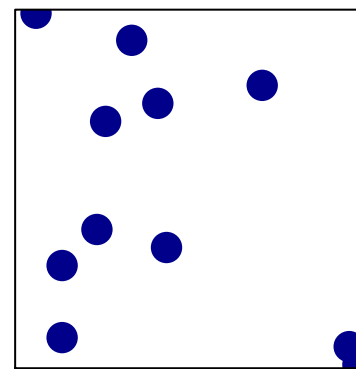
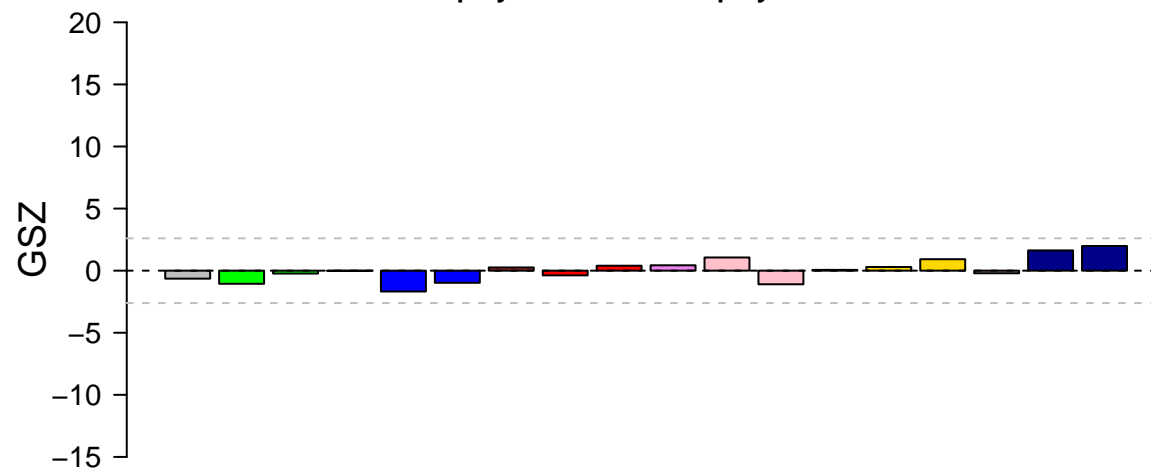
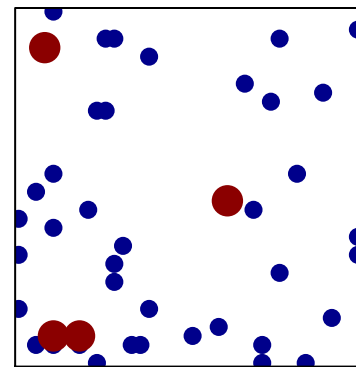
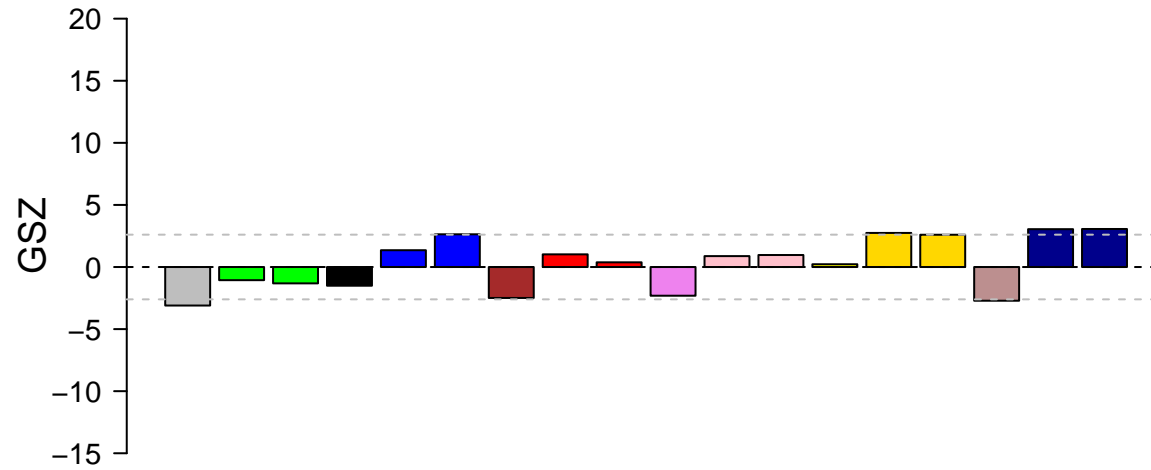


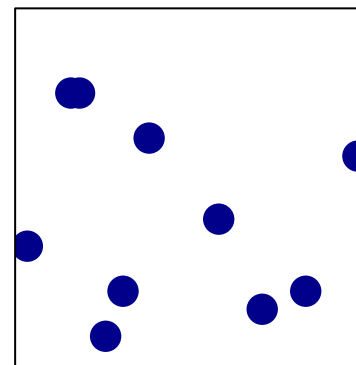
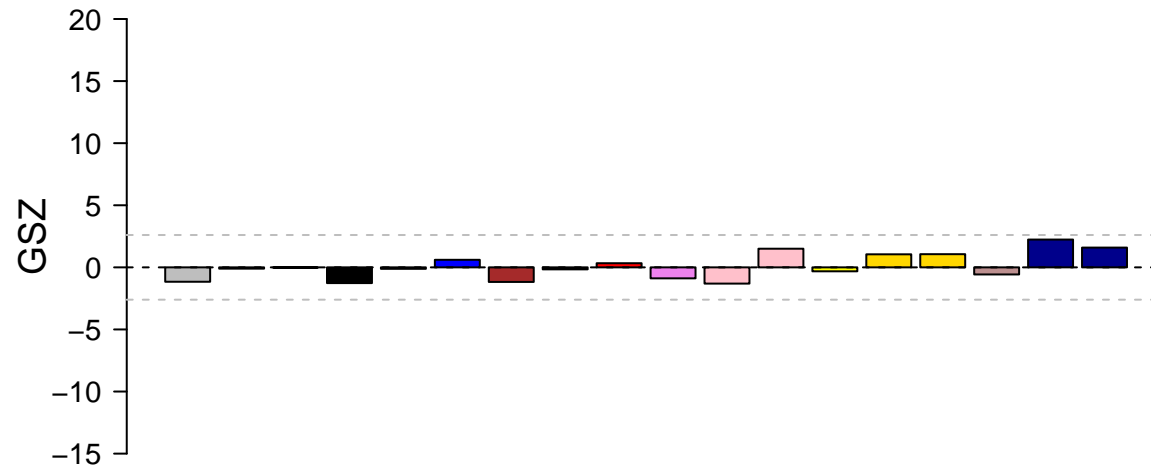
actin polymerization or depolymerization



