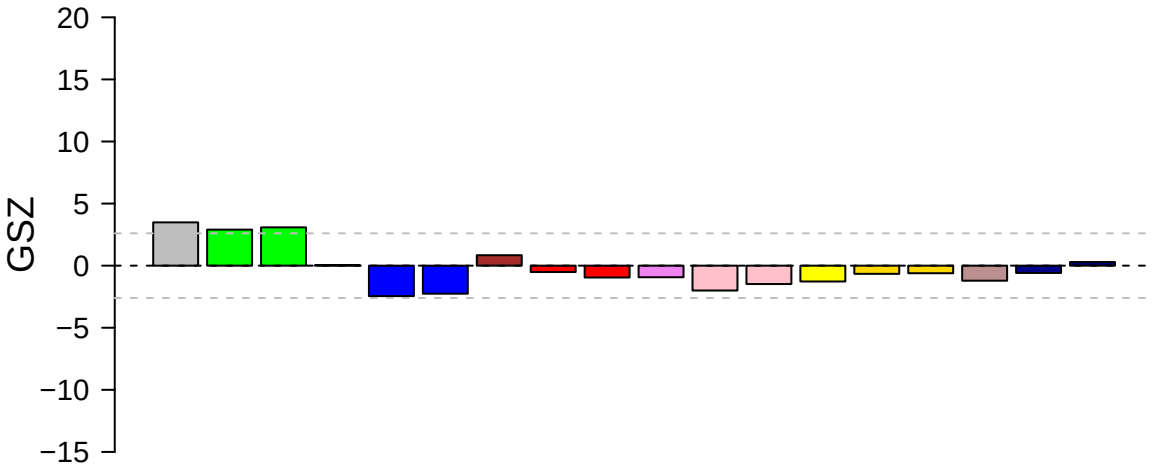
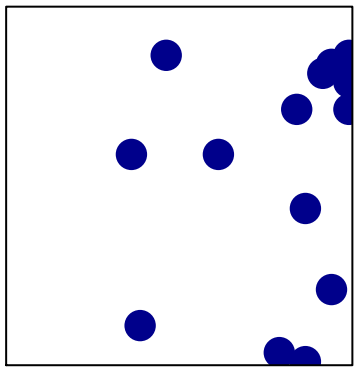
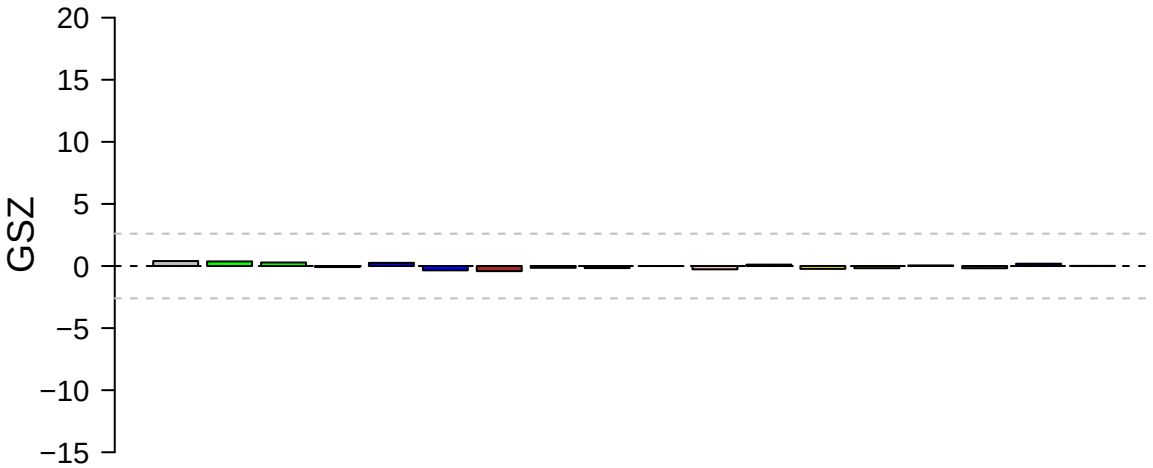
3'−5' exonuclease activity

