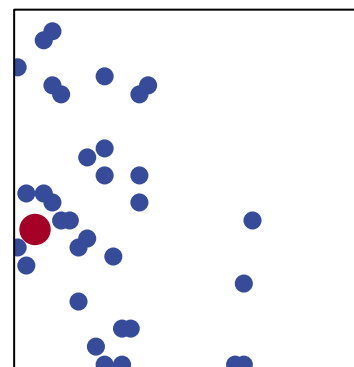
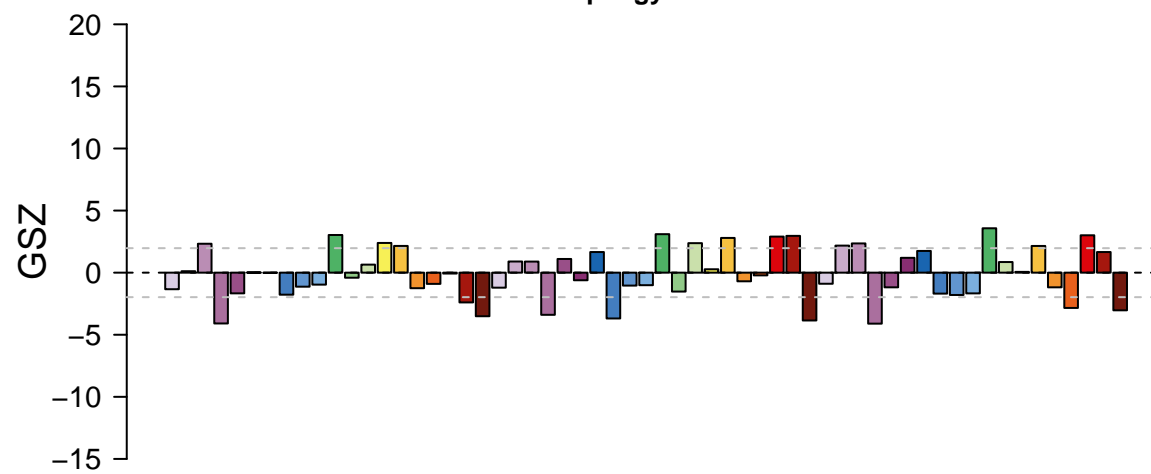
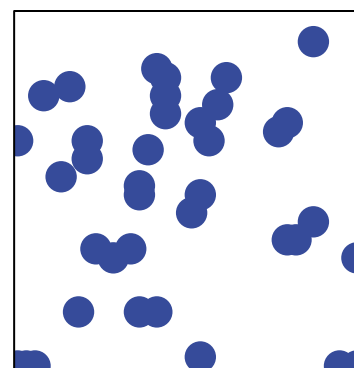
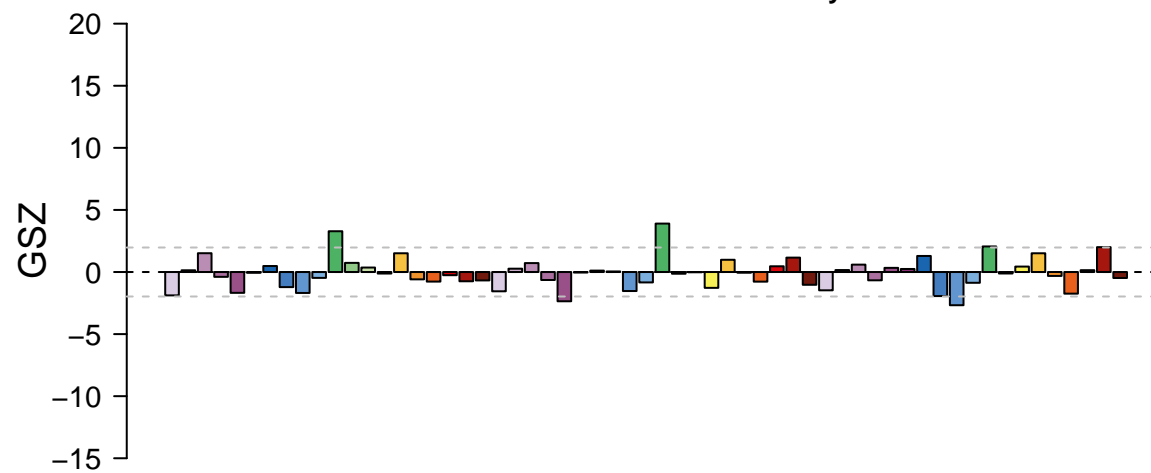


### Mitophagy factors



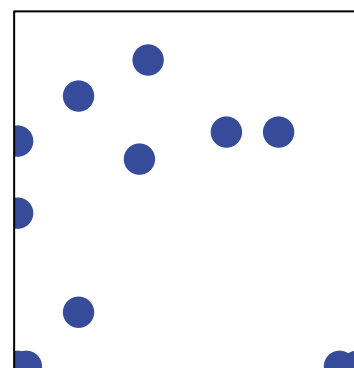
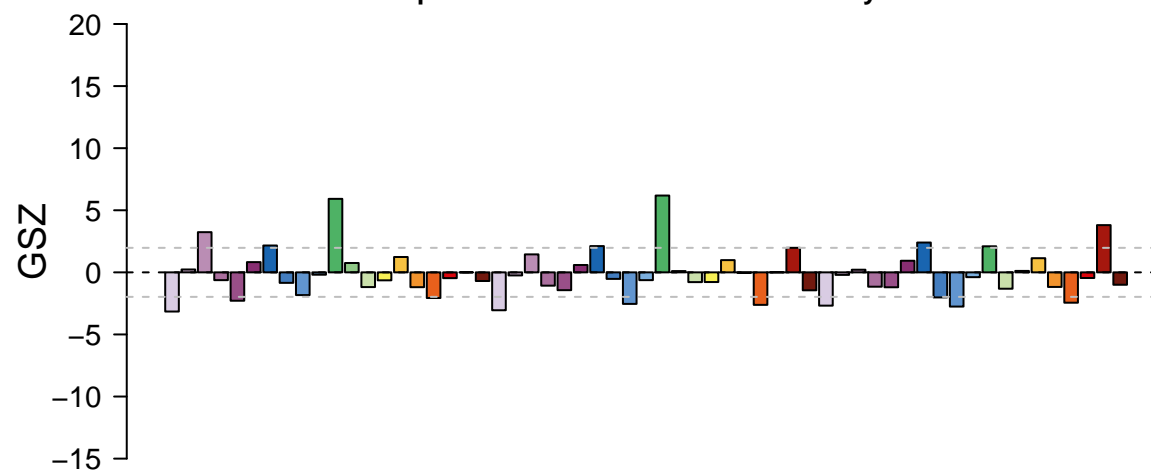
# features = 35 , max = 2