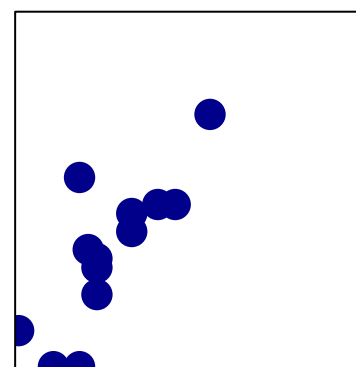
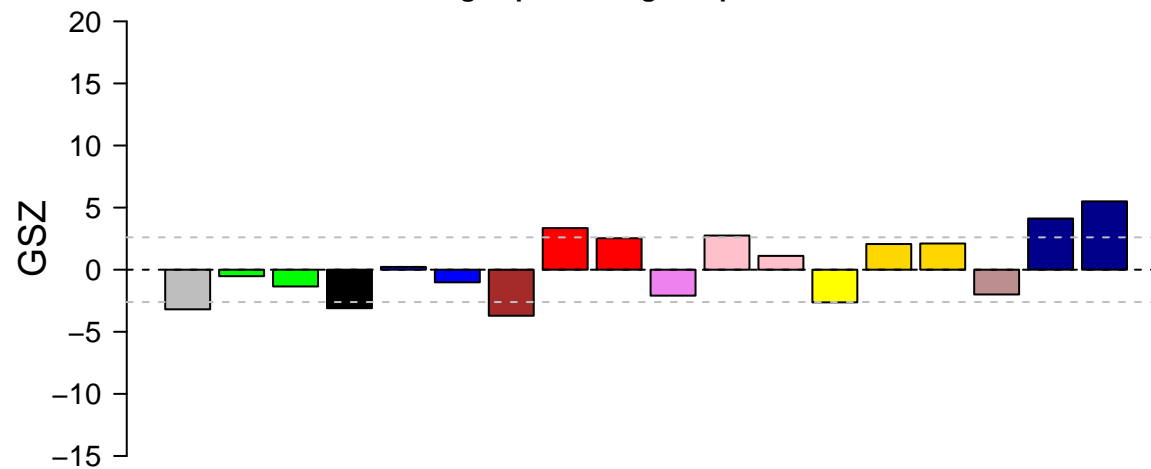
activation of GTPase activity



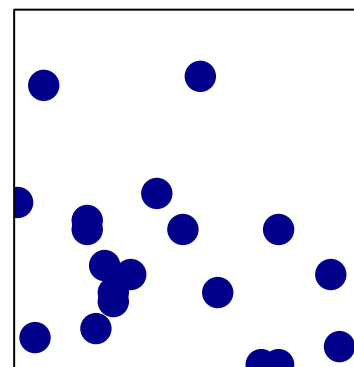
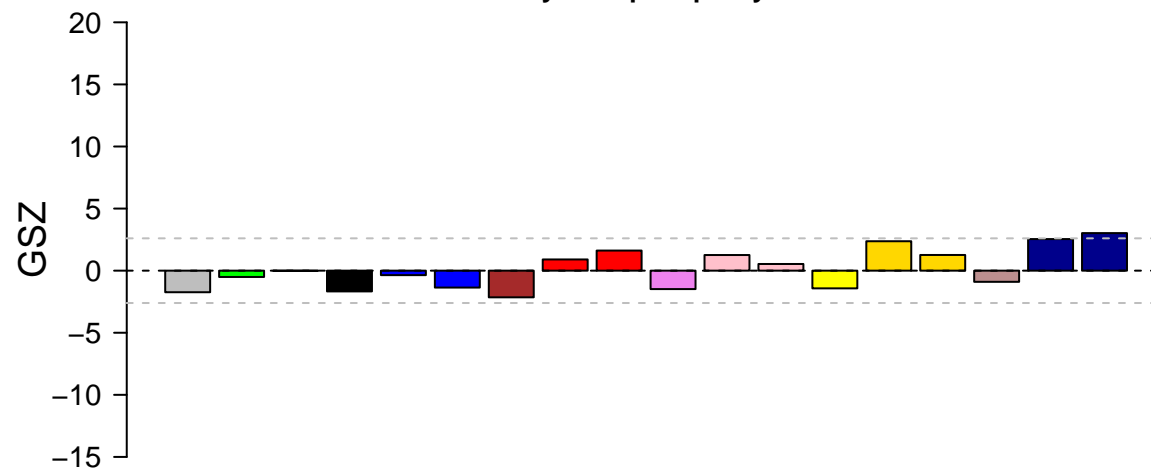
activation of MAPK activity



