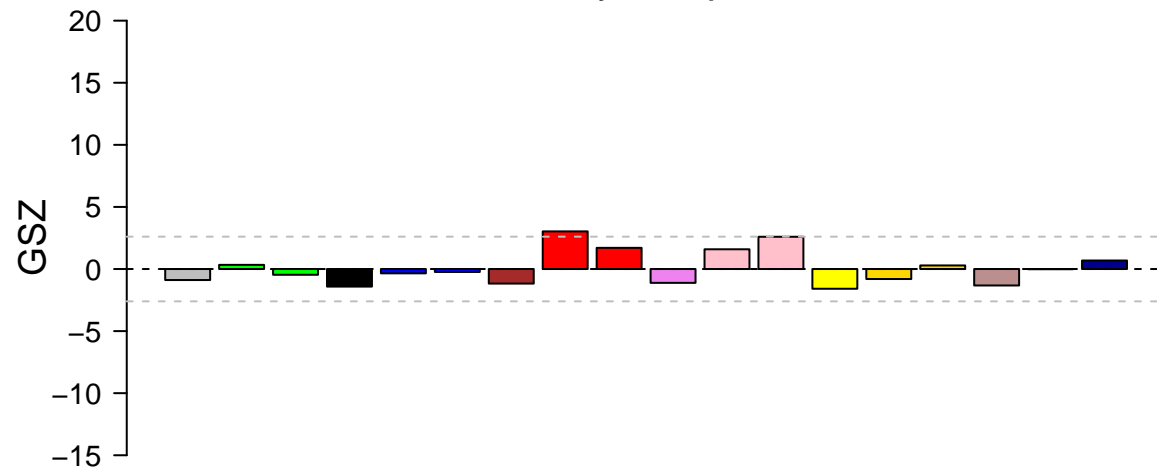
# features = 16 , max = 1

cellular calcium ion homeostasis



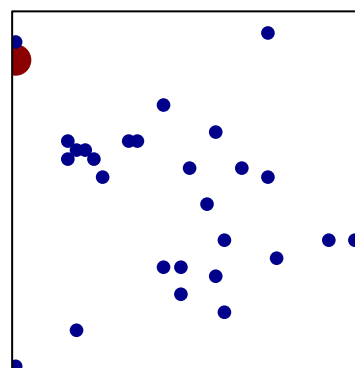
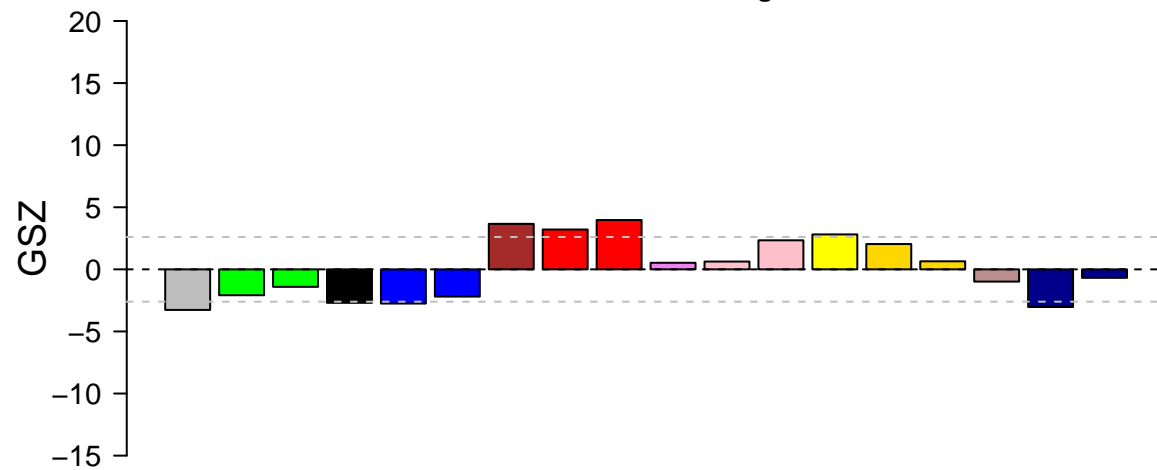
# features = 35 , max = 2