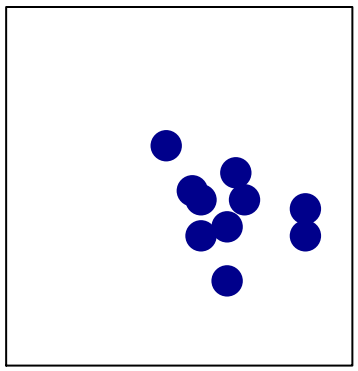




features = 15 , max = 1

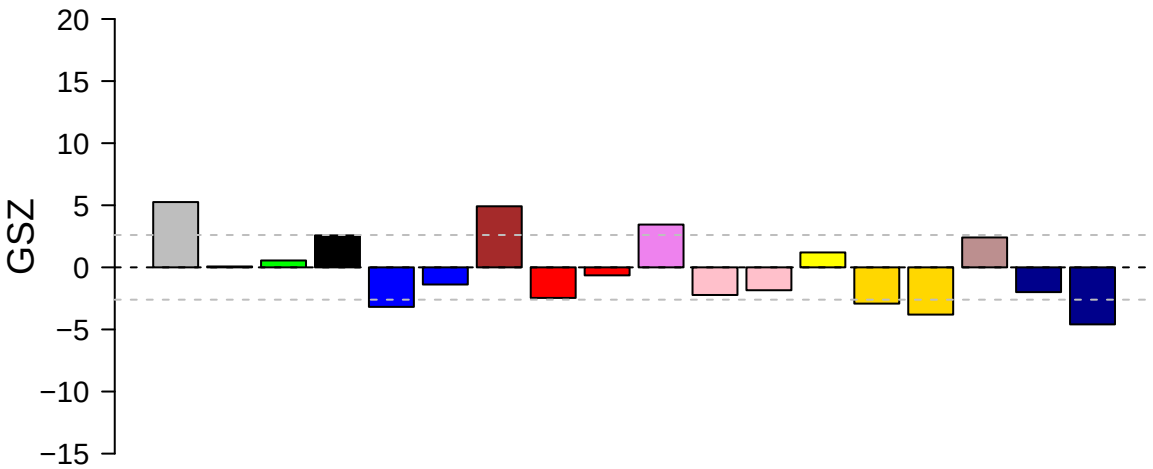
acrosin binding

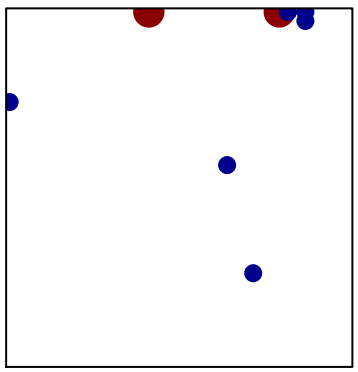




features = 10 , max = 1

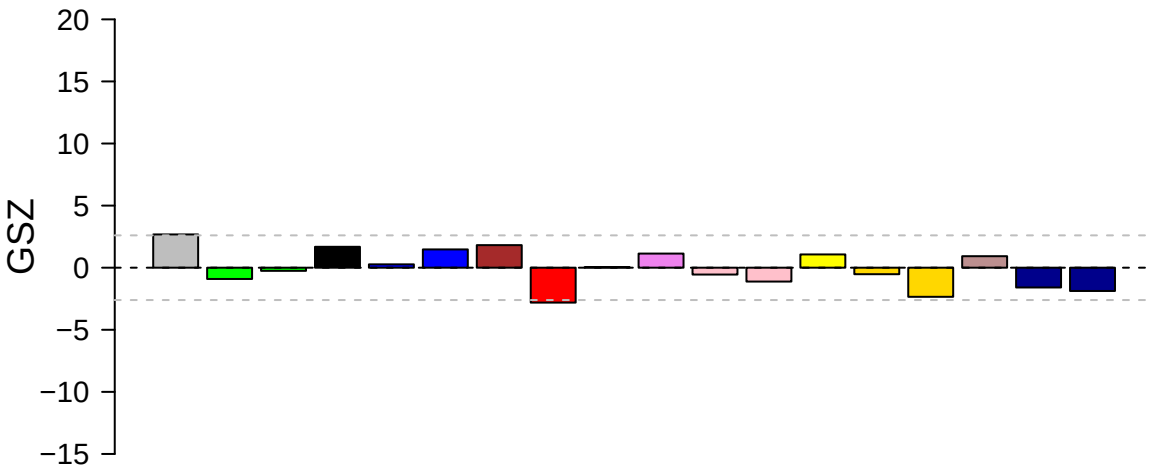
adenohypophysis development

