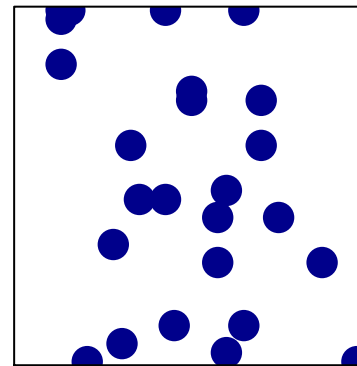
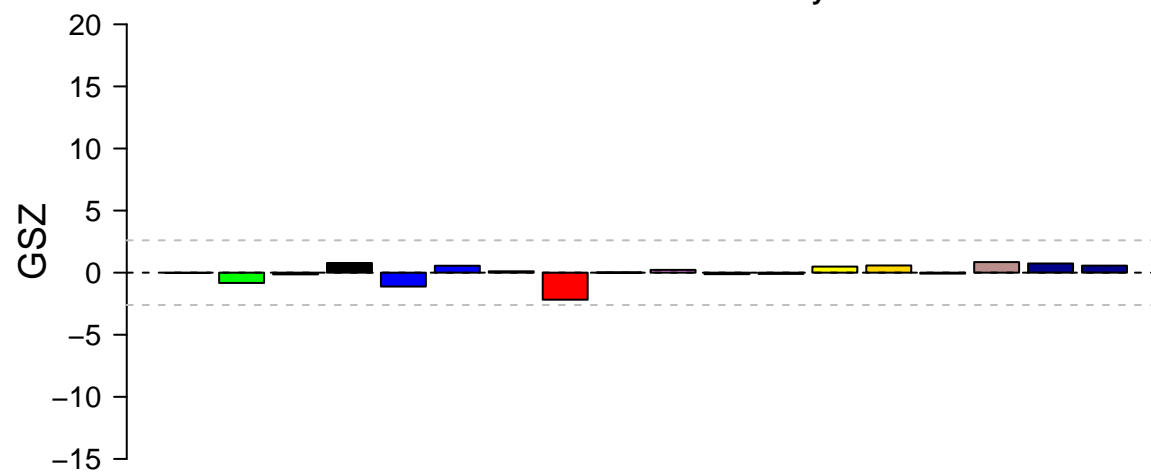
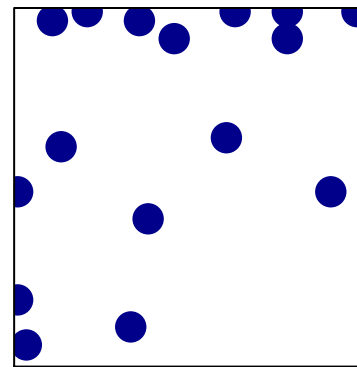
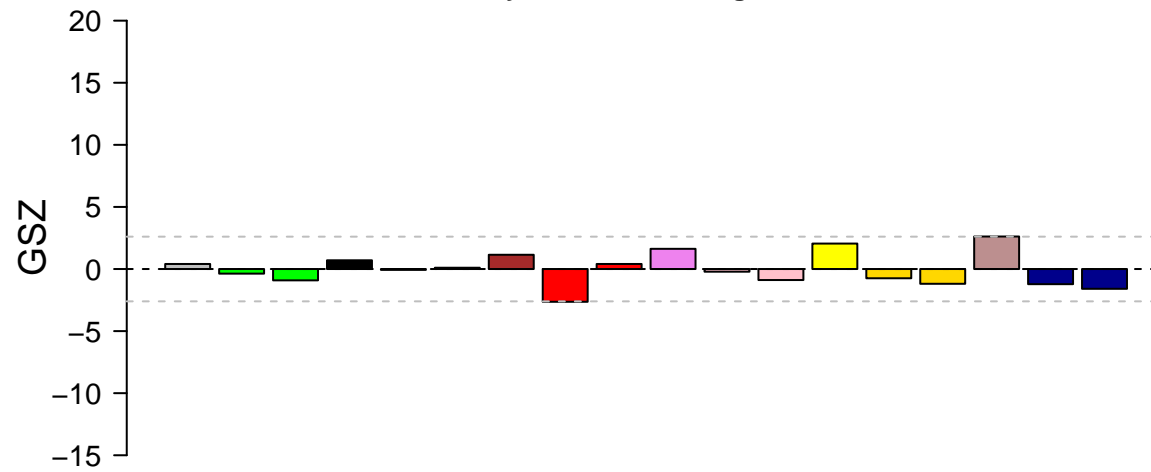


activation of MAPKK activity



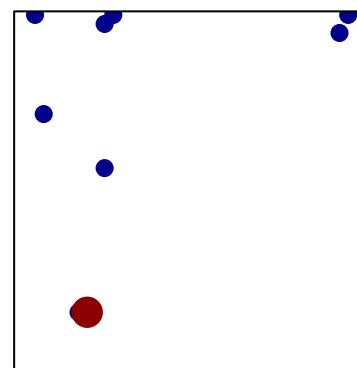
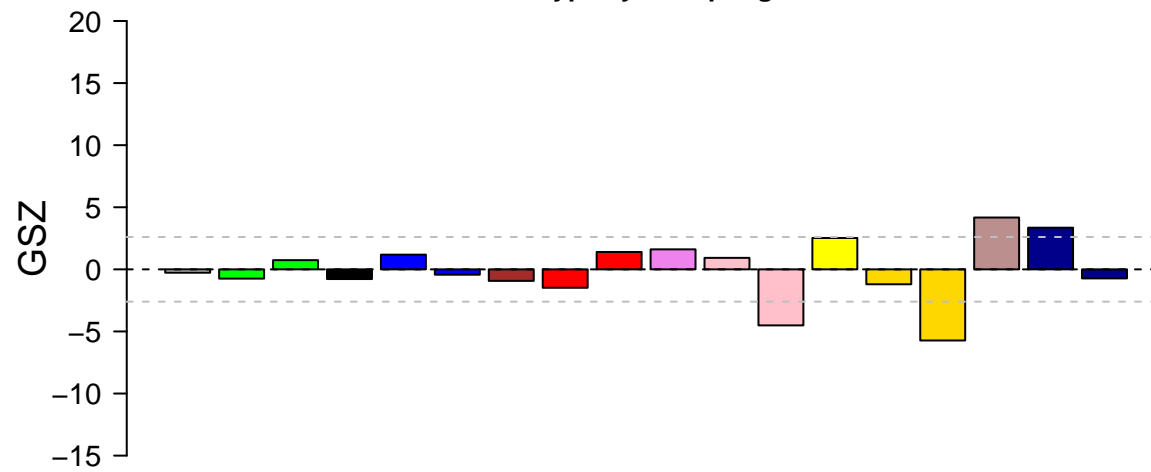
features = 25 , max = 1

