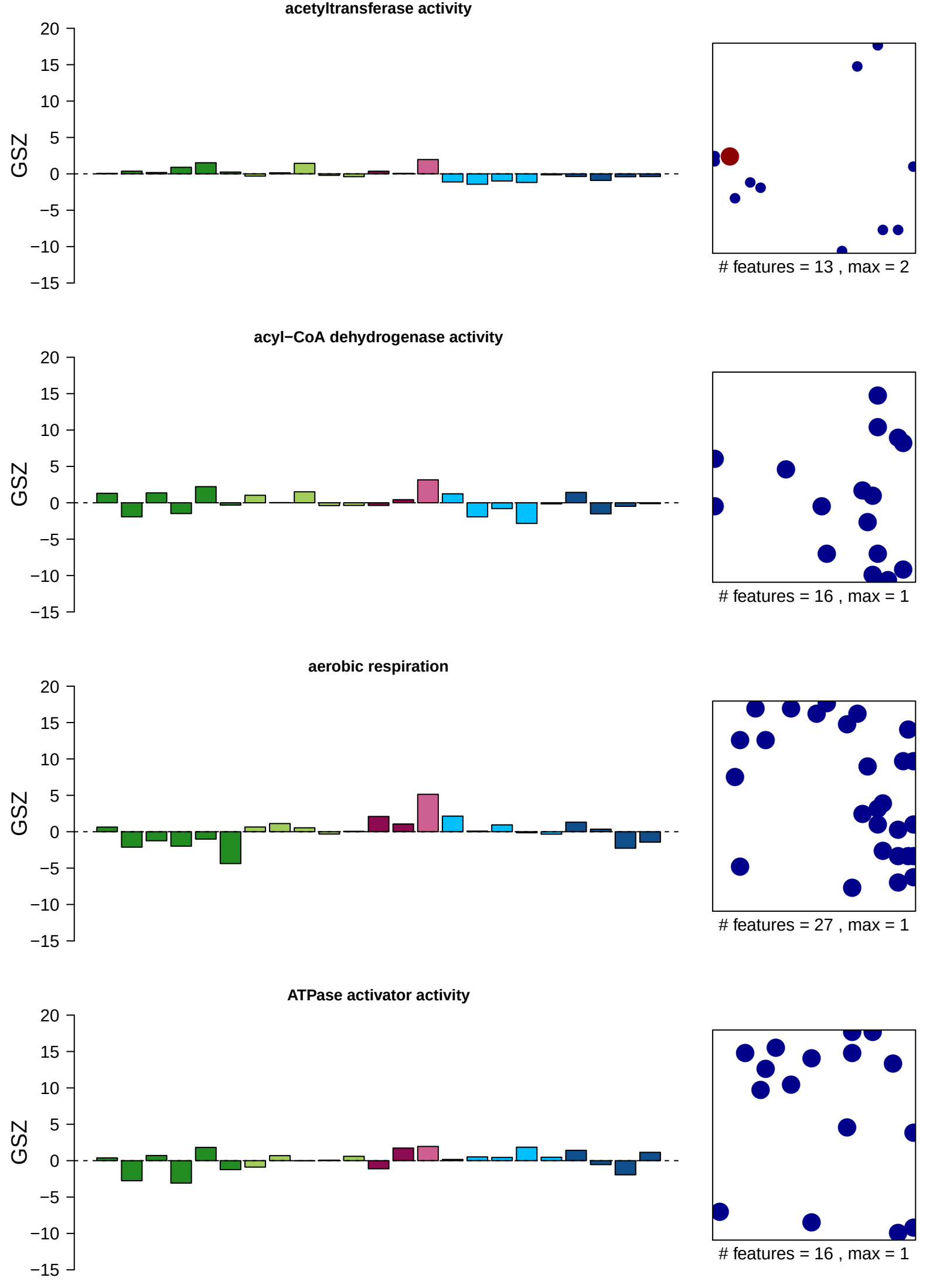
