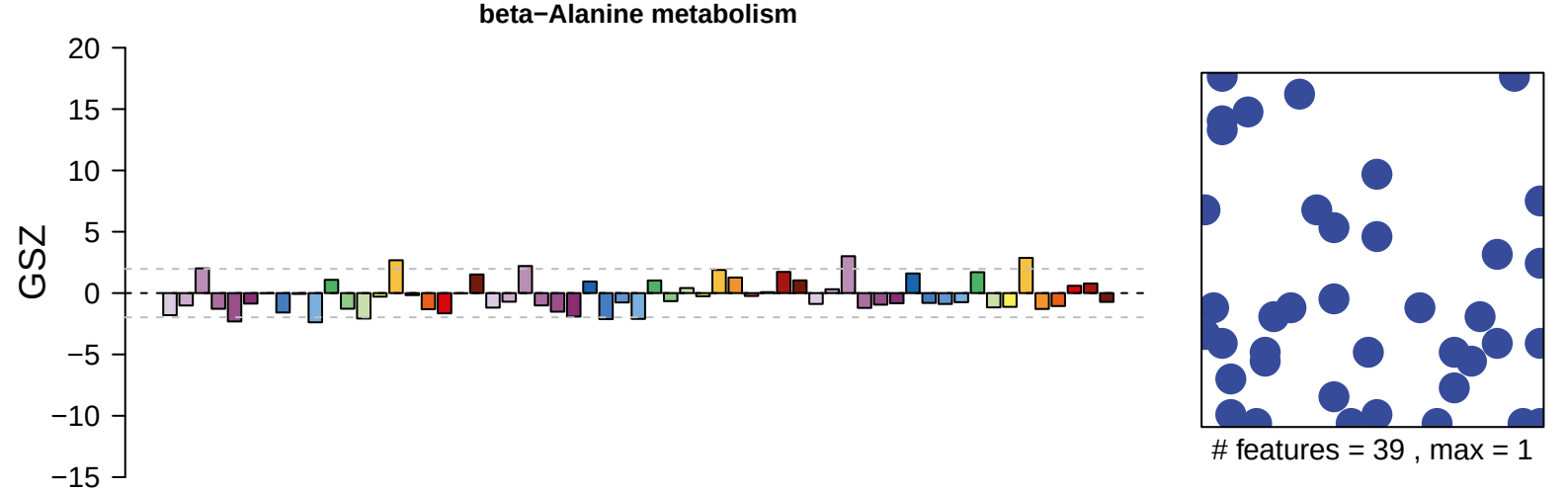
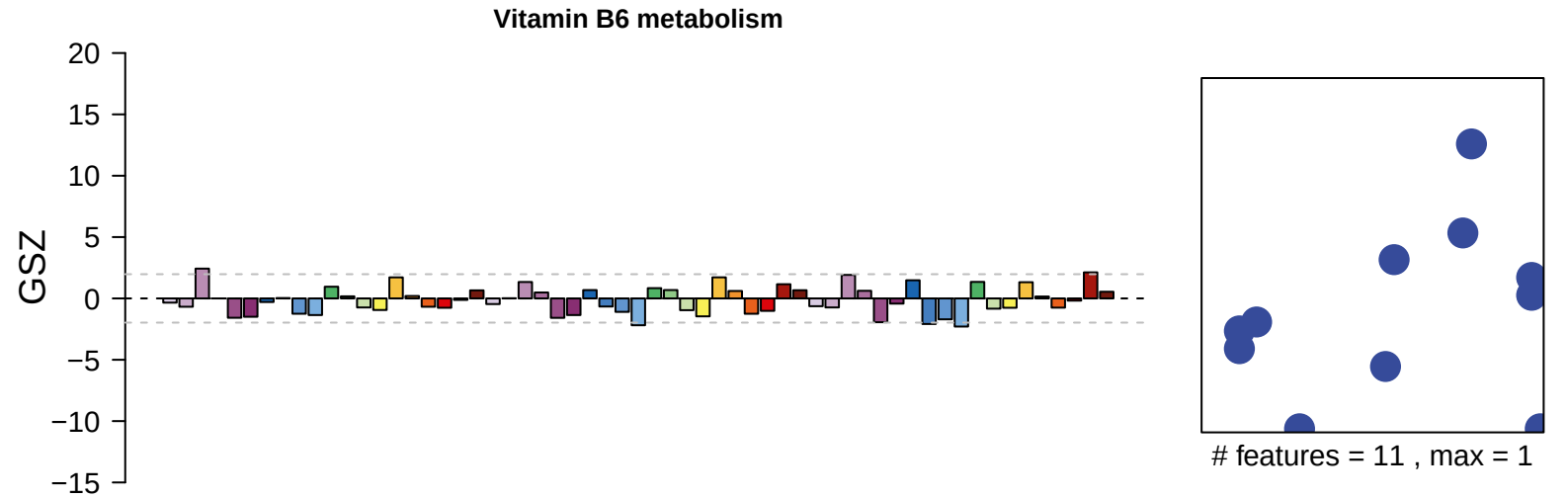