### Protein – Clathrin-mediated endocytosis



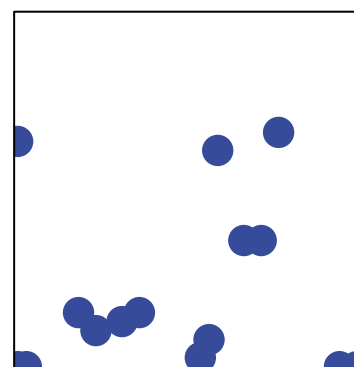
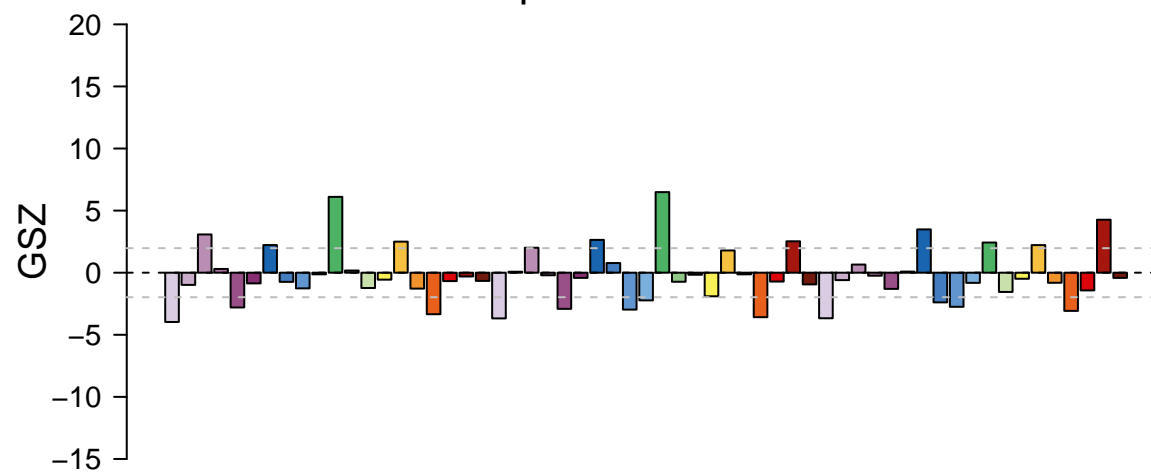
# features = 38 , max = 1

### Endoplasmic reticulum membrane and cytosol



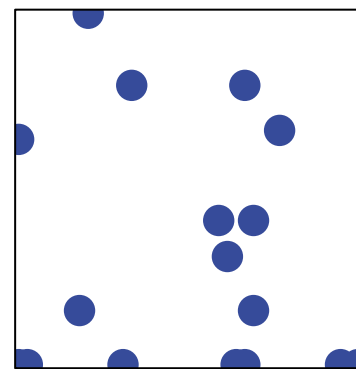
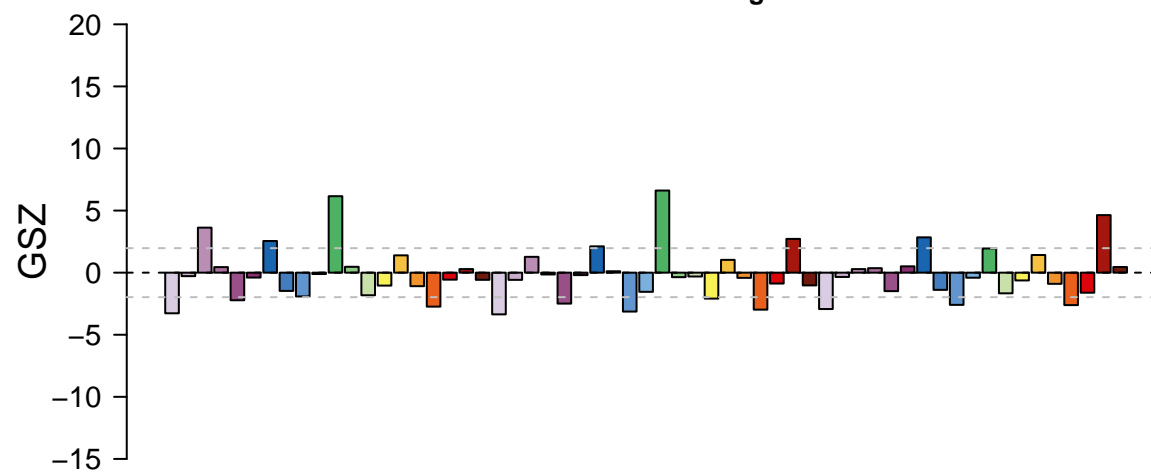
# features = 12 , max = 1