antigen processing and presentation

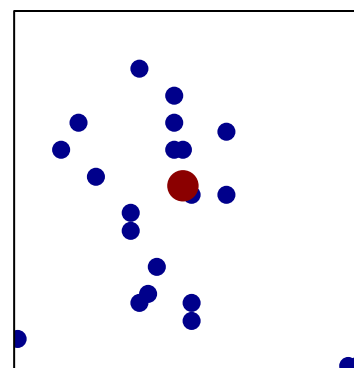
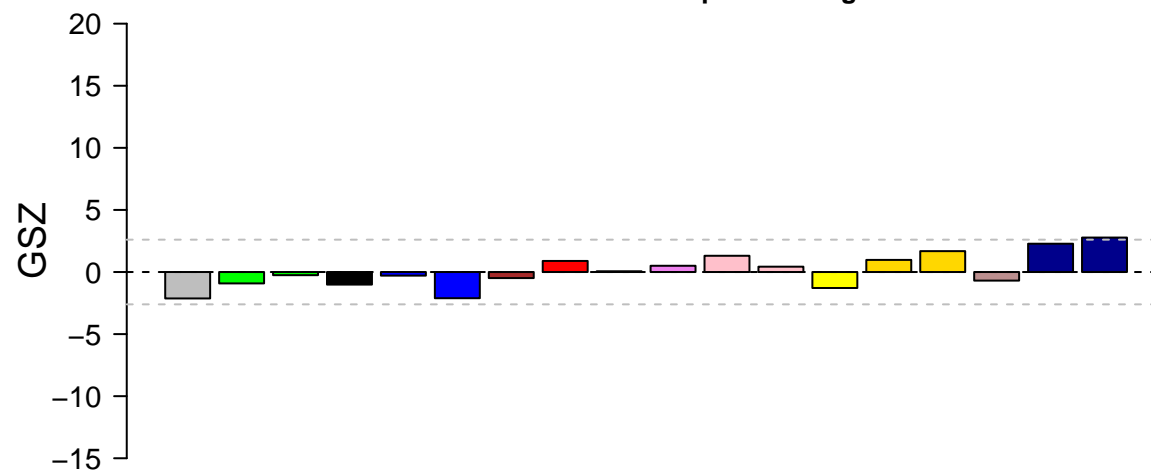


carbohydrate phosphorylation



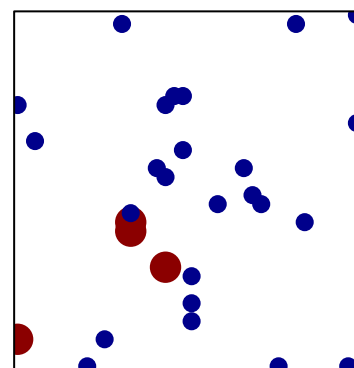
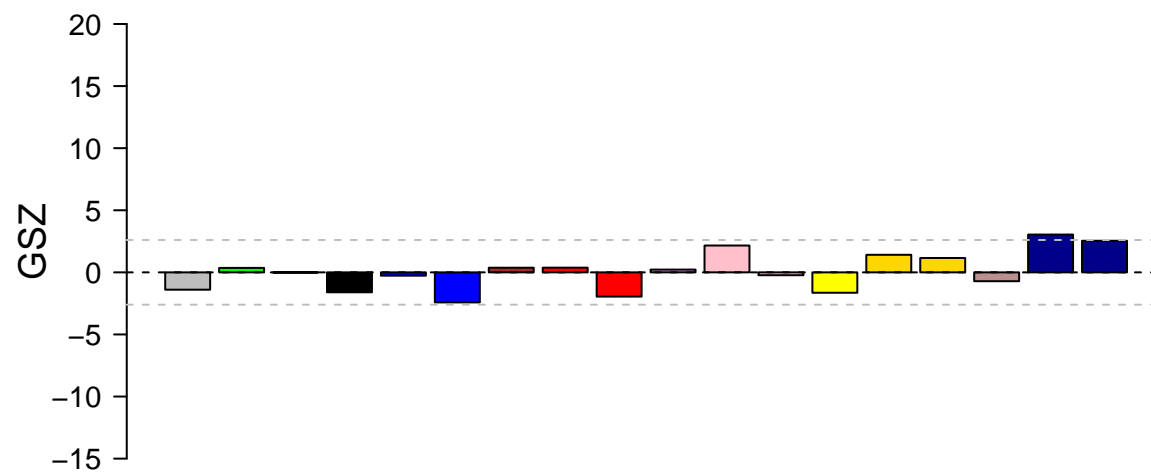
features = 19 , max = 1

CCR chemokine receptor binding



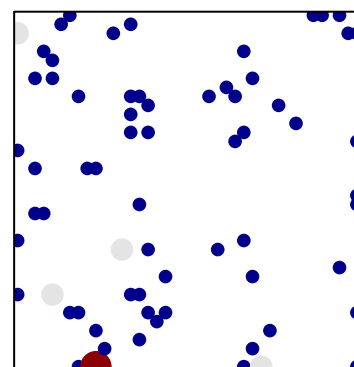
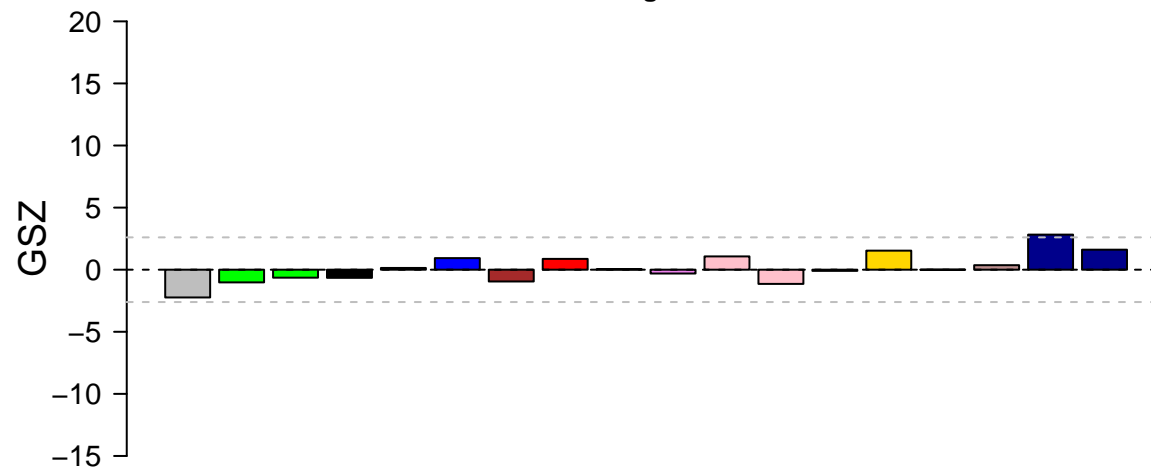
features = 23 , max = 2