actomyosin structure organization



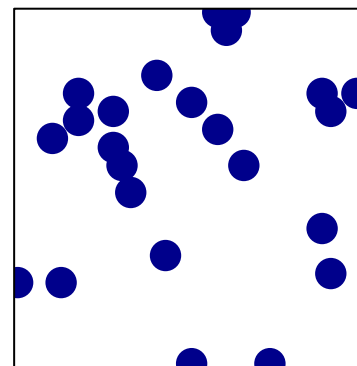
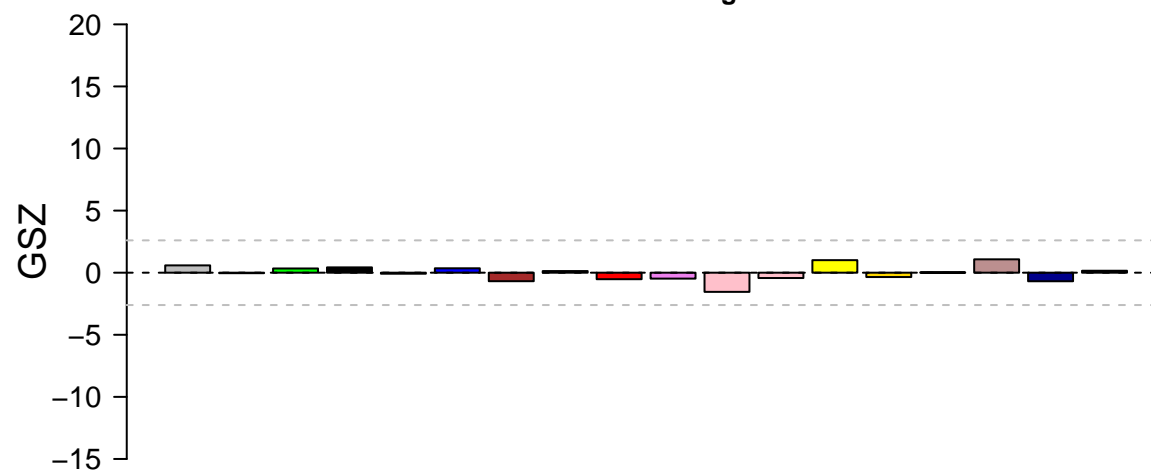
features = 16 , max = 1

camera-type eye morphogenesis



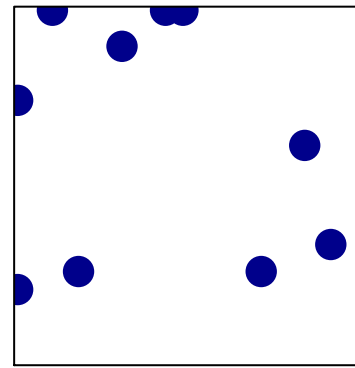
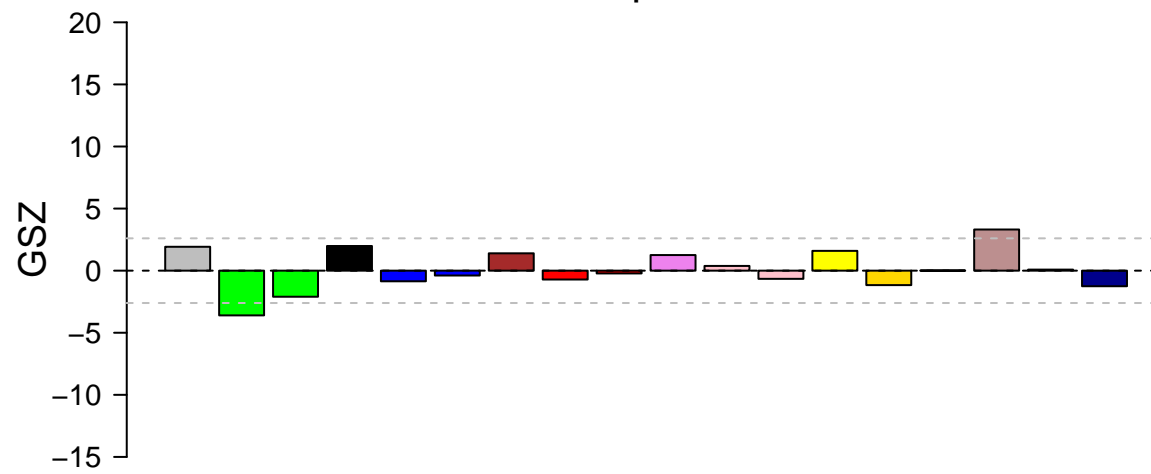
features = 10 , max = 2

