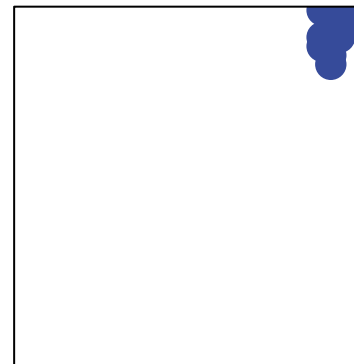
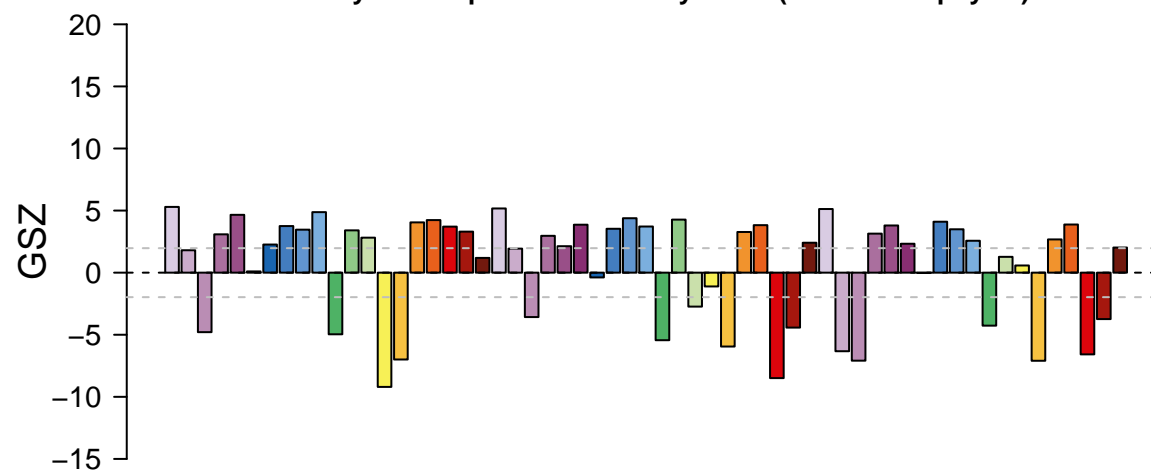
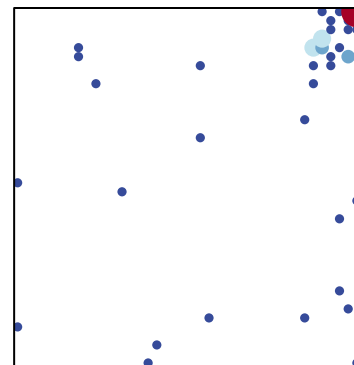
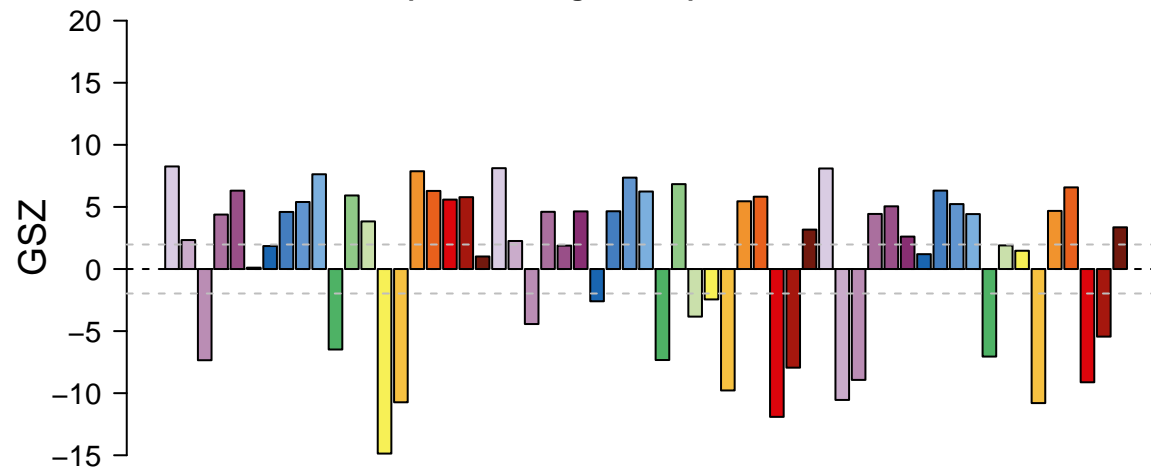


Photosynthesis protein – Photosystem I (P700 chlorophyll a)



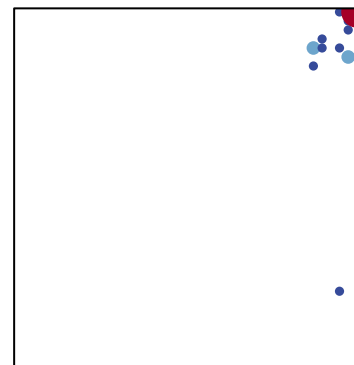
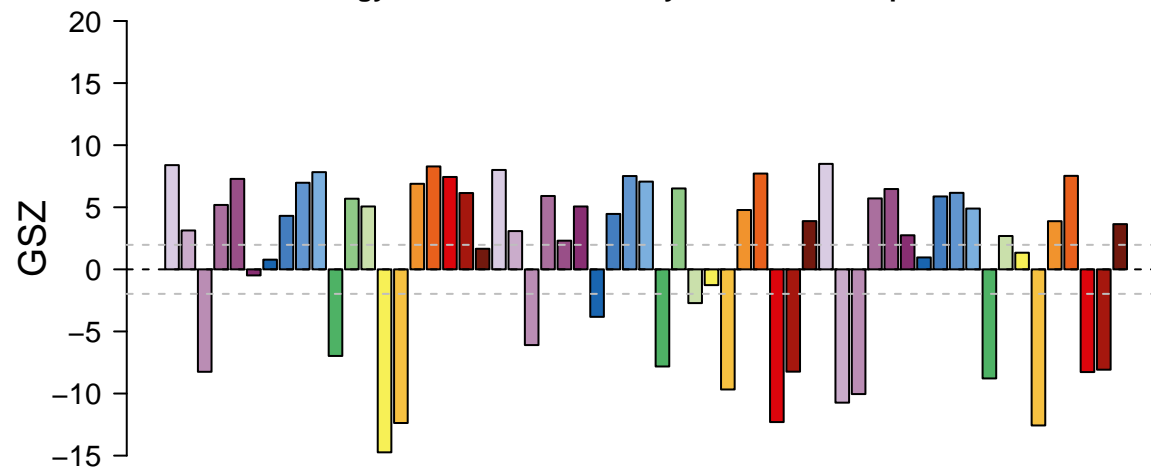
features = 10 , max = 1