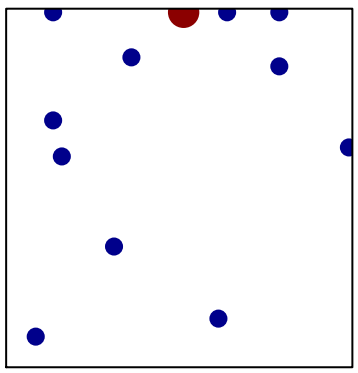




features = 10 , max = 2

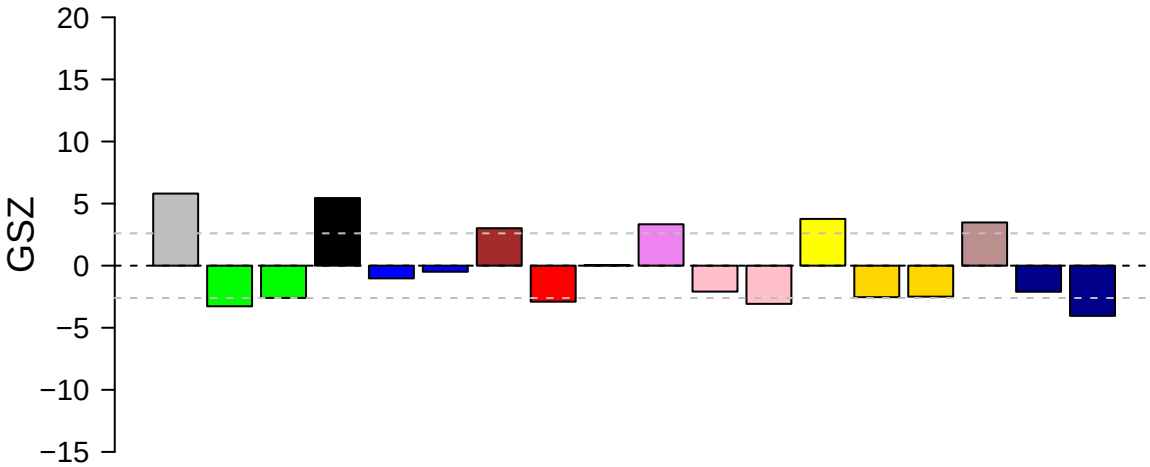
adherens junction

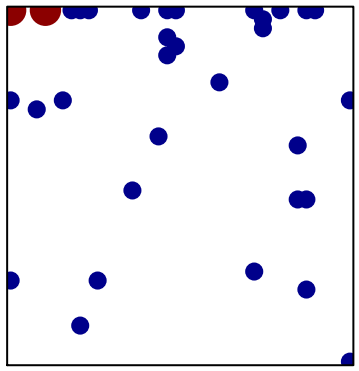




features = 13 , max = 2

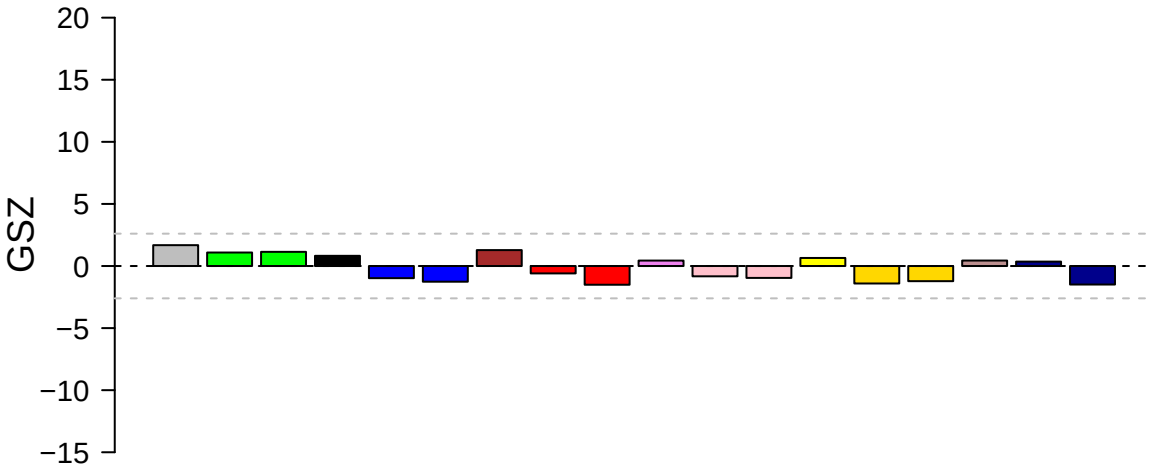
anterior/posterior pattern specification

