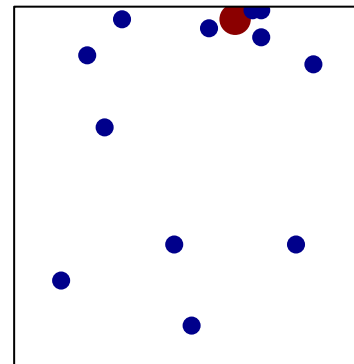
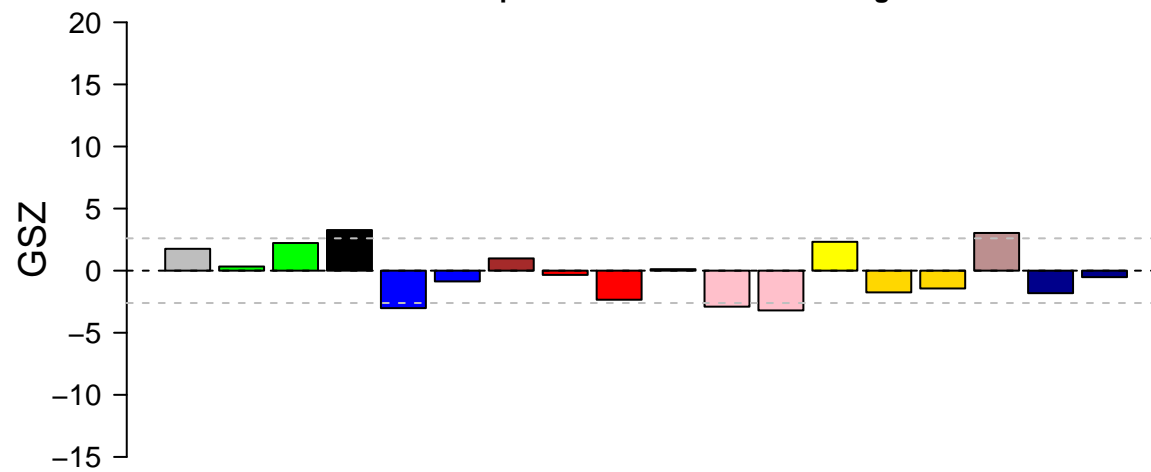
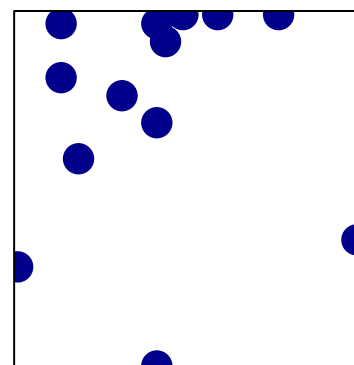
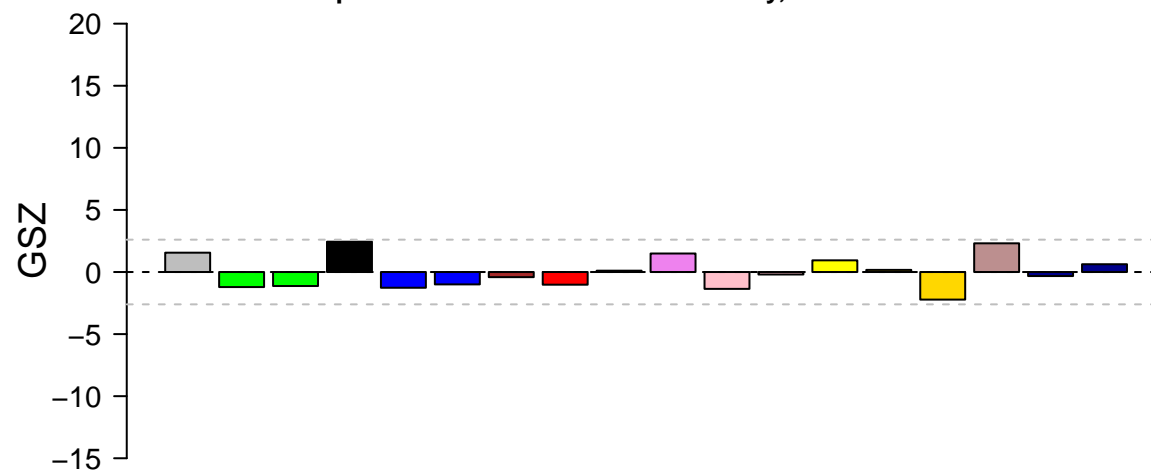


