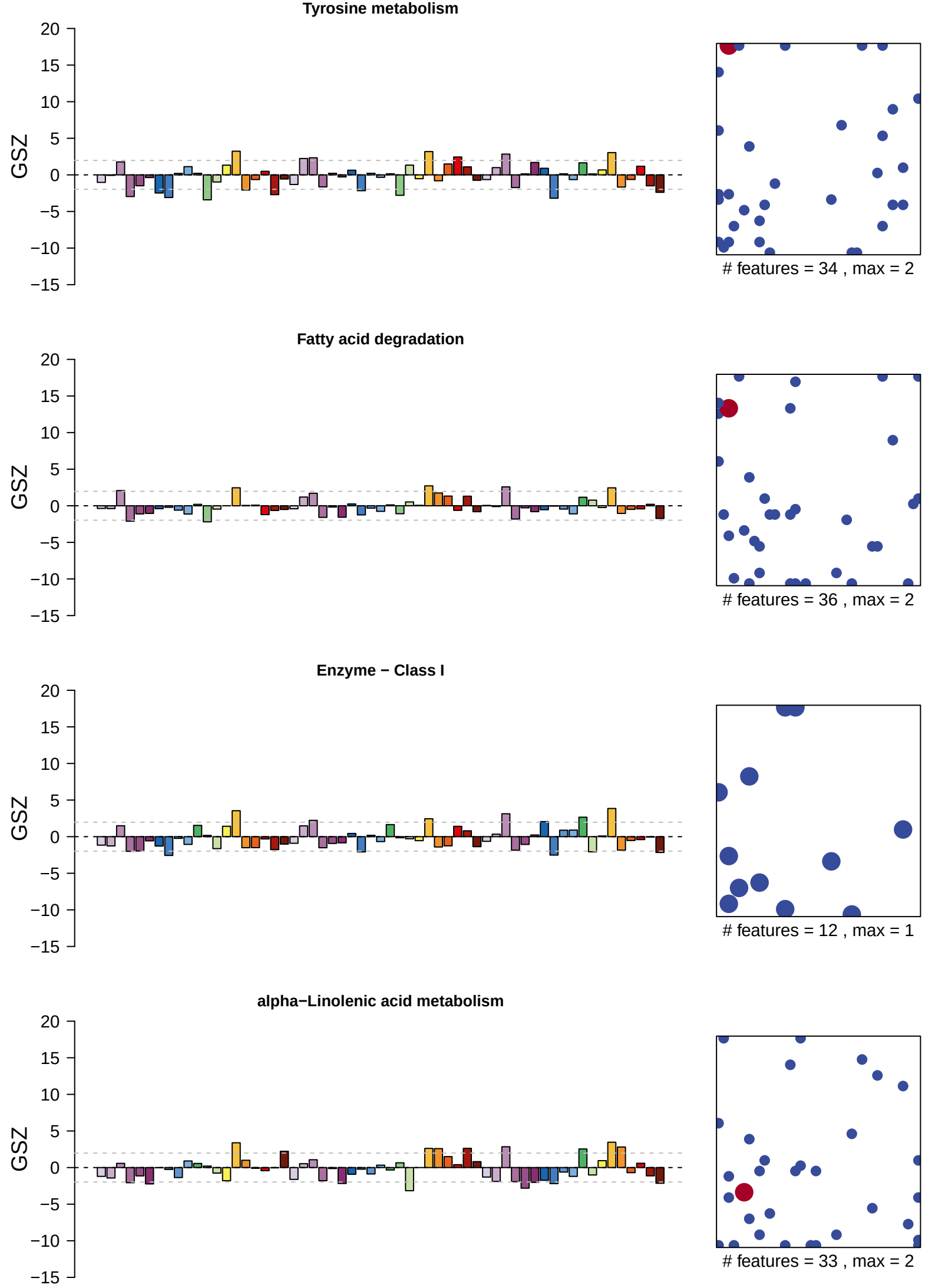