Transporter catalog – Transport electron carriers



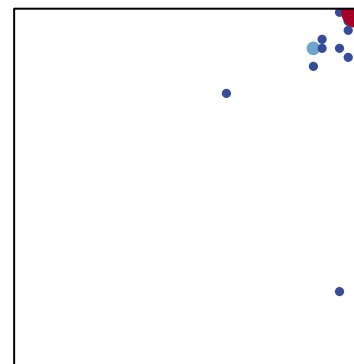
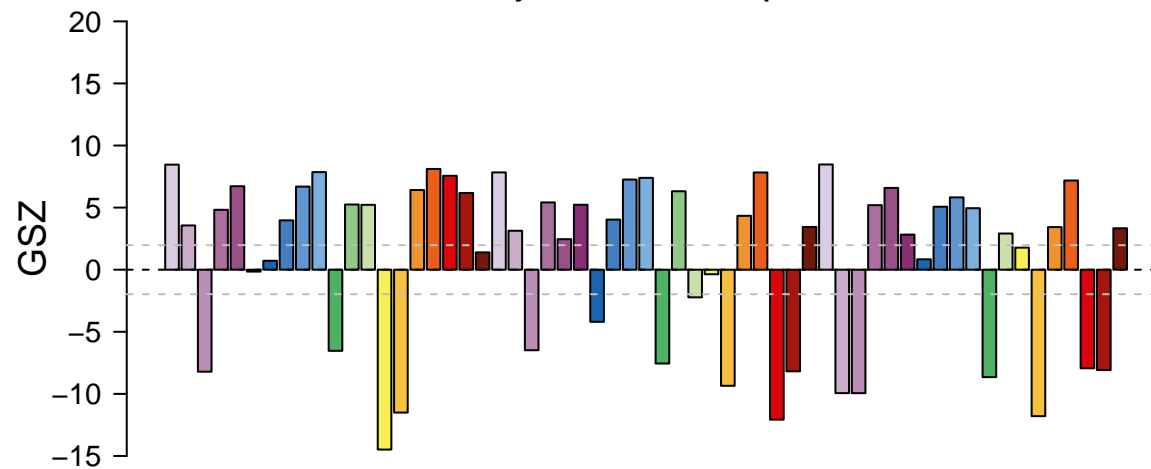
features = 47 , max = 6

Energy metabolism – Photosynthesis antenna proteins



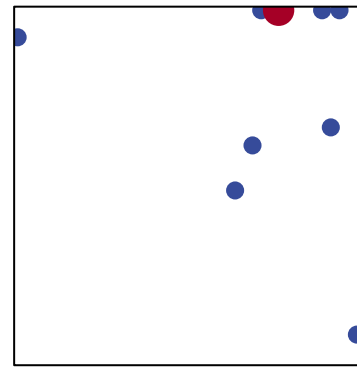
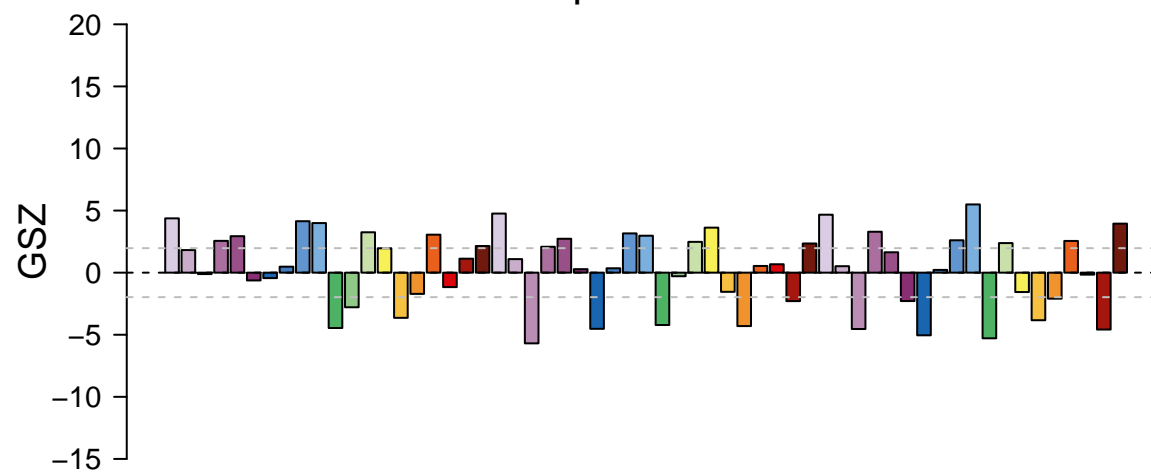
features = 18 , max = 6