cell chemotaxis



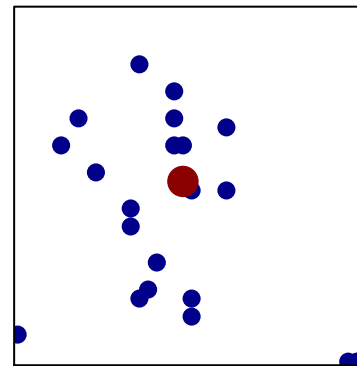
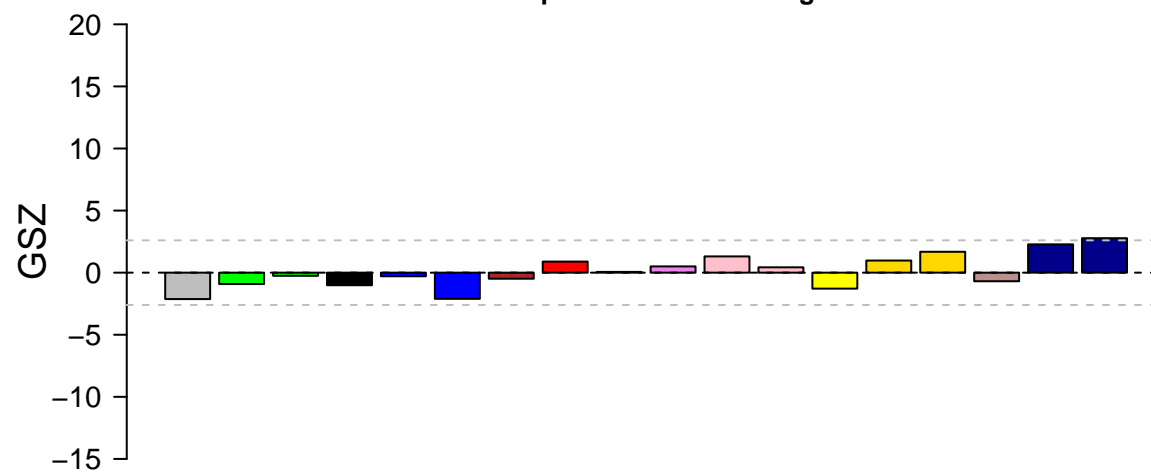
features = 33 , max = 2

cell migration



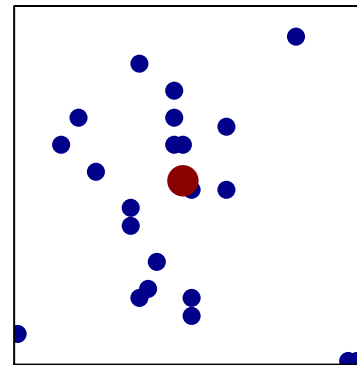
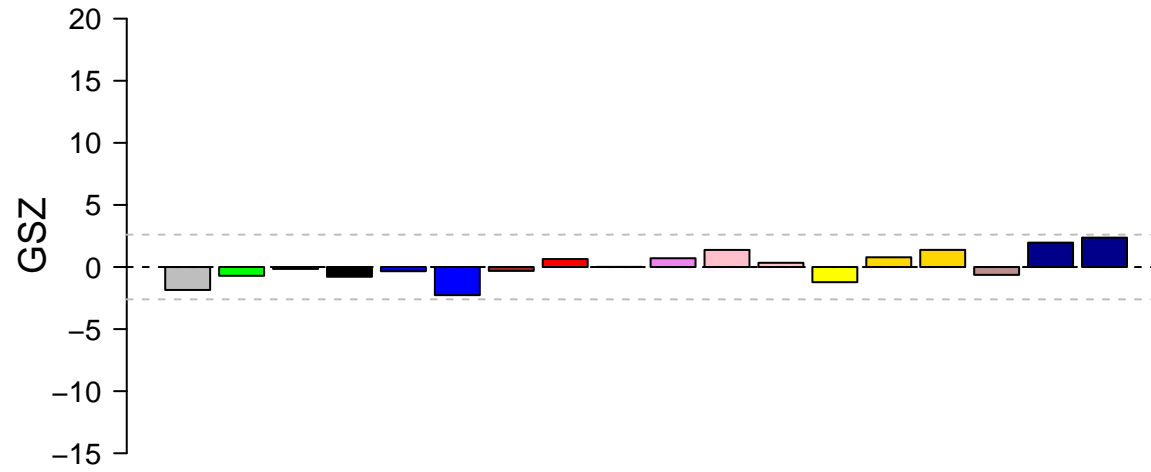
features = 74 , max = 3