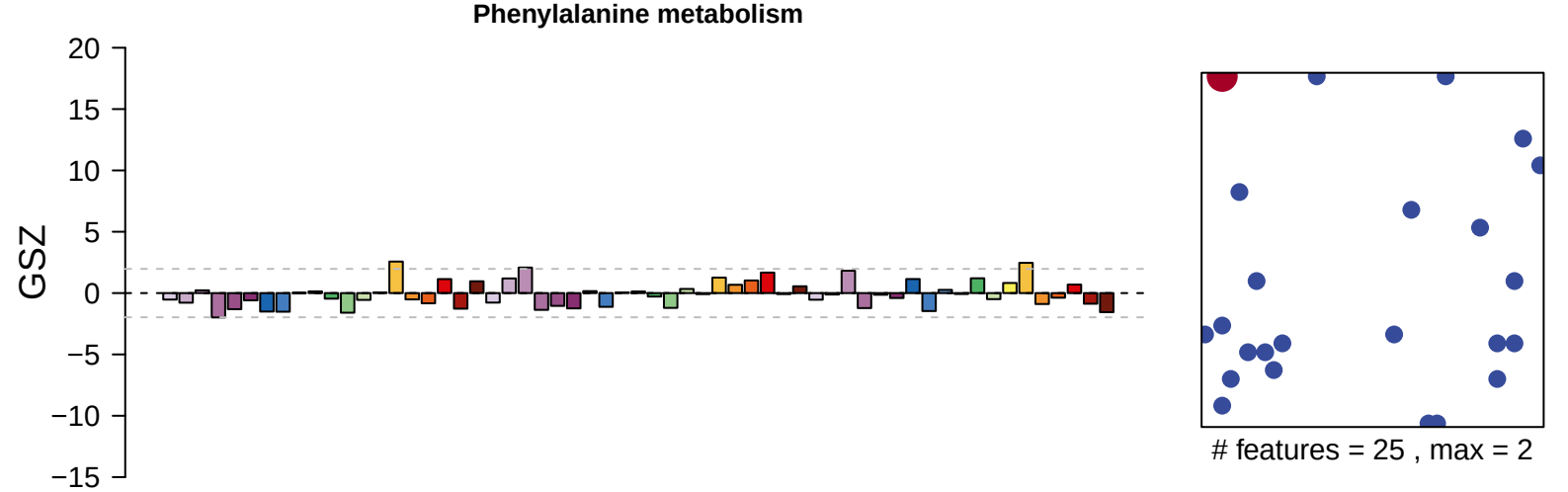
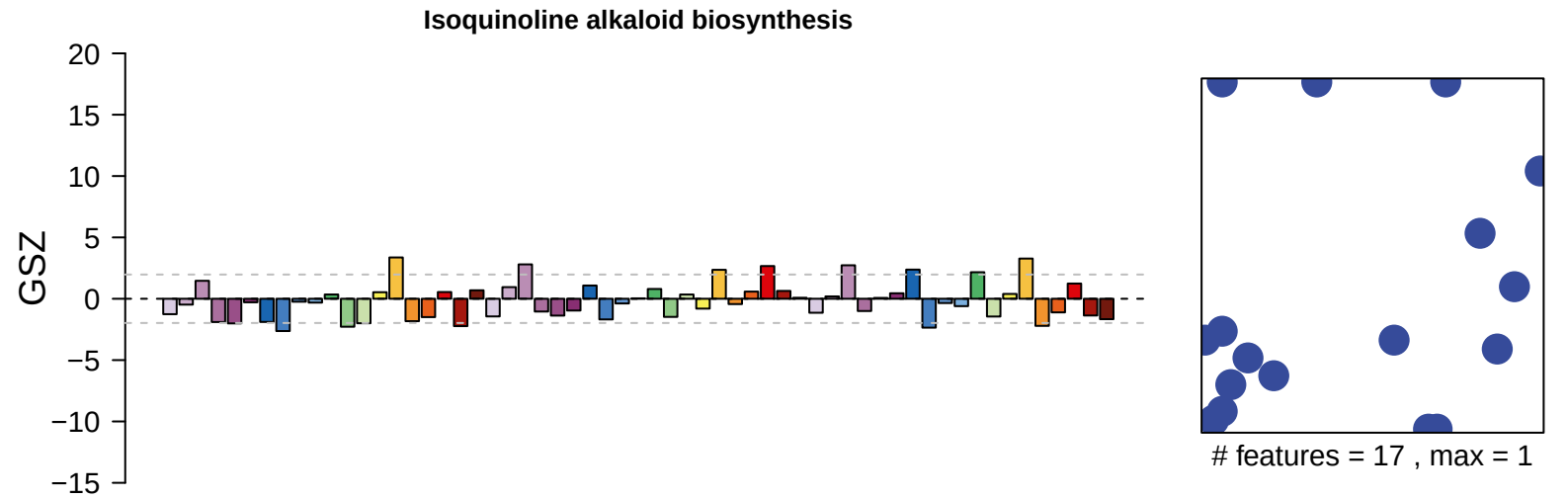
