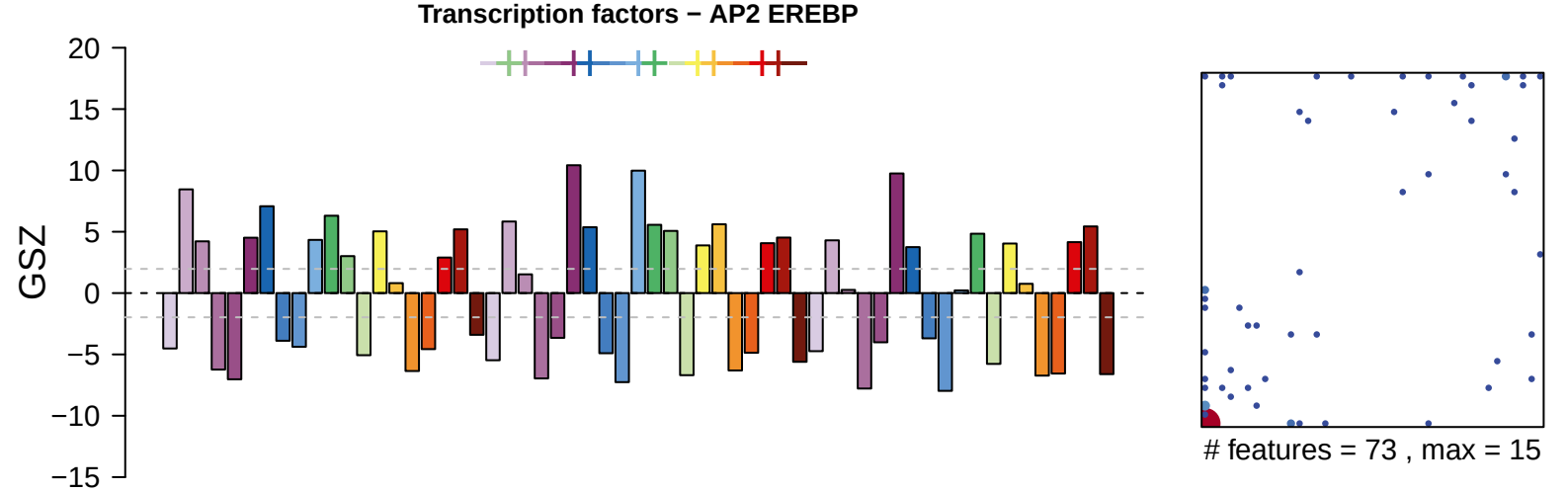
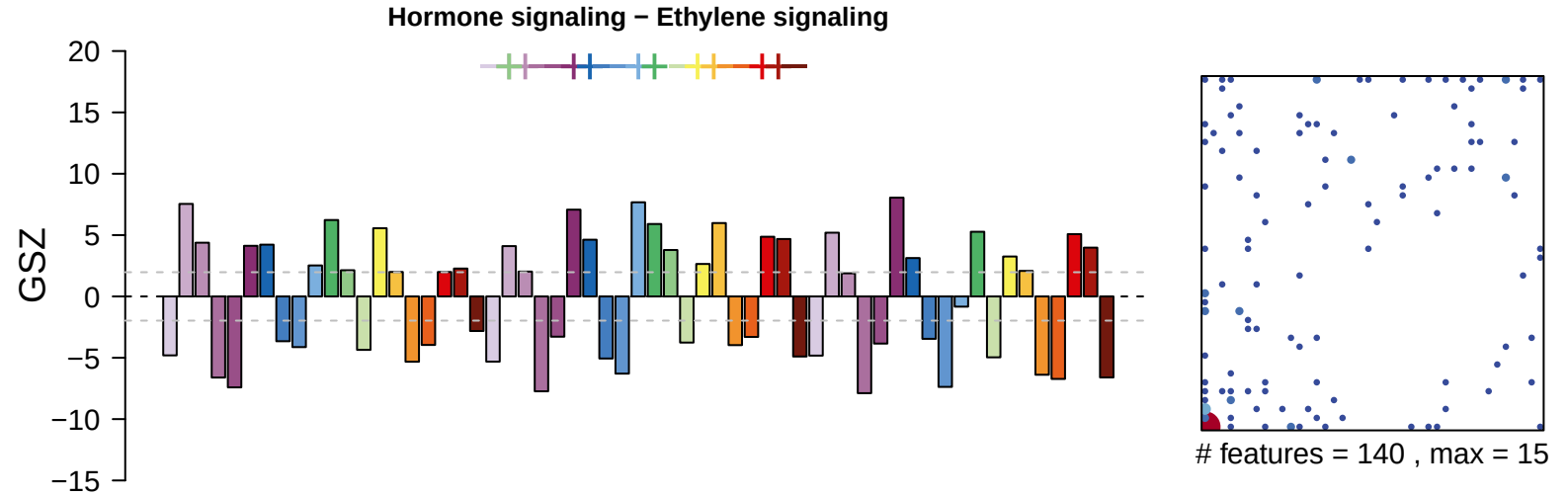
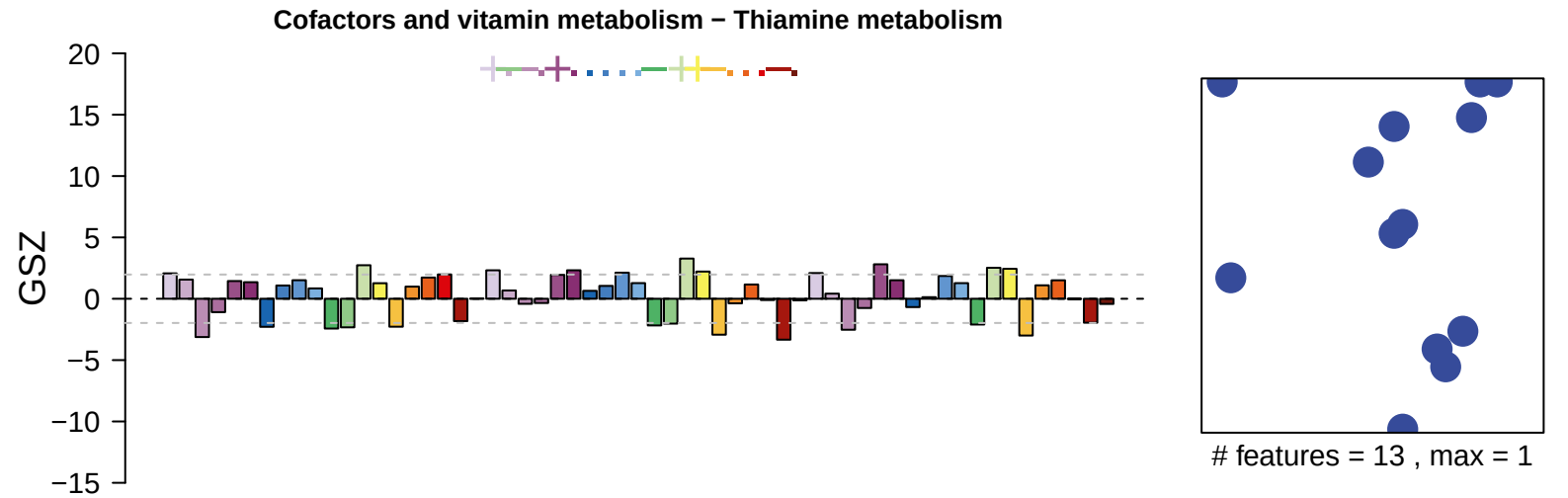