cation binding



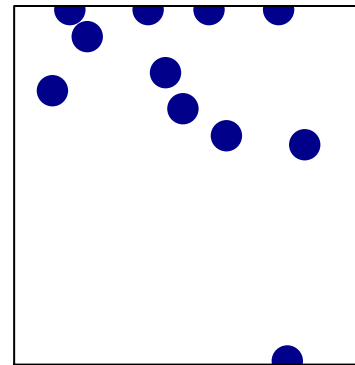
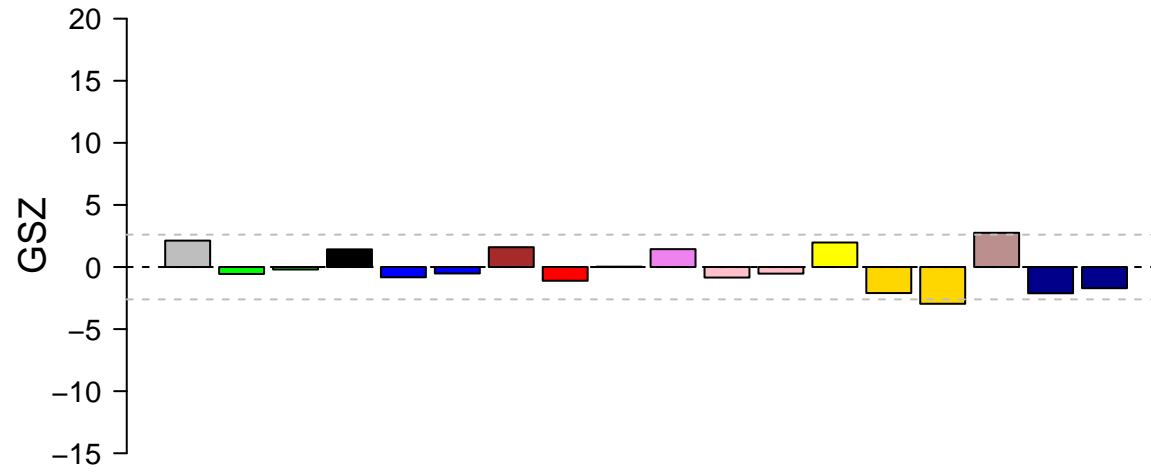
features = 24 , max = 1

cell fate specification



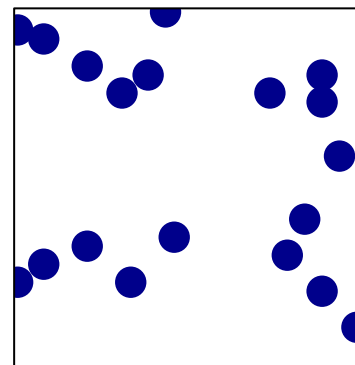
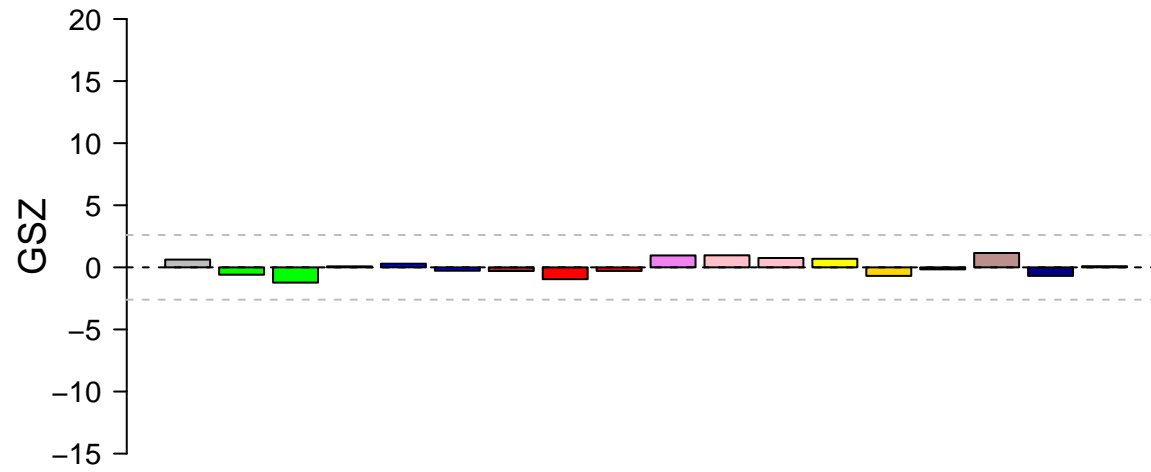
features = 10 , max = 1

determination of dorsal identity



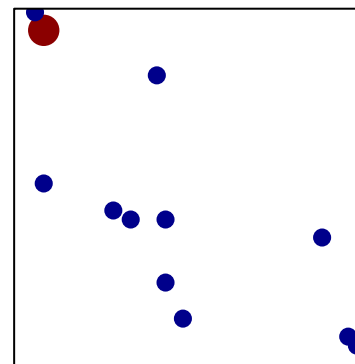
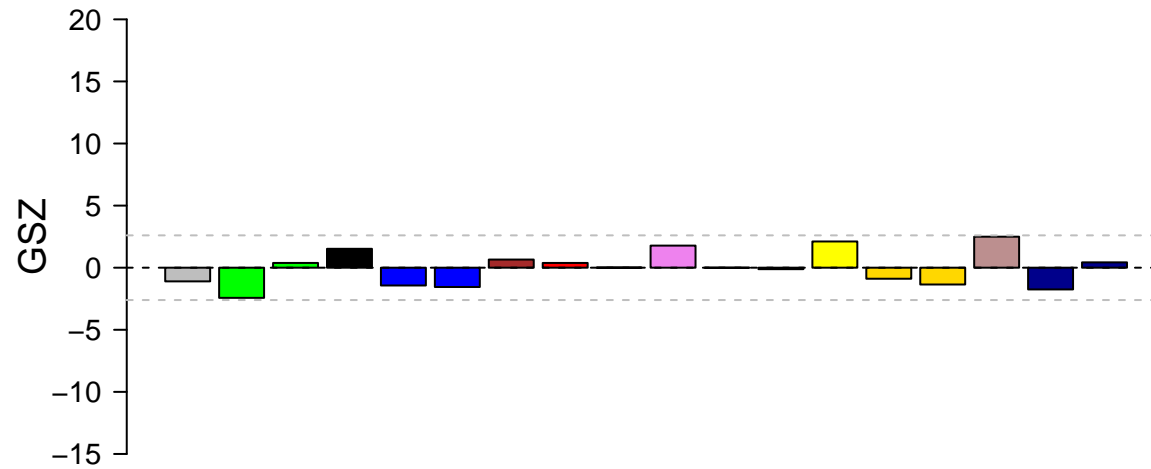
features = 11 , max = 1