### Chaperone – HSP70 / DNAK



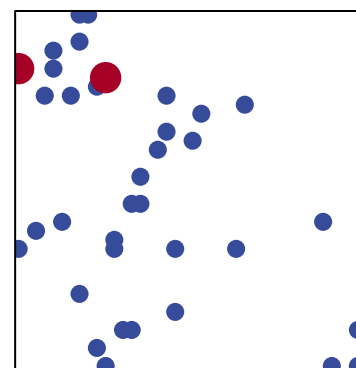
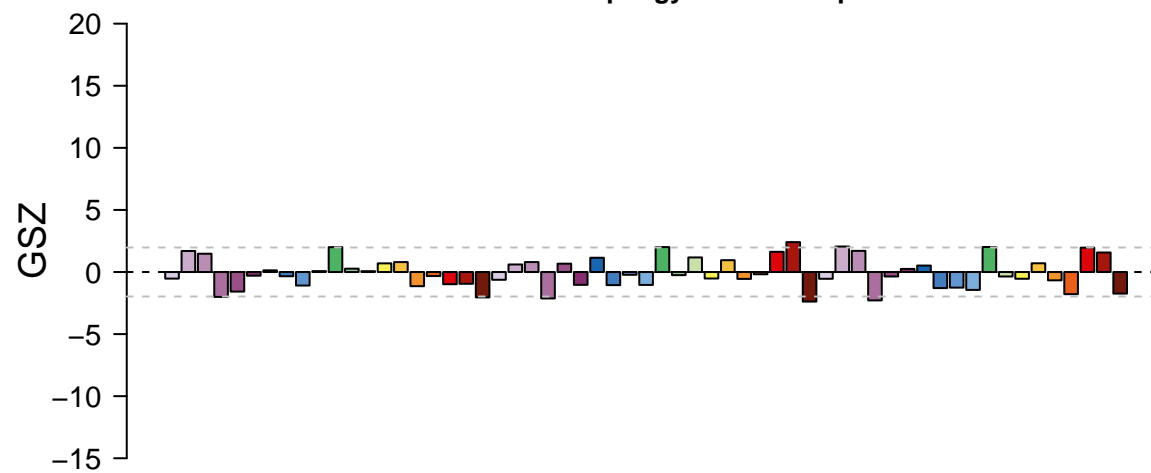
# features = 15 , max = 1

### Proteasome – Assembling factors



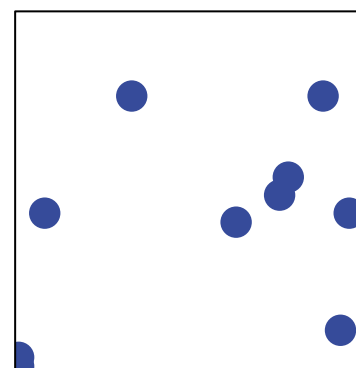
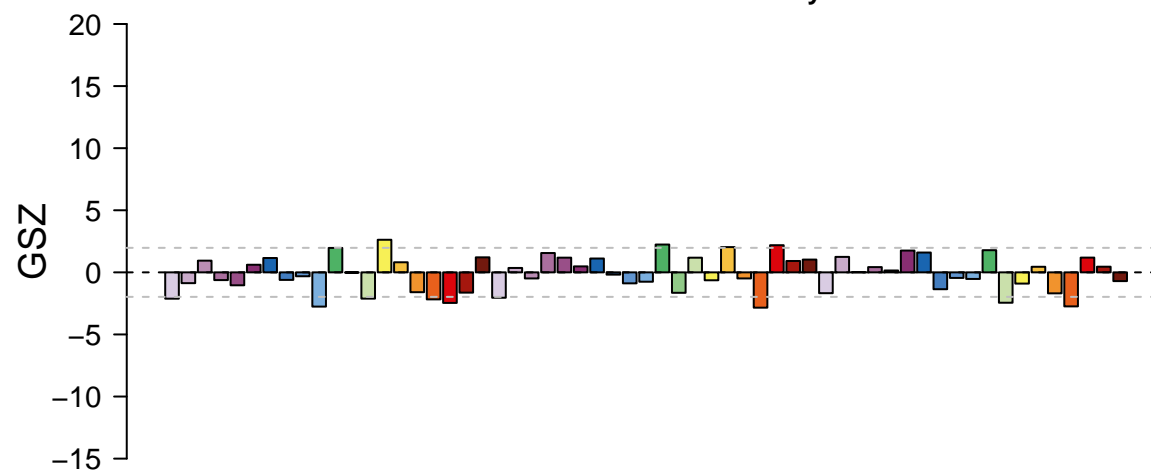
# features = 17 , max = 1

### Protein – Other autophagy associated proteins



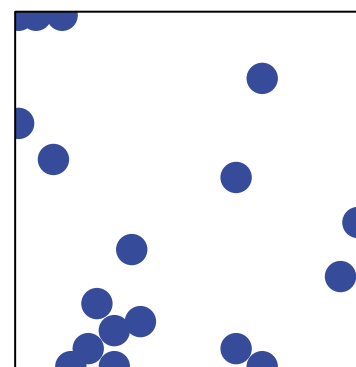
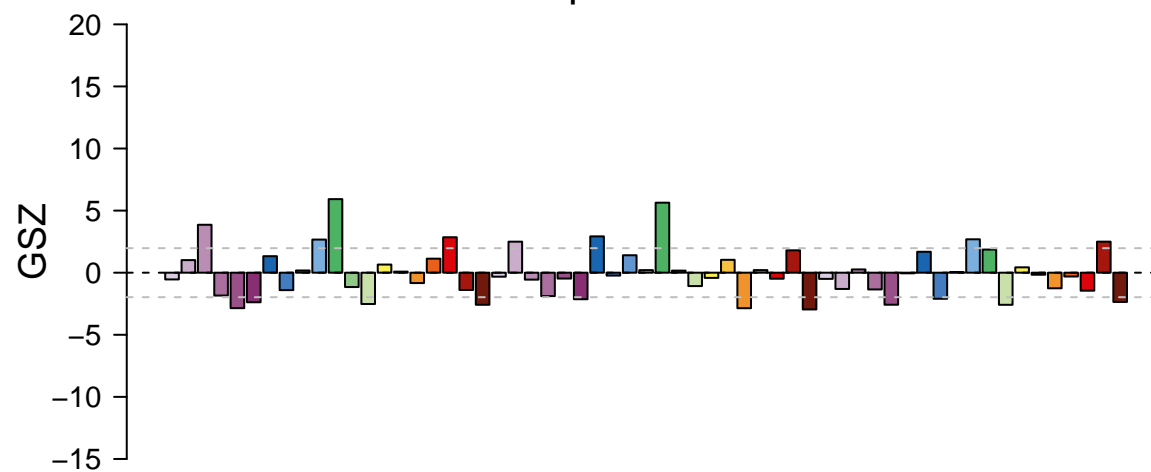
# features = 38 , max = 2