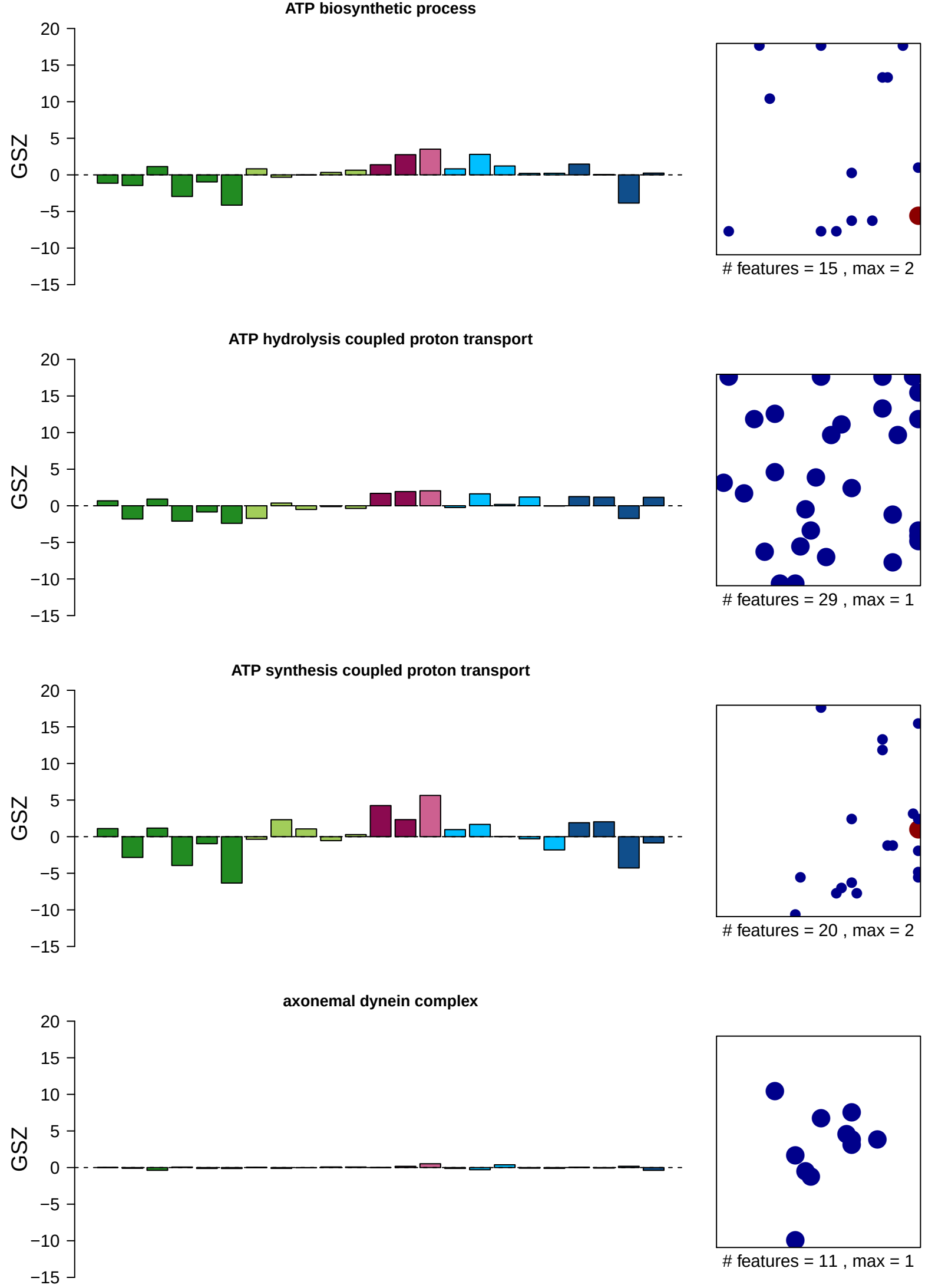
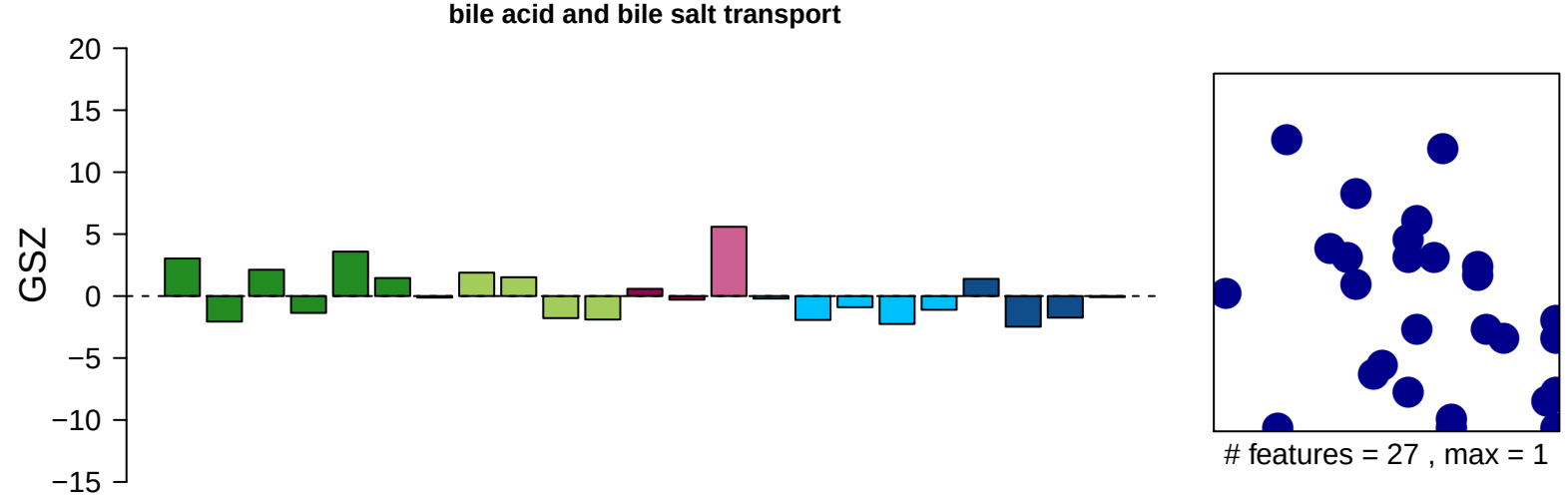