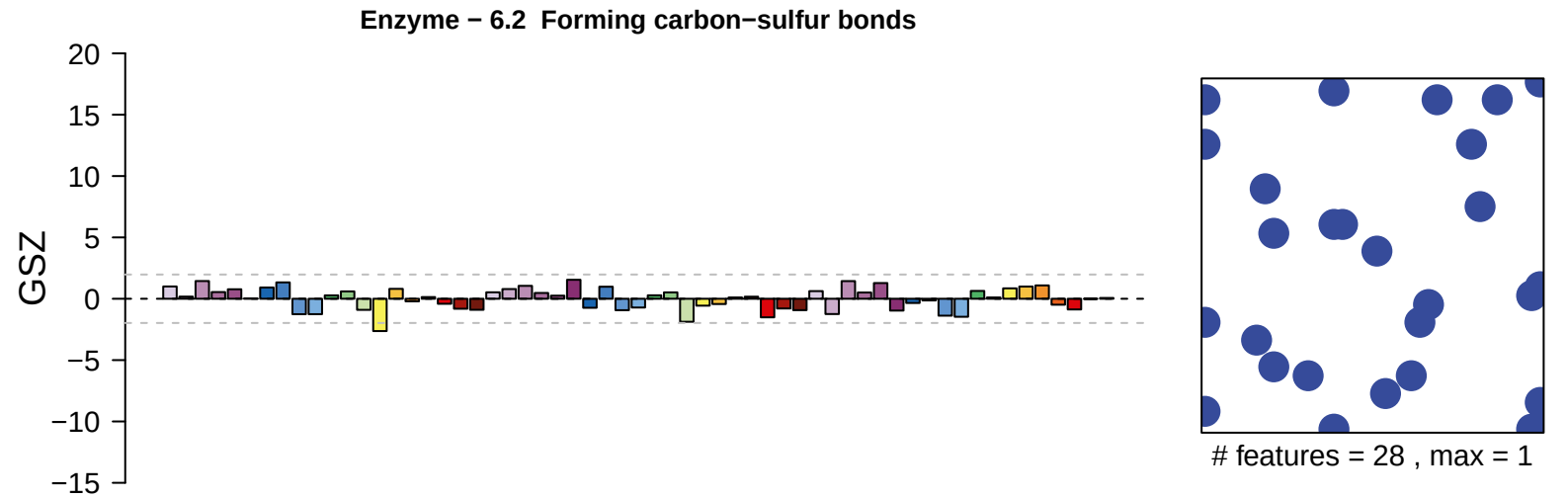