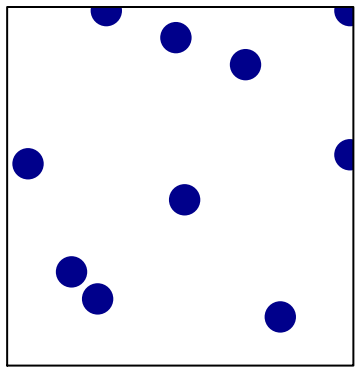




features = 35 , max = 2

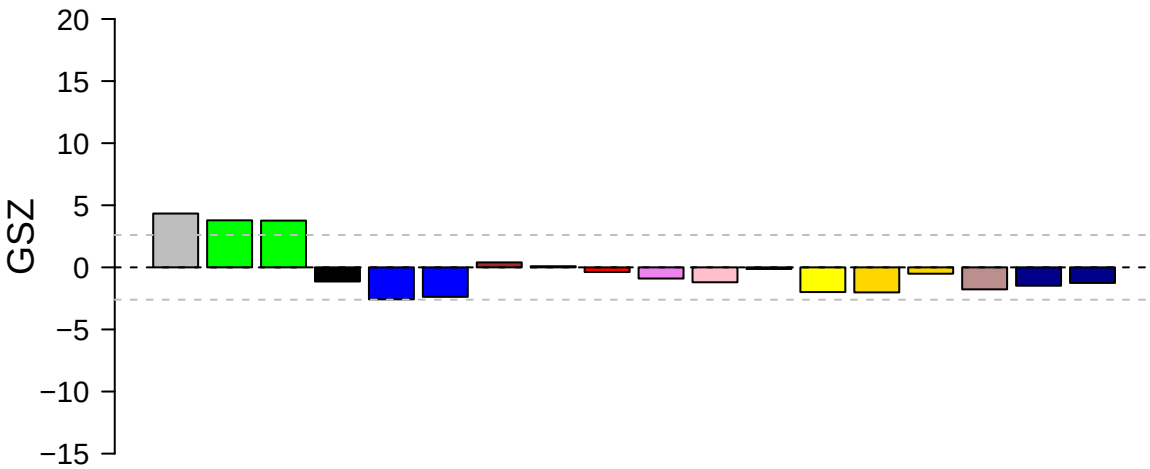
artery morphogenesis

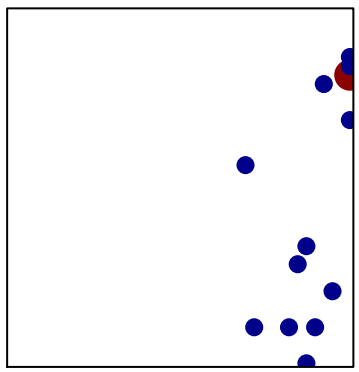




features = 10 , max = 1

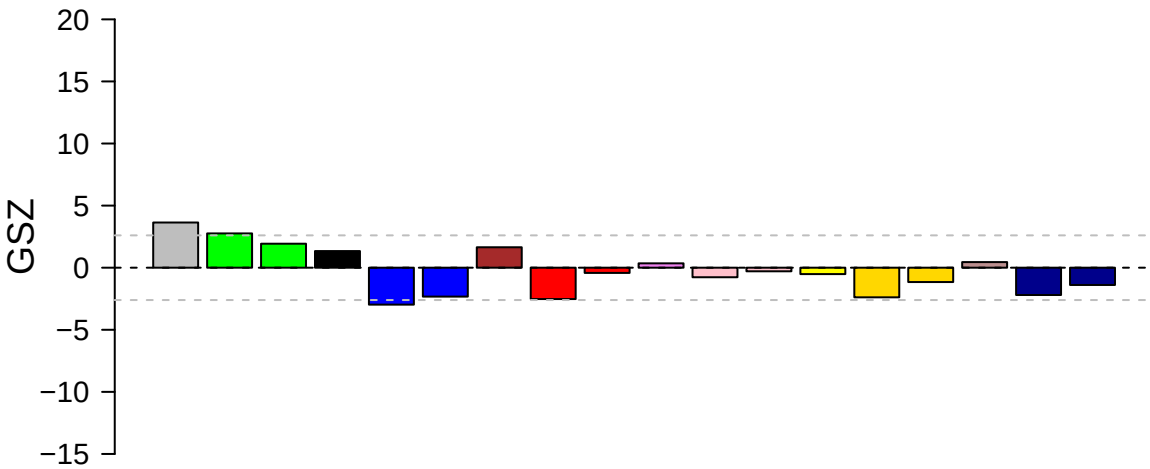