Photosynthesis – antenna proteins



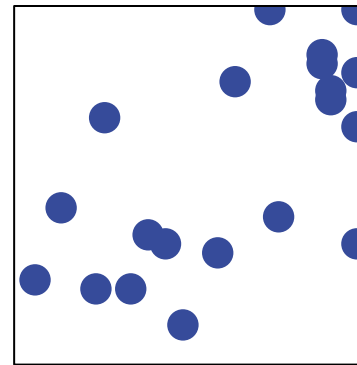
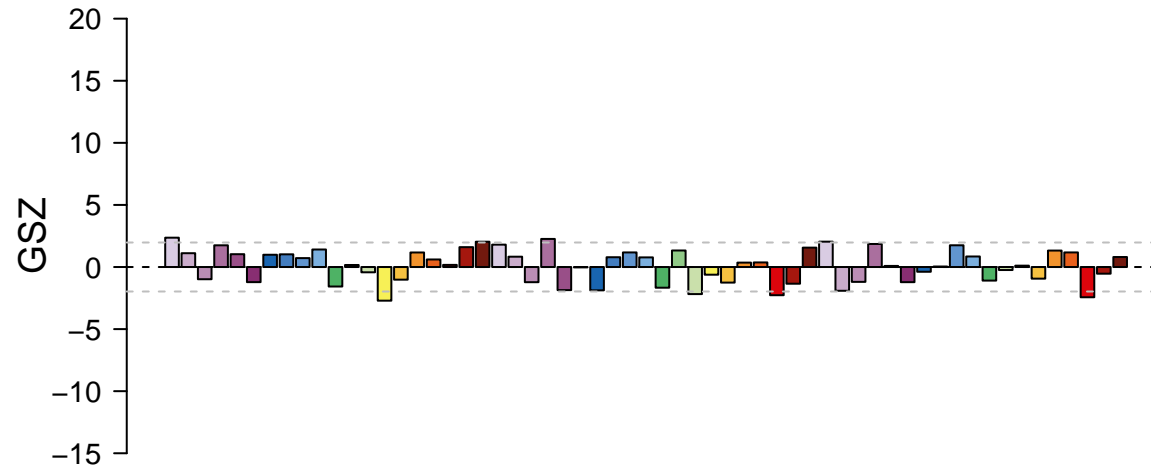
features = 18 , max = 6

Transcription factors – GRF



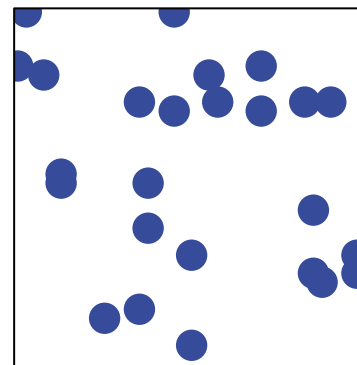
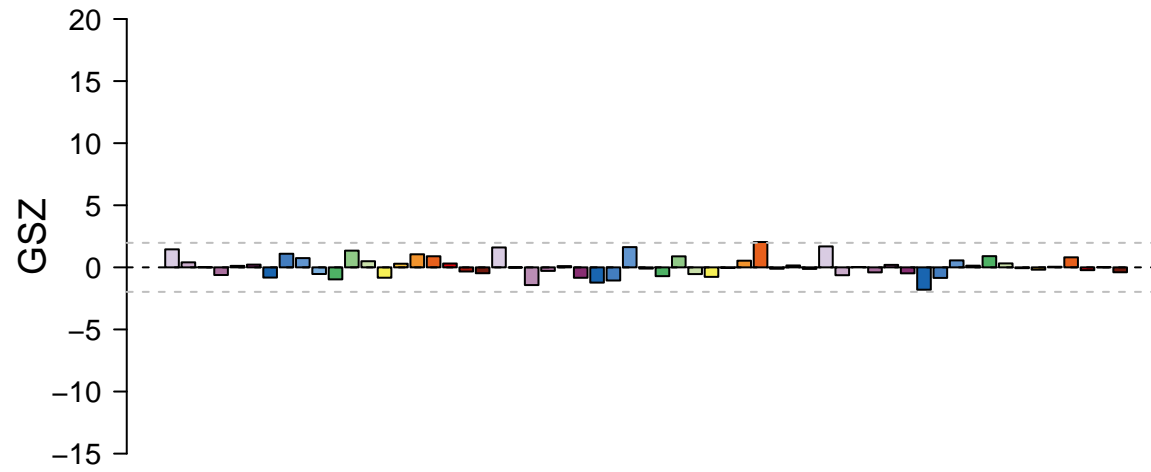
features = 11 , max = 2

Lipid metabolism – Biosynthesis of unsaturated fatty acids



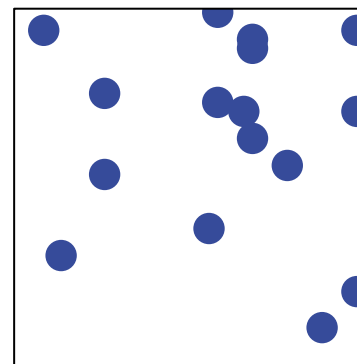
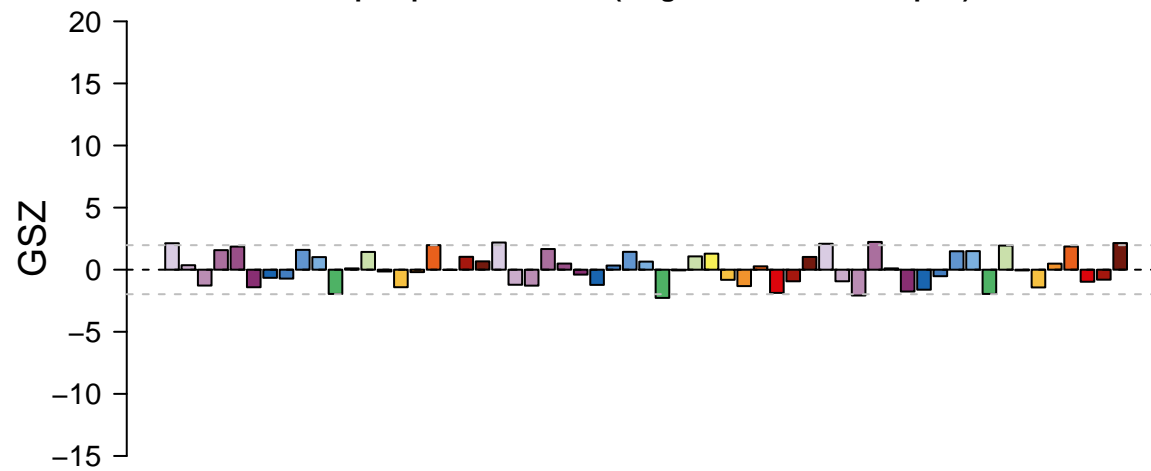
features = 20 , max = 1