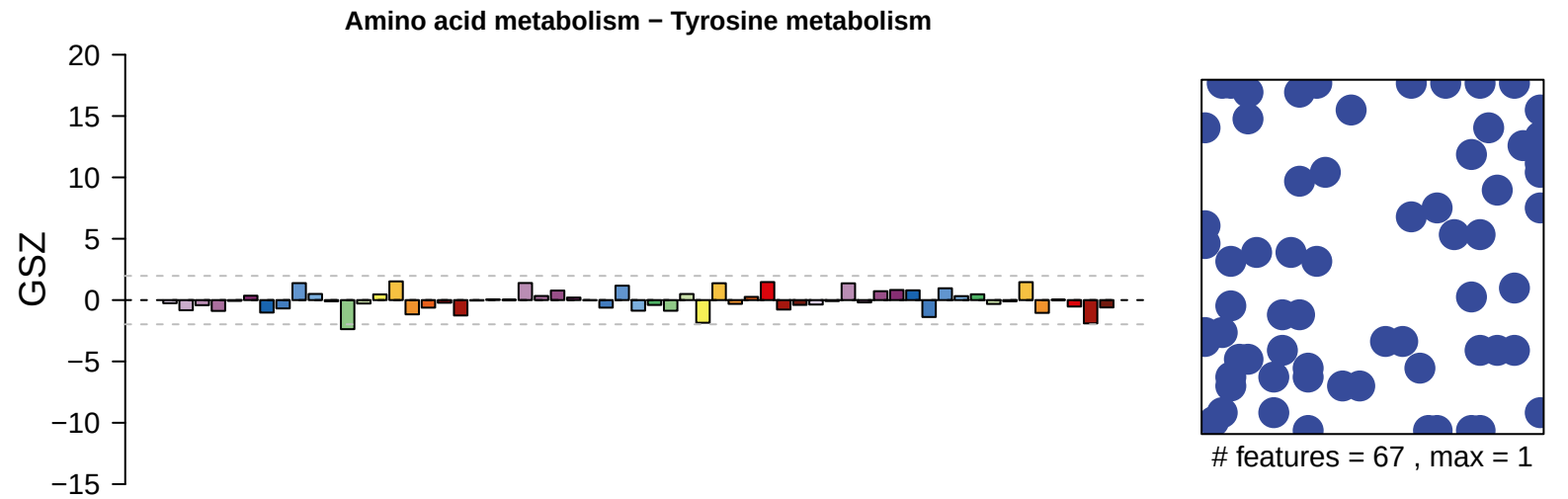
