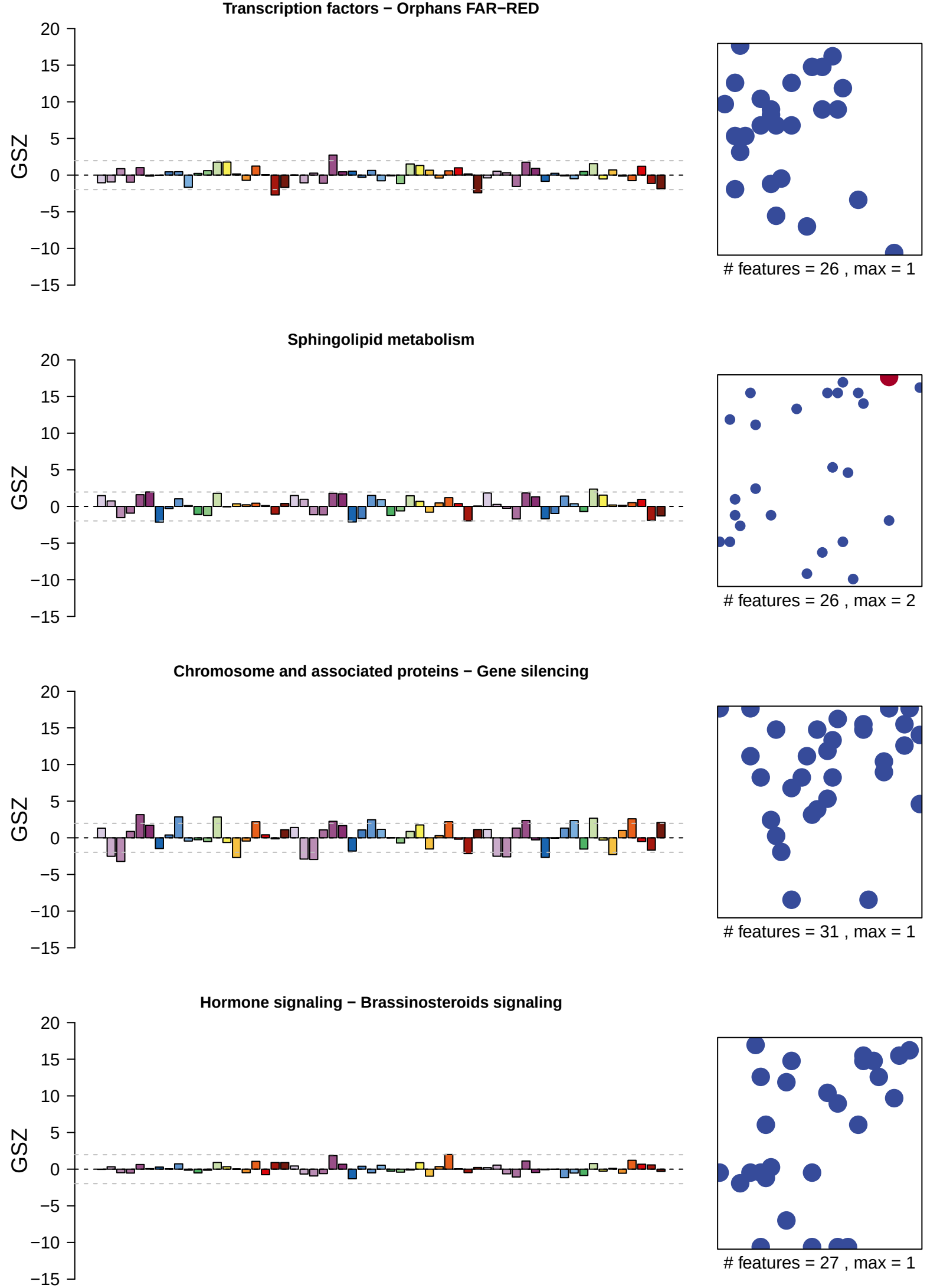
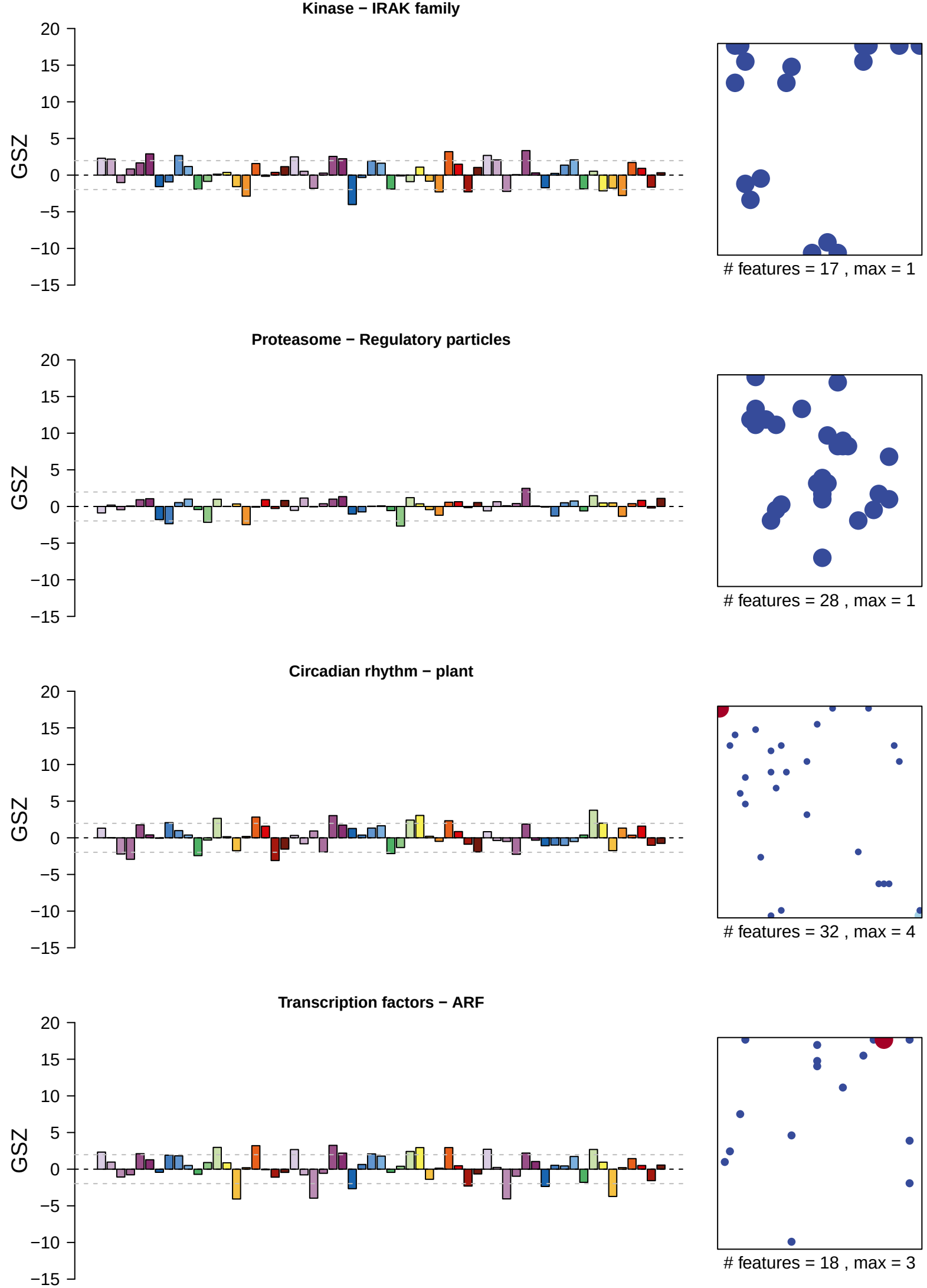
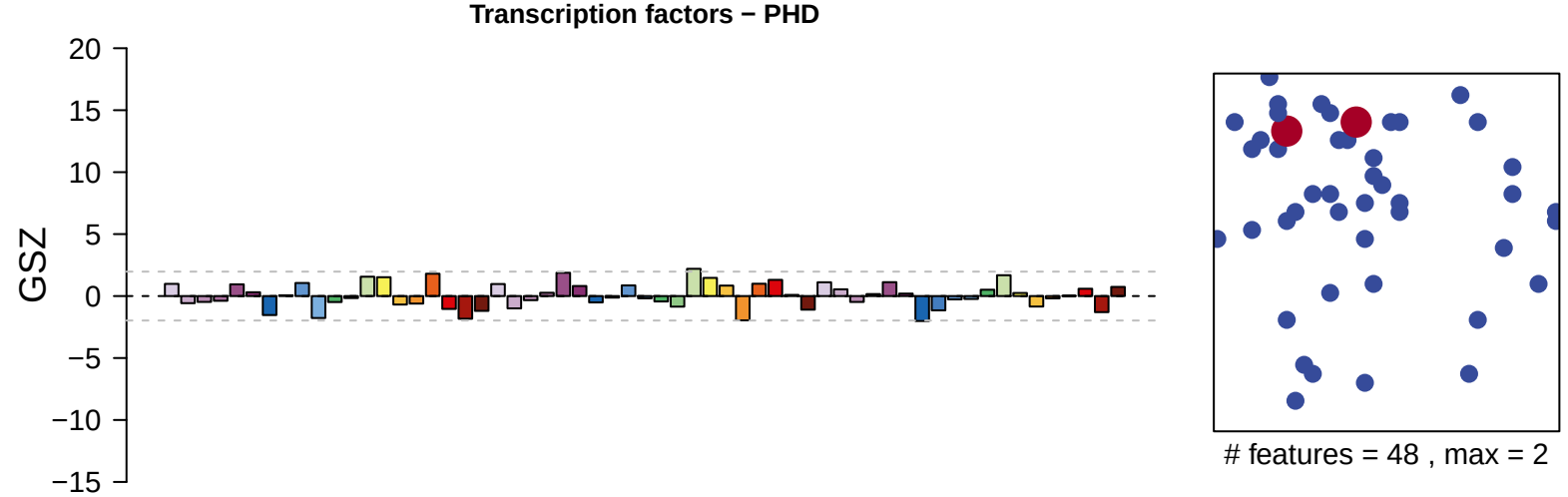
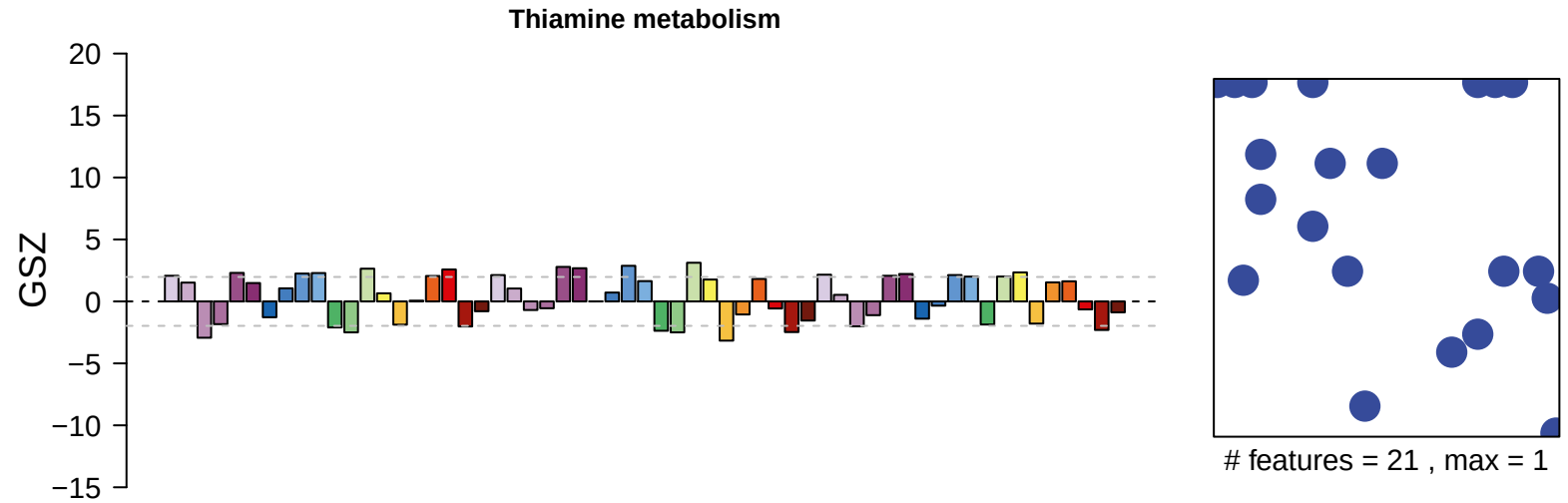

Transcription factors − Orphans FAR−RED
GSZ
20
15
10
5
0
−5
−10
−15
features = 26 , max = 1