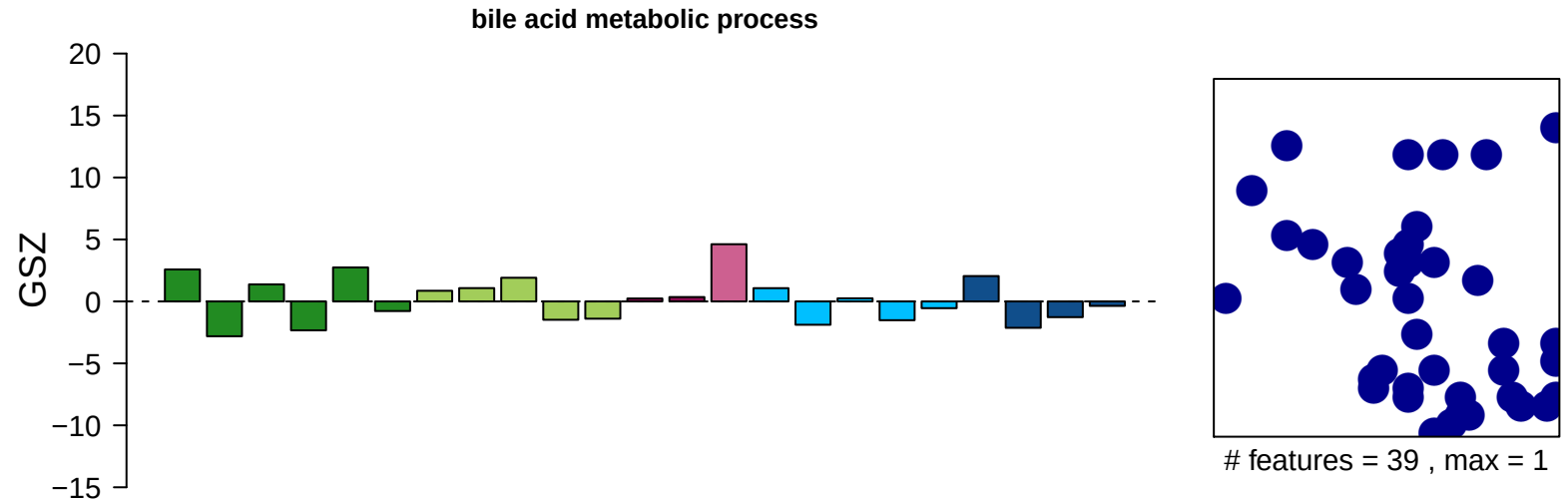