### Valine leucine and isoleucine biosynthesis



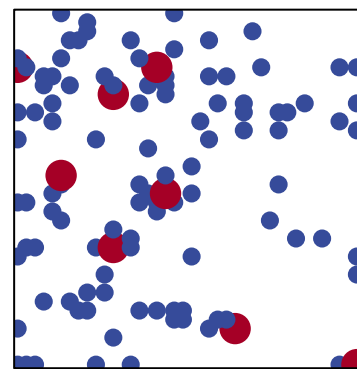
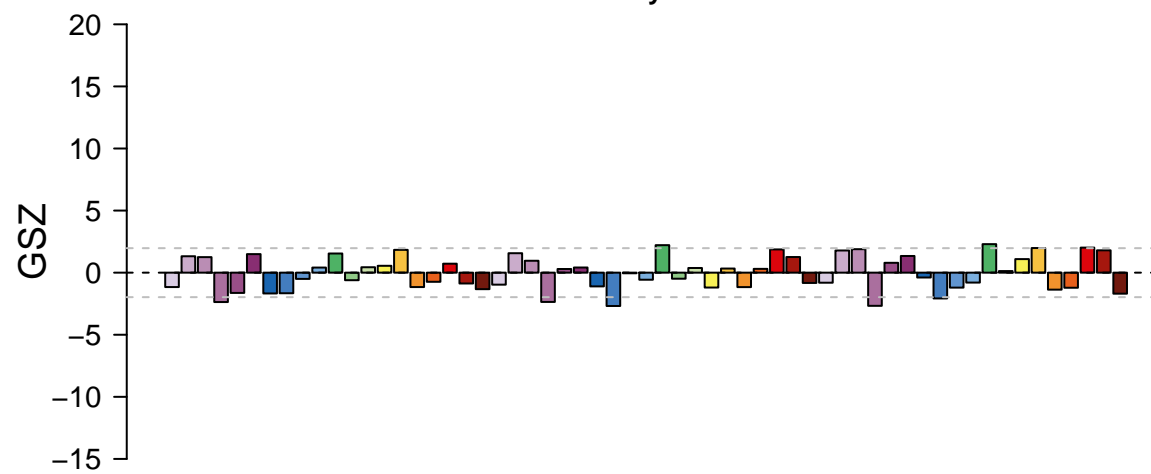
# features = 10 , max = 1

### Chaperone – HSP20



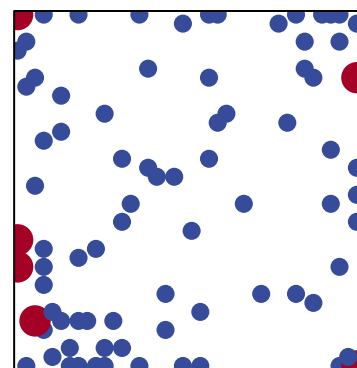
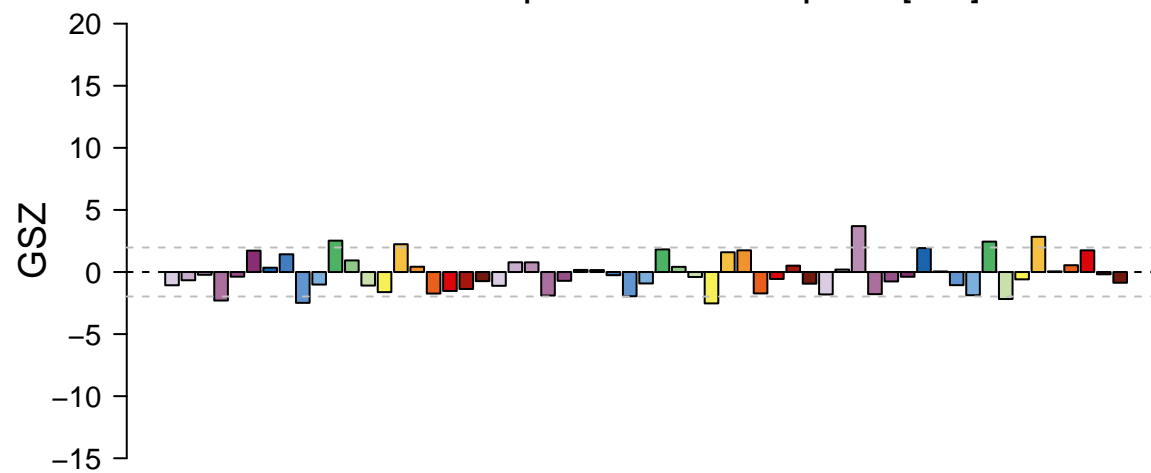
# features = 18 , max = 1