Protein – Calcium ion-dependent exocytosis



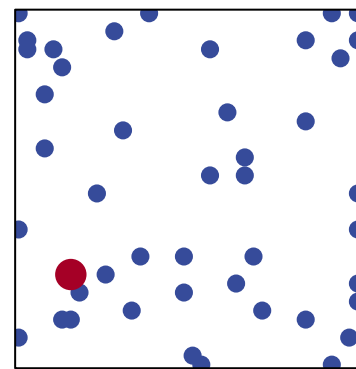
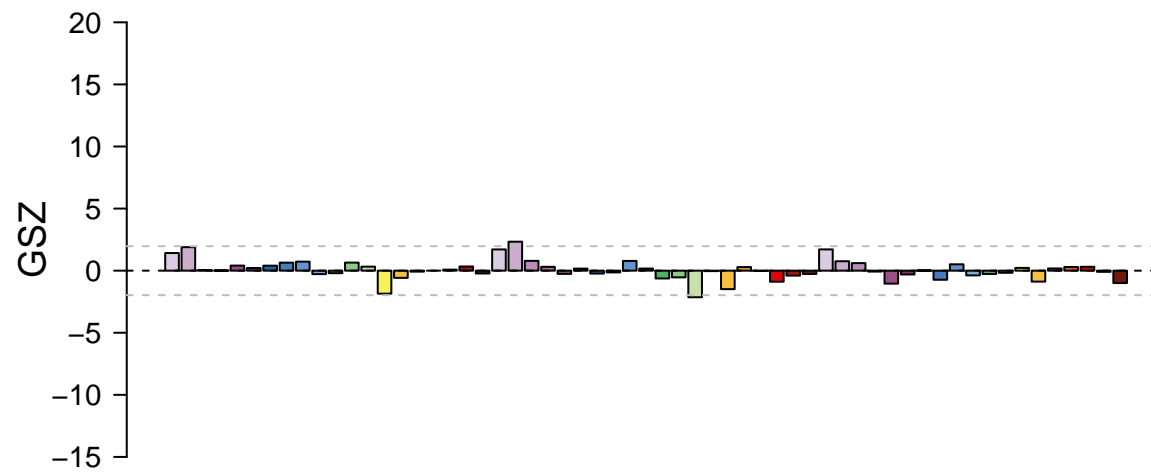
features = 25 , max = 1

Repair protein – SSBR (single strand breaks repair)



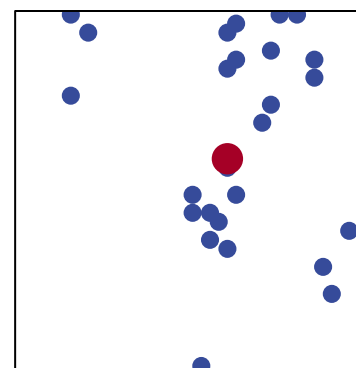
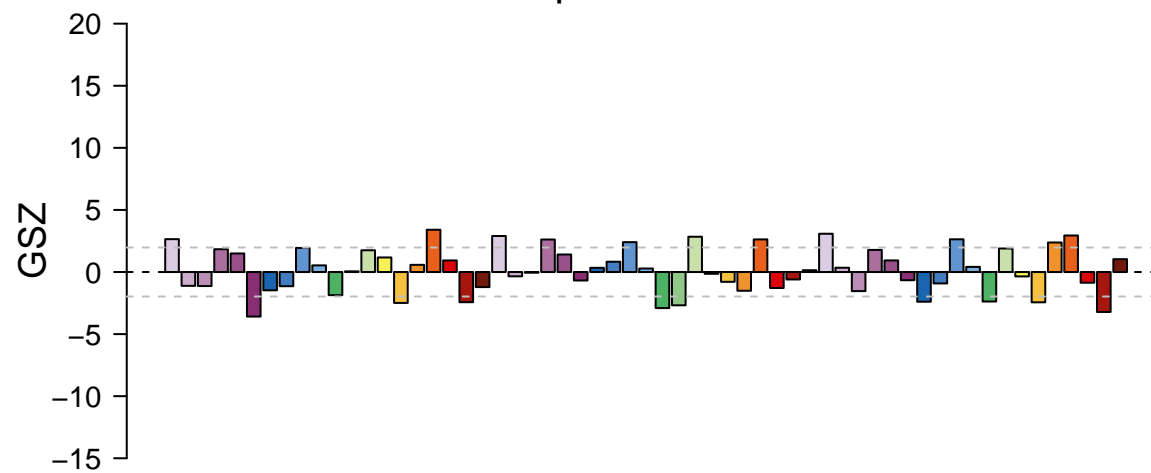
features = 16 , max = 1