developmental process



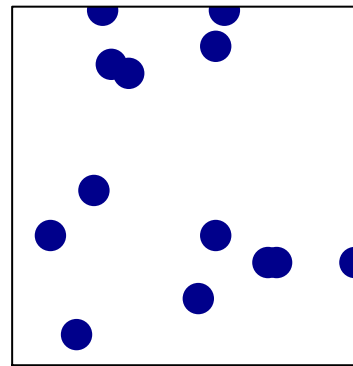
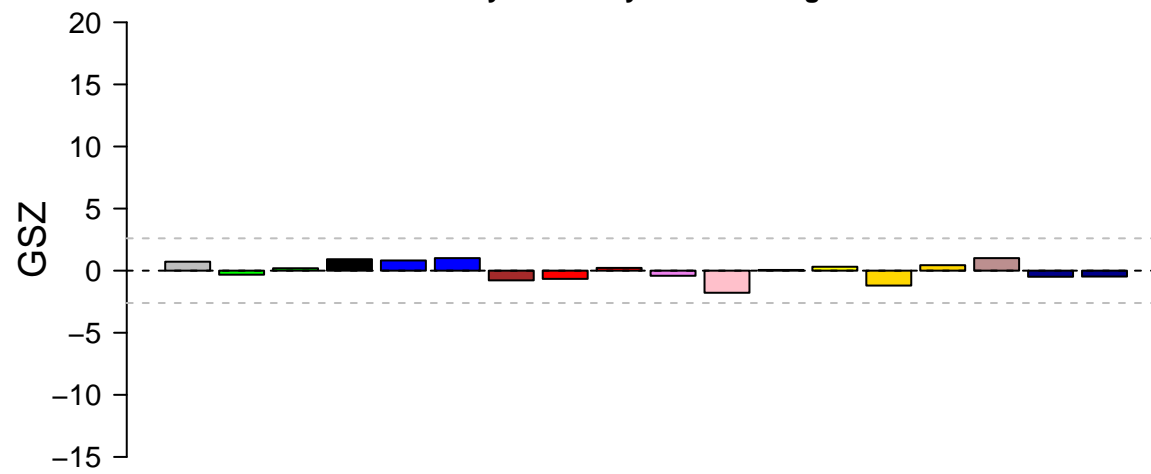
features = 19 , max = 1

dynamins family protein polymerization involved in mitochondrial fission



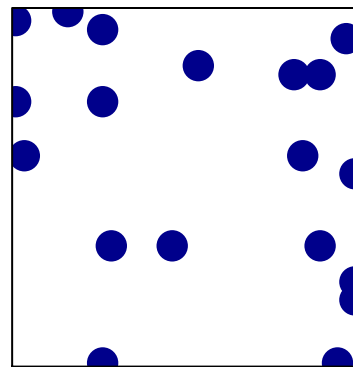
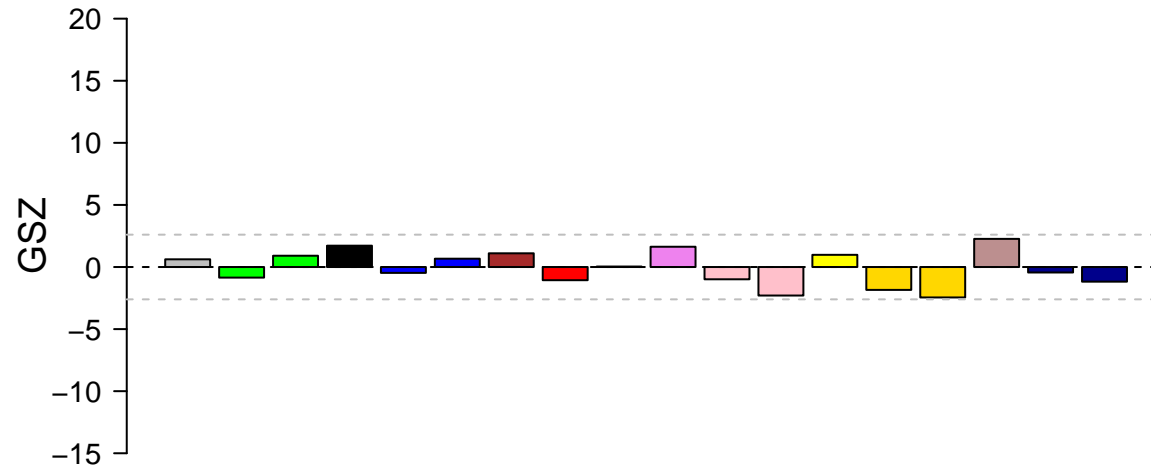
features = 13 , max = 2