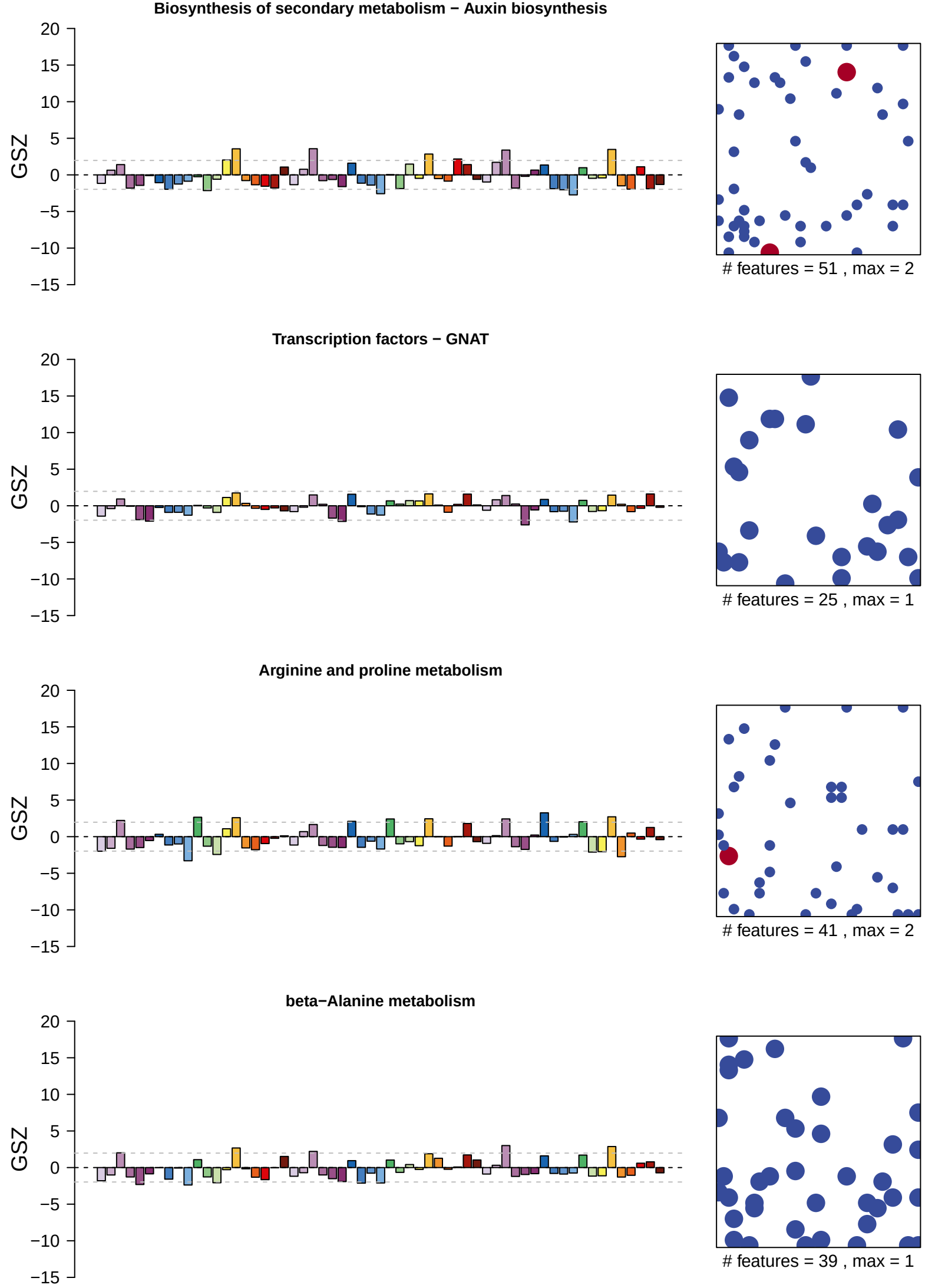