ATP−dependent DNA helicase activity

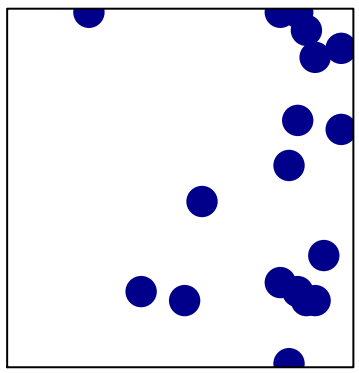




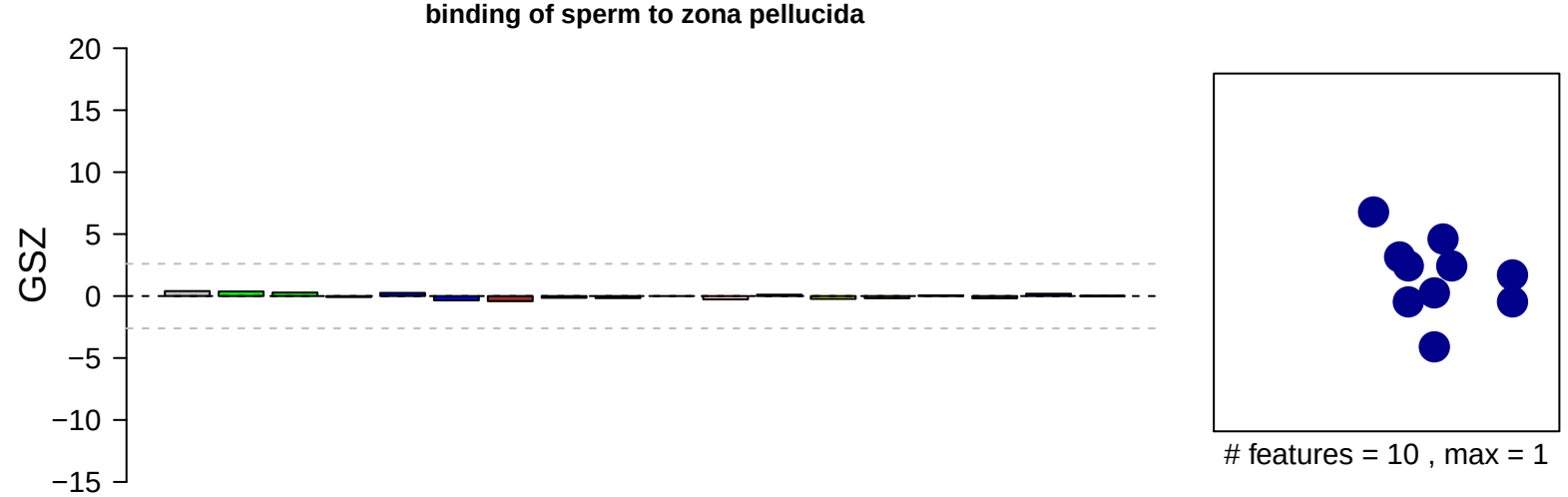
features = 14 , max = 2

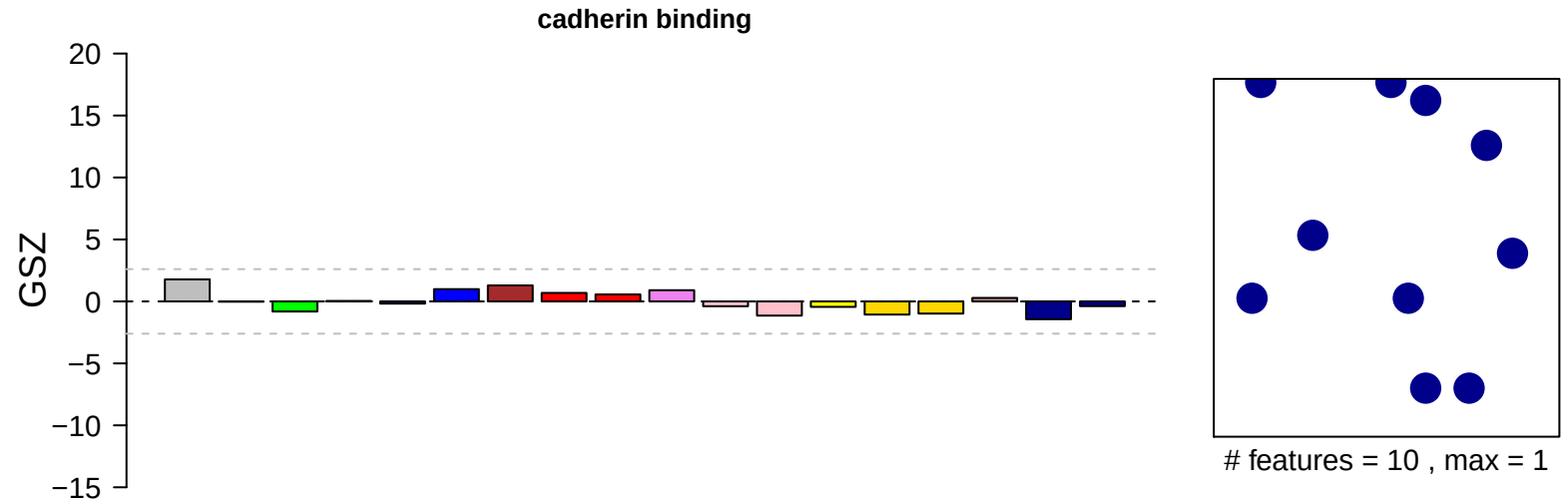
base−excision repair

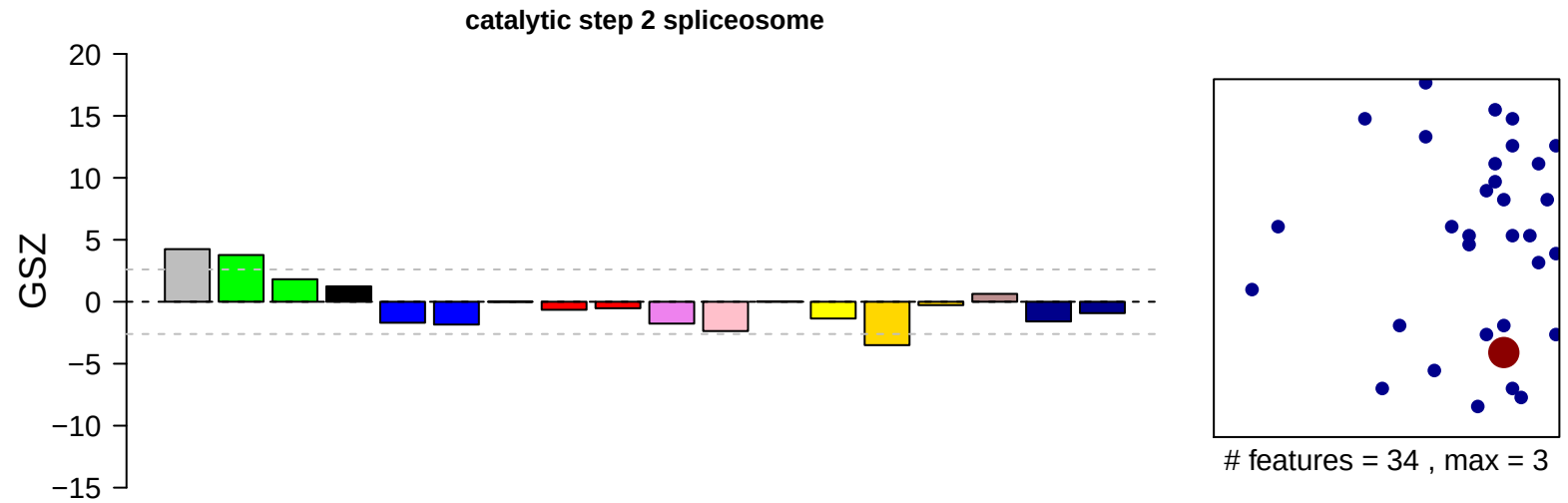