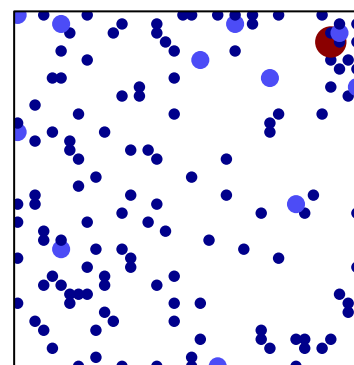
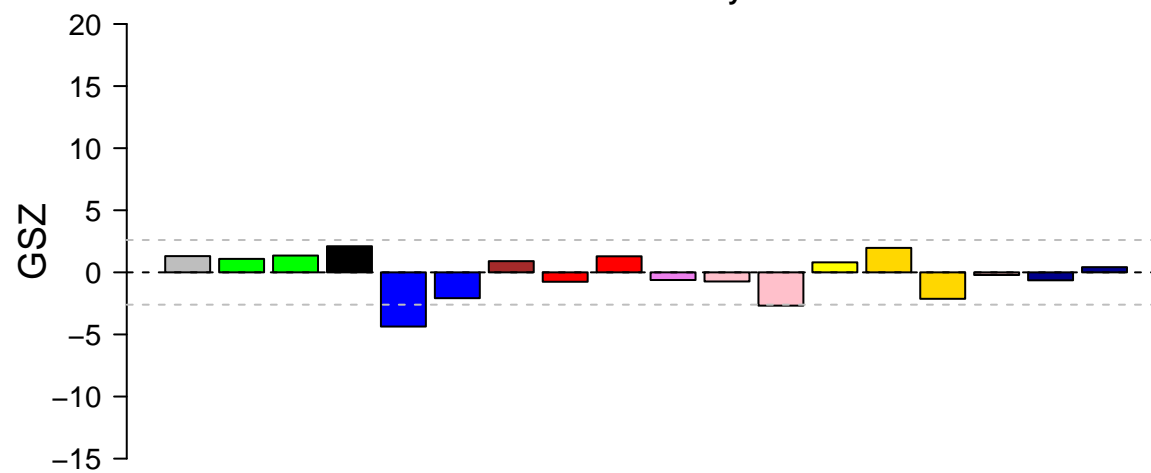
ATP-dependent chromatin remodeling



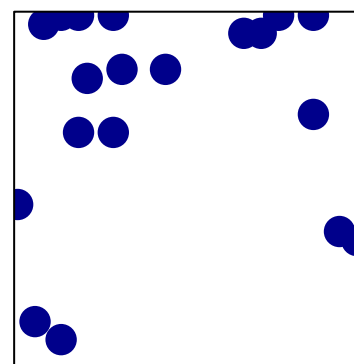
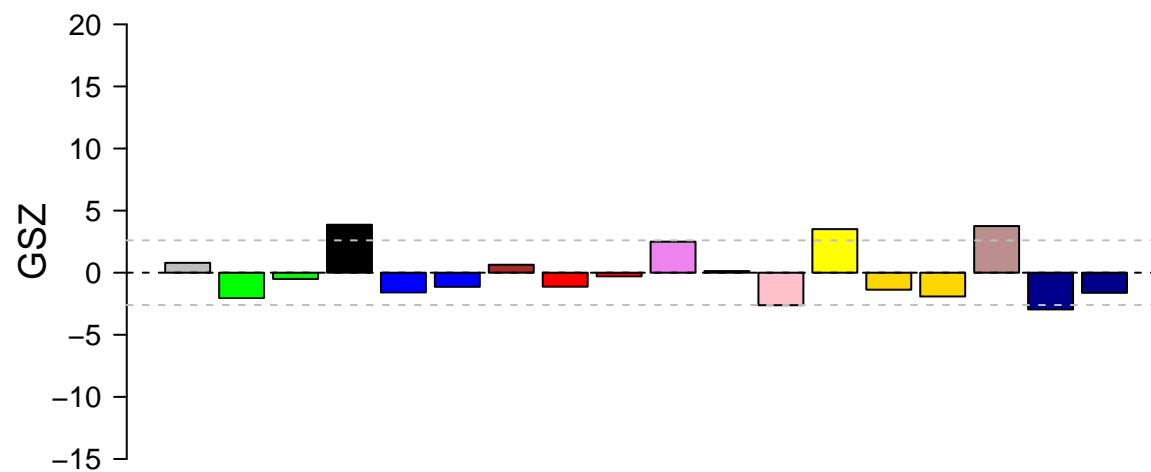
ATP-dependent microtubule motor activity, minus-end-directed