Tyrosine metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 34 , max = 2
Fatty acid degradation
GSZ
20
15
10
5
0
−5
−10
−15
features = 36 , max = 2
Enzyme − Class I
GSZ
20
15
10
5
0
−5
−10
−15
features = 12 , max = 1
alpha−Linolenic acid metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 33 , max = 2

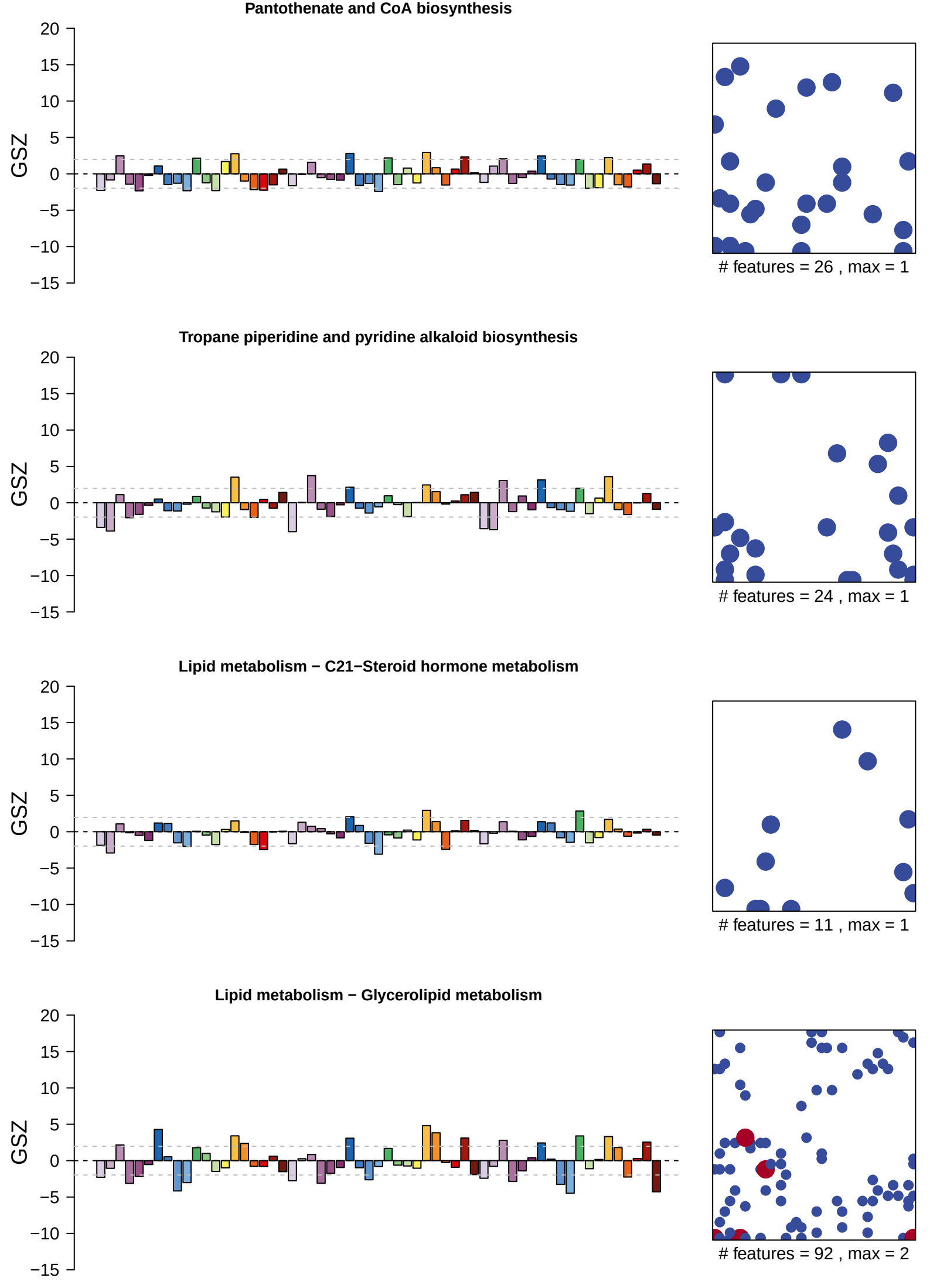

Phenylalanine metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 25 , max = 2