Amino acid metabolism – Glutamate metabolism



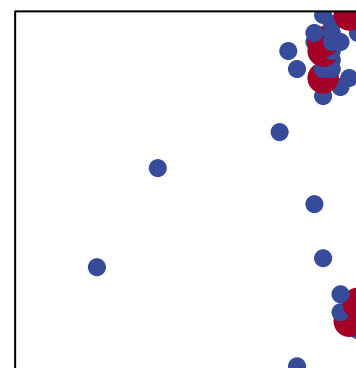
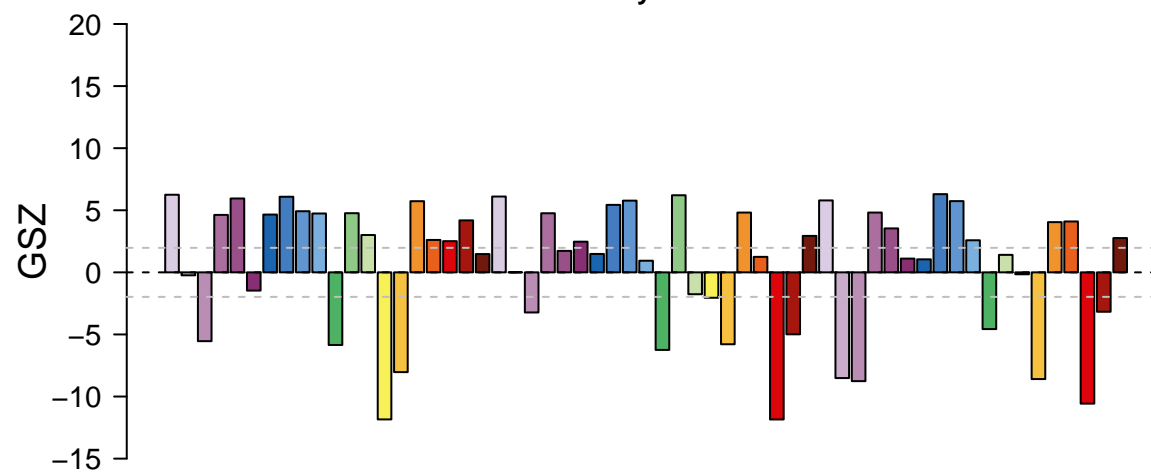
features = 46 , max = 2

Transcription factors – MTERF



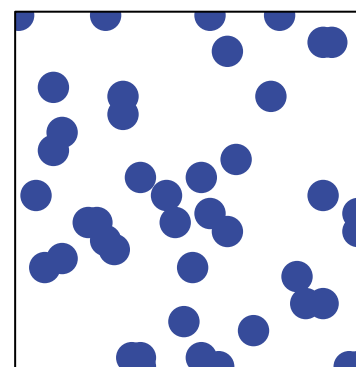
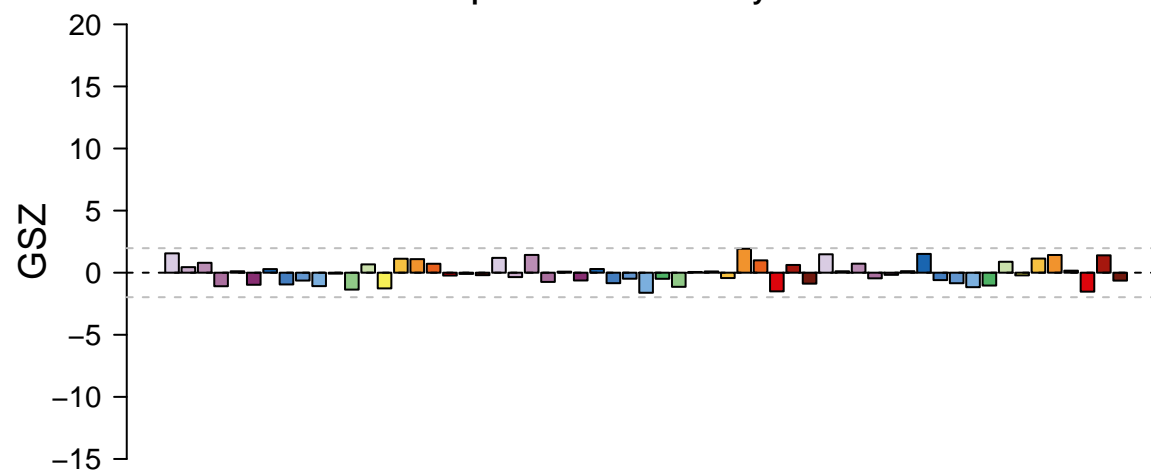
features = 28 , max = 2

Photosynthesis



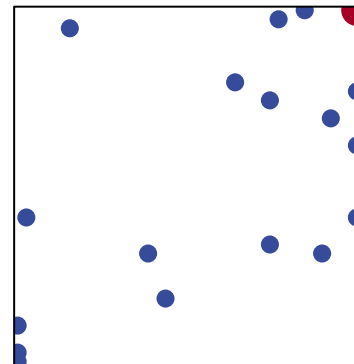
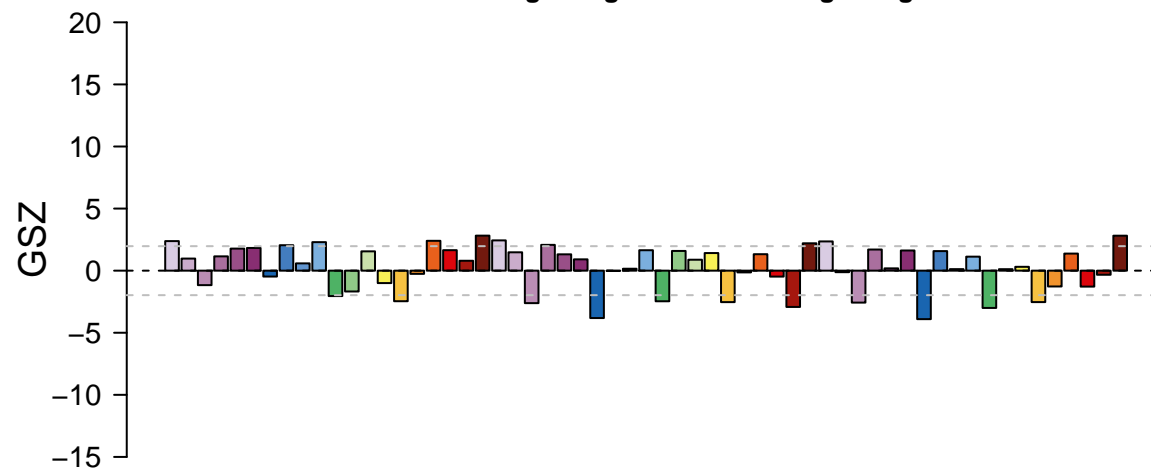
features = 38 , max = 2