acetyltransferase activity
GSZ
20
15
10
5
0
−5
−10
−15
features = 13 , max = 2
acyl−CoA dehydrogenase activity
GSZ
20
15
10
5
0
−5
−10
−15
features = 16 , max = 1
aerobic respiration
GSZ
20
15
10
5
0
−5
−10
−15
features = 27 , max = 1
ATPase activator activity
GSZ
20
15
10
5
0
−5
−10
−15
features = 16 , max = 1

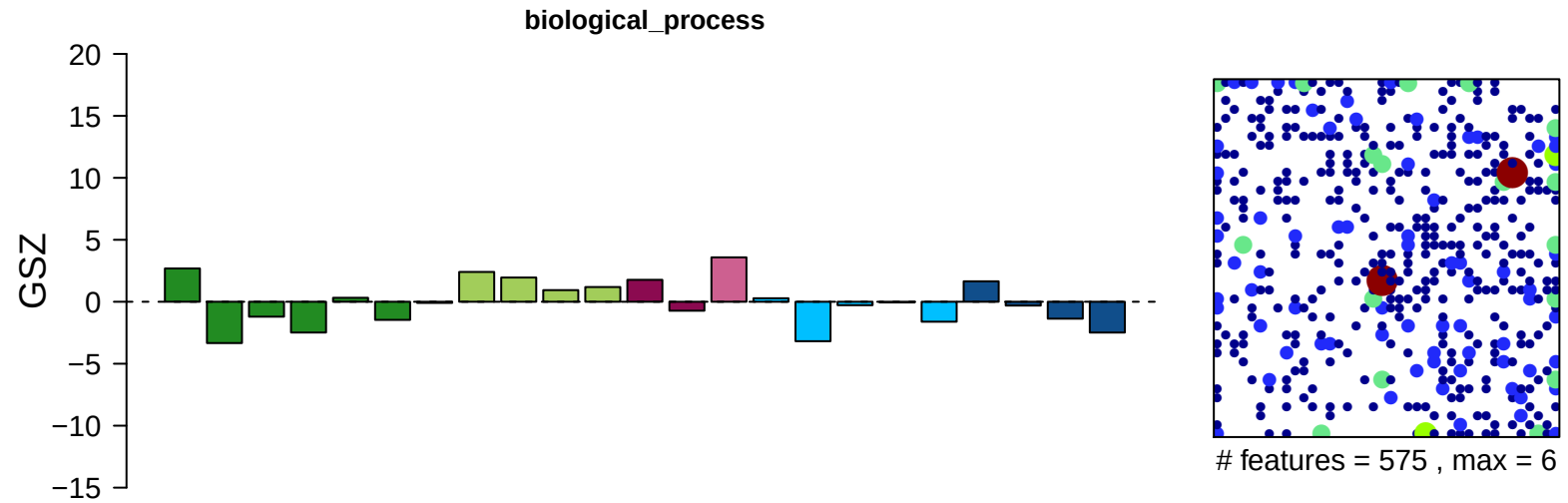

ATP biosynthetic process
GSZ
20
15
10
5
0
−5
−10
−15
features = 15 , max = 2
ATP hydrolysis coupled proton transport
GSZ
20
15
10
5
0
−5
−10
−15
features = 29 , max = 1
ATP synthesis coupled proton transport
GSZ
20
15
10
5
0
−5
−10
−15
features = 20 , max = 2
axonemal dynein complex
GSZ
20
15
10
5
0
−5
−10
−15
features = 11 , max = 1