Sphingolipid metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 26 , max = 2
Chromosome and associated proteins − Gene silencing
GSZ
20
15
10
5
0
−5
−10
−15
features = 31 , max = 1
Hormone signaling − Brassinosteroids signaling
GSZ
20
15
10
5
0
−5
−10
−15
features = 27 , max = 1

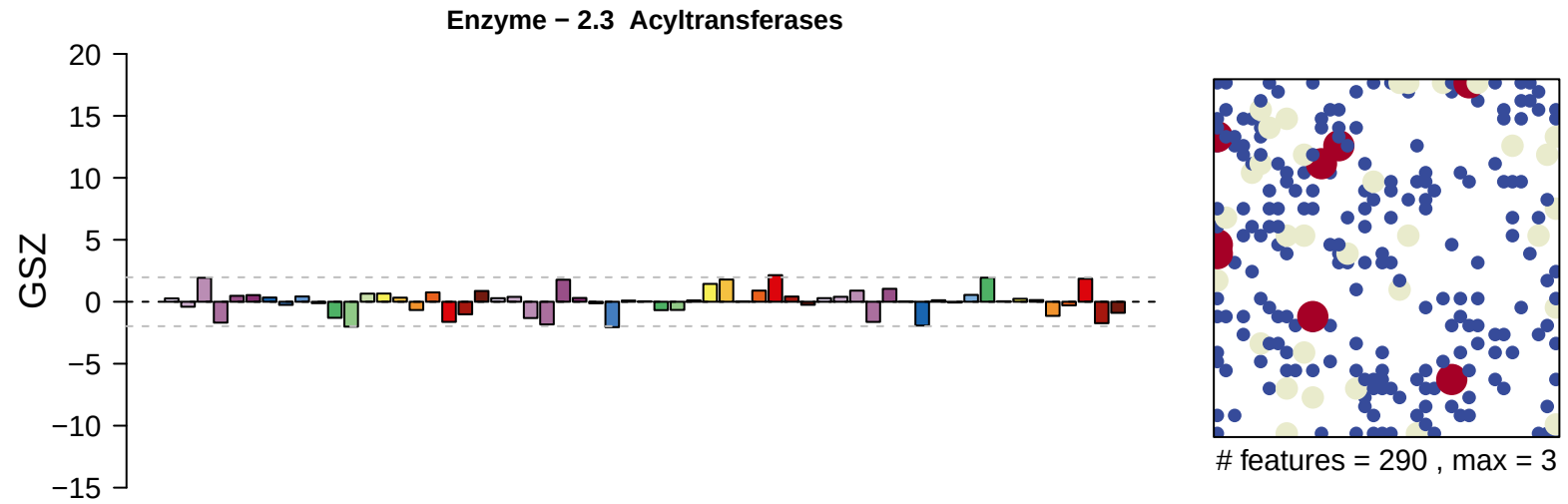

Kinase − IRAK family
GSZ
20
15
10
5
0
−5
−10
−15
features = 17 , max = 1
Proteasome − Regulatory particles
GSZ
20
15
10
5
0
−5
−10
−15
features = 28 , max = 1