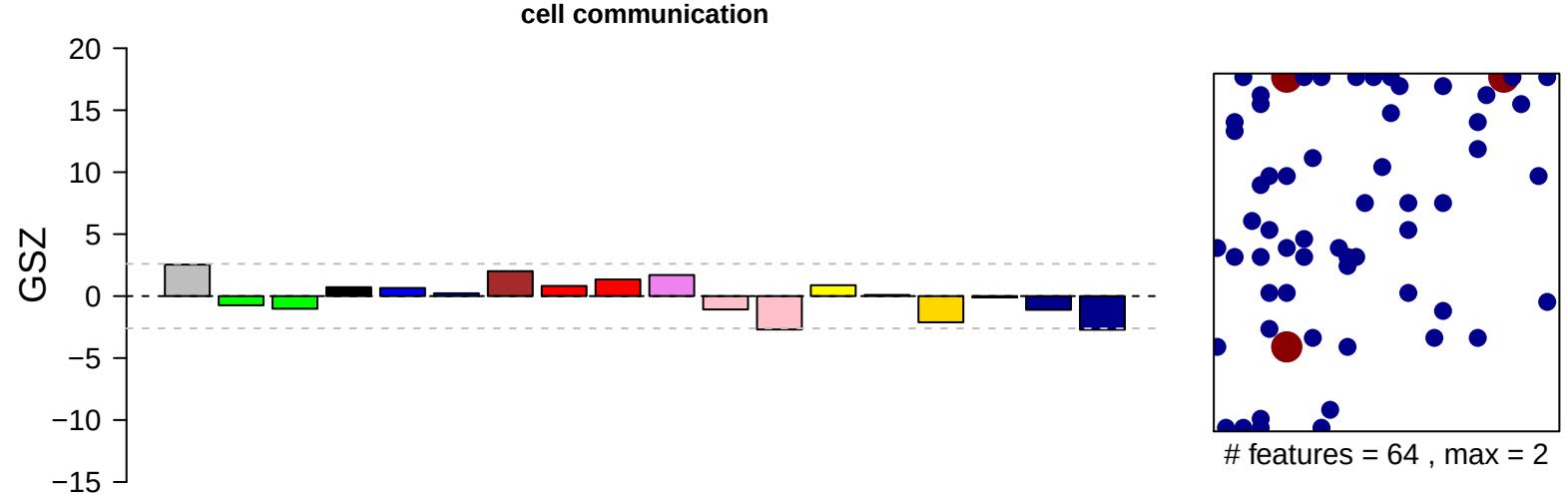


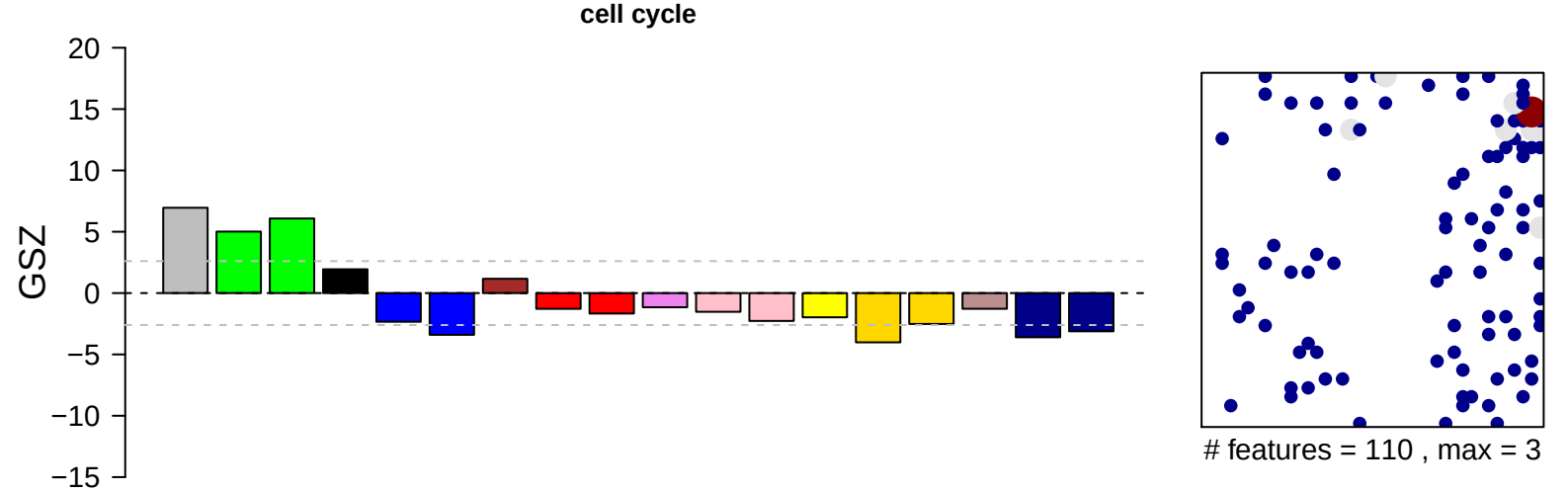
features = 18 , max = 1

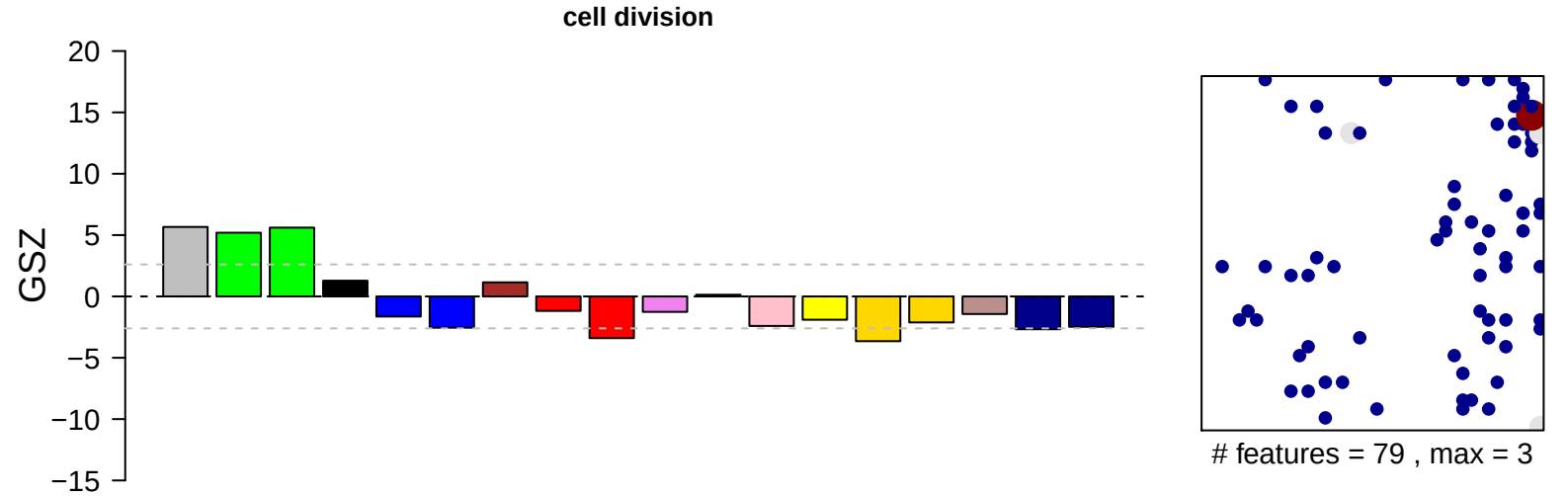

binding of sperm to zona pellucida
GSZ
20
15
10
5
0
−5
−10
−15
features = 10 , max = 1