dynein heavy chain binding



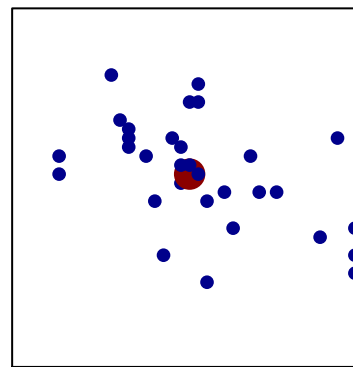
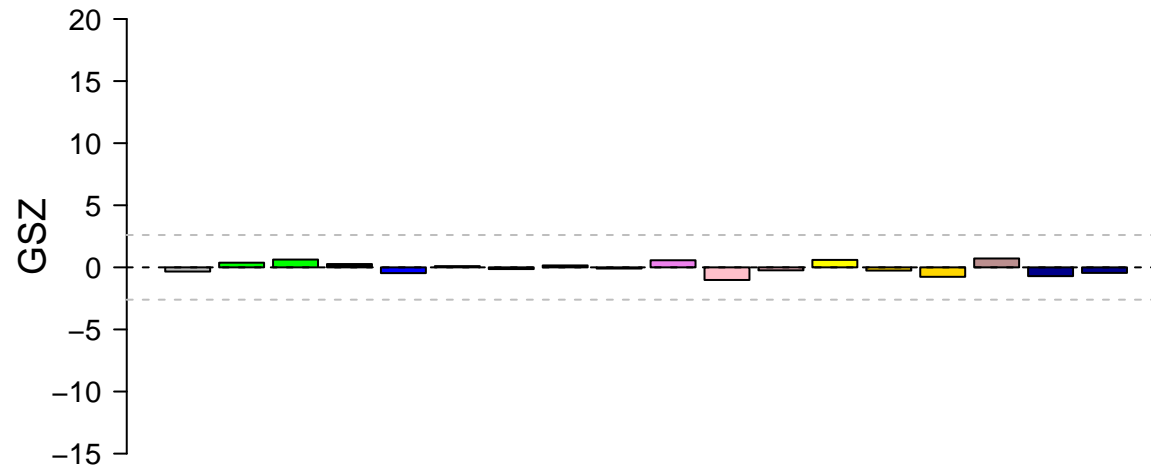
features = 13 , max = 1