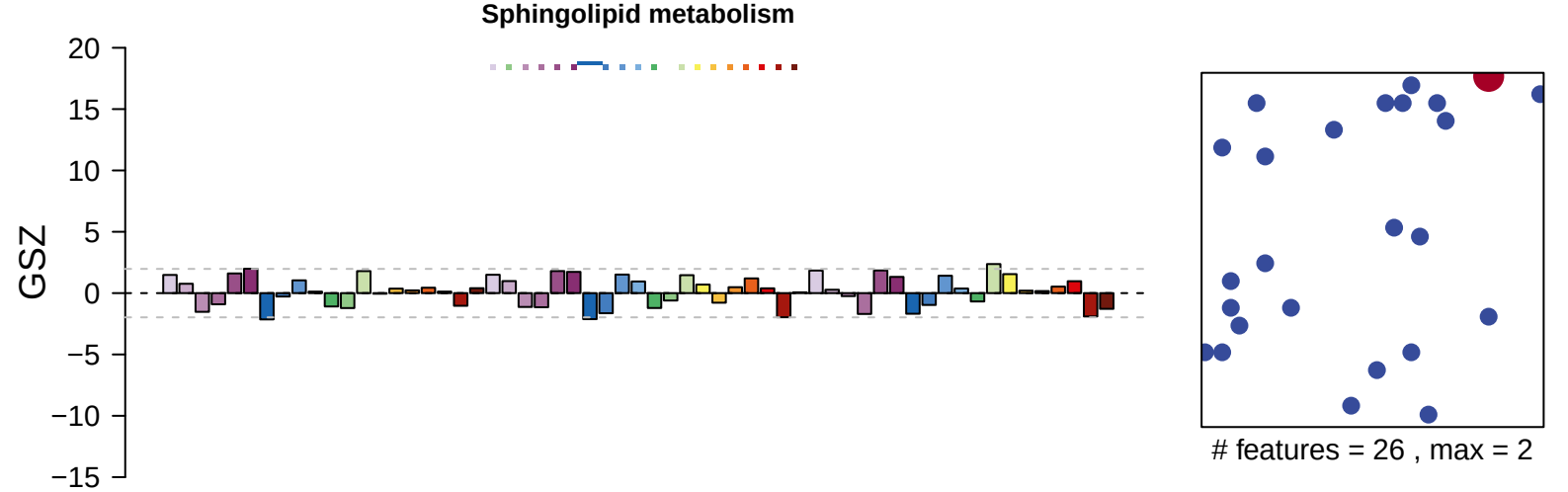
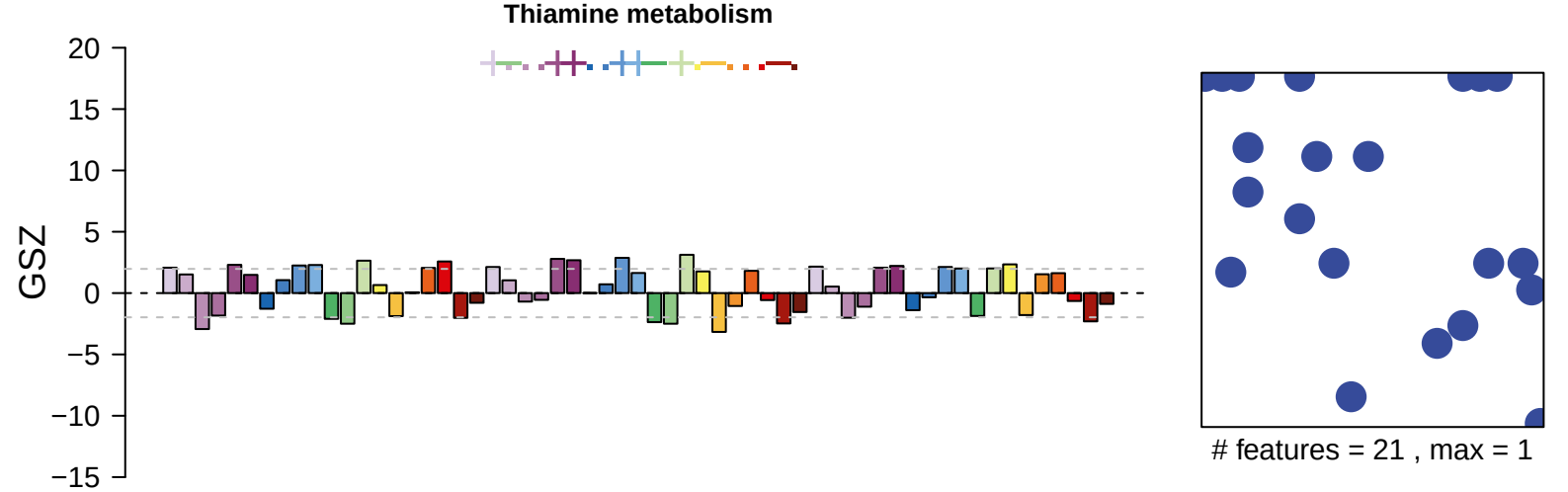