cellular response to interferon- γ



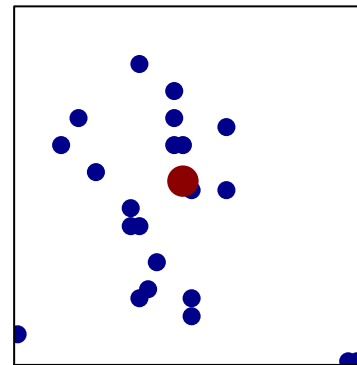
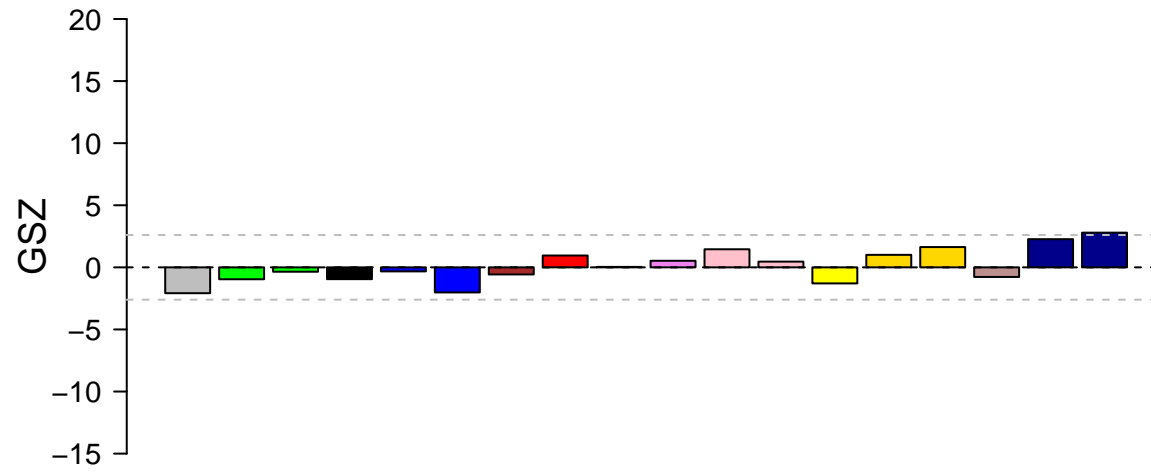
features = 23 , max = 2

cellular response to interleukin-1



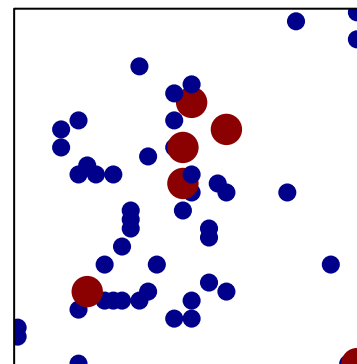
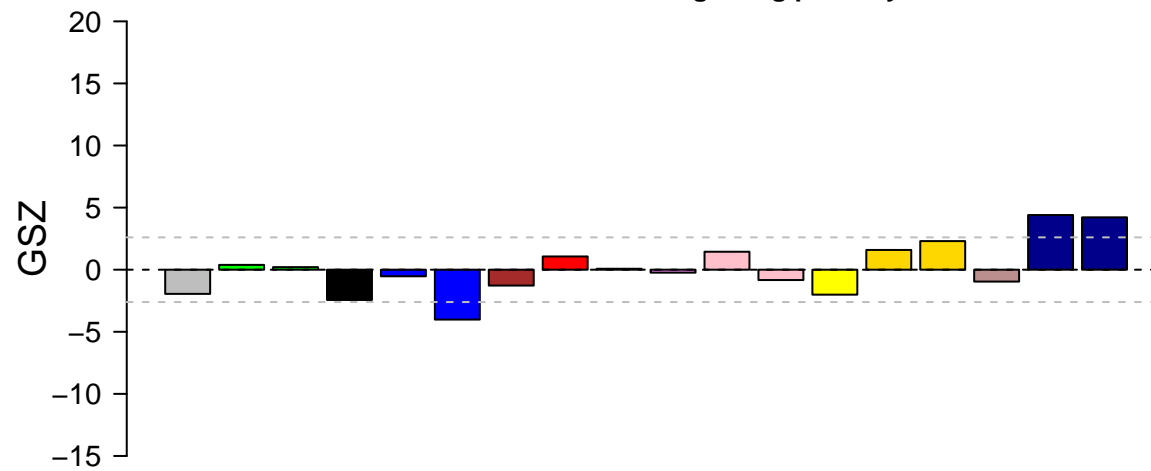
features = 24 , max = 2

cellular response to tumor necrosis factor



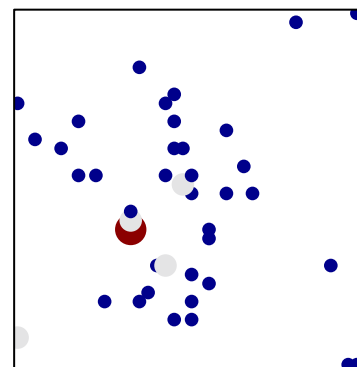
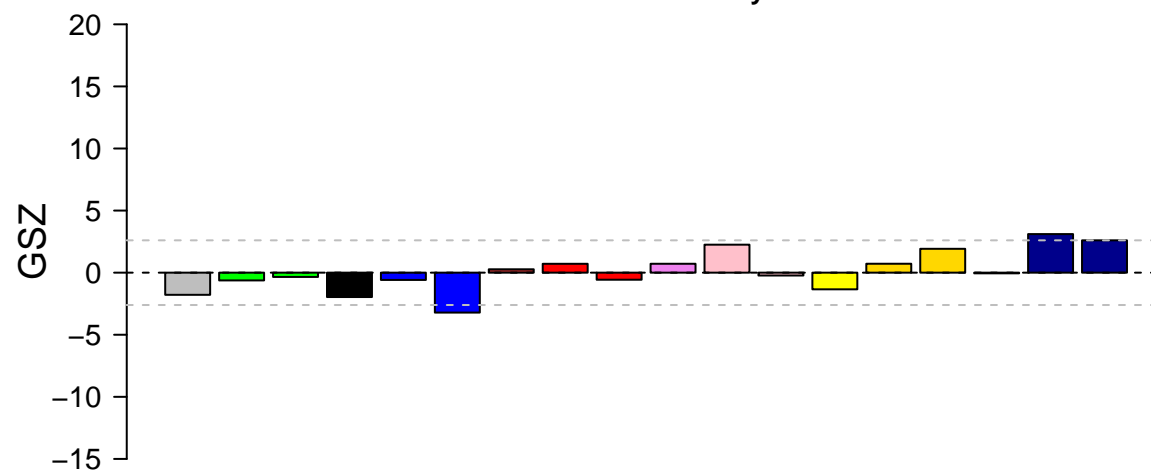
features = 24 , max = 2