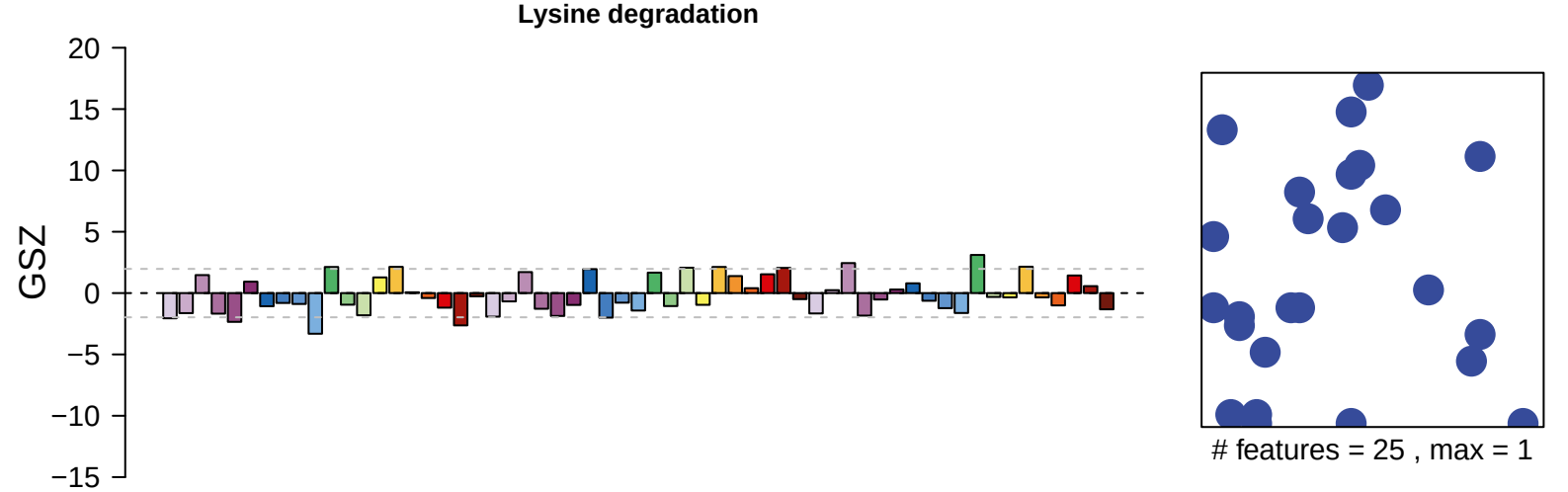

Other amino acids metabolism − Glutathione metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 58 , max = 3