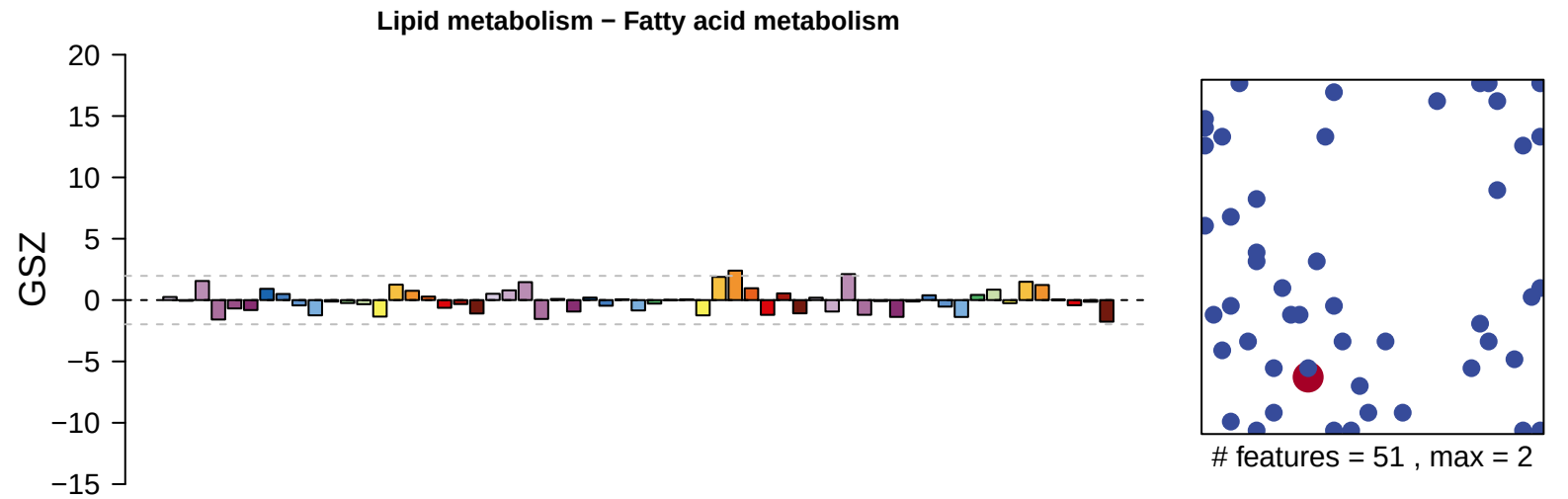
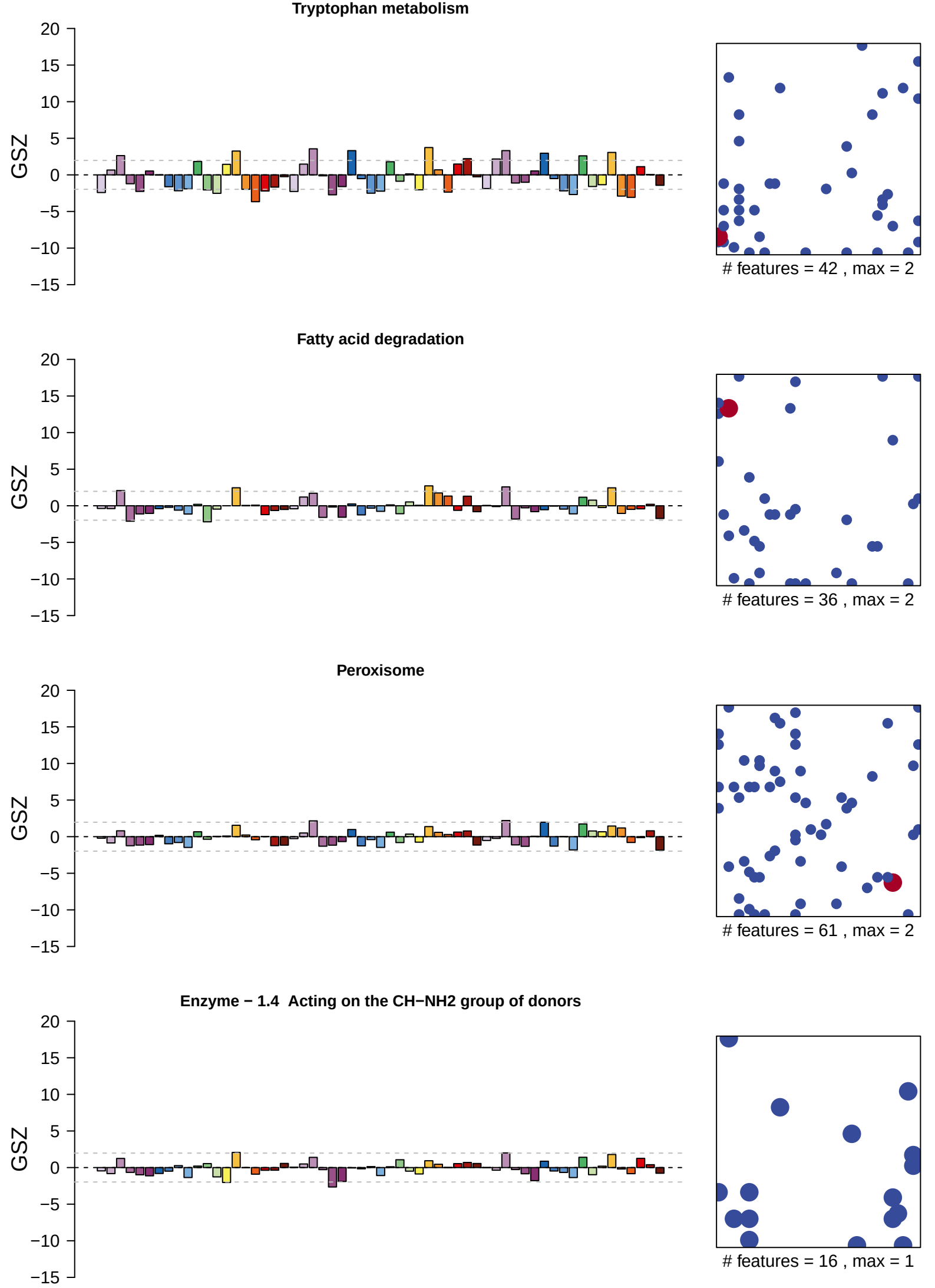