Terpenoid backbone biosynthesis



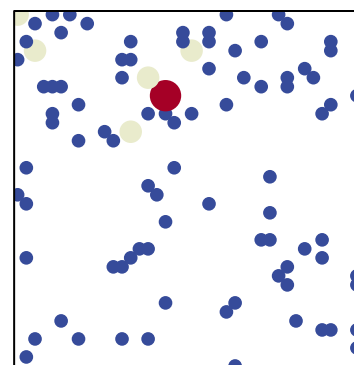
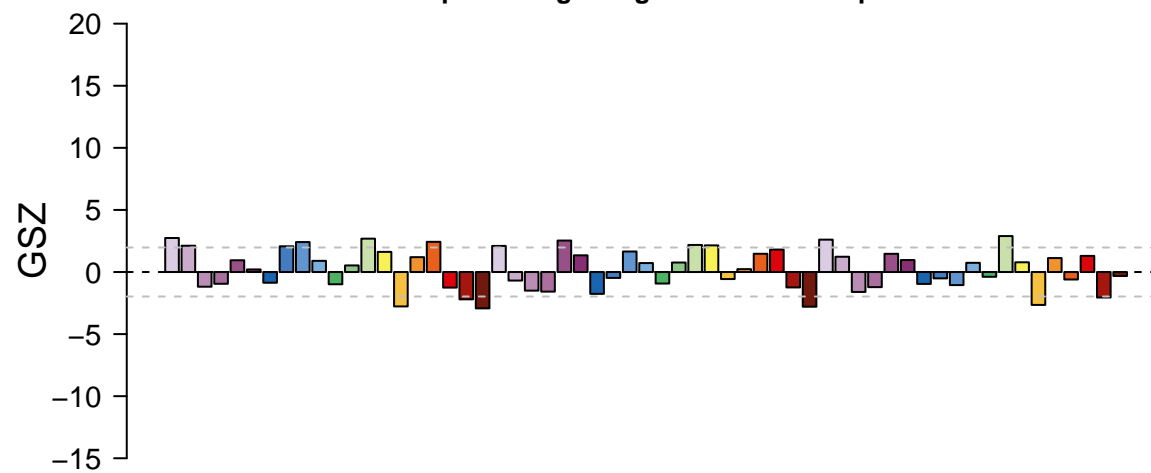
features = 43 , max = 1

Hormone signaling – Gibberellin signaling



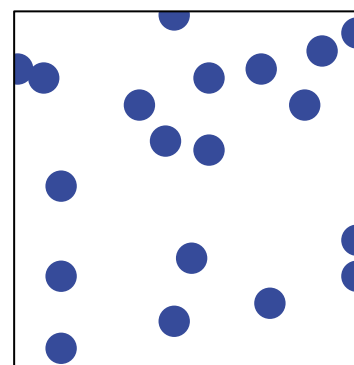
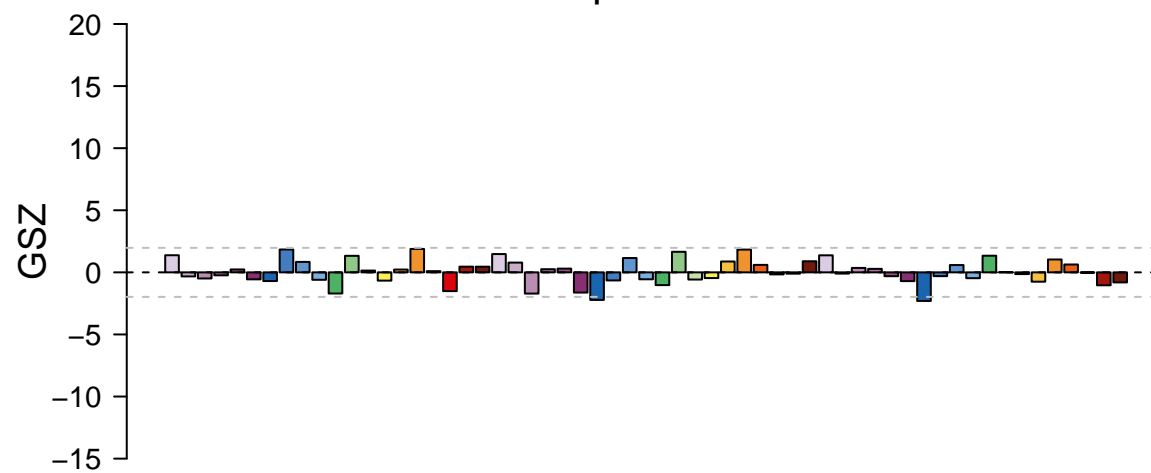
features = 19 , max = 2