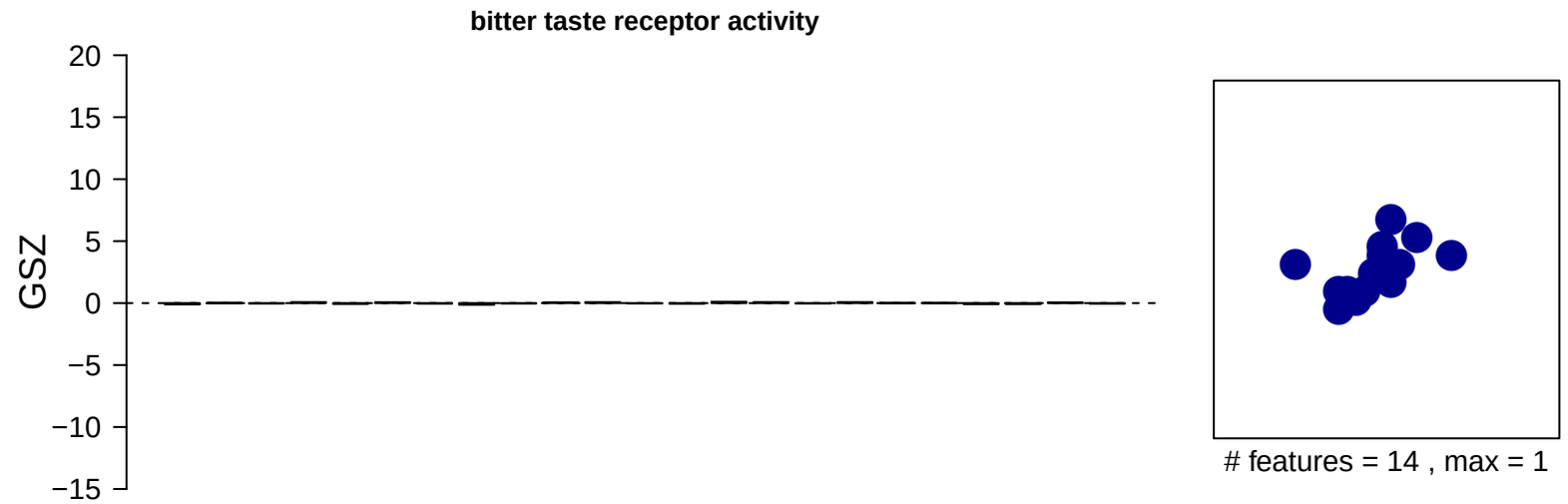
features = 27 , max = 1



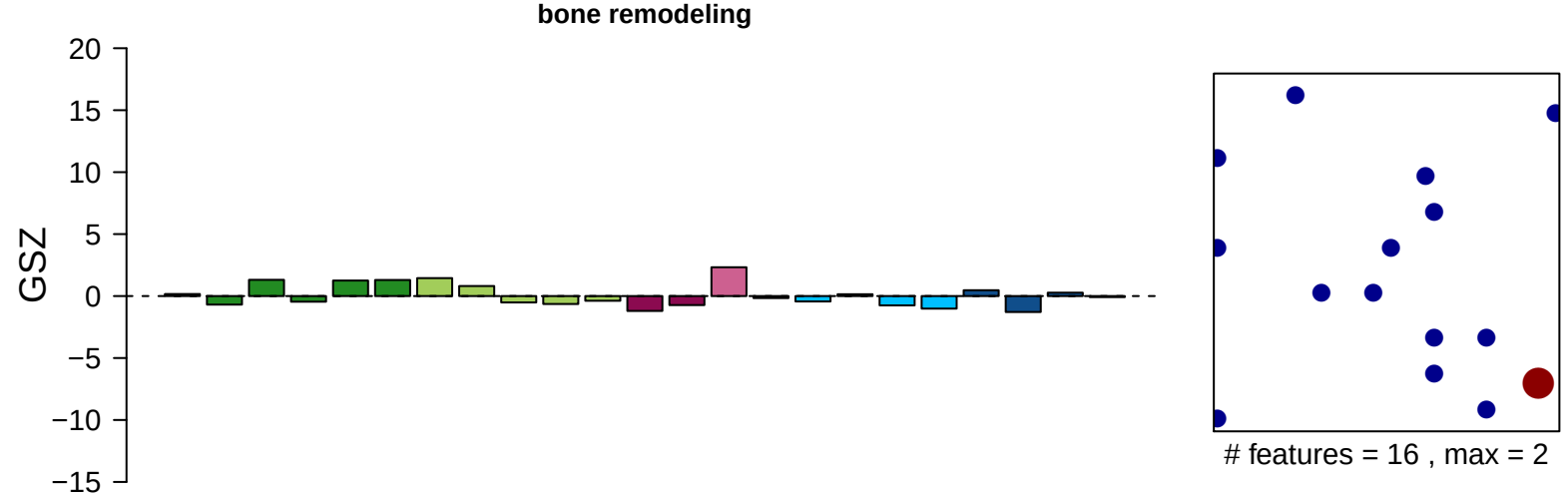
features = 39 , max = 1

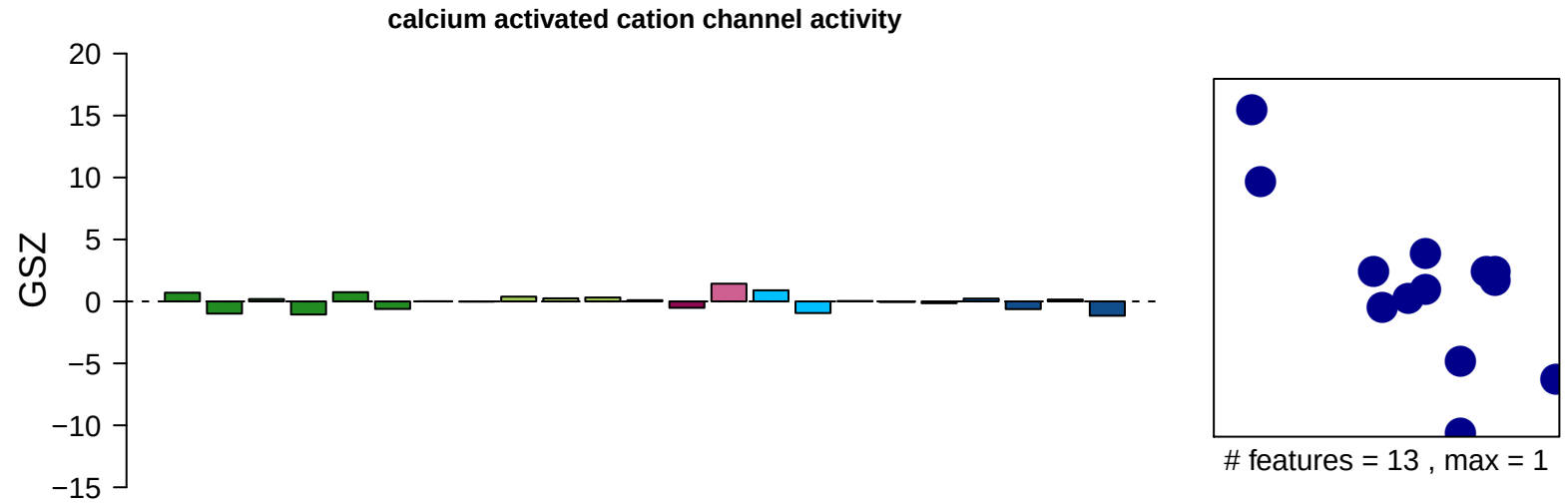


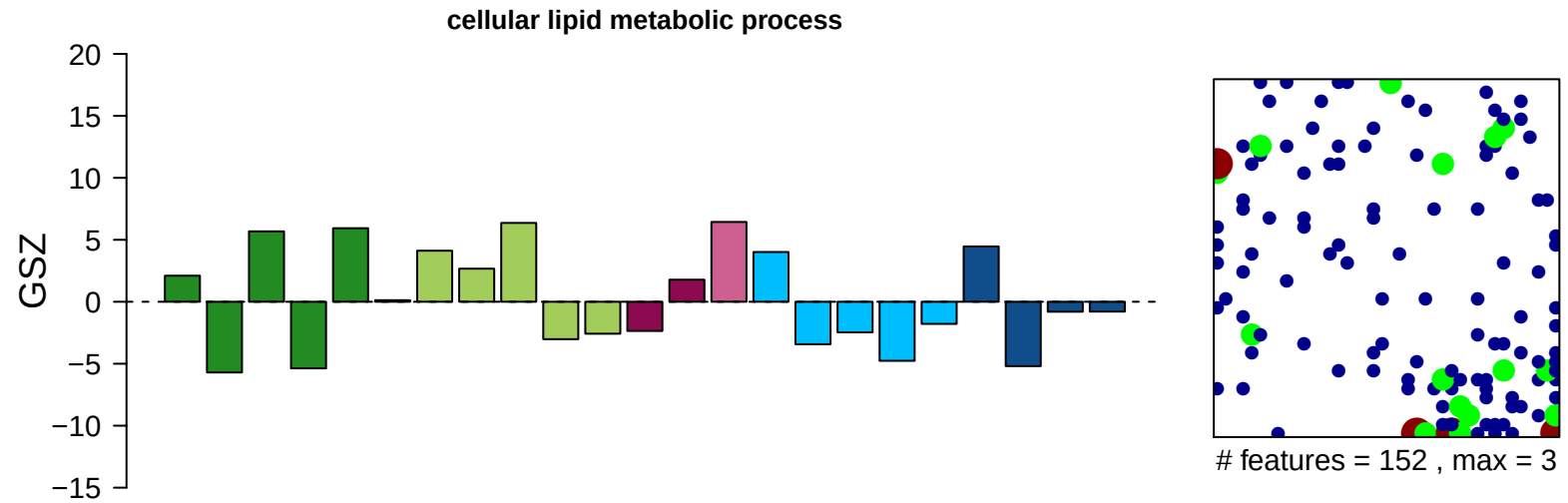
features = 575 , max = 6

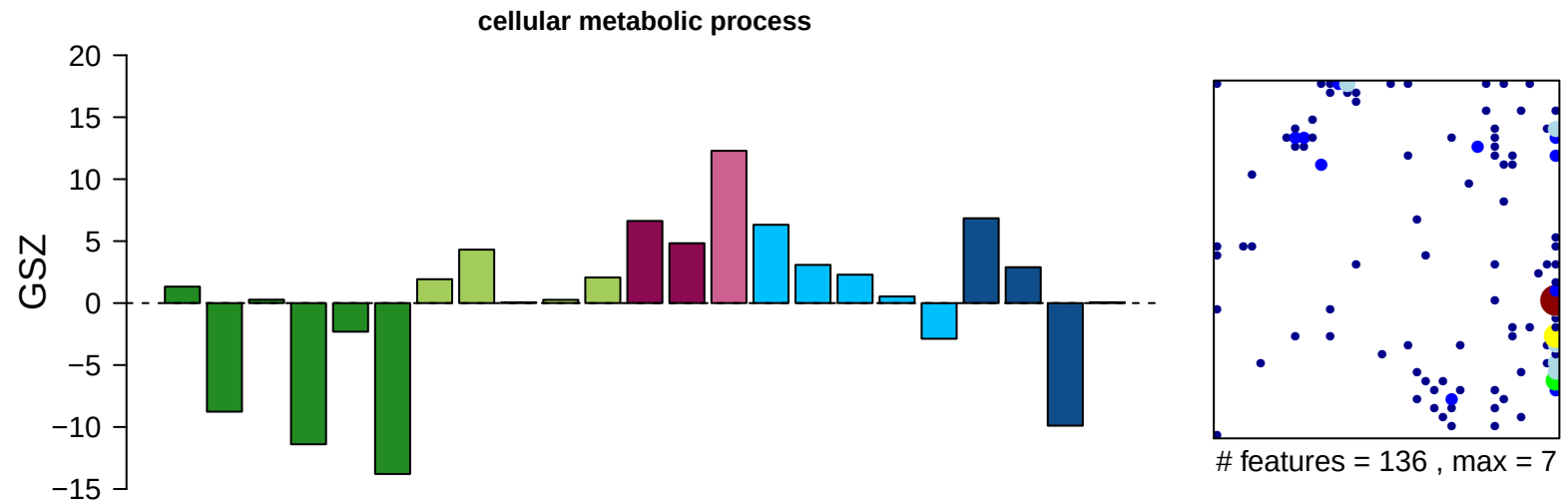