eye photoreceptor cell development



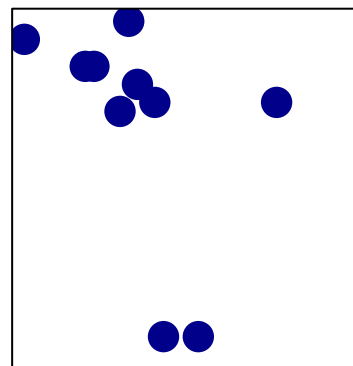
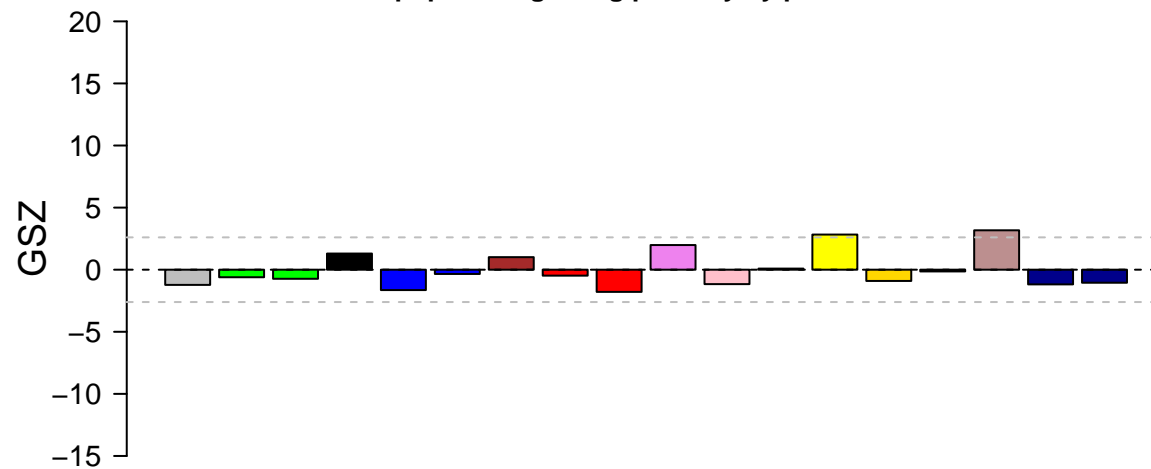
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# features = 19 , max = 1
```

G-protein coupled serotonin receptor activity



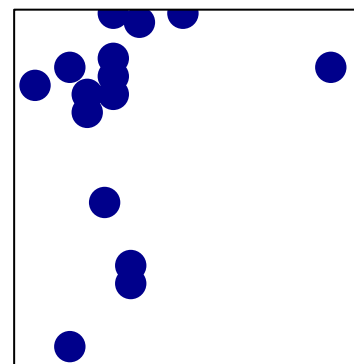
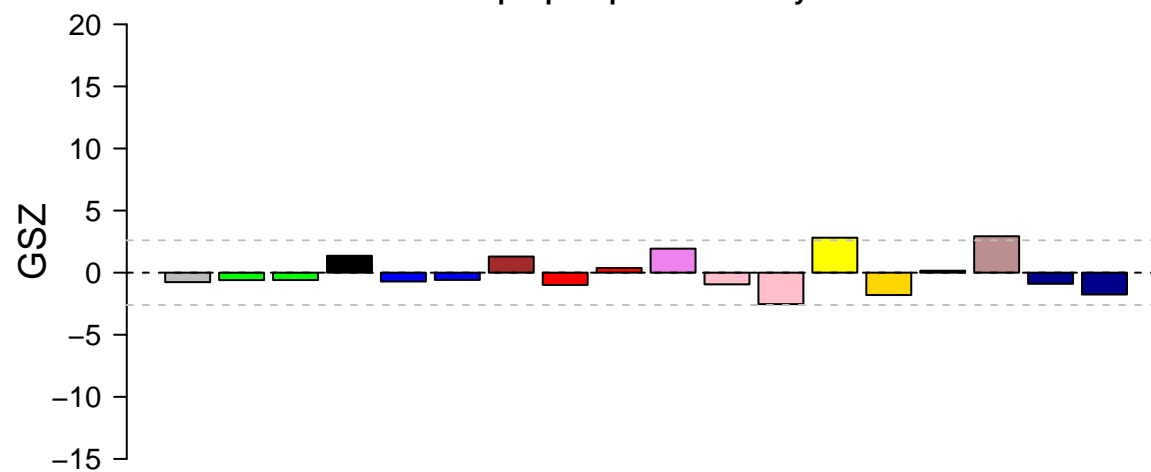
features = 34 , max = 3

intrinsic apoptotic signaling pathway by p53 class mediator



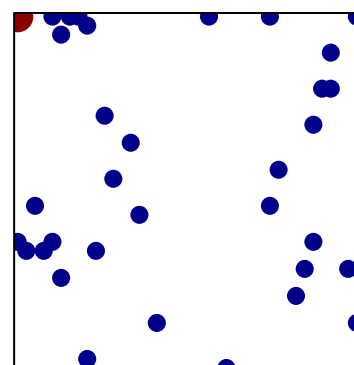
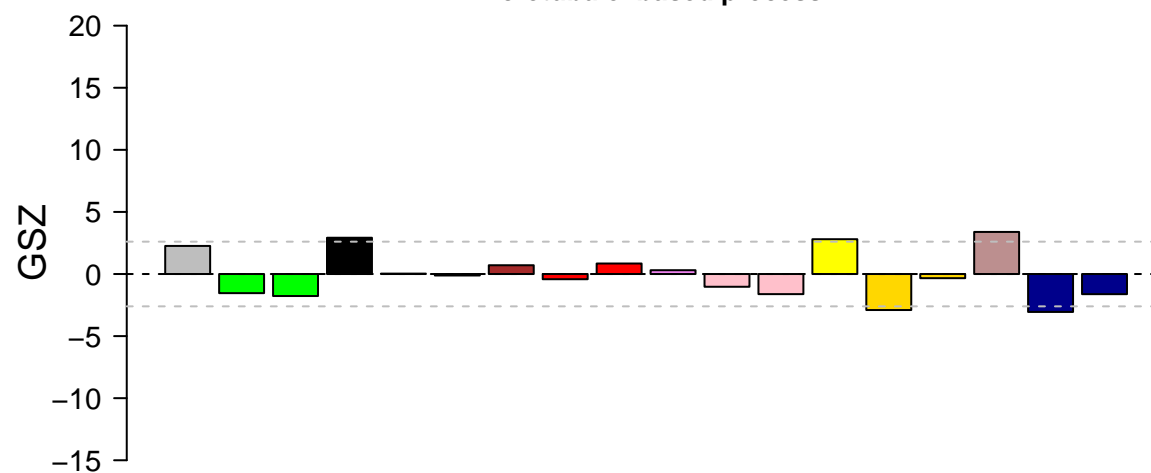