CTP biosynthetic process



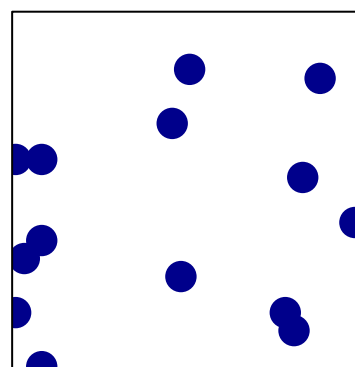
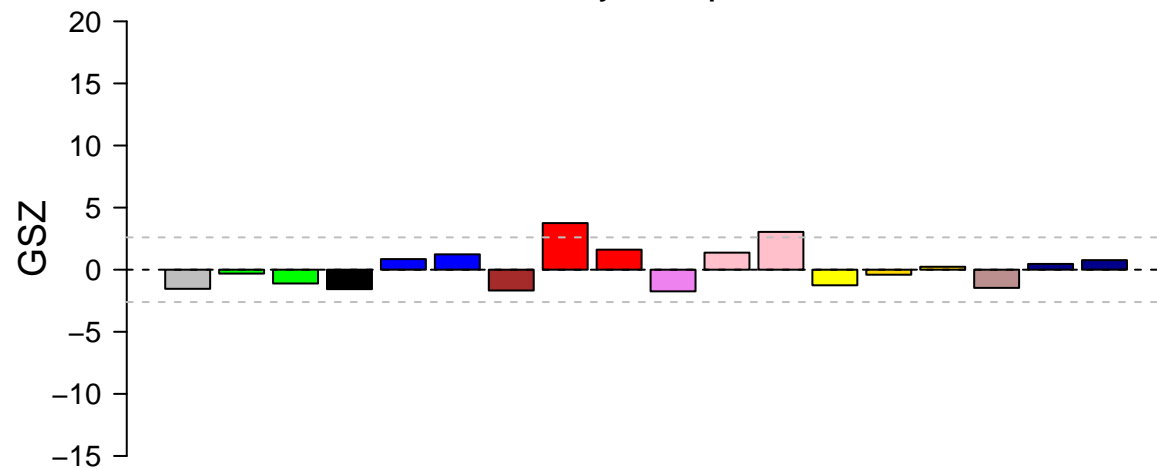
# features = 13 , max = 1

detection of visible light



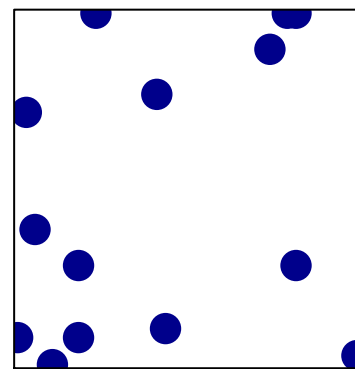
# features = 30 , max = 3

GTP biosynthetic process



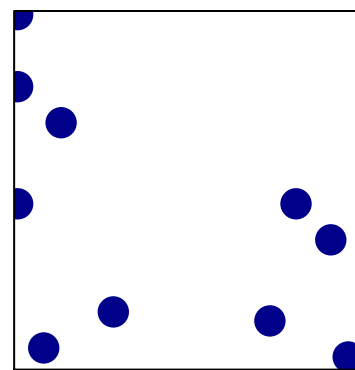
# features = 14 , max = 1

## iron ion homeostasis



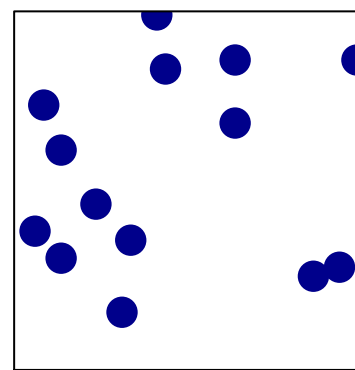
```
# features = 14 , max = 1
```

**regulation of cell differentiation**



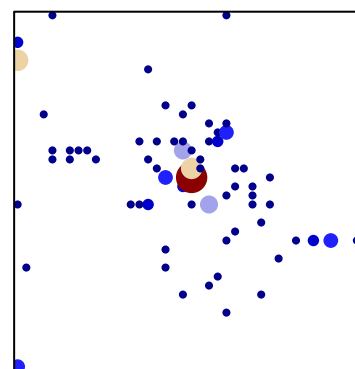
```
# features = 10 , max = 1
```

**response to endoplasmic reticulum stress**



```
# features = 14 , max = 1
```

**response to stimulus**



```
# features = 102 , max = 8
```