features = 14 , max = 1

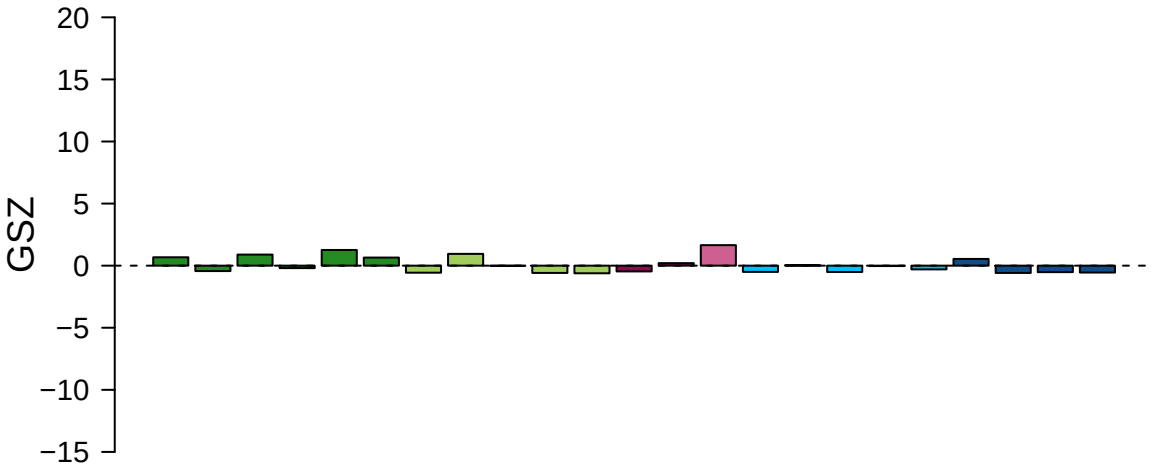

bone remodeling
GSZ
20
15
10
5
0
−5
−10
−15
features = 16 , max = 2

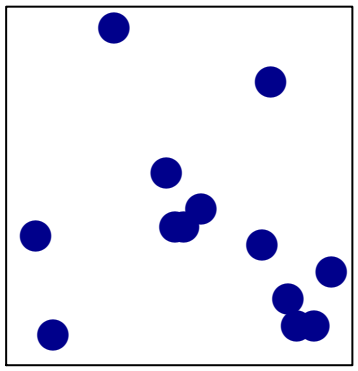

calcium activated cation channel activity
GSZ
20
15
10
5
0
−5
−10
−15