features = 10 , max = 1

lipid phosphatase activity



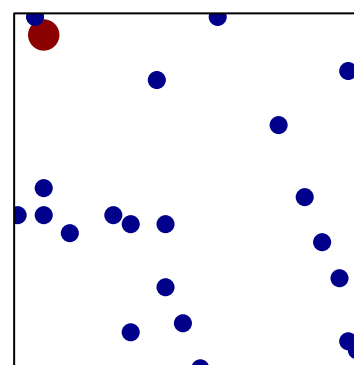
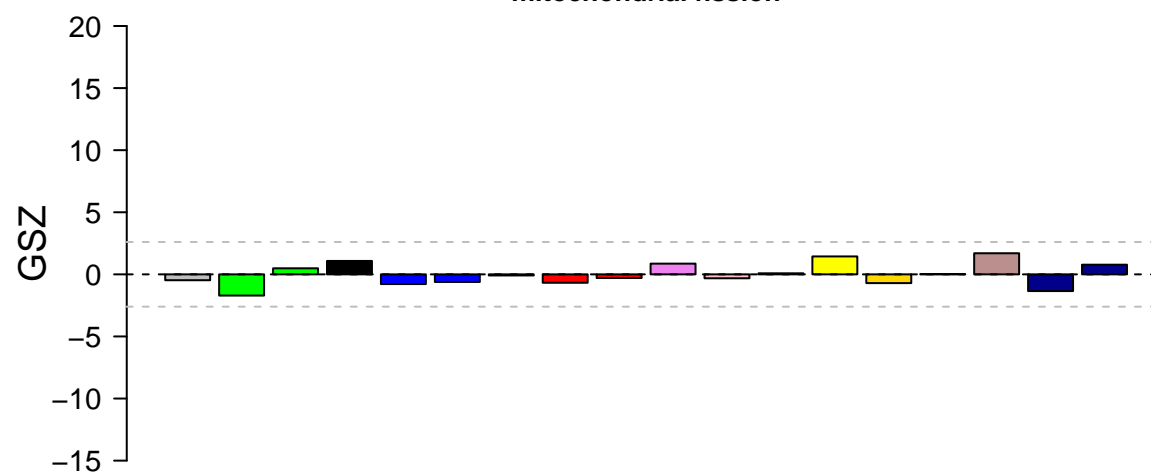
features = 15 , max = 1

microtubule-based process



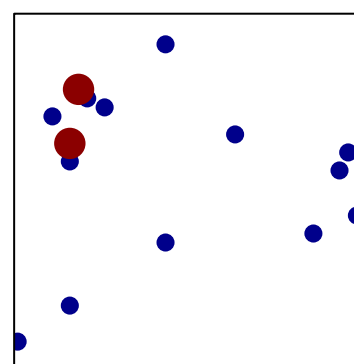
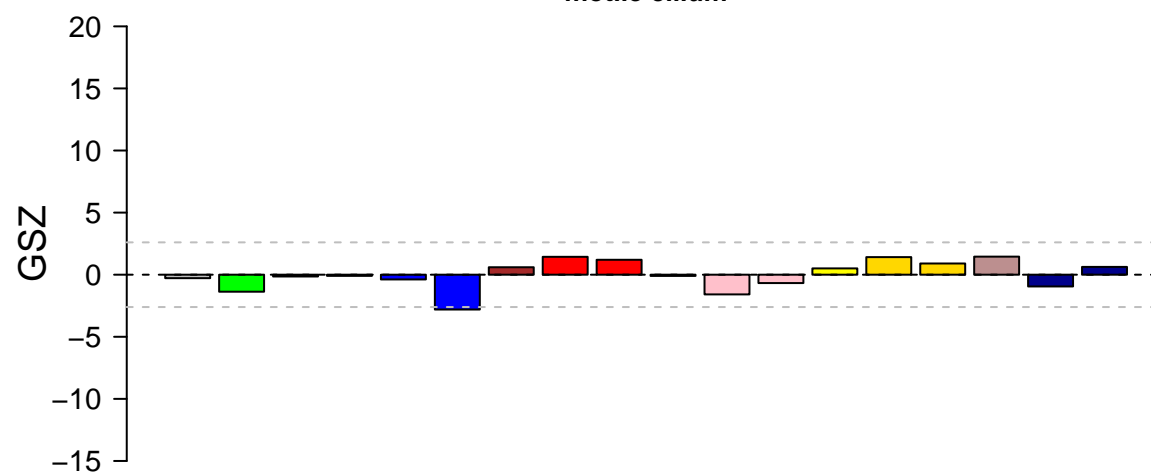
features = 35 , max = 2

mitochondrial fission



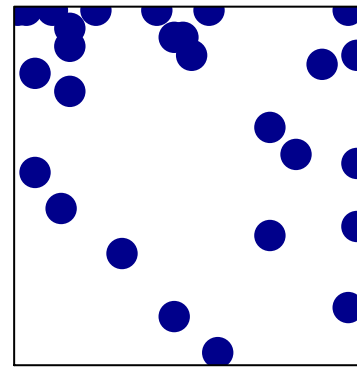
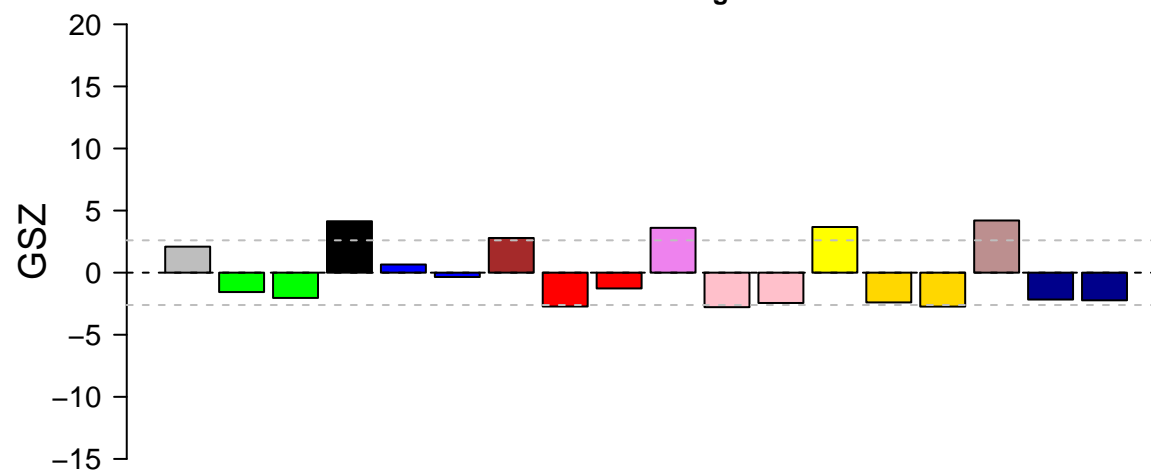
features = 23 , max = 2

motile cilium



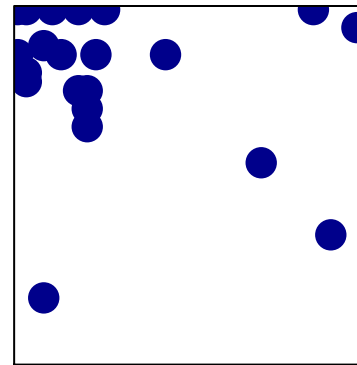