Transcription factors − AP2 EREBP
GSZ
20
15
10
5
0
−5
−10
−15
features = 73 , max = 15

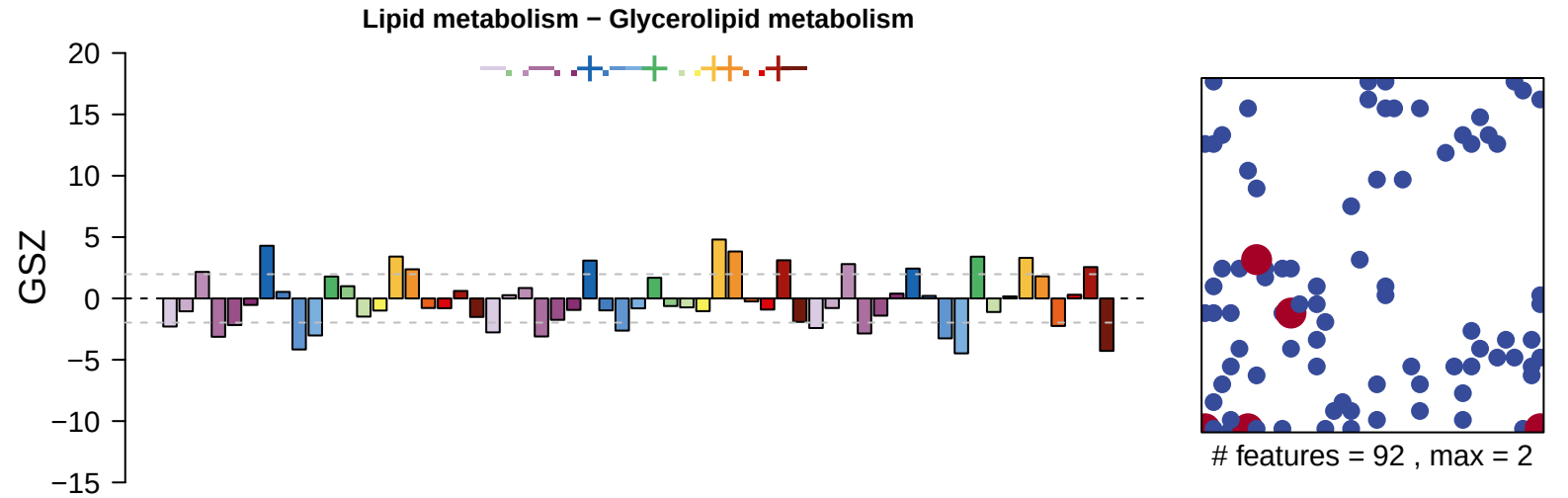

Hormone signaling − Ethylene signaling
GSZ
20
15
10
5
0
−5
−10
−15
features = 140 , max = 15


Cofactors and vitamin metabolism − Thiamine metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 13 , max = 1