Plant specific signaling – Flower development



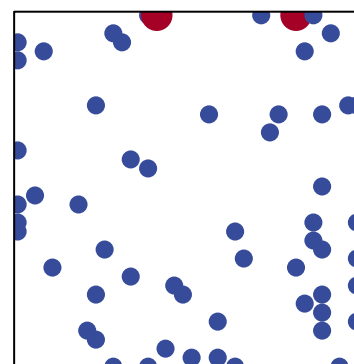
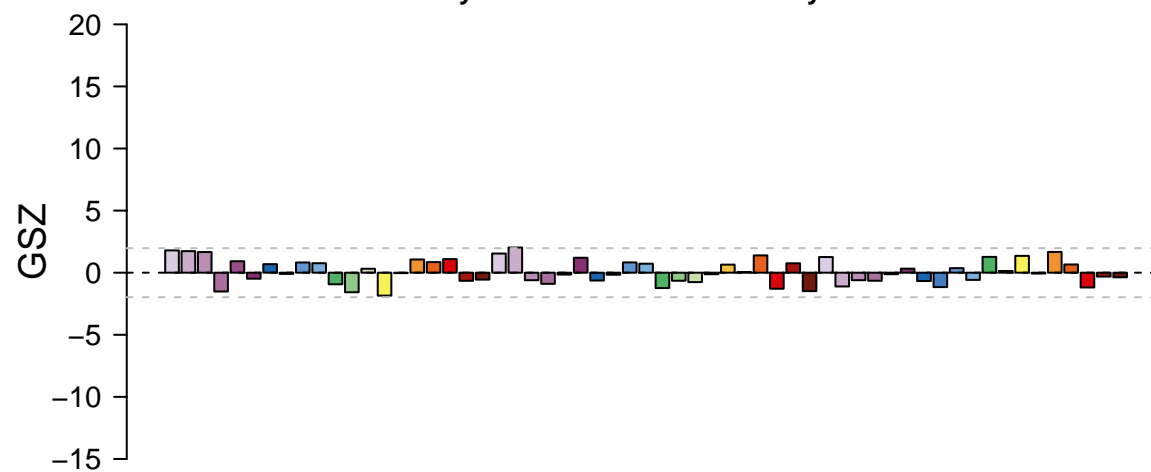
features = 100 , max = 3

Ether lipid metabolism



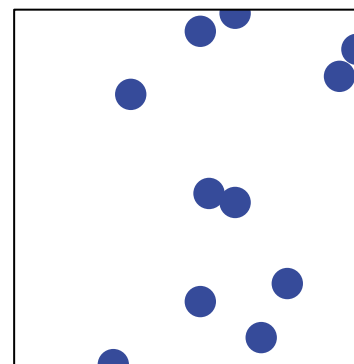
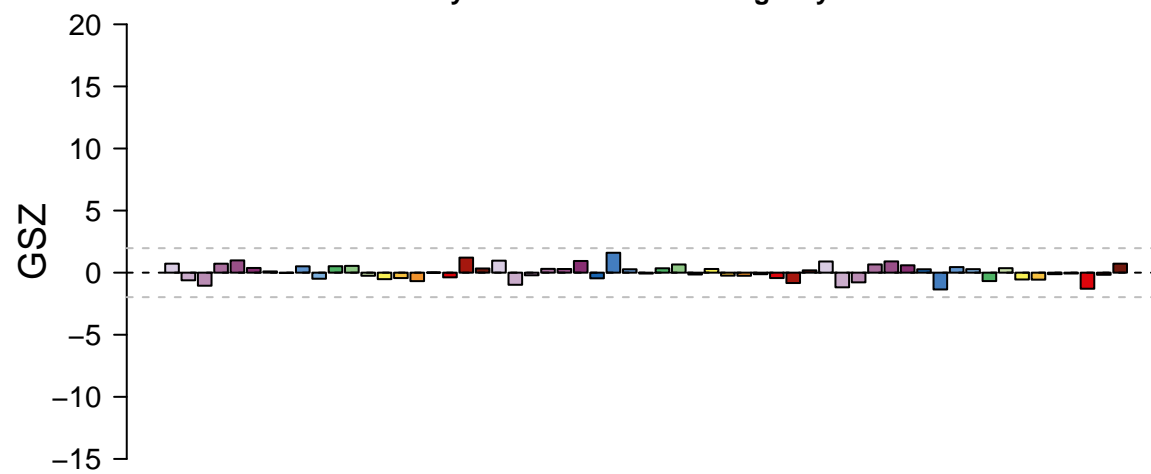
features = 19 , max = 1

Enzyme – 4.1 Carbon-carbon lyases



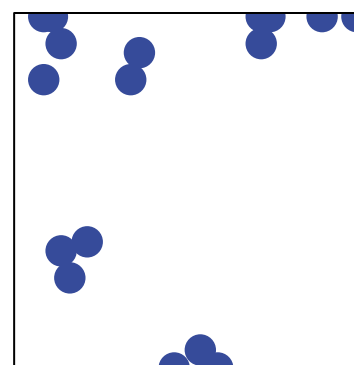
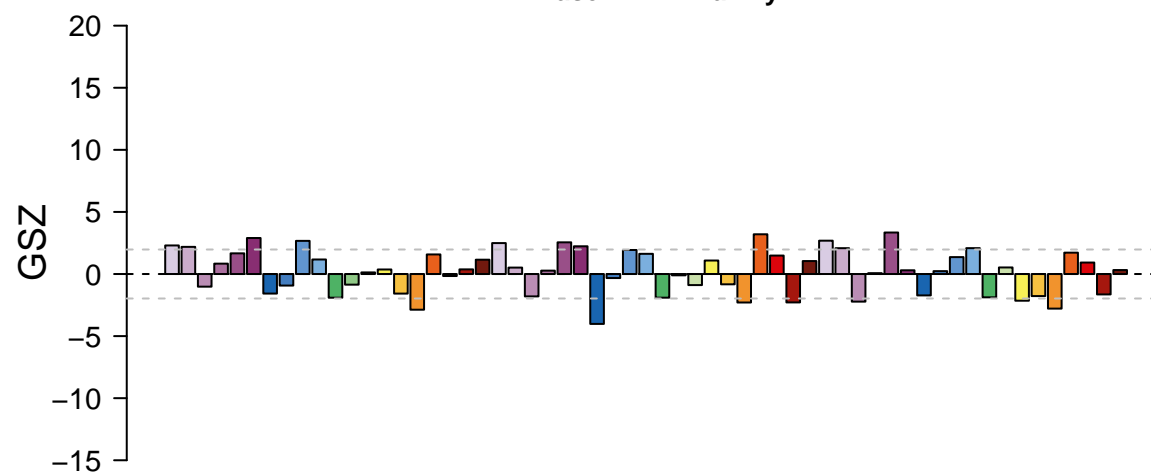
features = 60 , max = 2