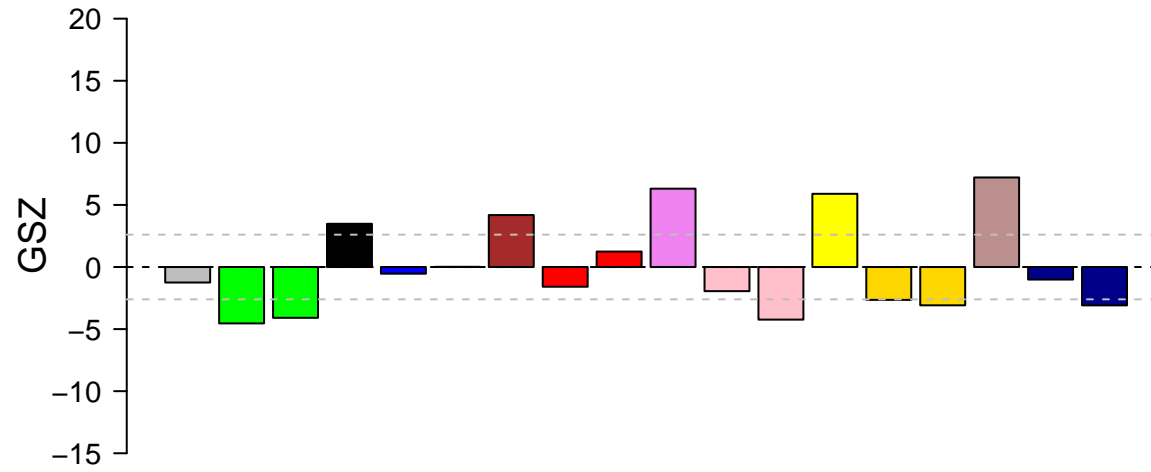
features = 18 , max = 2

motor neuron axon guidance



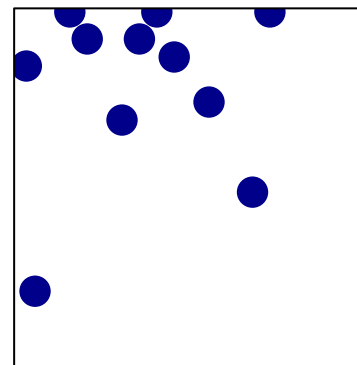
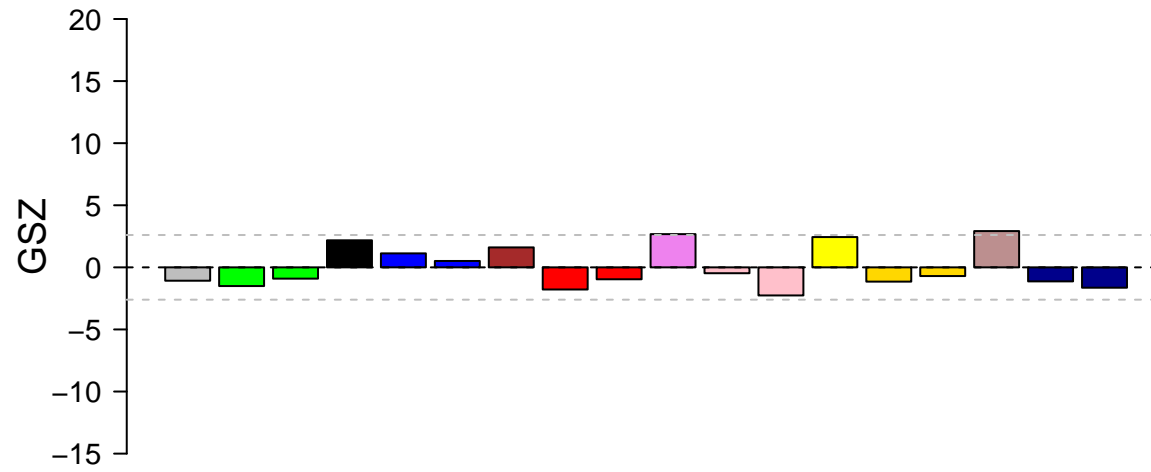
features = 27 , max = 1

negative regulation of angiogenesis



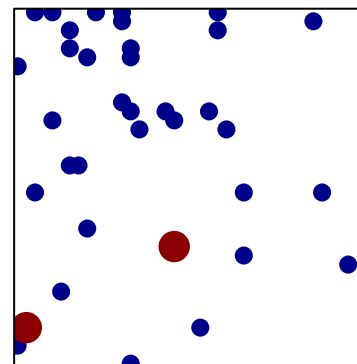
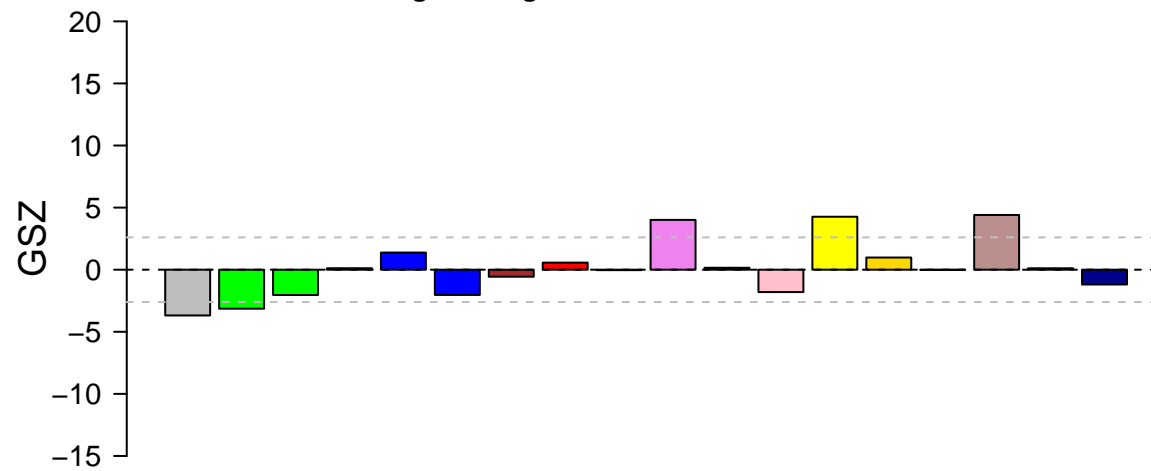
features = 21 , max = 1

negative regulation of cell differentiation



features = 11 , max = 1

negative regulation of JAK-STAT cascade



features = 38 , max = 2