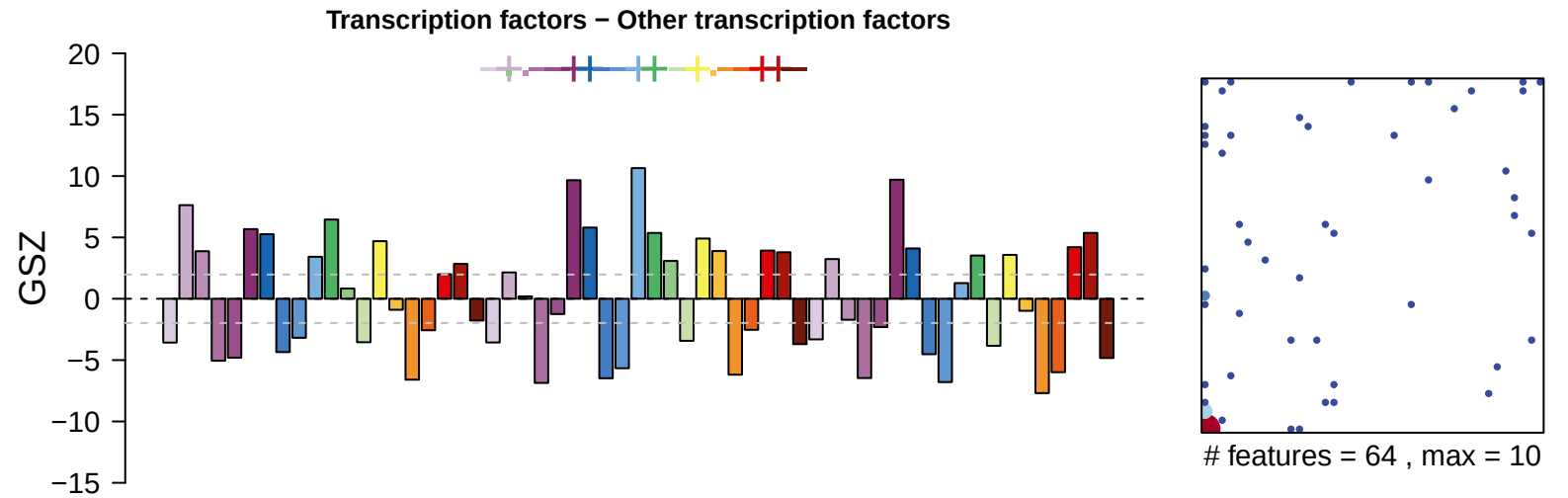

Sphingolipid metabolism
GSZ
20
15
10
5
0
−5
−10
−15
features = 26 , max = 2



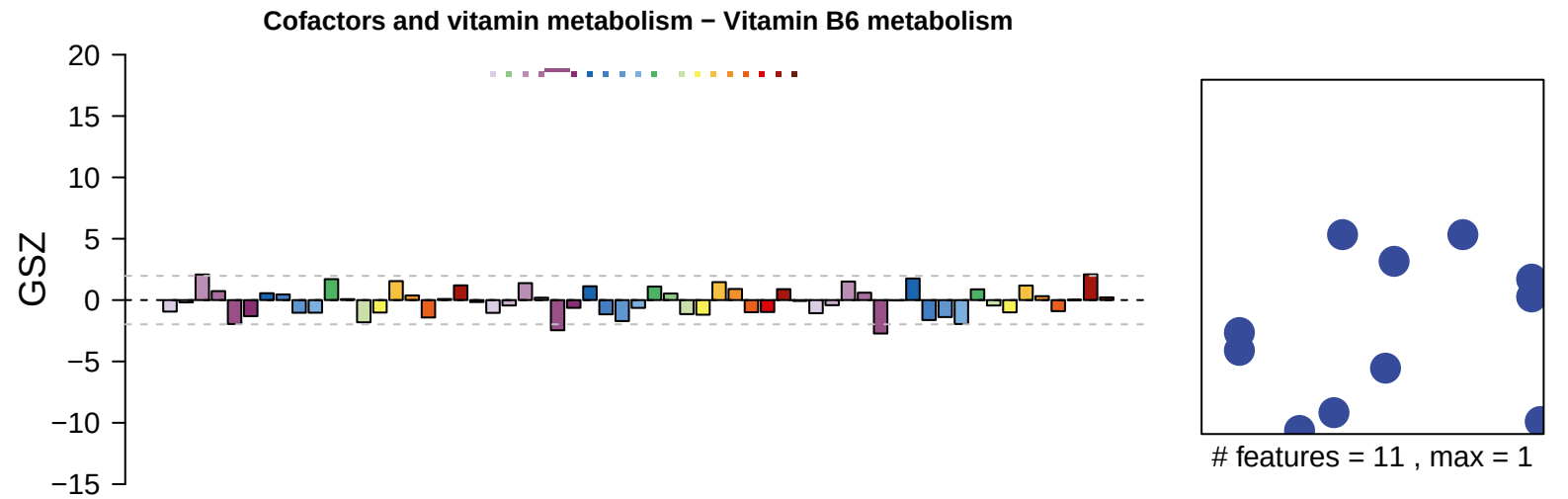
features = 21 , max = 1