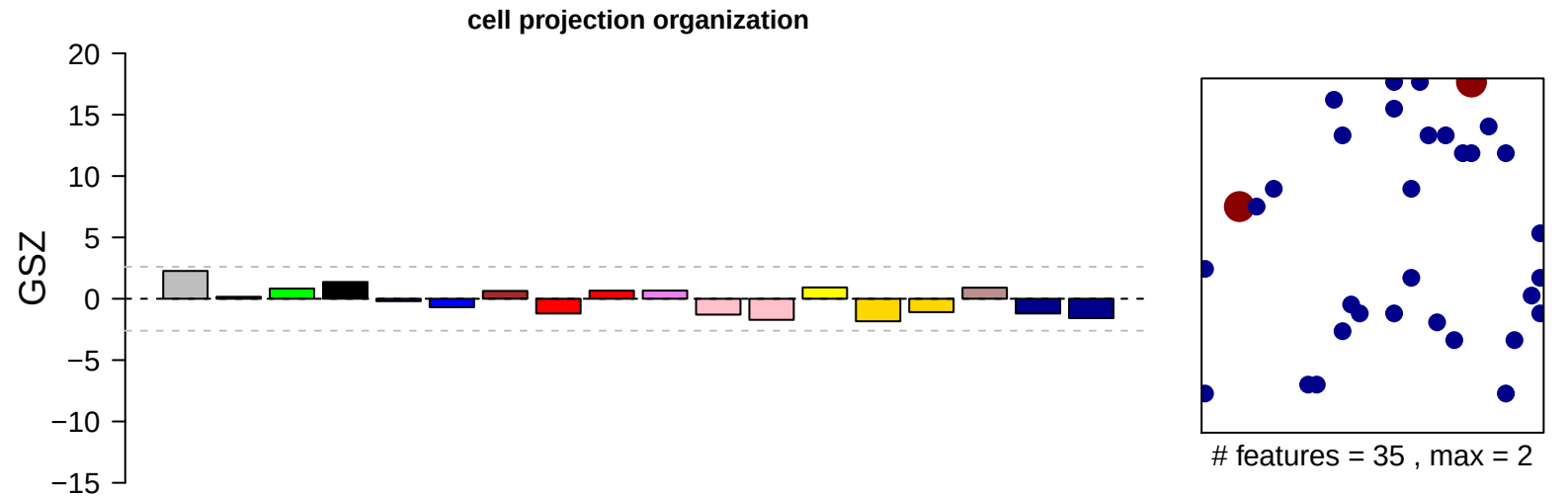

cadherin binding
GSZ
20
15
10
5
0
−5
−10
−15
features = 10 , max = 1

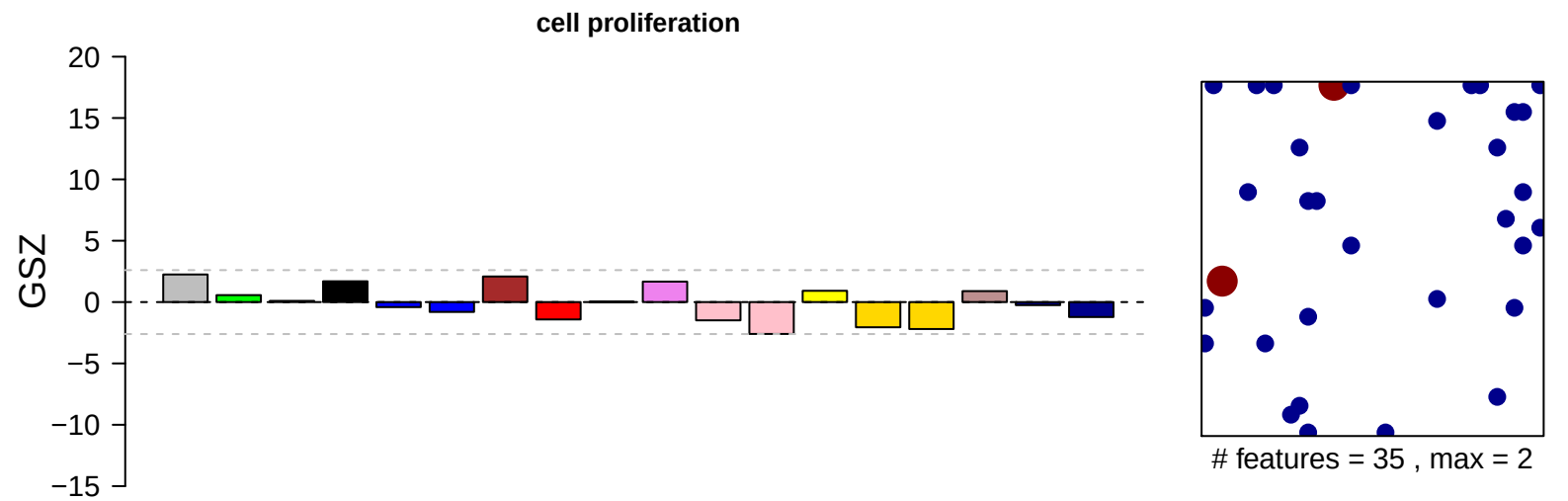

catalytic step 2 spliceosome
GSZ
20
15
10
5
0
−5
−10
−15
features = 34 , max = 3


cell communication
GSZ
20
15
10
5
0
−5
−10
−15
features = 64 , max = 2

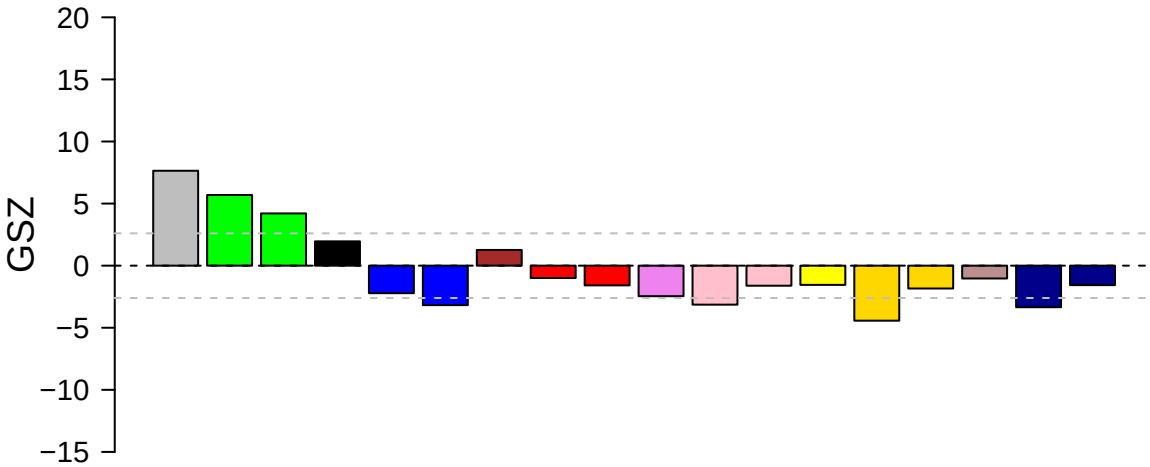

cell cycle
GSZ
20
15
10
5
0
−5
−10
−15
features = 110 , max = 3

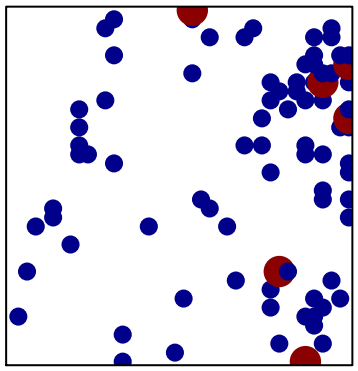

cell division
GSZ
20
15
10
5
0
−5
−10
−15
features = 79 , max = 3


cell projection organization
GSZ