chemokine-mediated signaling pathway



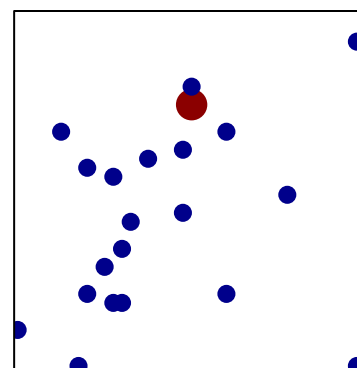
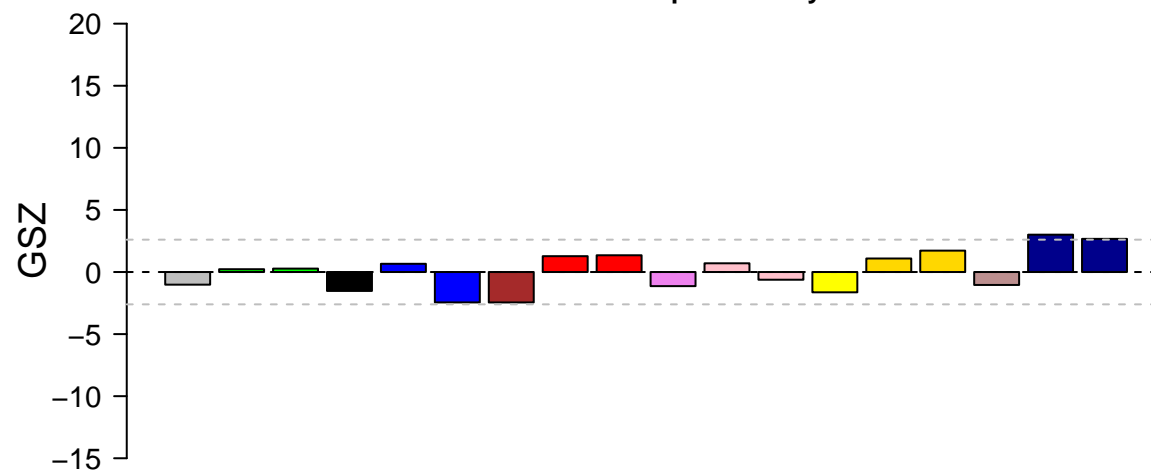
features = 58 , max = 2

chemokine activity



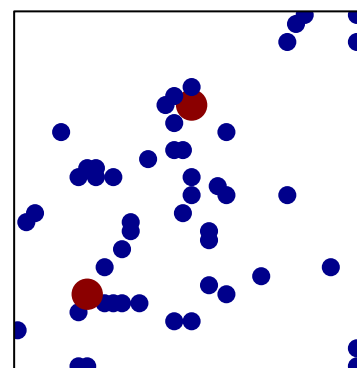
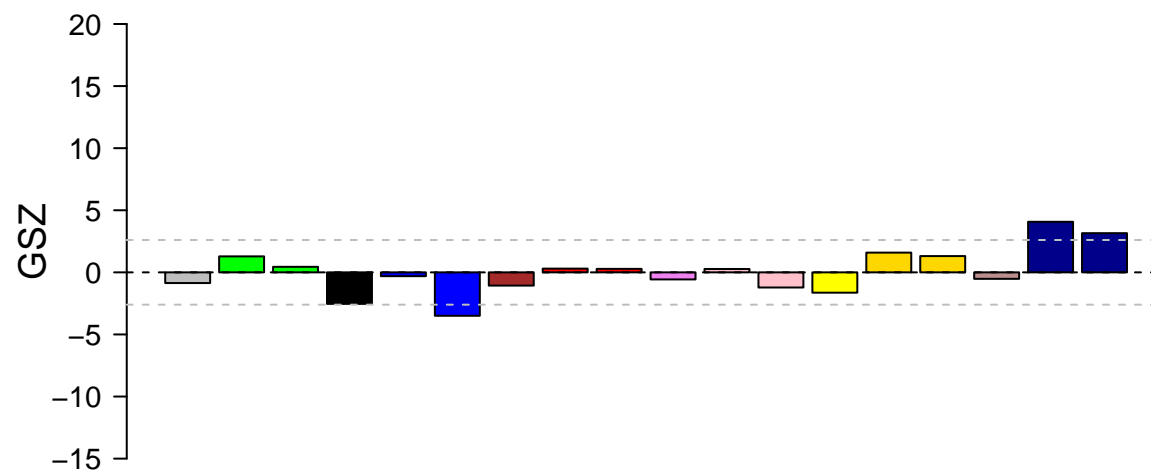
features = 47 , max = 3