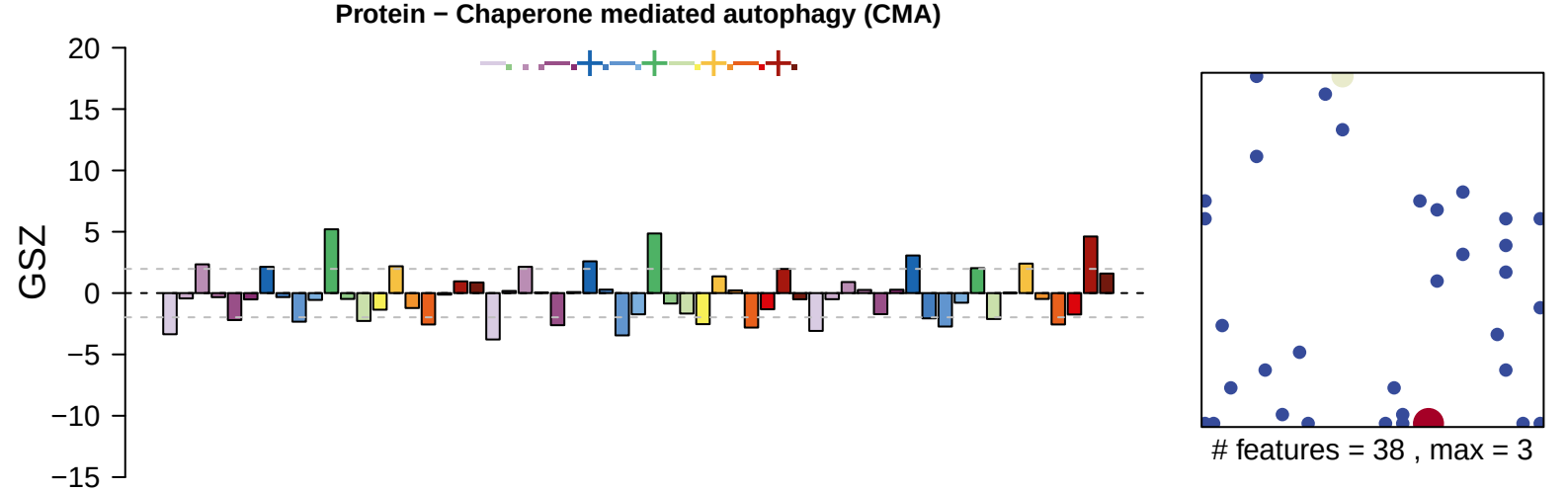
features = 92 , max = 2

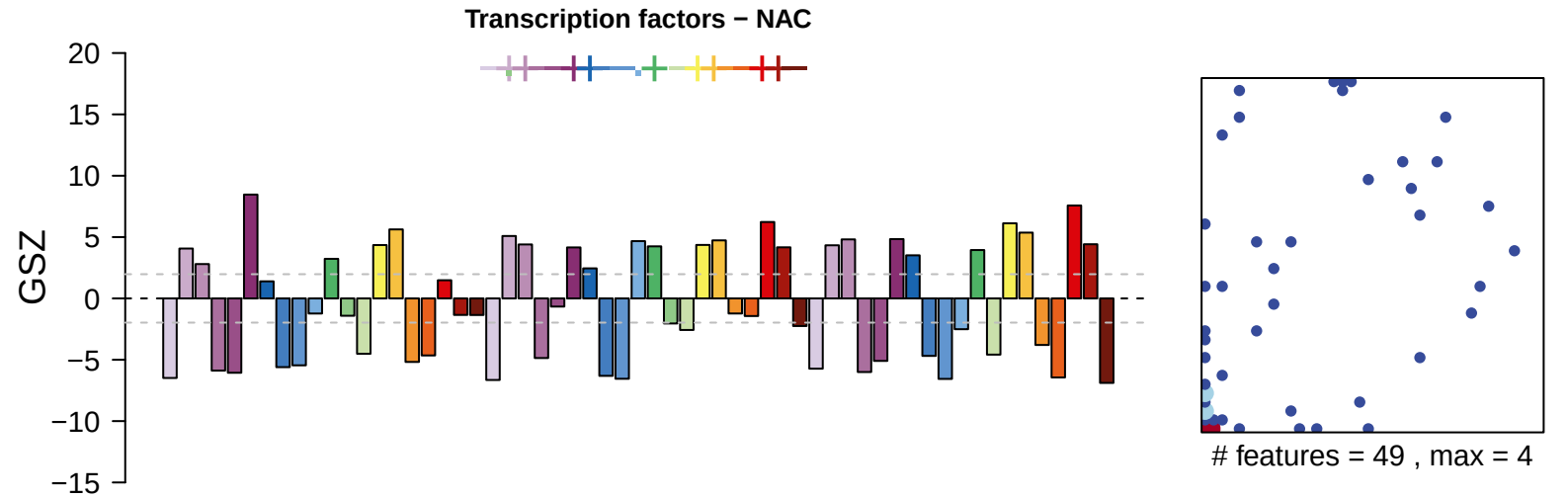


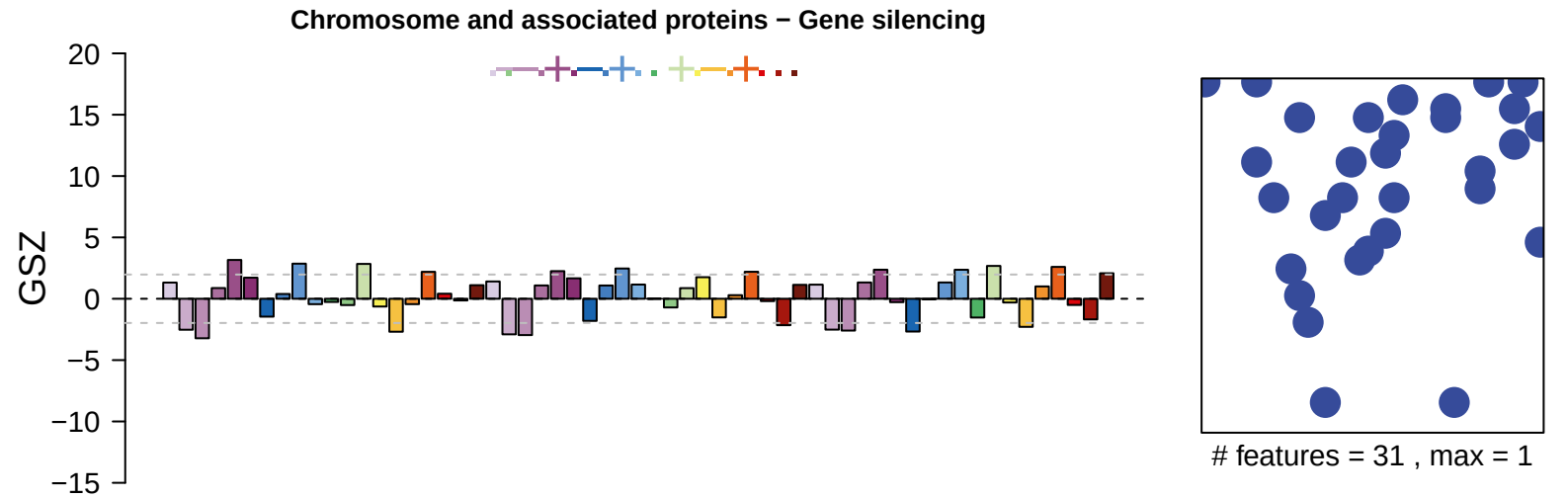
features = 64 , max = 10

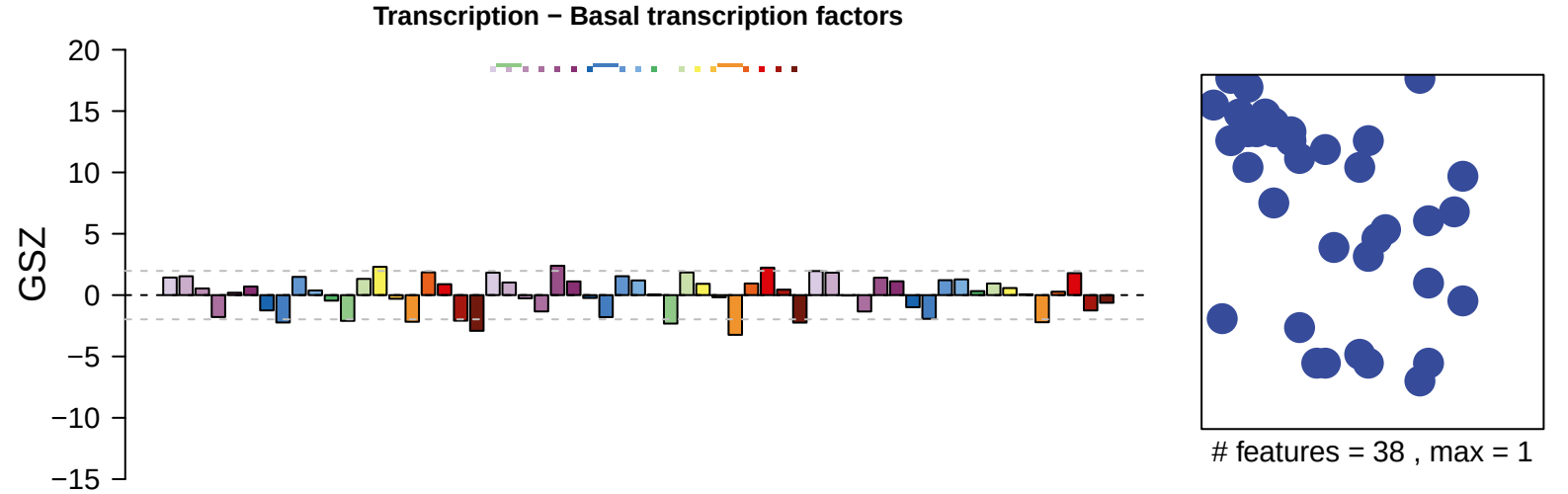