beta−Alanine metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 39 , max = 1

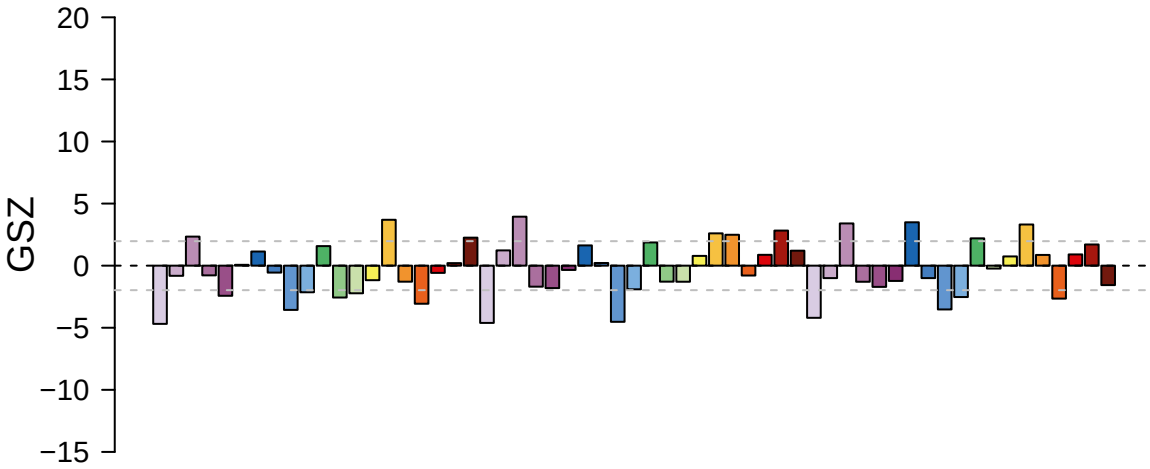

Vitamin B6 metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 11 , max = 1

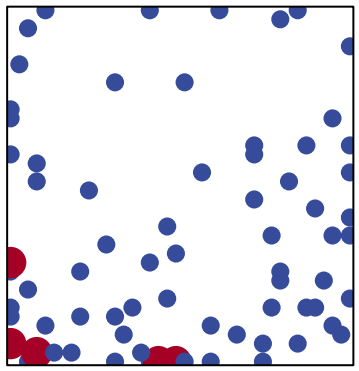

Enzyme − 6.2 Forming carbon−sulfur bonds
GSZ
20
15
10
5
0
−5
−10
−15
features = 28 , max = 1


Lipid metabolism − Fatty acid metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 51 , max = 2


Tryptophan metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 42 , max = 2
Fatty acid degradation
GSZ
20
15
10
5
0
−5
−10
−15
features = 36 , max = 2
Peroxisome
GSZ
20
15
10
5
0
−5
−10
−15
features = 61 , max = 2
Enzyme − 1.4 Acting on the CH−NH2 group of donors
GSZ
20
15
10
5
0
−5
−10
−15
features = 16 , max = 1

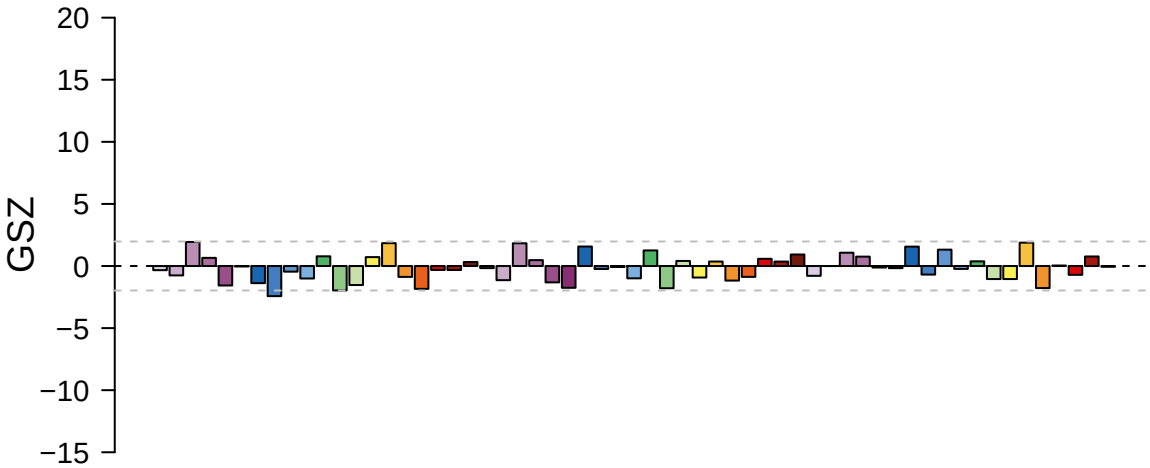
Pores ion channels [TC:1]