chemokine receptor activity



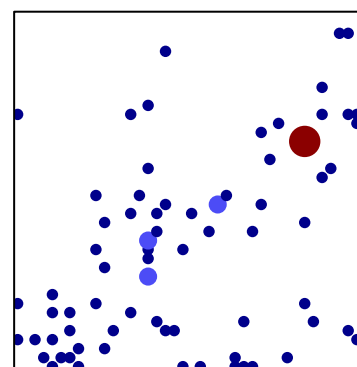
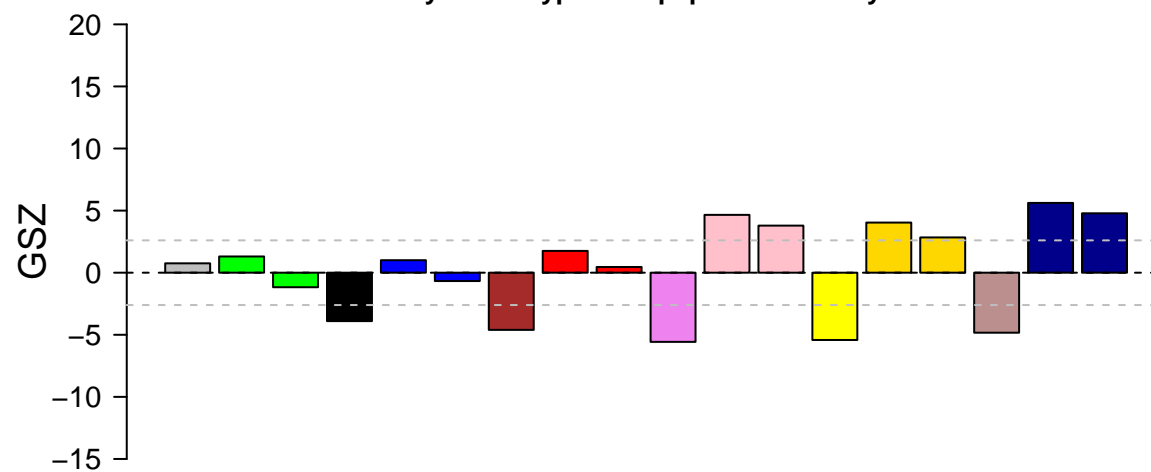
features = 22 , max = 2

chemotaxis



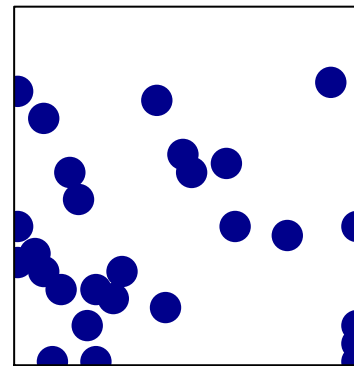
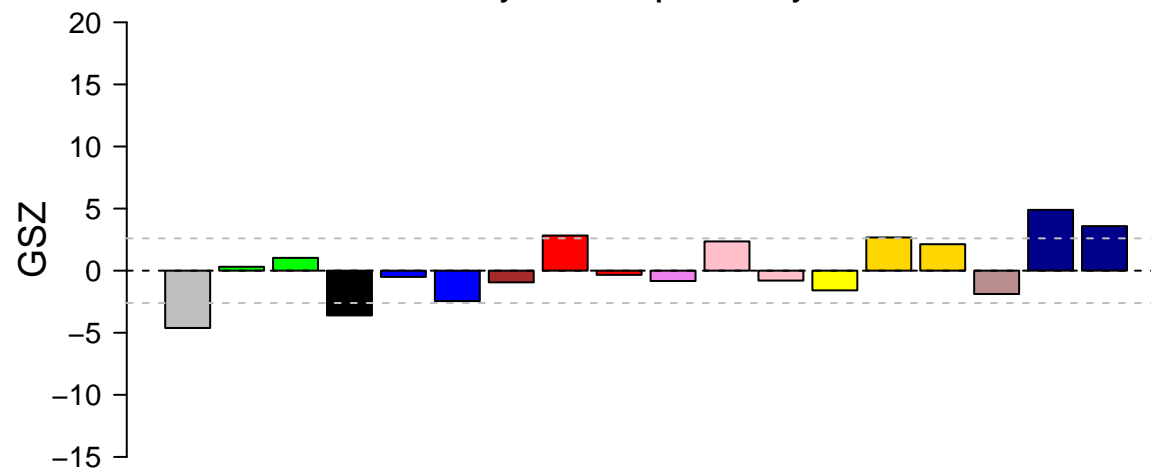
features = 53 , max = 2

cysteine-type endopeptidase activity



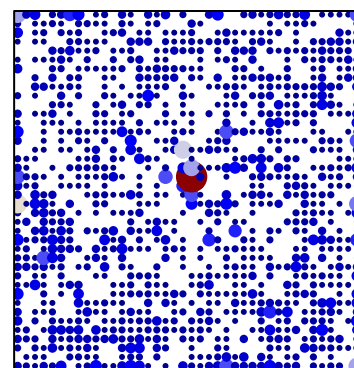
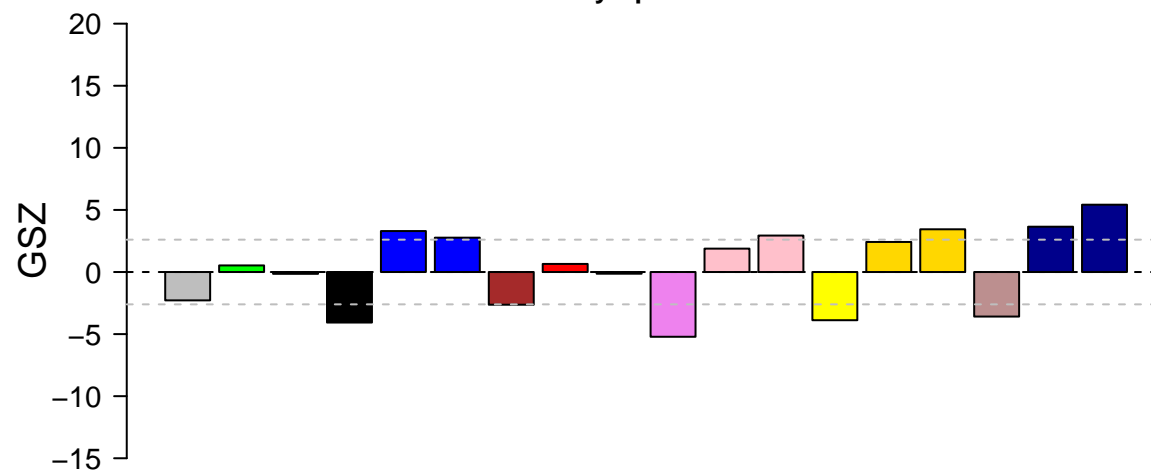
features = 78 , max = 4