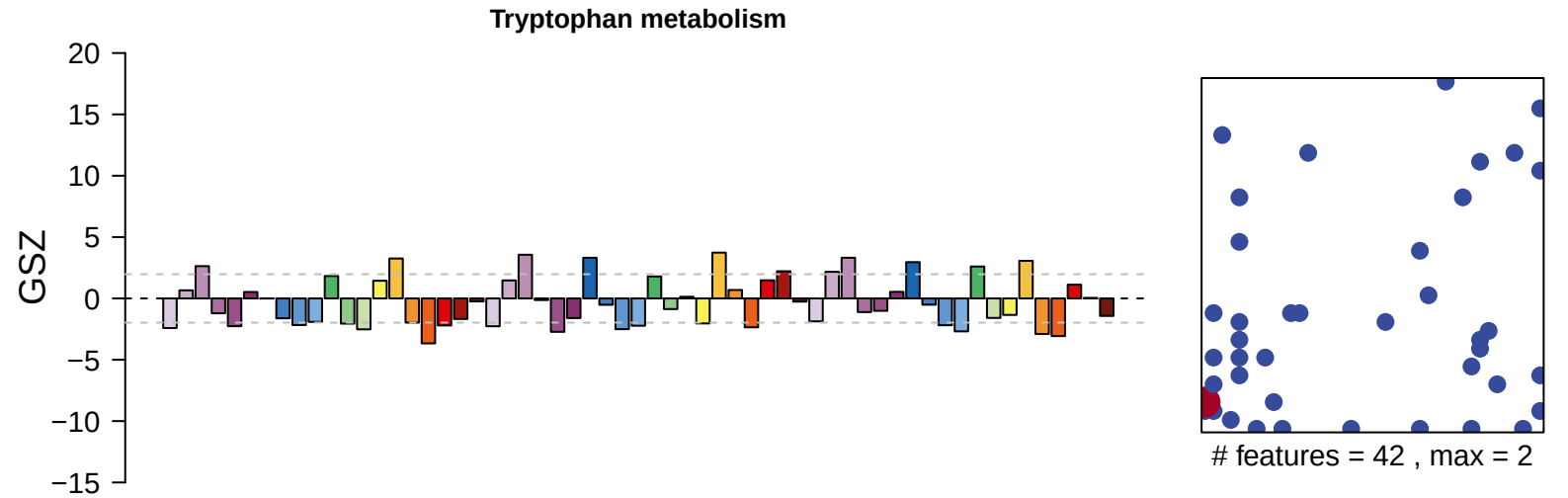

Isoquinoline alkaloid biosynthesis
GSZ
20
15
10
5
0
−5
−10
−15
features = 17 , max = 1

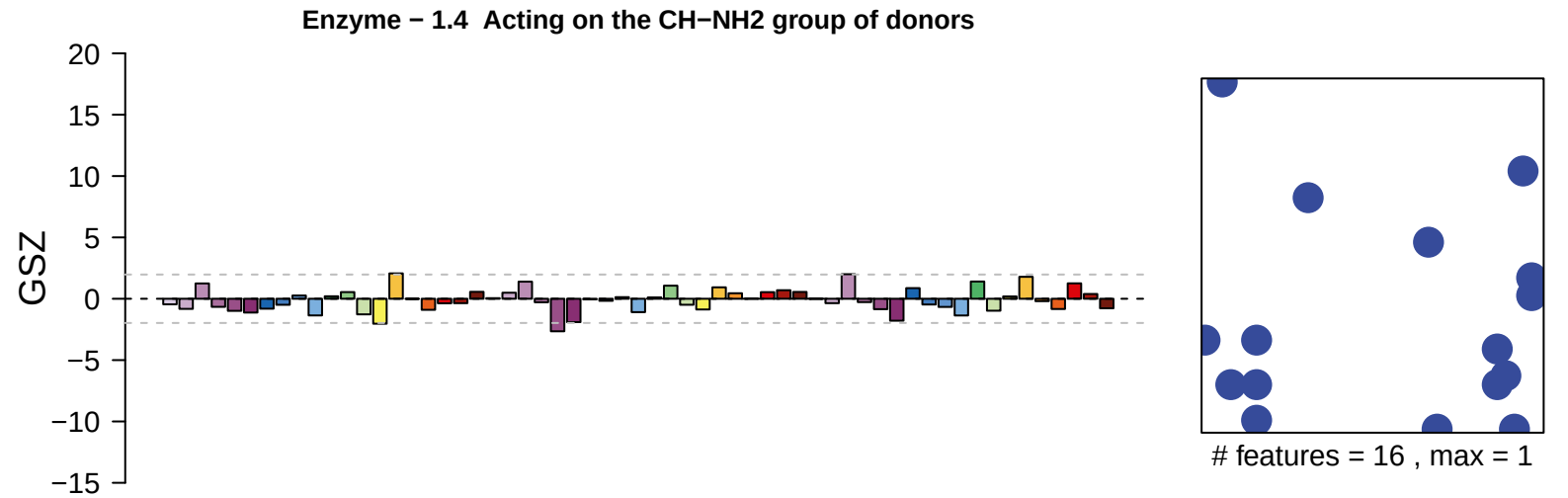

Amino acid metabolism − Tyrosine metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 67 , max = 1