Circadian rhythm − plant
GSZ
20
15
10
5
0
−5
−10
−15
features = 32 , max = 4
Transcription factors − ARF
GSZ
20
15
10
5
0
−5
−10
−15
features = 18 , max = 3



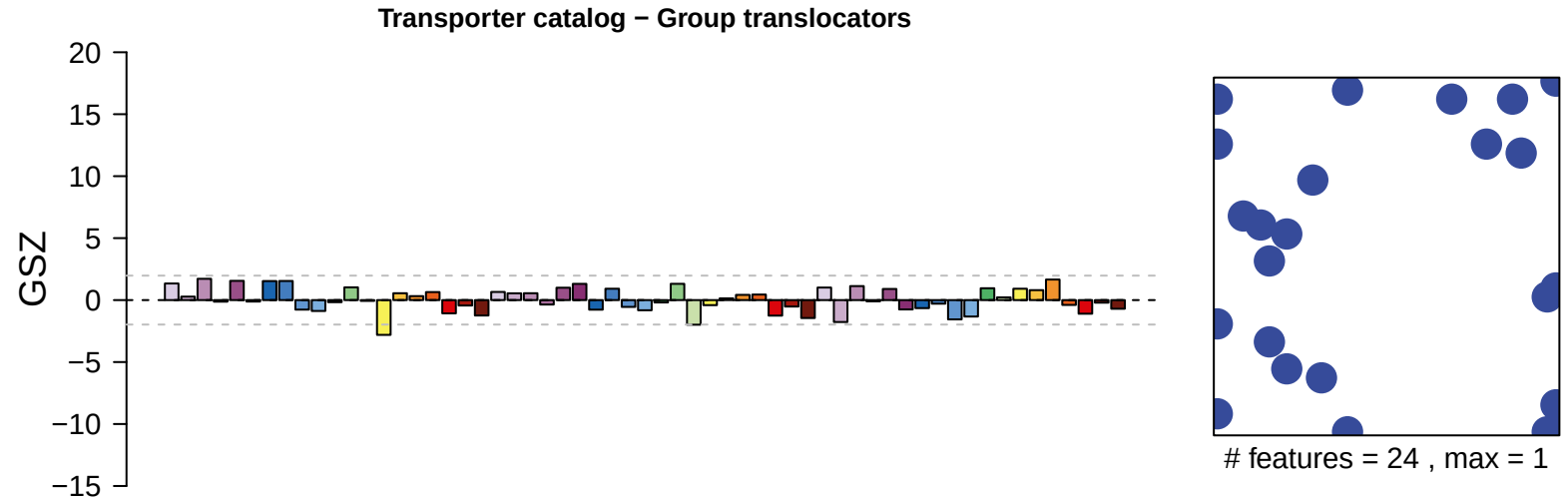
features = 48 , max = 2



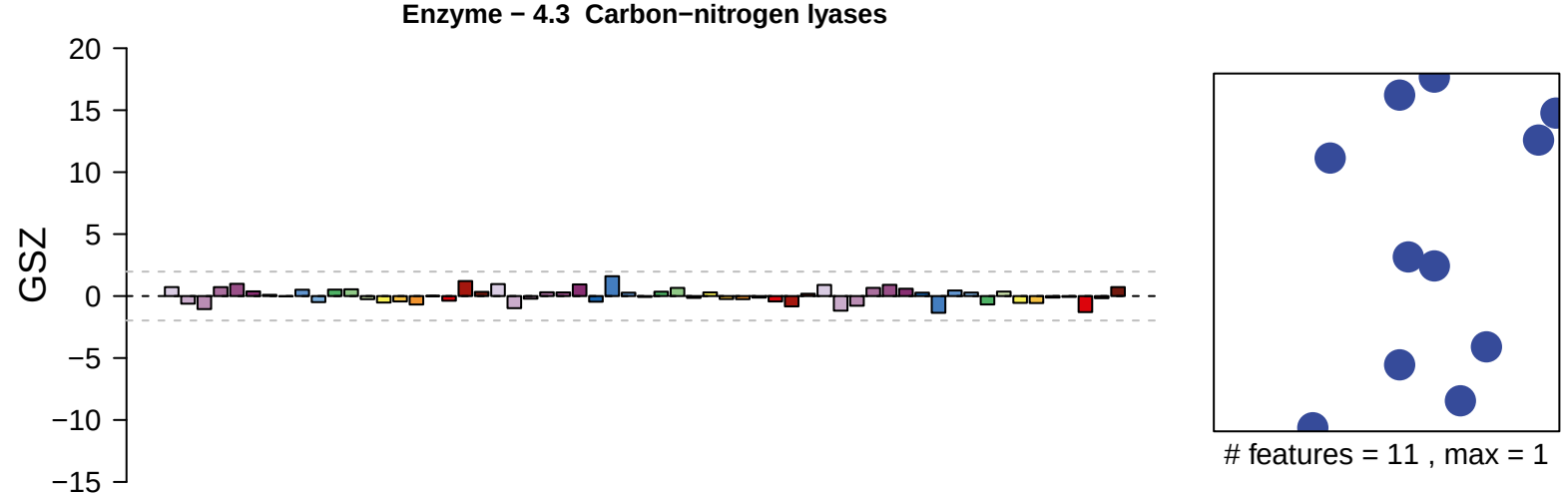