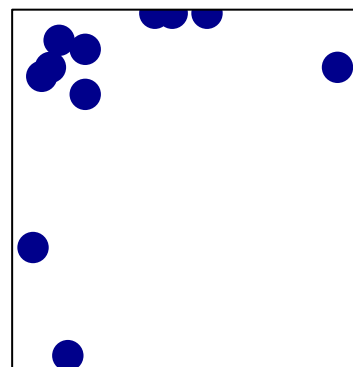
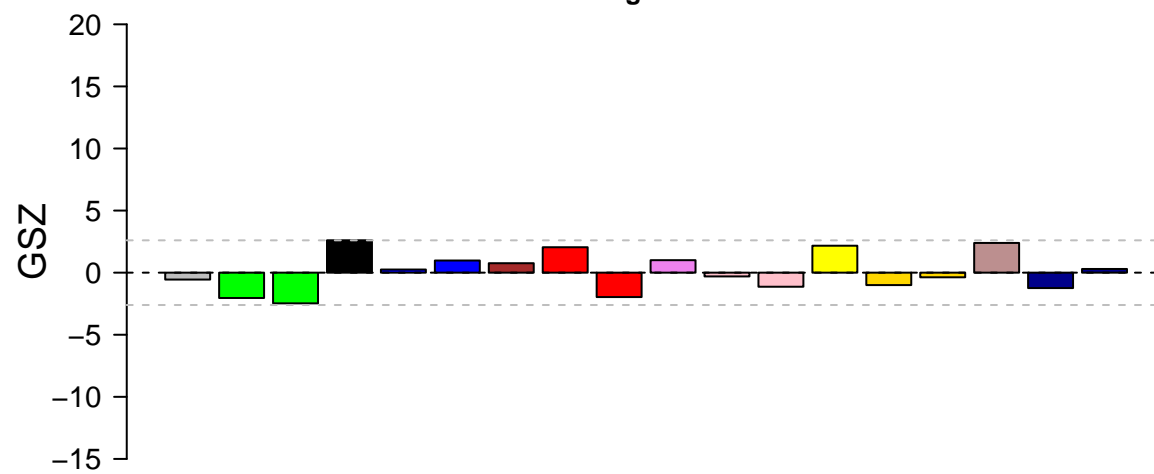
ATPase activity



axon

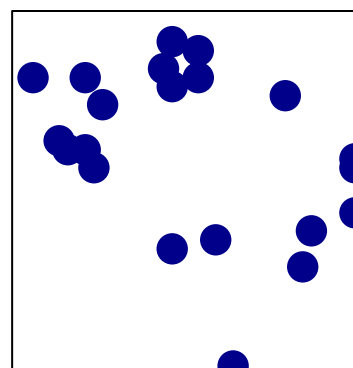
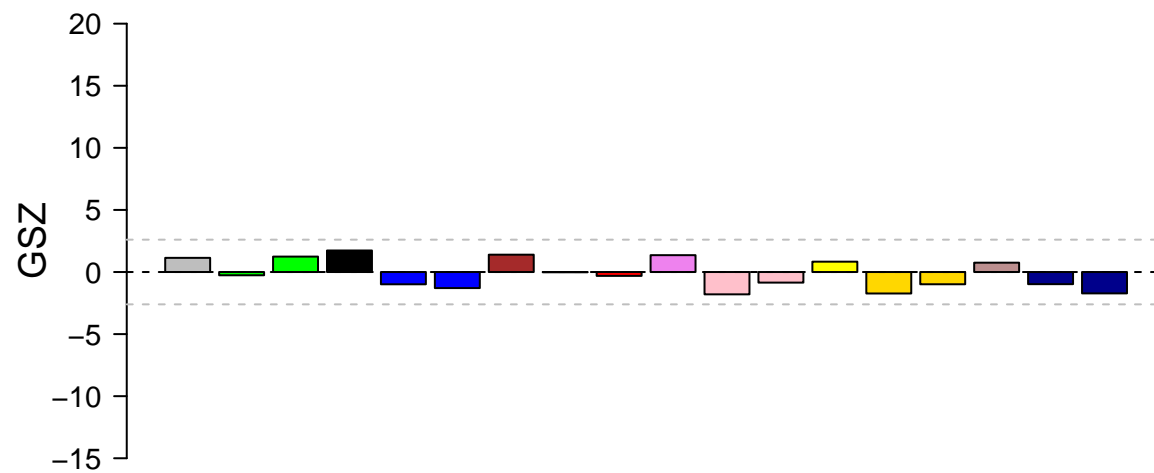