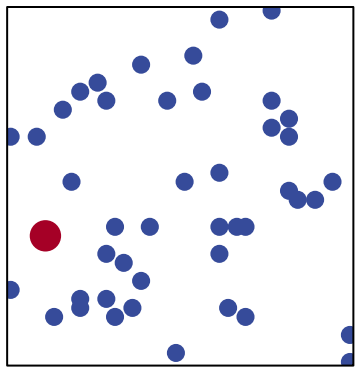




features = 77 , max = 2

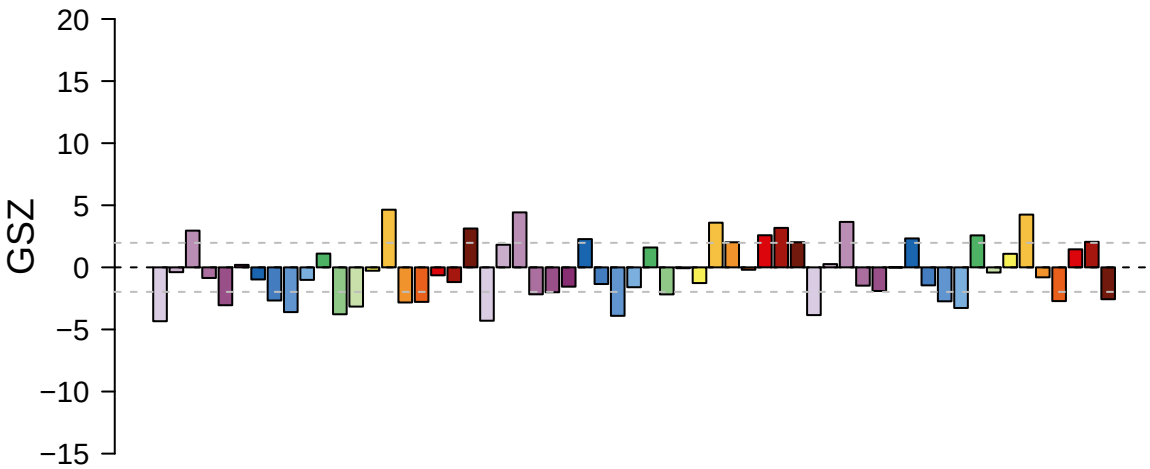
Folding sorting degradation − Protein export

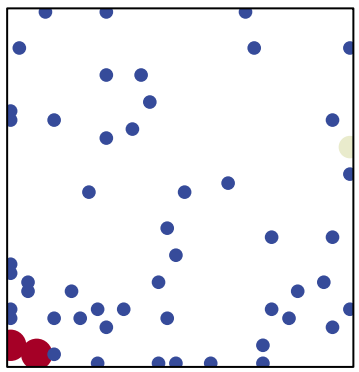




features = 46 , max = 2

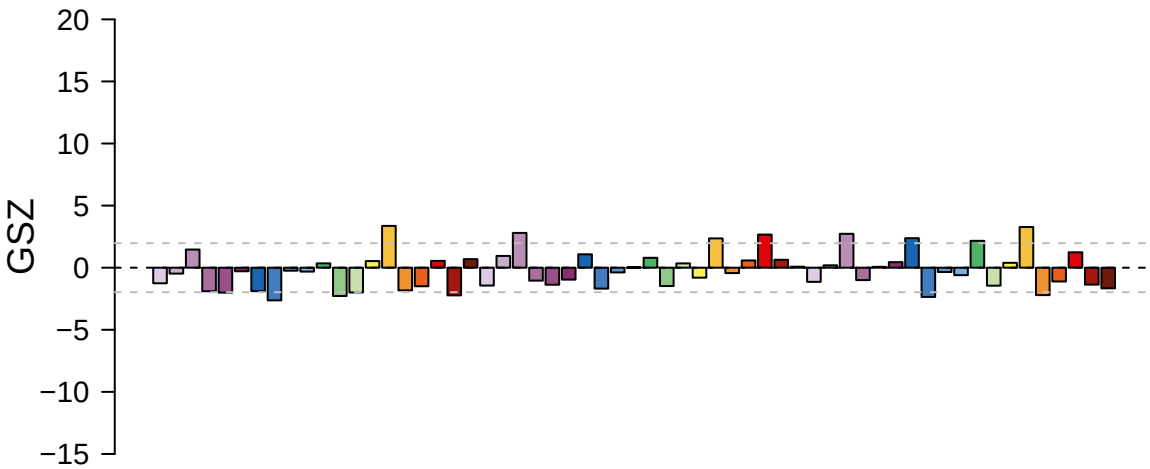
Other amino acids metabolism − Glutathione metabolism