Enzyme – 4.3 Carbon–nitrogen lyases



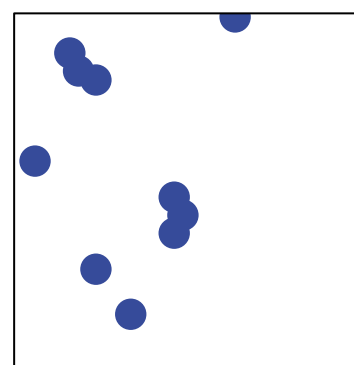
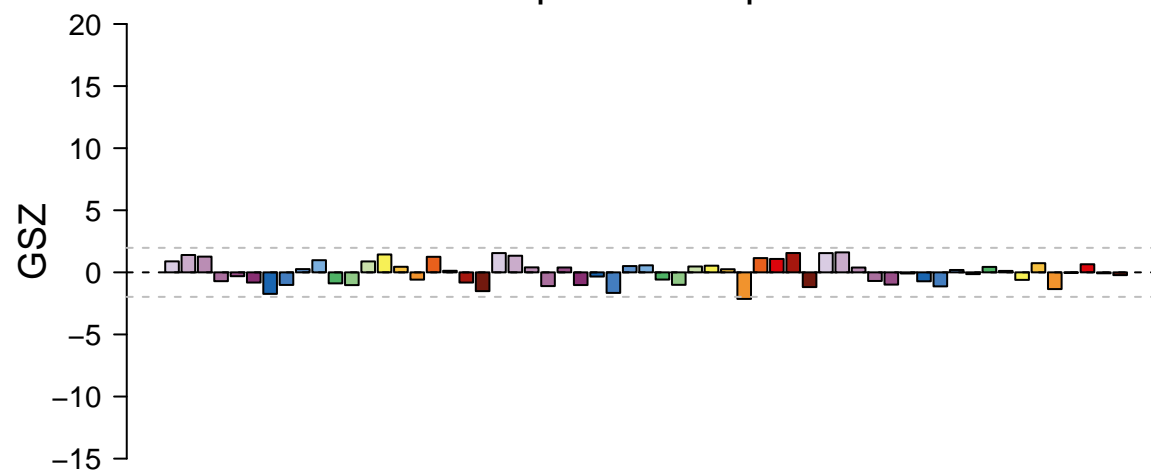
features = 11 , max = 1

Kinase – IRAK family



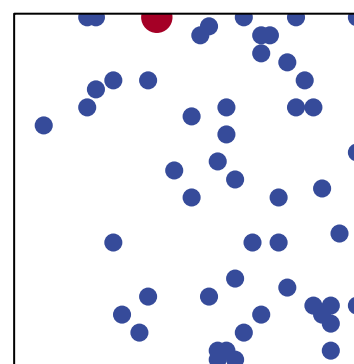
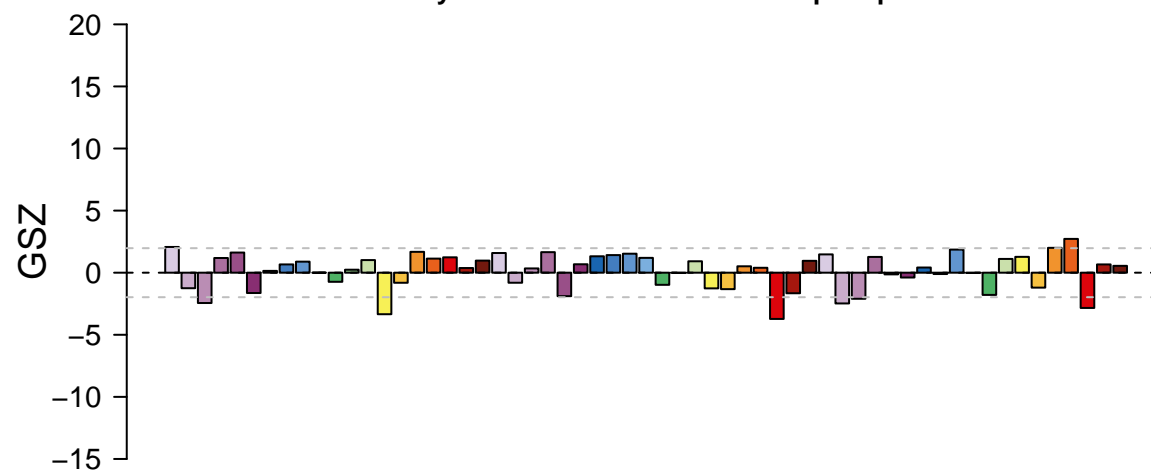
features = 17 , max = 1

Minor spliceosome components



features = 10 , max = 1

Carbohydrate metabolism – Pentose phosphate



features = 54 , max = 2