axon regeneration



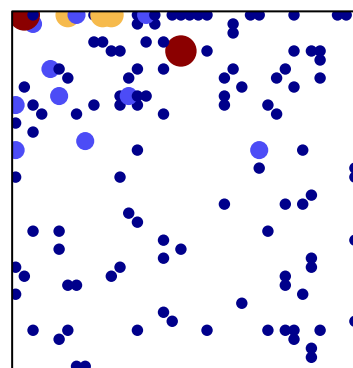
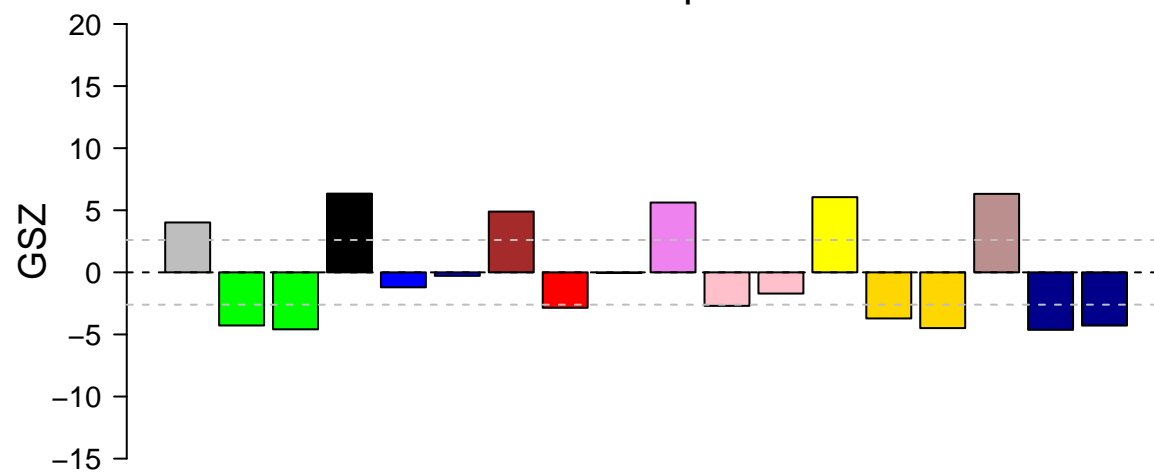
# features = 11 , max = 1