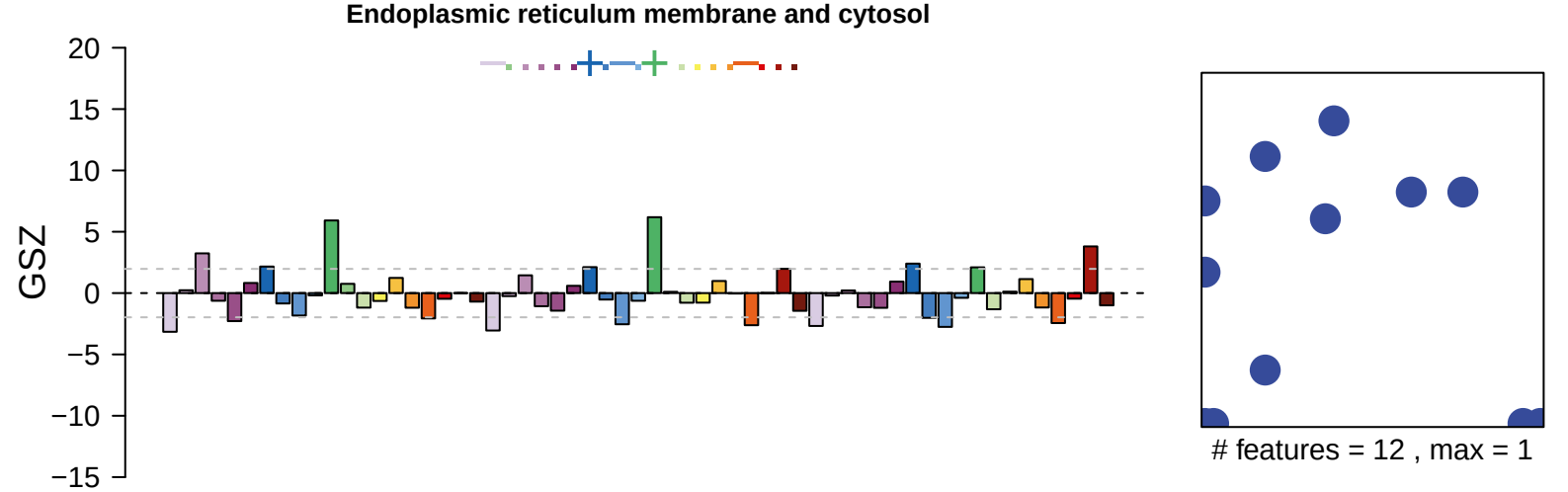
features = 11 , max = 1

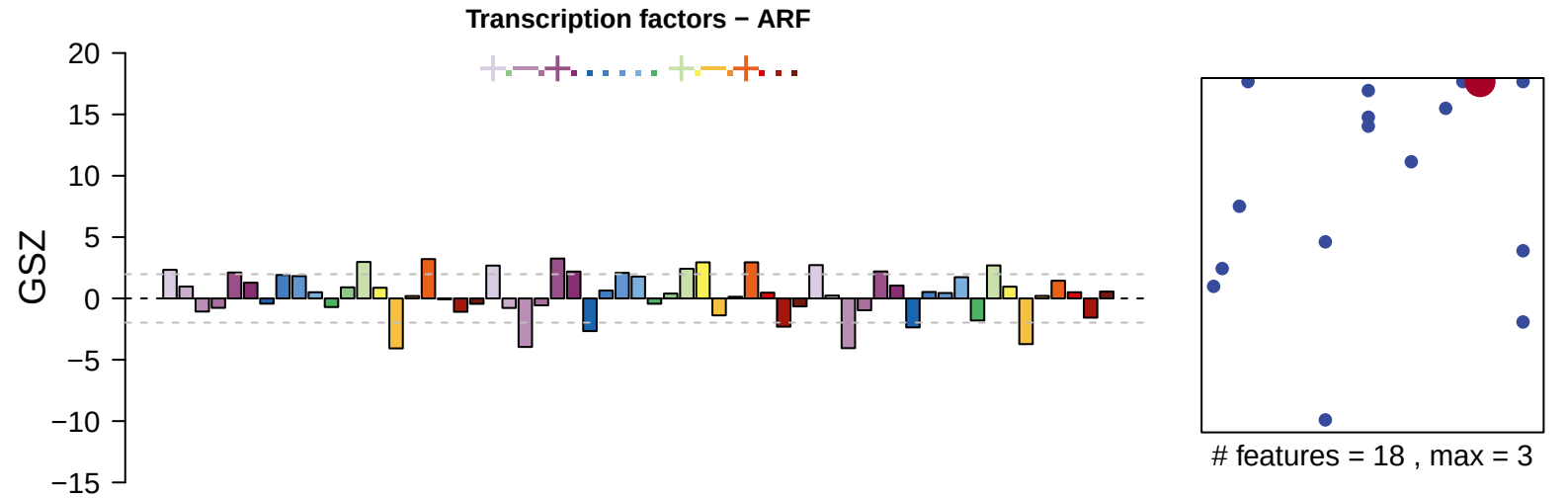

Protein − Chaperone mediated autophagy (CMA)
GSZ
20
15
10
5
0
−5
−10
−15
features = 38 , max = 3