cytokine receptor activity



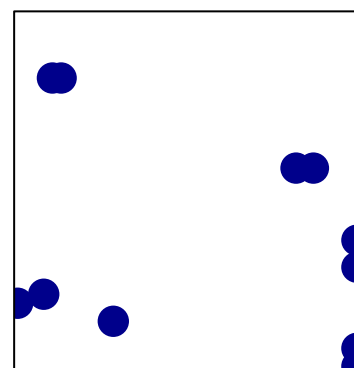
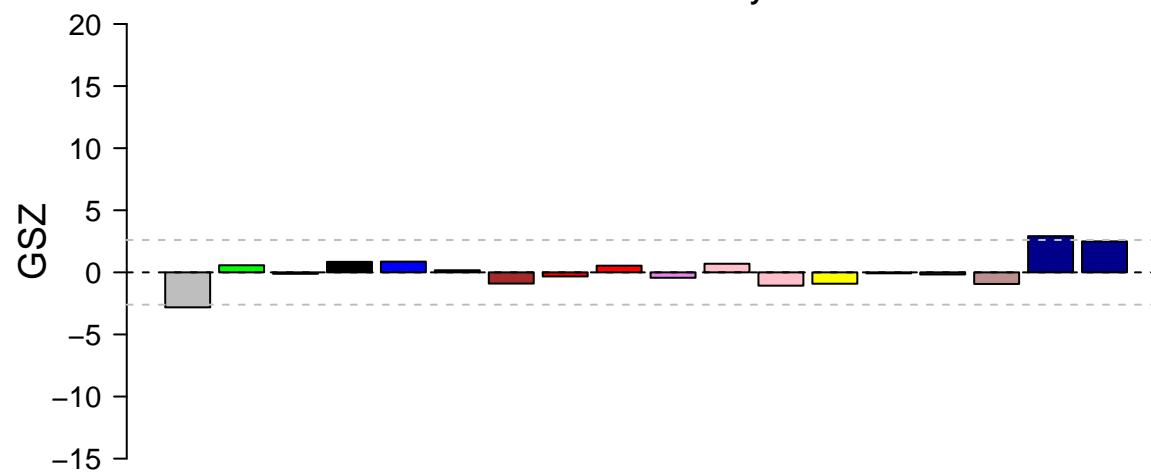
features = 27 , max = 1

cytoplasm



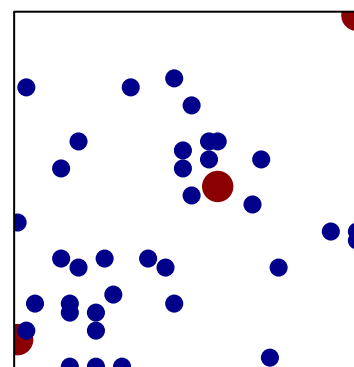
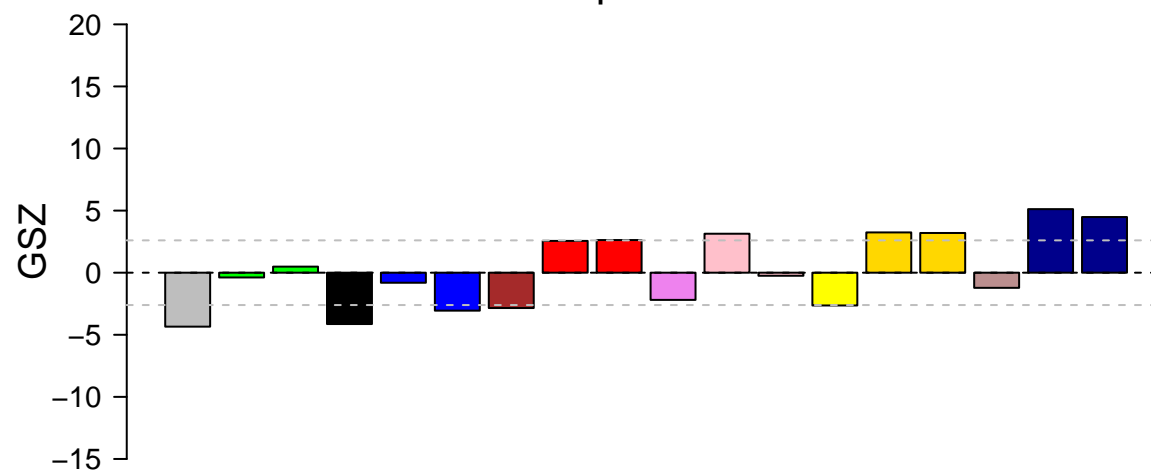
features = 2189 , max = 22

deaminase activity



features = 11 , max = 1

defense response to bacterium



features = 42 , max = 2