features = 13 , max = 1


cellular lipid metabolic process
GSZ
20
15
10
5
0
−5
−10
−15
features = 152 , max = 3


cellular metabolic process
GSZ
20
15
10
5
0
−5
−10
−15
features = 136 , max = 7

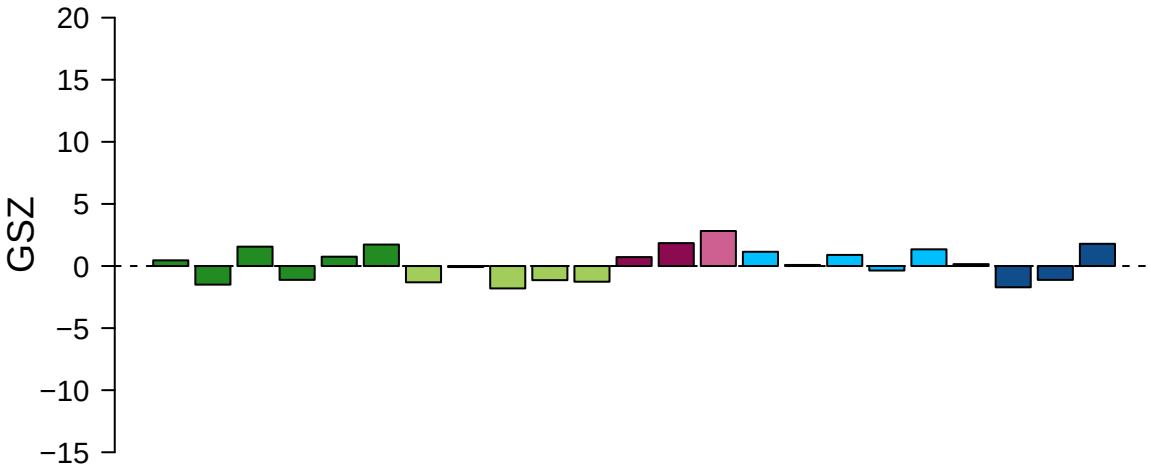
cellular response to electrical stimulus