features = 21 , max = 1

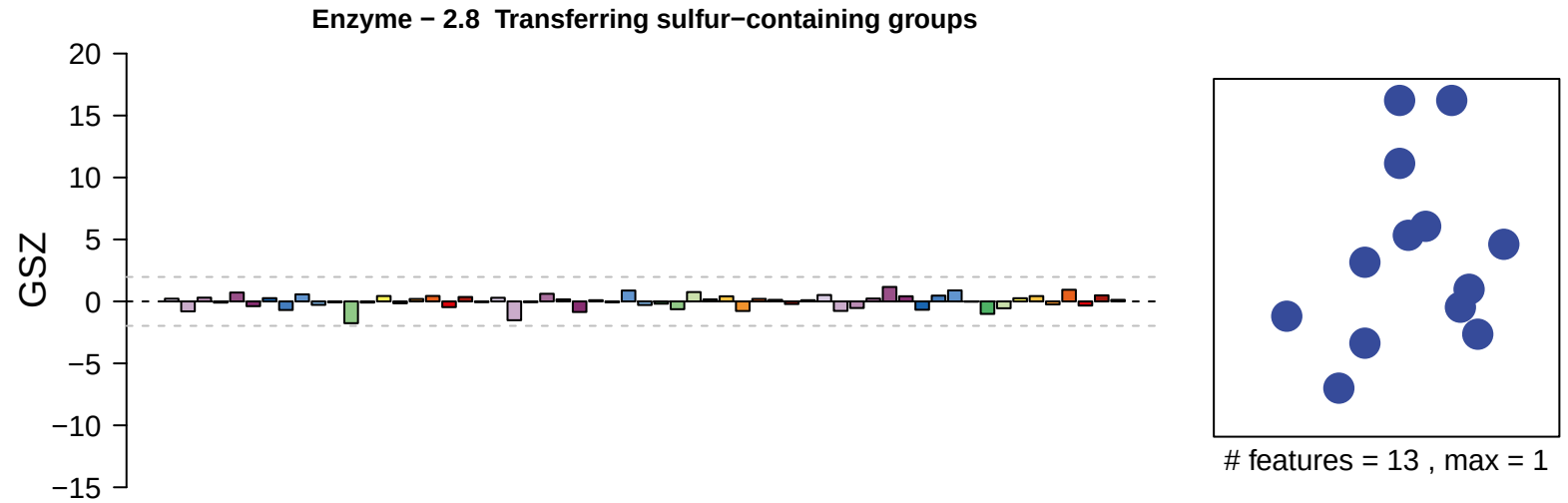


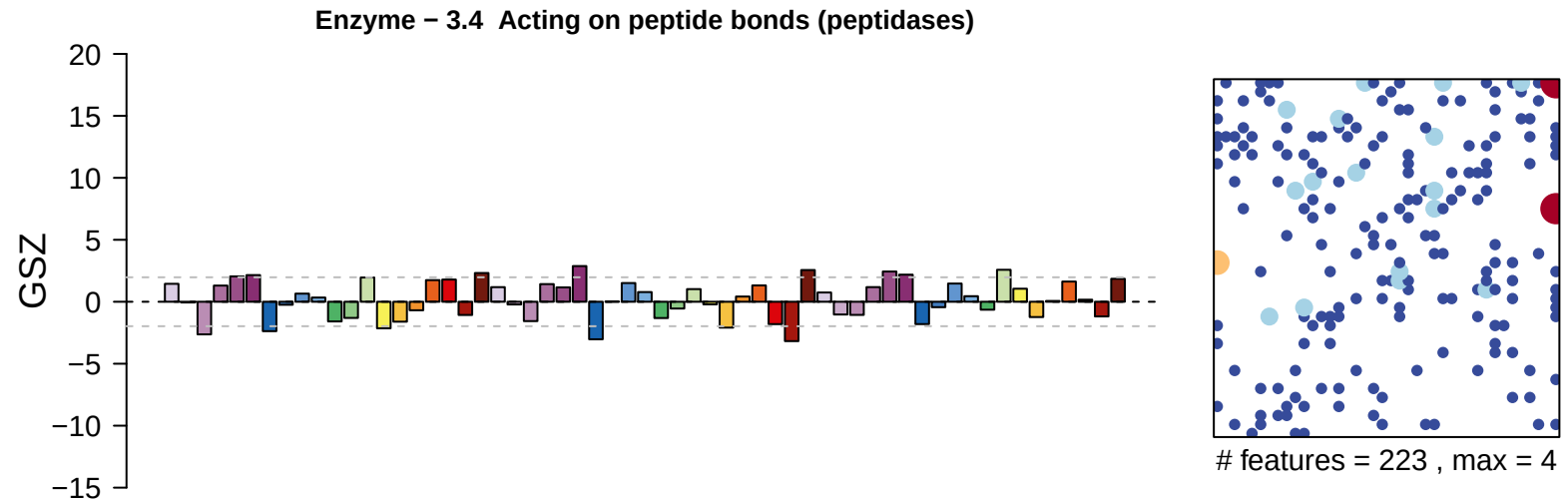
features = 290 , max = 3

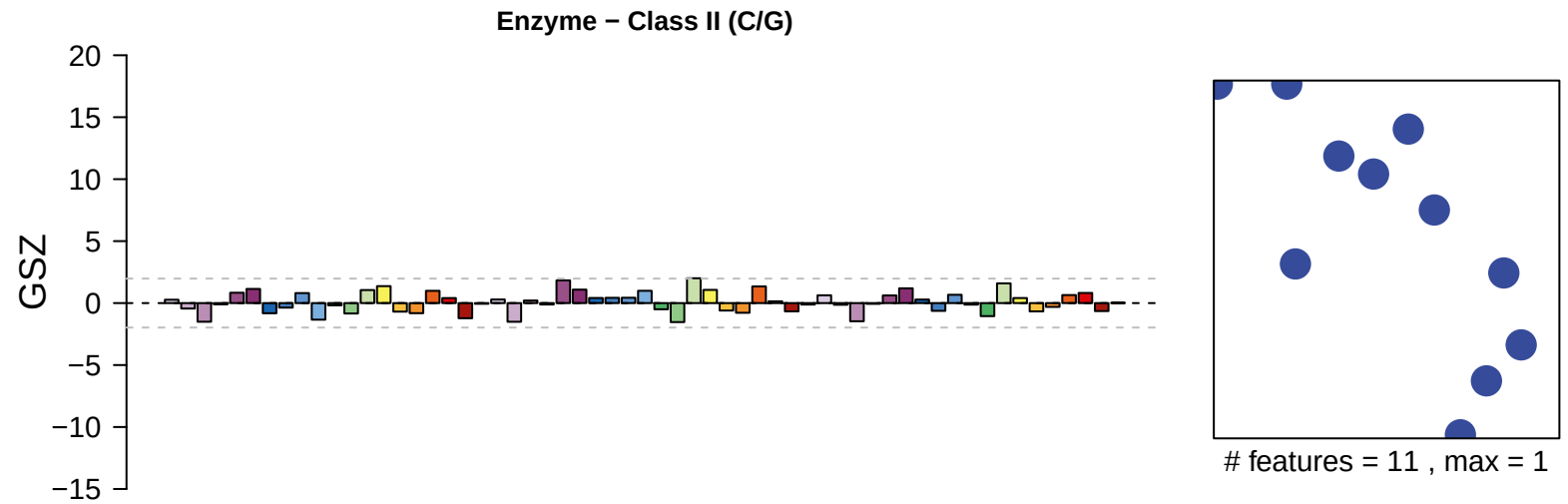


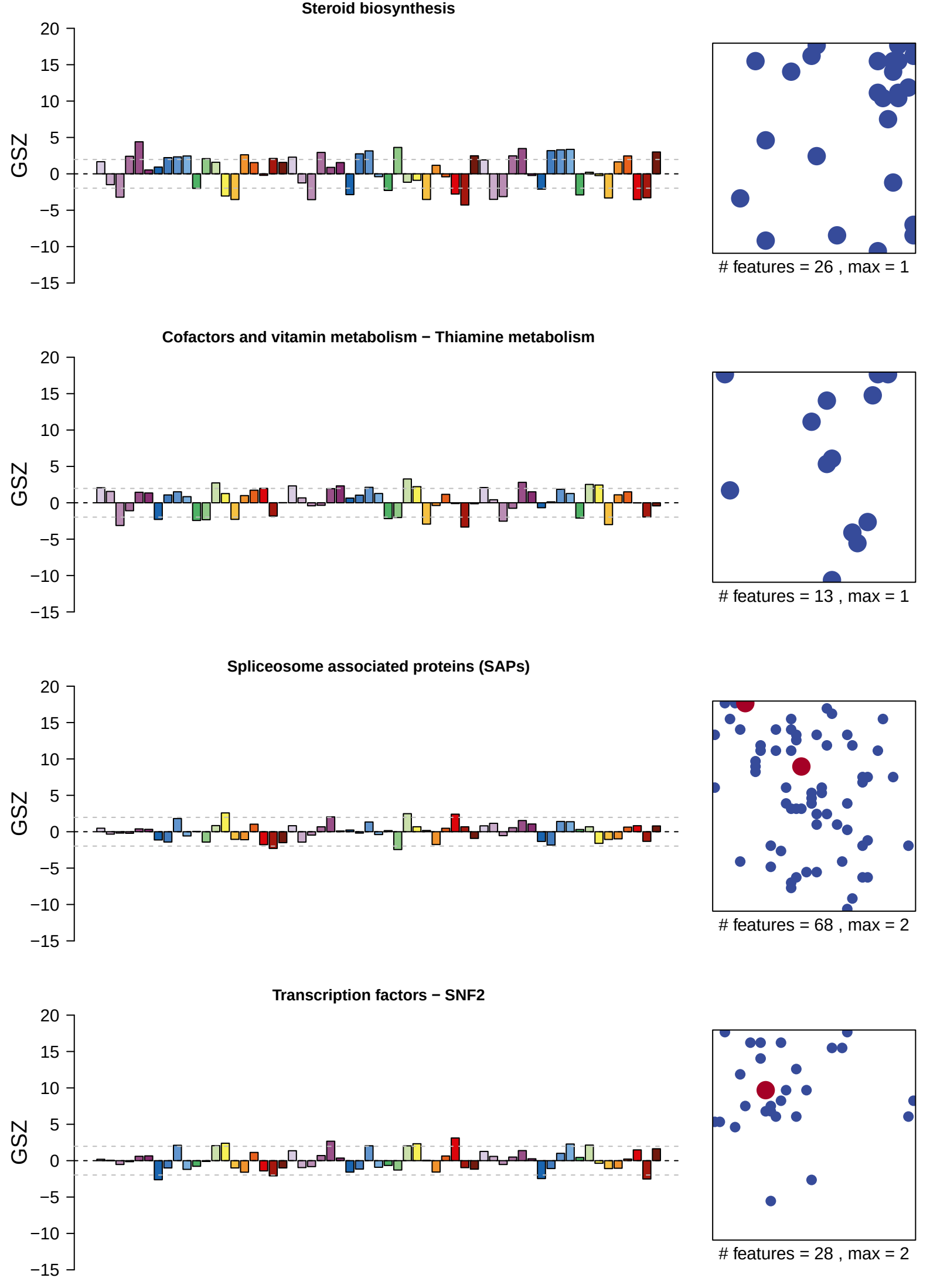
features = 24 , max = 1


Enzyme − 4.3 Carbon−nitrogen lyases
GSZ
20
15
10
5
0
−5
−10
−15
features = 11 , max = 1


Enzyme − 2.8 Transferring sulfur−containing groups
GSZ
20
15
10
5
0
−5
−10
−15
features = 13 , max = 1


Enzyme − 3.4 Acting on peptide bonds (peptidases)
GSZ
20
15
10
5
0
−5
−10
−15
features = 223 , max = 4


Enzyme − Class II (C/G)
GSZ
20
15
10
5
0
−5
−10
−15
features = 11 , max = 1


Steroid biosynthesis
GSZ
20
15
10
5
0
−5
−10
−15
features = 26 , max = 1
Cofactors and vitamin metabolism − Thiamine metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 13 , max = 1
Spliceosome associated proteins (SAPs)
GSZ
20
15
10
5
0
−5
−10
−15
features = 68 , max = 2
Transcription factors − SNF2
GSZ
20
15
10
5
0
−5
−10
−15
features = 28 , max = 2