### Endocytosis



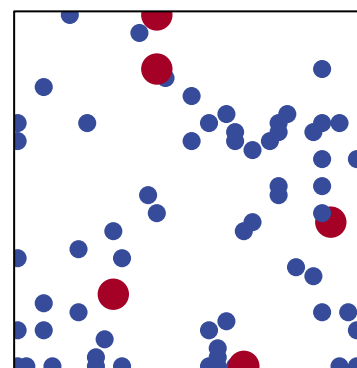
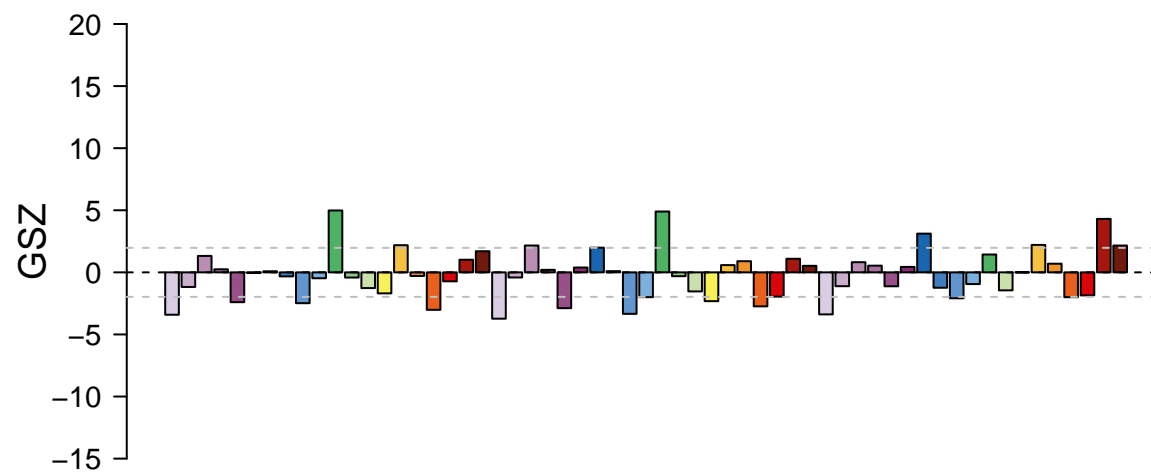
# features = 119 , max = 2

### Electrochemical potential-driven transporters [TC:2]



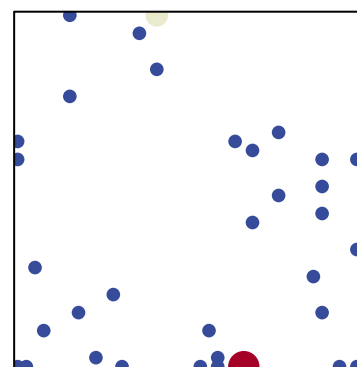
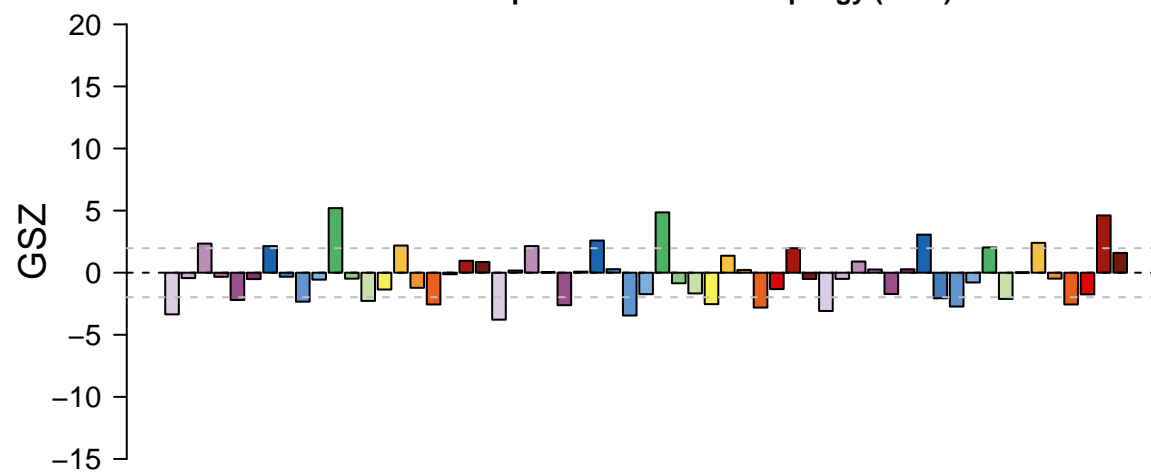
# features = 88 , max = 2

### Exosome – Proteins found in most exosomes



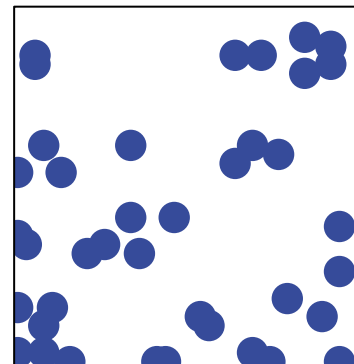
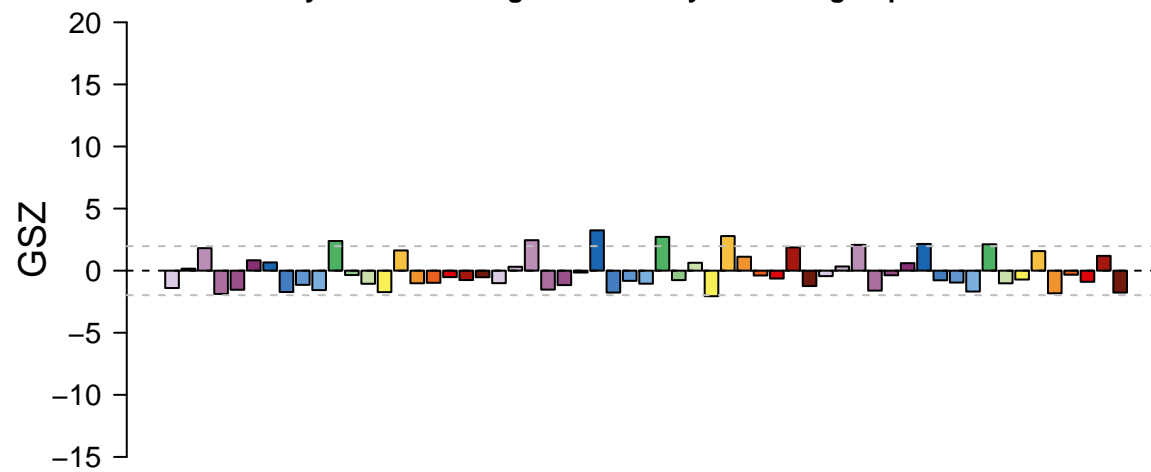
# features = 71 , max = 2