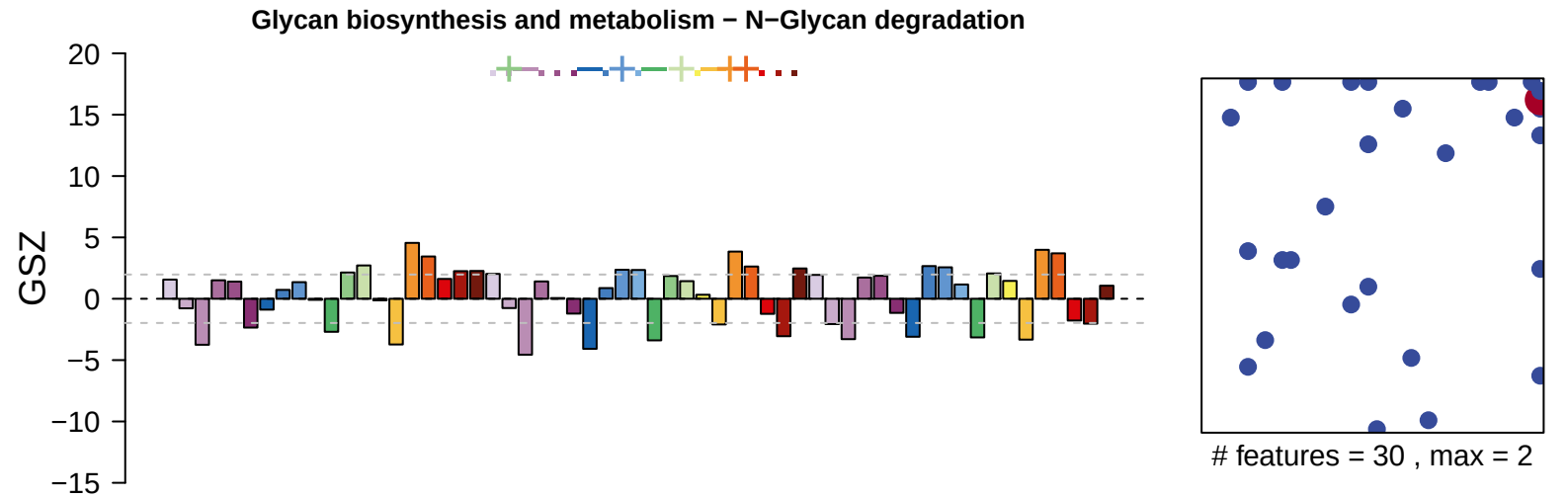

Transcription factors − NAC
GSZ
20
15
10
5
0
−5
−10
−15
features = 49 , max = 4

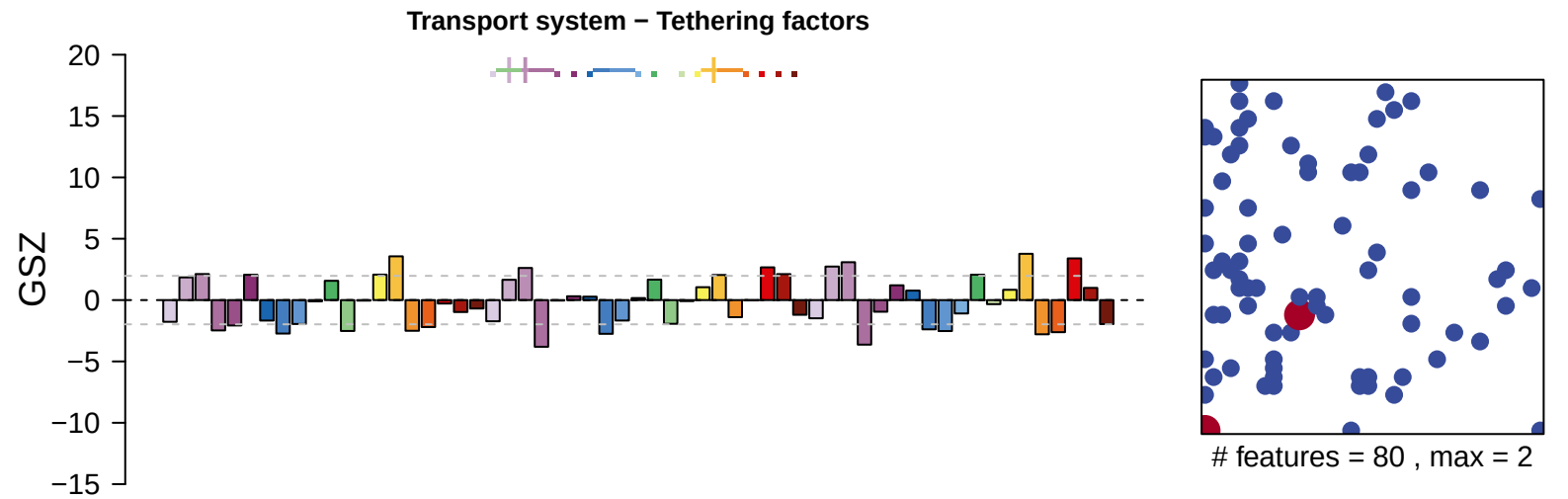

Chromosome and associated proteins − Gene silencing
GSZ
20
15
10
5
0
−5
−10
−15
features = 31 , max = 1

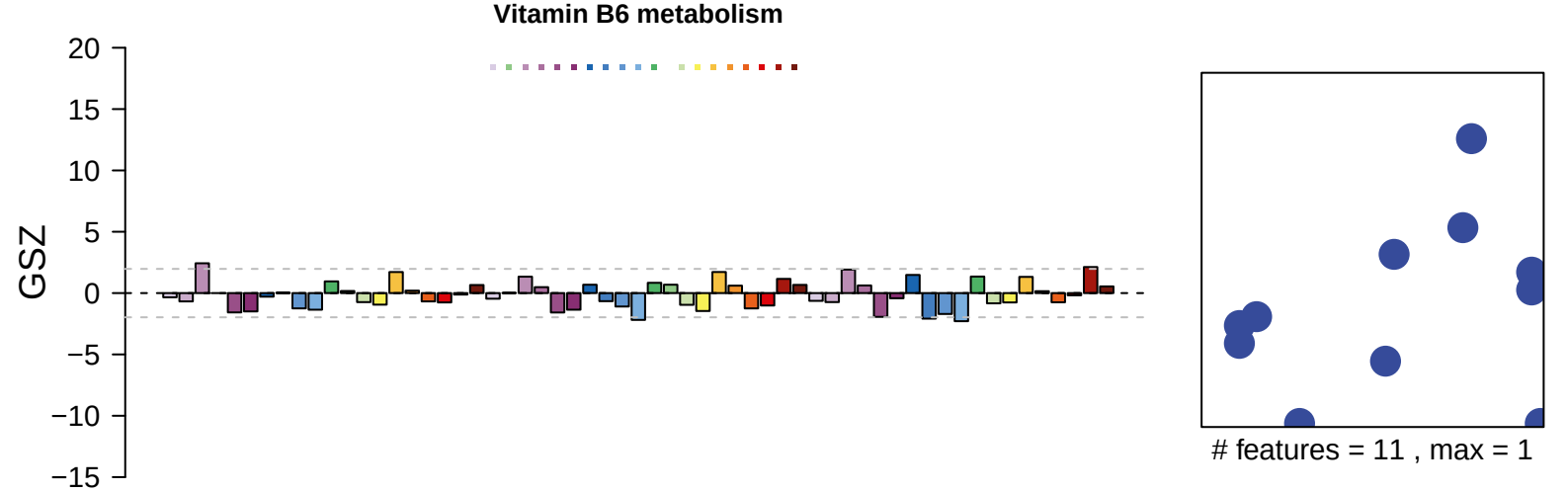

Transcription − Basal transcription factors
GSZ
20
15
10
5
0
−5
−10
−15
features = 38 , max = 1

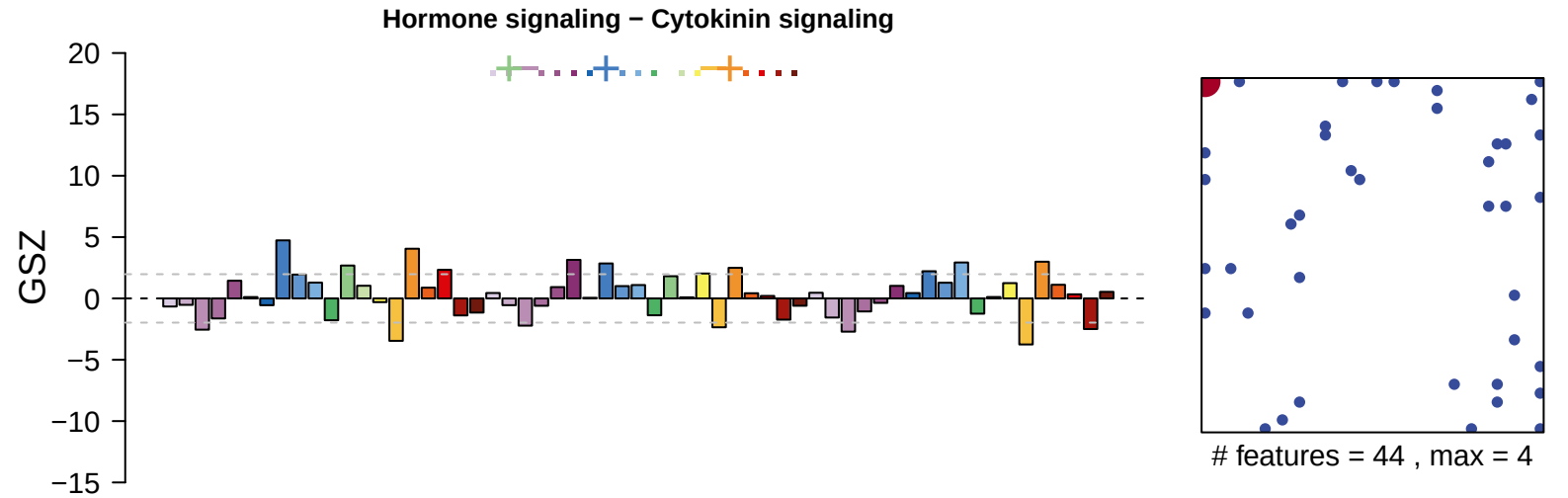

Endoplasmic reticulum membrane and cytosol
GSZ
20
15
10
5
0
−5
−10
−15
features = 12 , max = 1