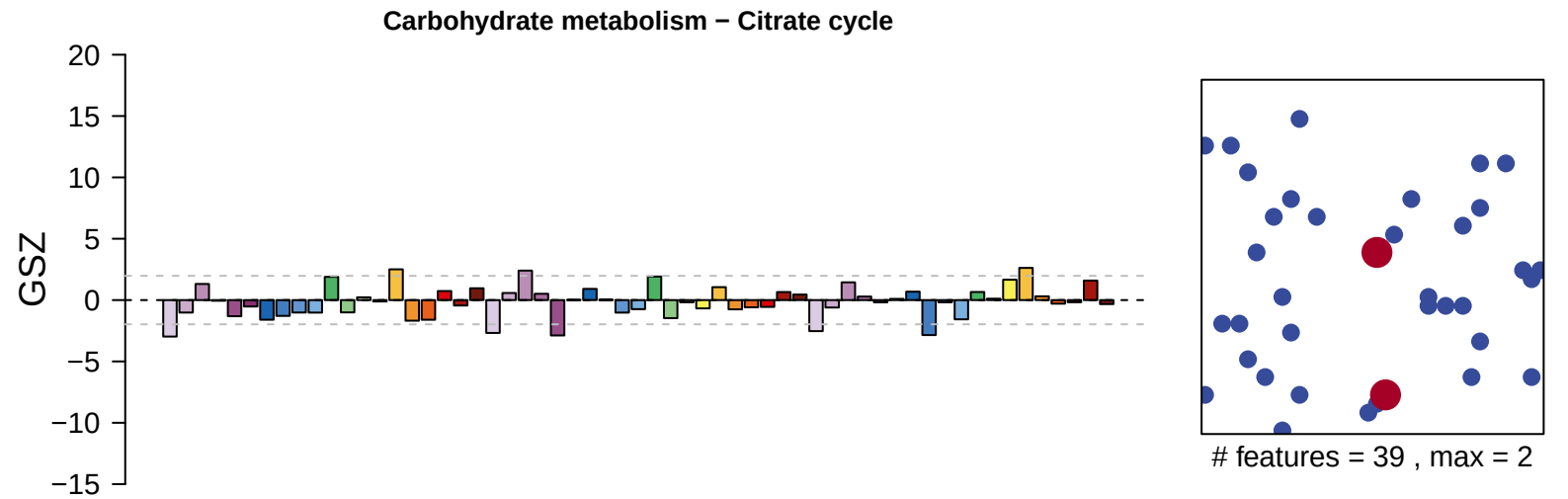

Biosynthesis of secondary metabolism − Auxin biosynthesis
GSZ
20
15
10
5
0
−5
−10
−15
features = 51 , max = 2
Transcription factors − GNAT
GSZ
20
15
10
5
0
−5
−10
−15
features = 25 , max = 1
Arginine and proline metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 41 , max = 2
beta−Alanine metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 39 , max = 1


Pantothenate and CoA biosynthesis
GSZ
20
15
10
5
0
−5
−10
−15
features = 26 , max = 1
Tropane piperidine and pyridine alkaloid biosynthesis
GSZ
20
15
10
5
0
−5
−10
−15
features = 24 , max = 1
Lipid metabolism − C21−Steroid hormone metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 11 , max = 1
Lipid metabolism − Glycerolipid metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 92 , max = 2


Lysine degradation
GSZ
20
15
10
5
0
−5
−10
−15
features = 25 , max = 1


Tryptophan metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 42 , 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


Carbohydrate metabolism − Citrate cycle
GSZ
20
15
10
5
0
−5
−10
−15
features = 39 , max = 2