### Protein – Chaperone mediated autophagy (CMA)



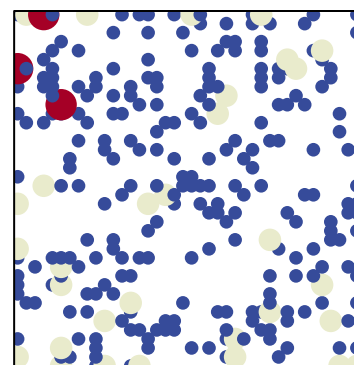
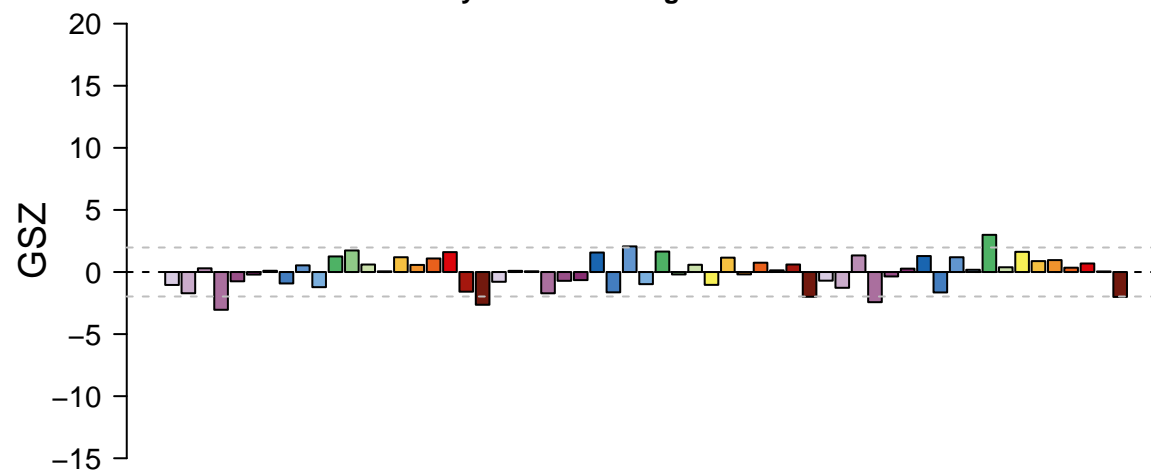
# features = 38 , max = 3

Enzyme – 1.2 Acting on the aldehyde or oxo group of donors



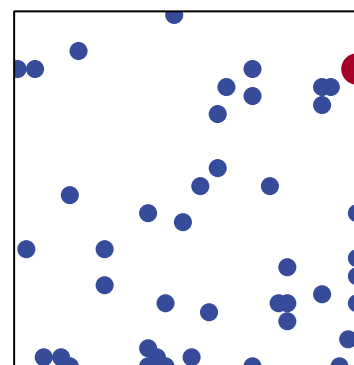
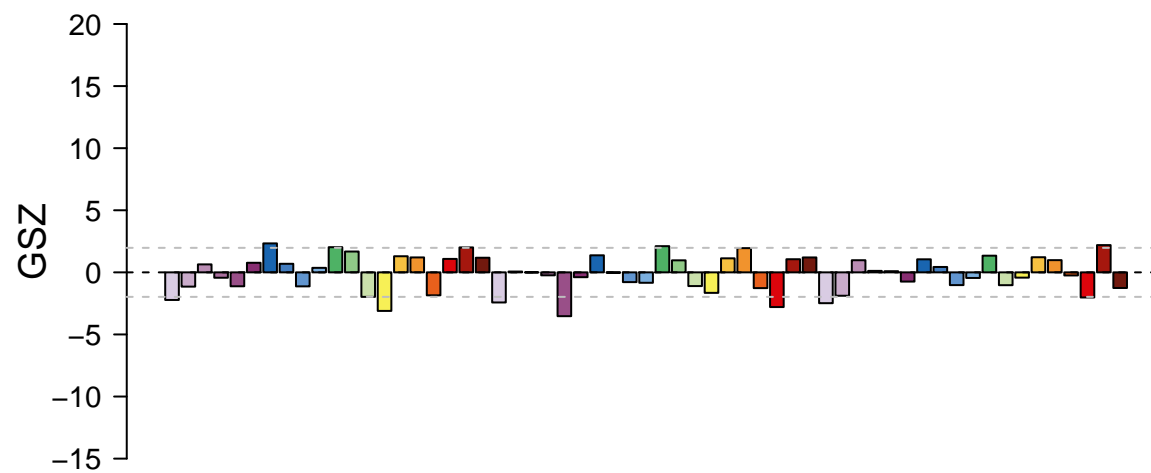
# features = 39 , max = 1