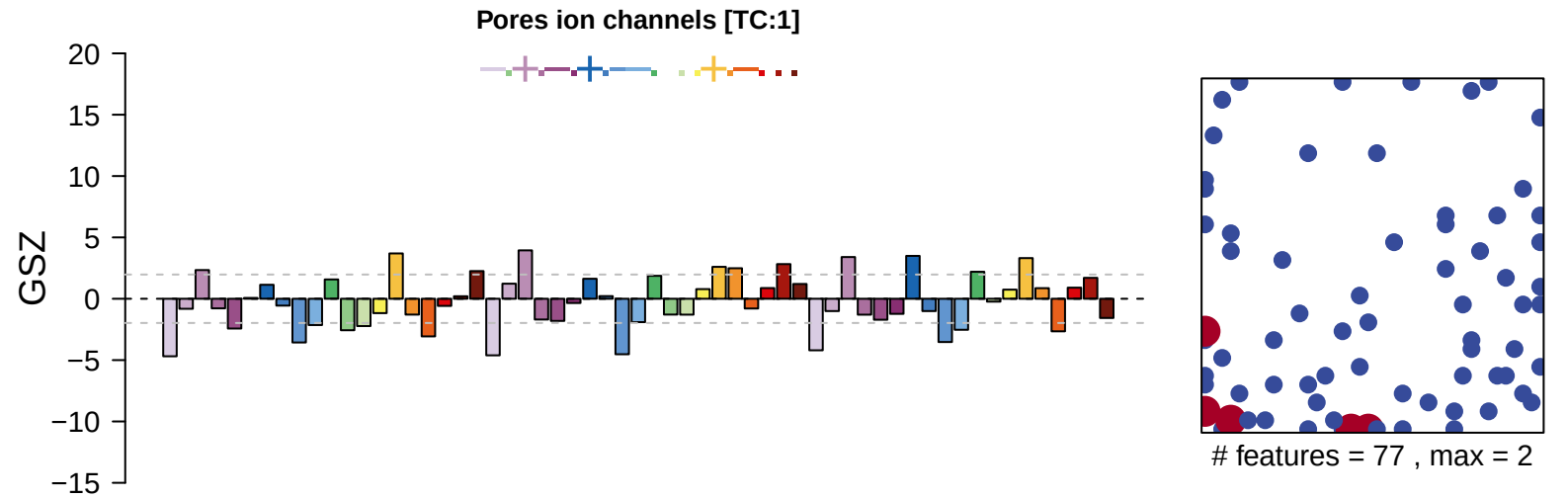

Transcription factors − ARF
GSZ
20
15
10
5
0
−5
−10
−15
features = 18 , max = 3

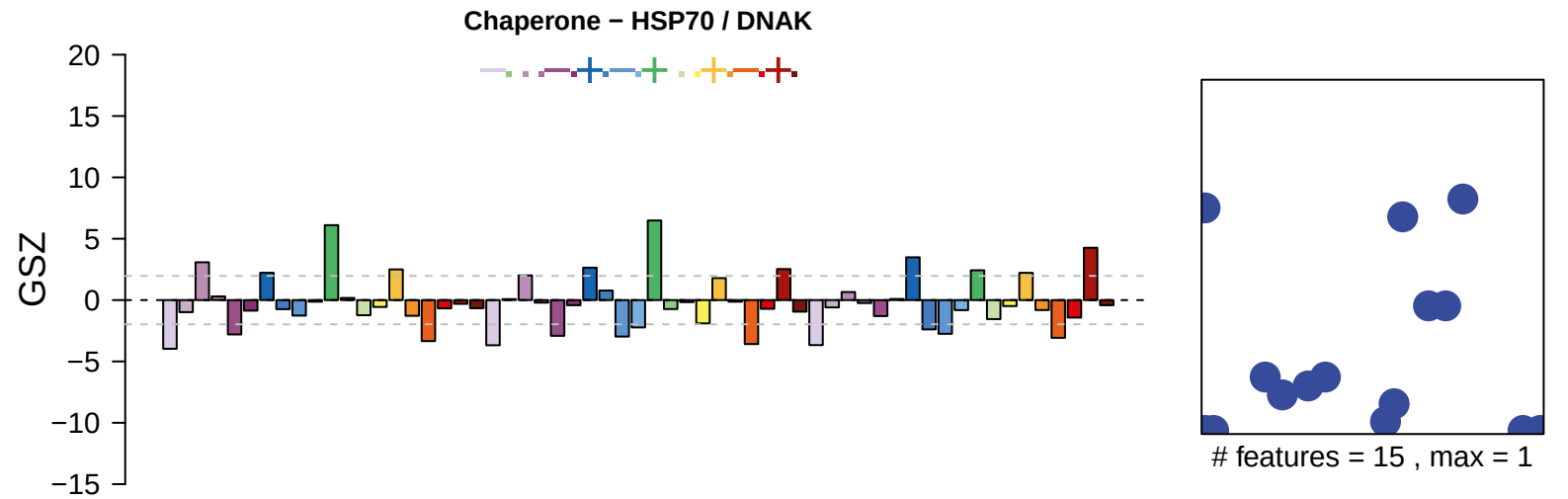

Glycan biosynthesis and metabolism − N−Glycan degradation
GSZ
20
15
10
5
0
−5
−10
−15
features = 30 , max = 2


Transport system − Tethering factors
GSZ
20
15
10
5
0
−5
−10
−15
features = 80 , max = 2


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


Hormone signaling − Cytokinin signaling
GSZ
20
15
10
5
0
−5
−10
−15
features = 44 , max = 4


Pores ion channels [TC:1]
GSZ
20
15
10
5
0
−5
−10
−15
features = 77 , max = 2


Chaperone − HSP70 / DNAK
GSZ
20
15
10
5
0
−5
−10
−15
features = 15 , max = 1