20
15
10
5
0
−5
−10
−15
features = 35 , max = 2


cell proliferation
GSZ
20
15
10
5
0
−5
−10
−15
features = 35 , max = 2

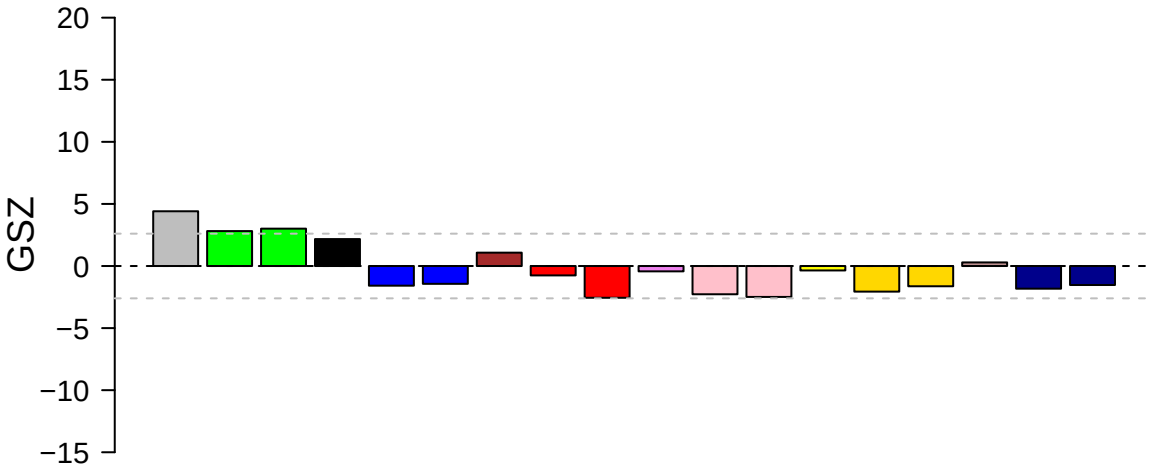
cellular response to DNA damage stimulus

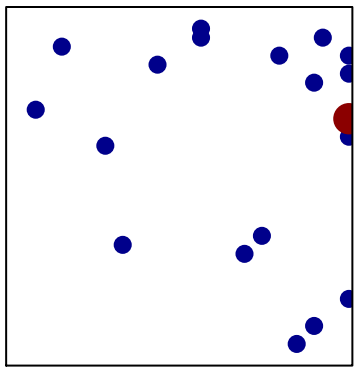




features = 91 , max = 2

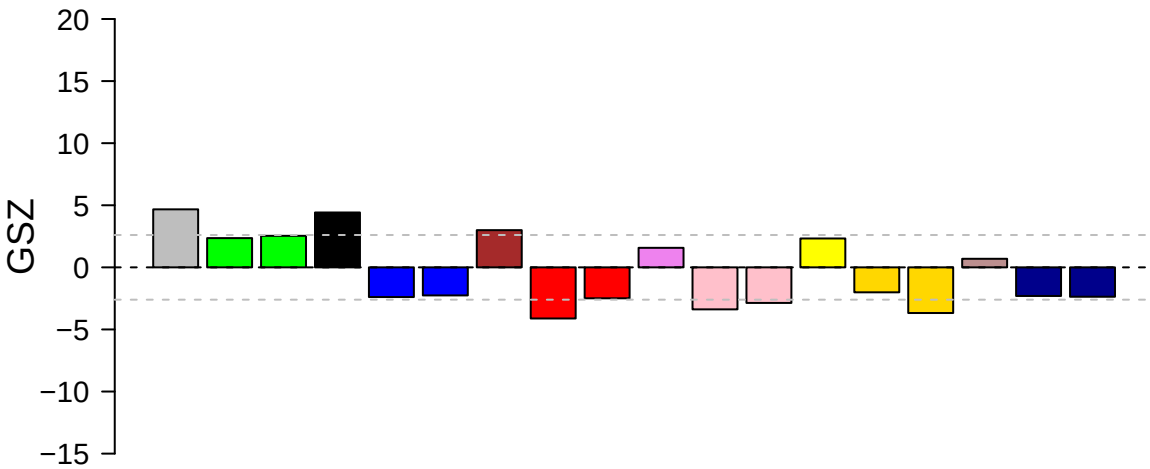
centriole

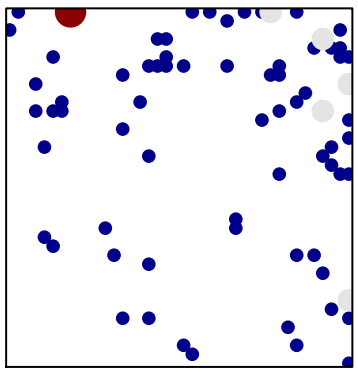




features = 20 , max = 2

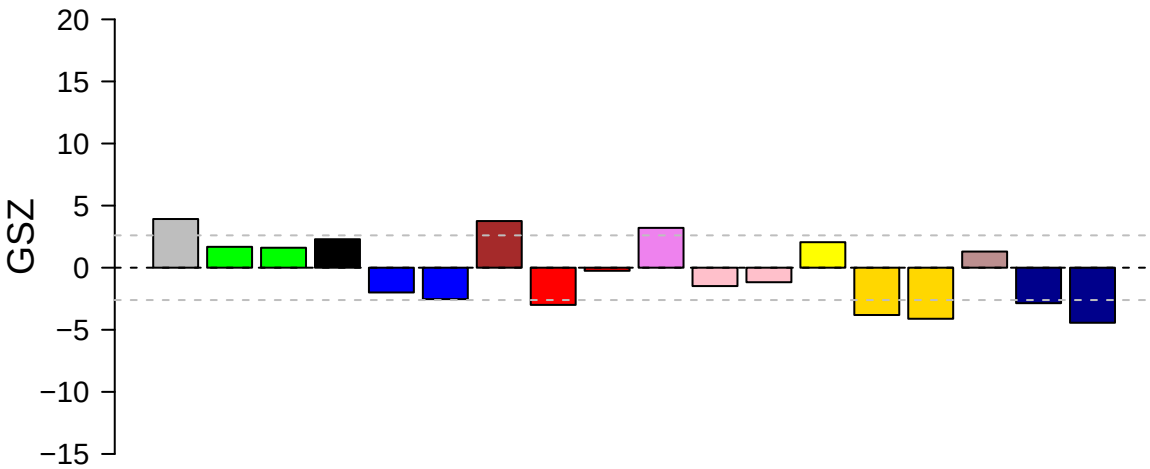
centrosome

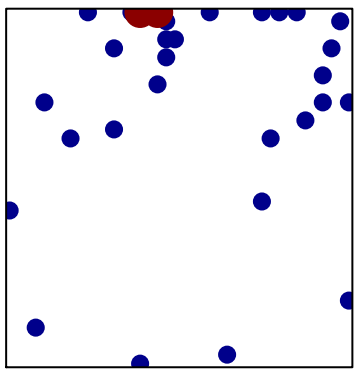




features = 80 , max = 3

chromatin





features = 32 , max = 2