Enzyme – 3.1 Acting on ester bonds



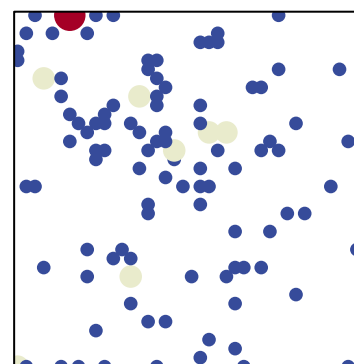
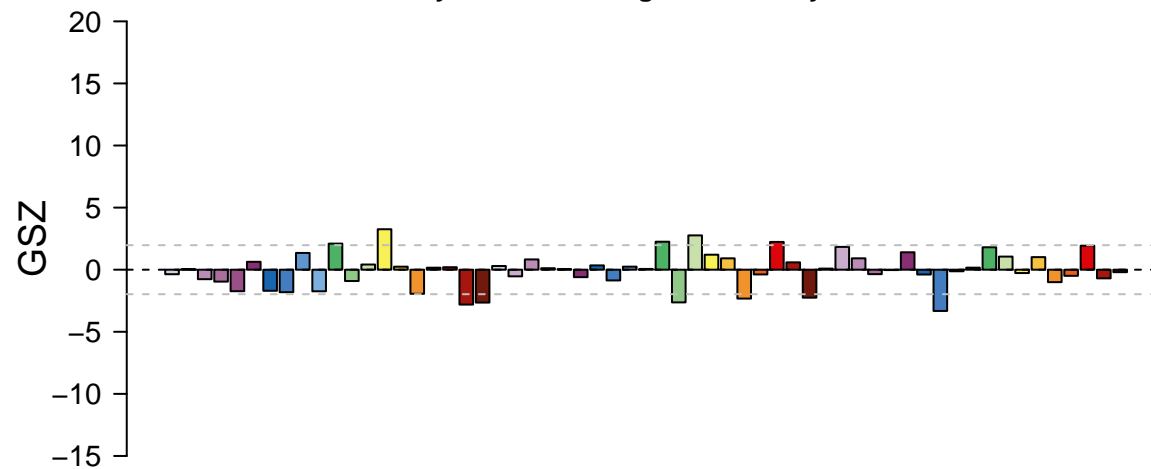
# features = 309 , max = 3

Ascorbate and aldarate metabolism



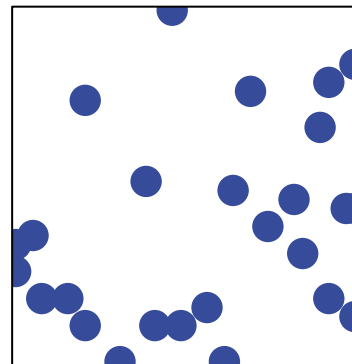
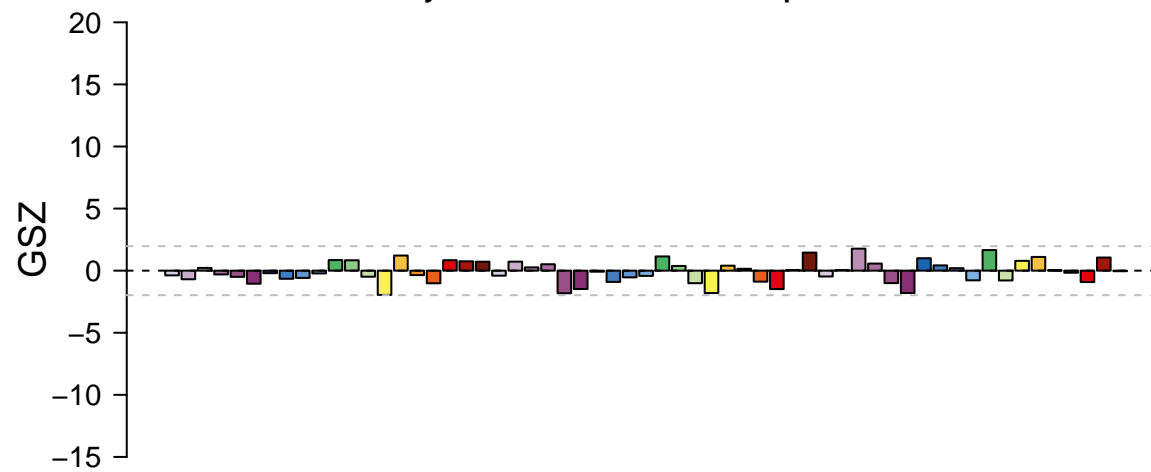
# features = 44 , max = 2

Enzyme – 3.6 Acting on acid anhydrides



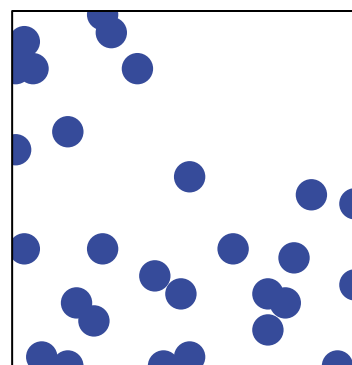
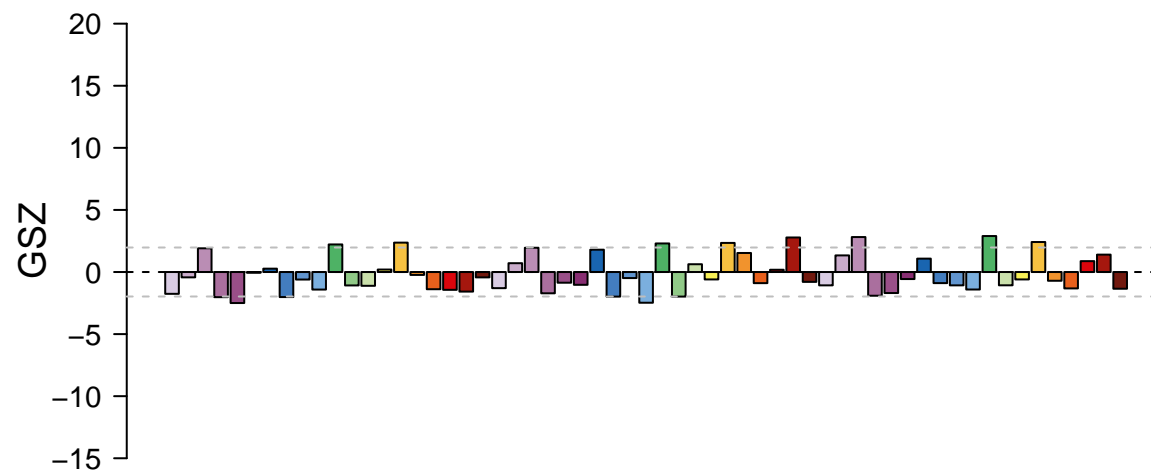
# features = 115 , max = 3