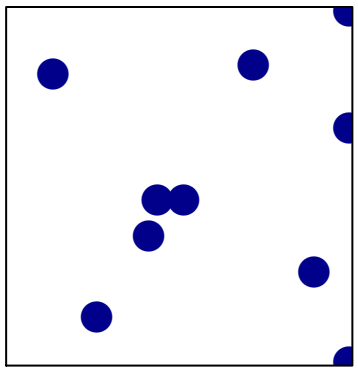




features = 13 , max = 1

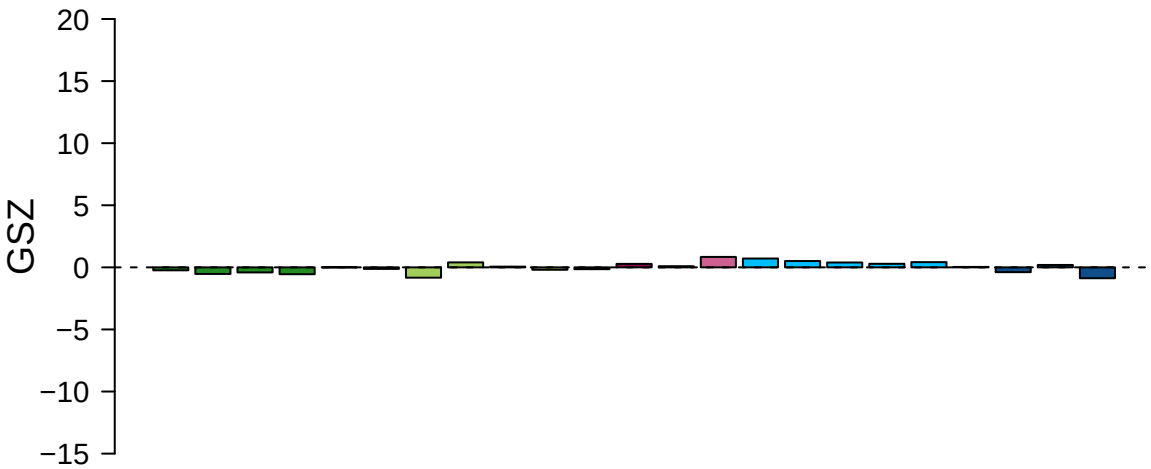
chloride ion homeostasis

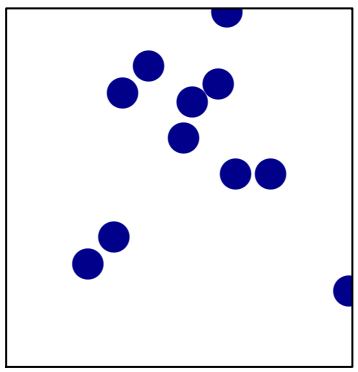




features = 10 , max = 1

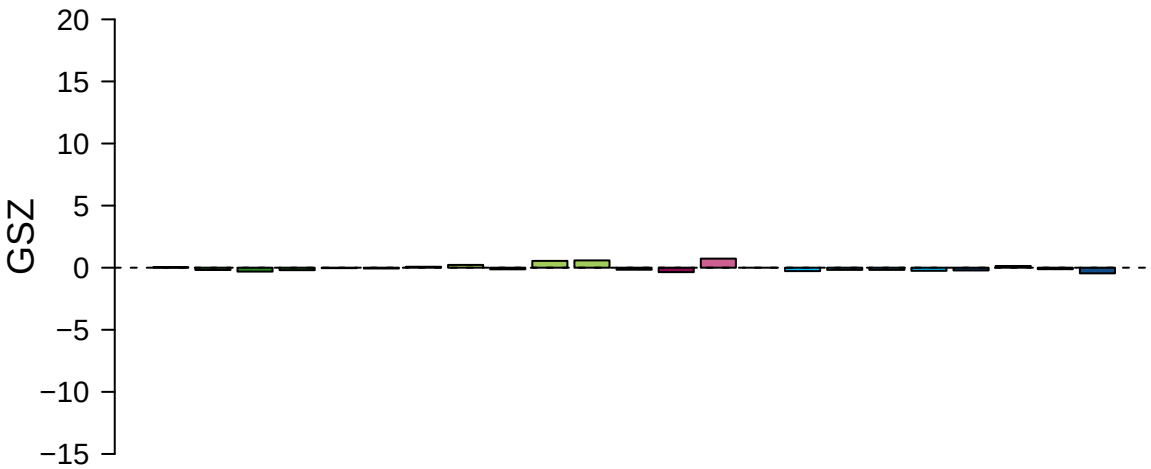
cilium organization

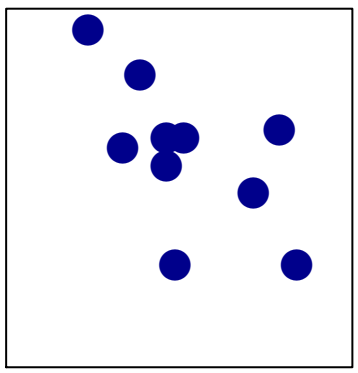




features = 11 , max = 1

cornea development in camera−type eye





features = 10 , max = 1