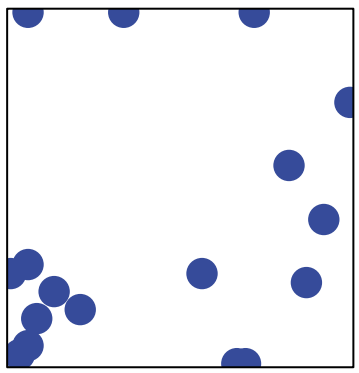




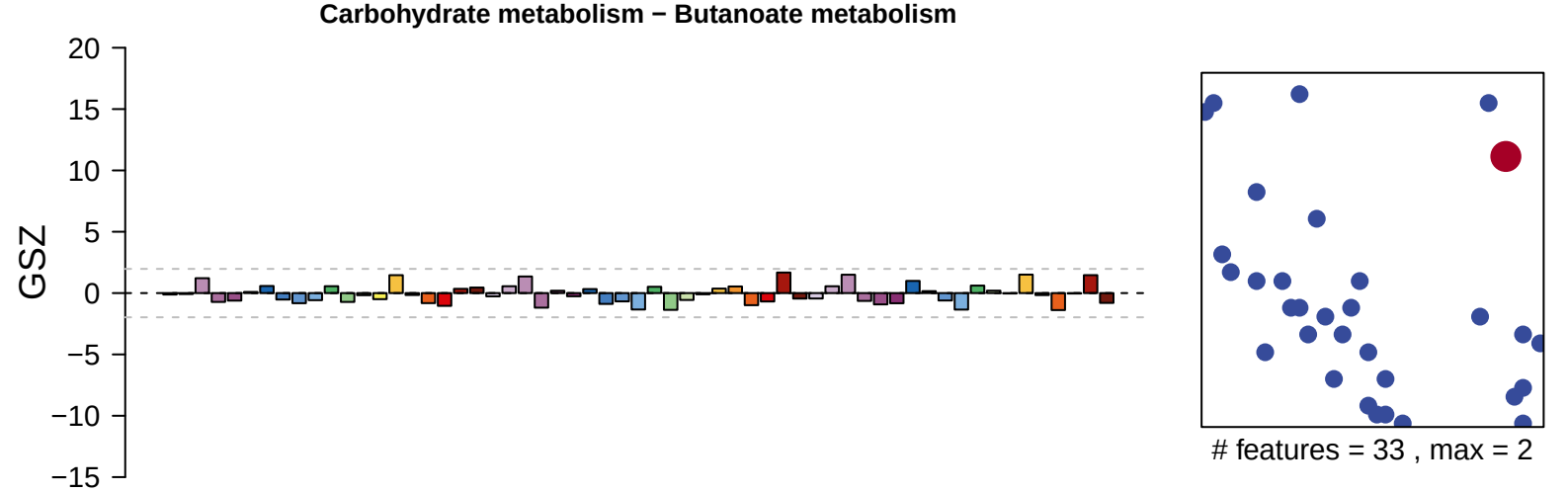
features = 58 , max = 3

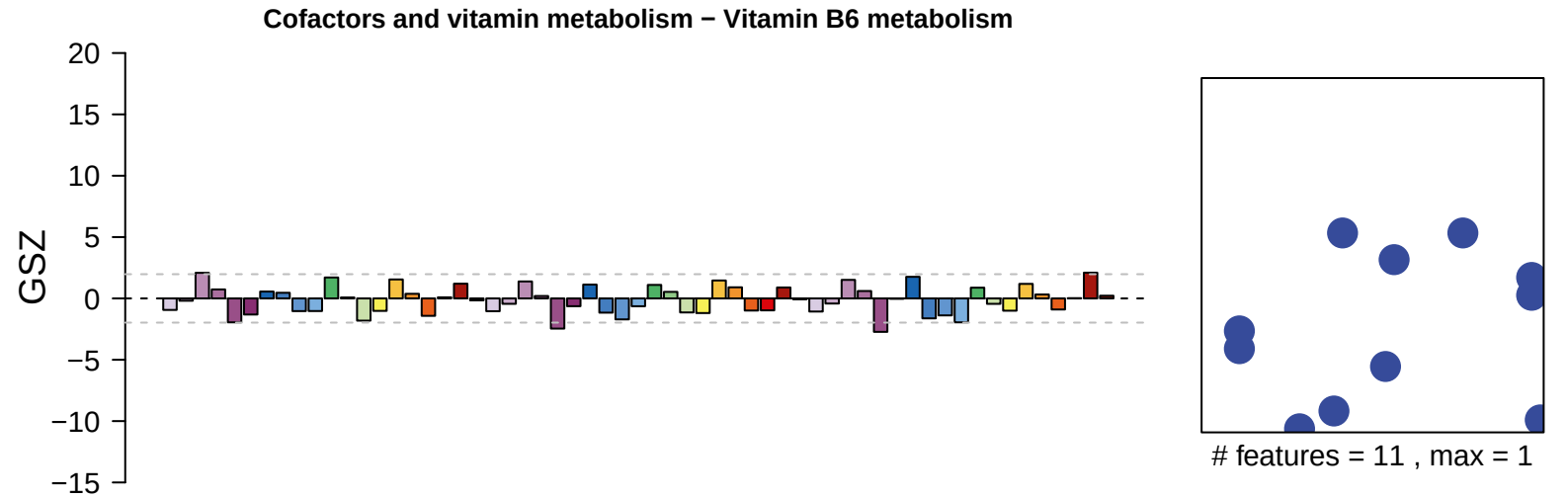
Isoquinoline alkaloid biosynthesis

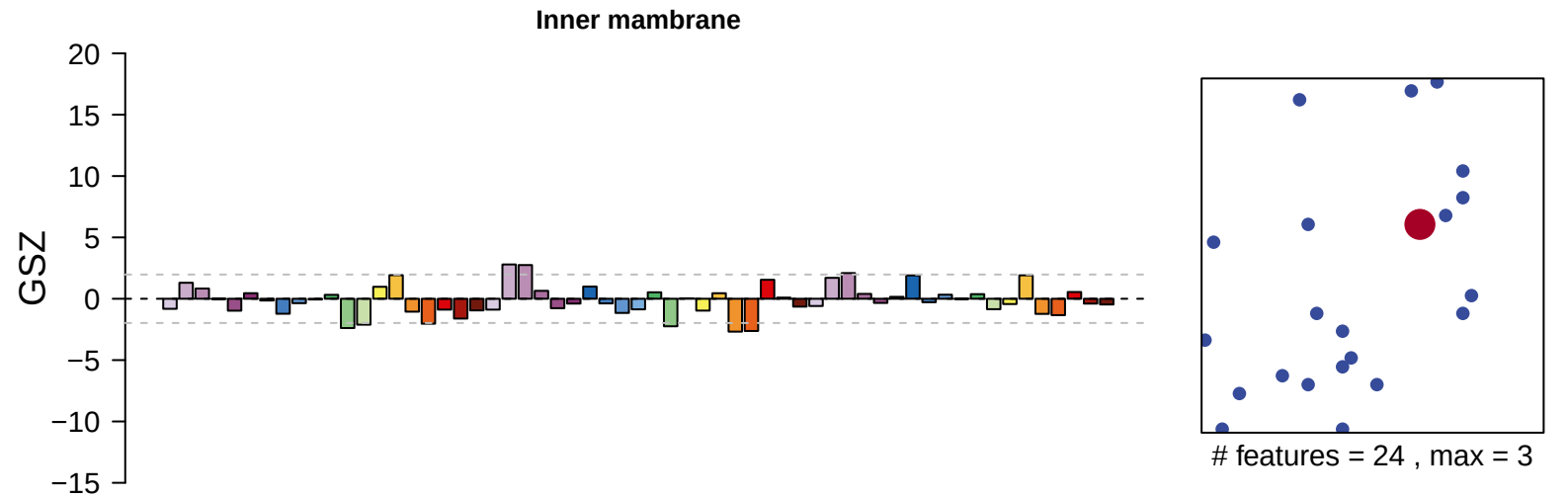


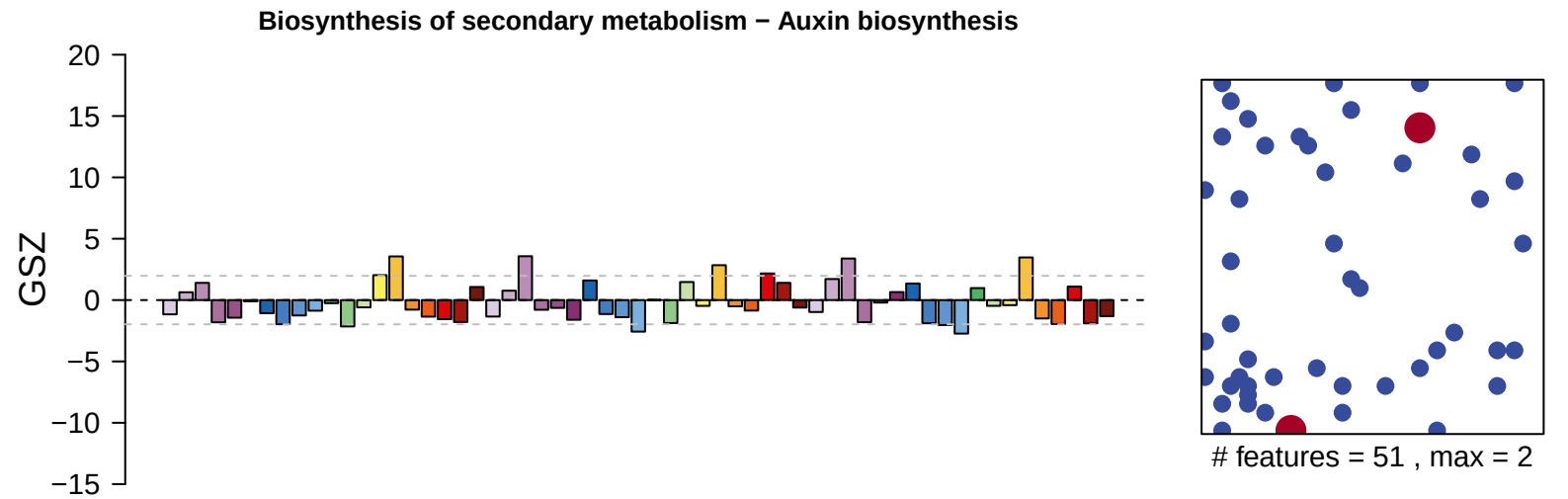


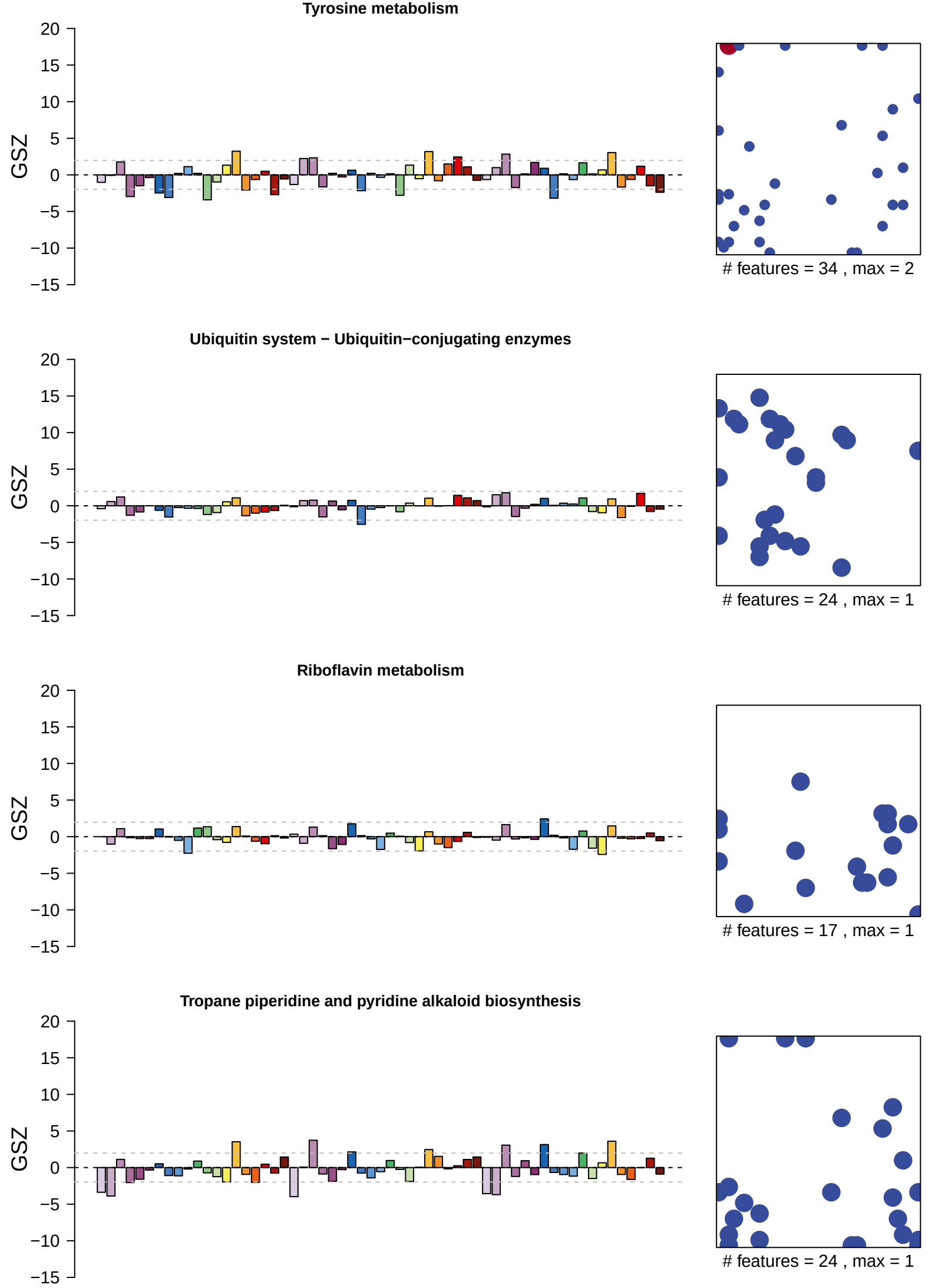
features = 17 , max = 1


Carbohydrate metabolism − Butanoate metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 33 , max = 2


Cofactors and vitamin metabolism − Vitamin B6 metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 11 , max = 1


Inner mambrane
GSZ
20
15
10
5
0
−5
−10
−15
features = 24 , max = 3


Biosynthesis of secondary metabolism − Auxin biosynthesis
GSZ
20
15
10
5
0
−5
−10
−15
features = 51 , max = 2


Tyrosine metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 34 , max = 2
Ubiquitin system − Ubiquitin−conjugating enzymes
GSZ
20
15
10
5
0
−5
−10
−15
features = 24 , max = 1
Riboflavin metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 17 , max = 1
Tropane piperidine and pyridine alkaloid biosynthesis
GSZ
20
15
10
5
0
−5
−10
−15
features = 24 , max = 1