## Enzyme – 5.1 Racemases and epimerases



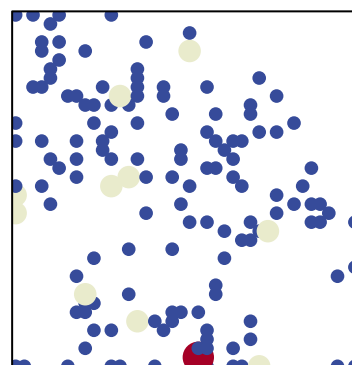
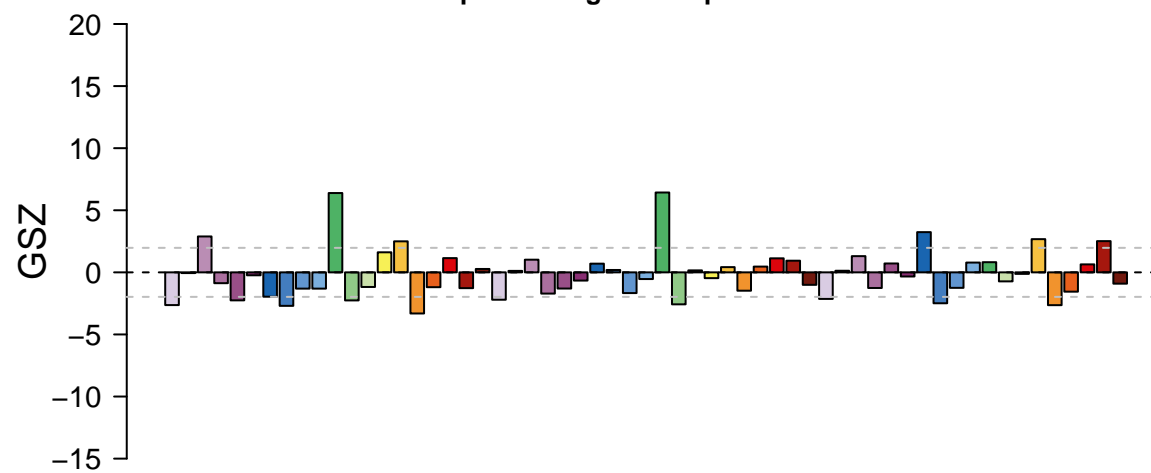
# features = 26 , max = 1

## Other amino acids metabolism – Beta-alanine metabolism



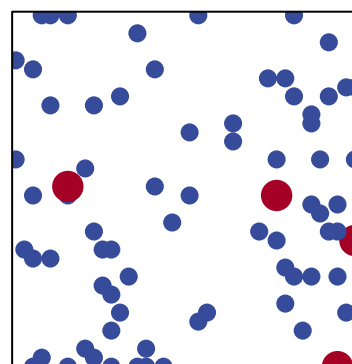
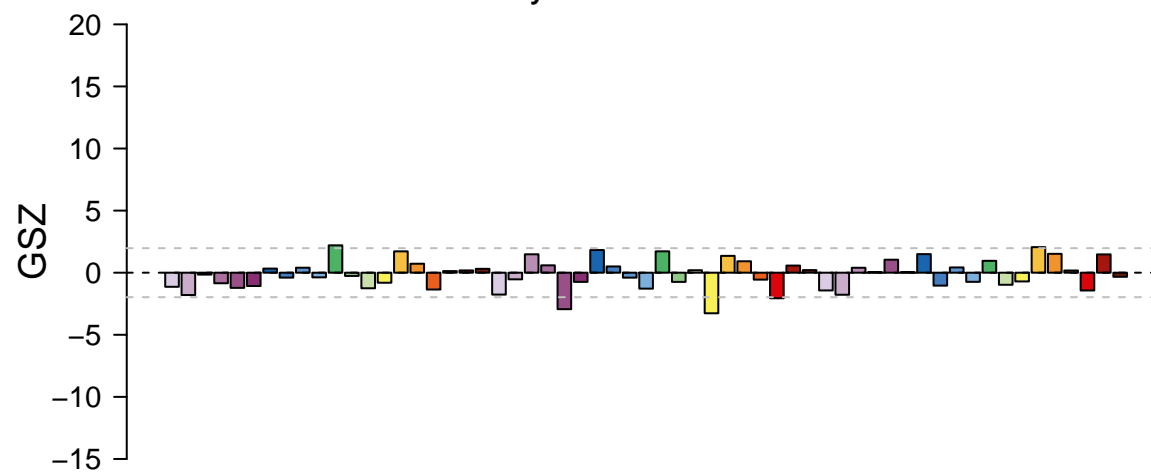
# features = 29 , max = 1

## Protein processing in endoplasmic reticulum



# features = 157 , max = 3

## Pyruvate metabolism



# features = 79 , max = 2