axoneme



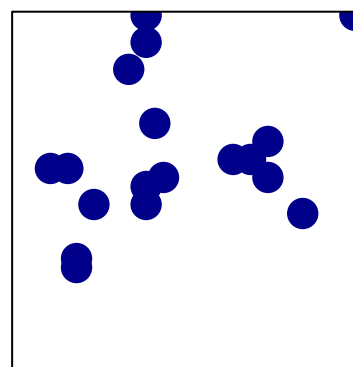
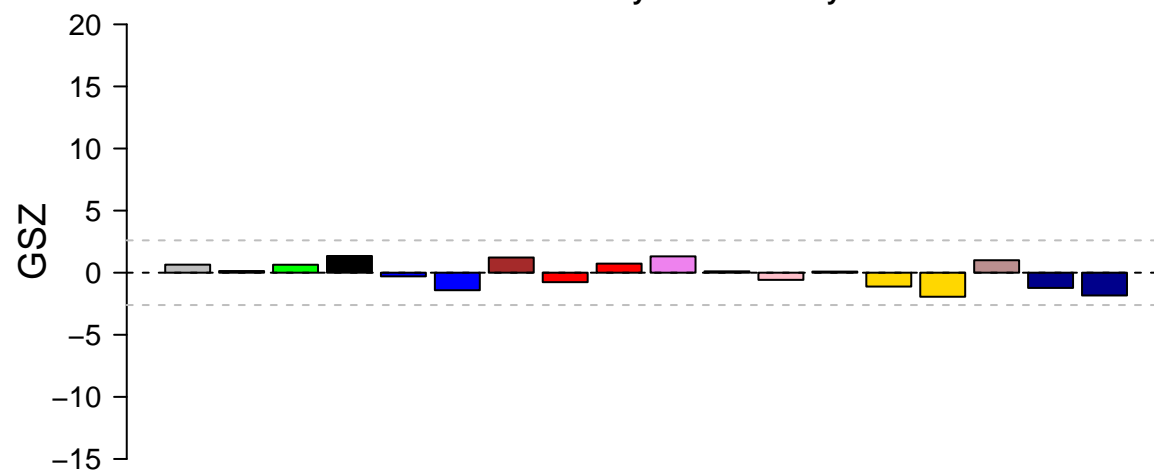
# features = 21 , max = 1

brain development



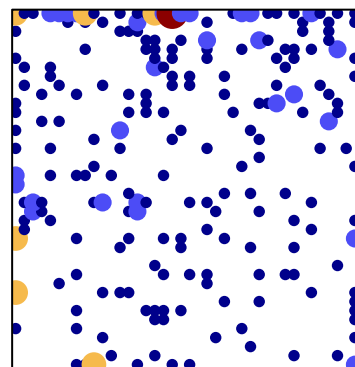
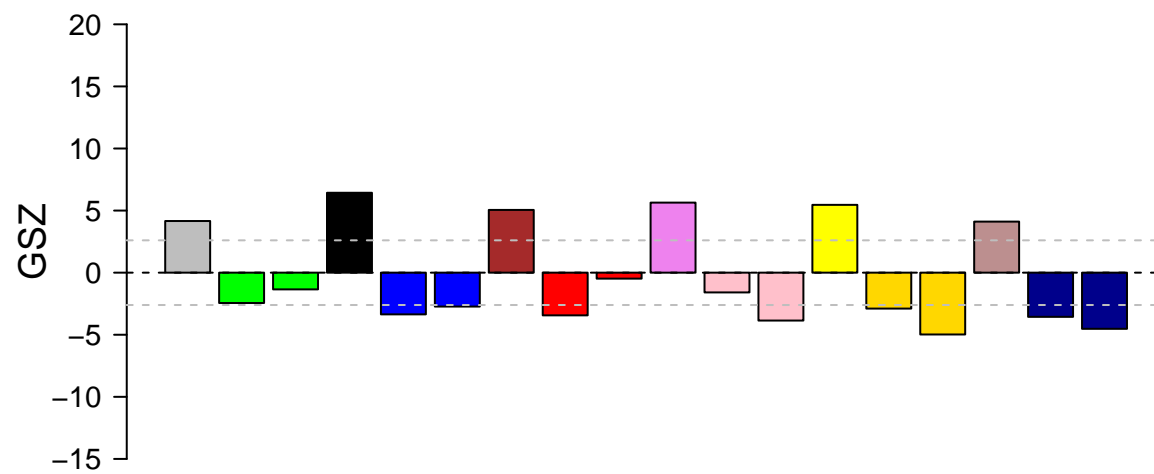
# features = 147 , max = 4

carbonate dehydratase activity



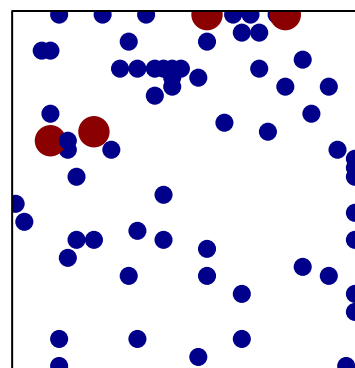
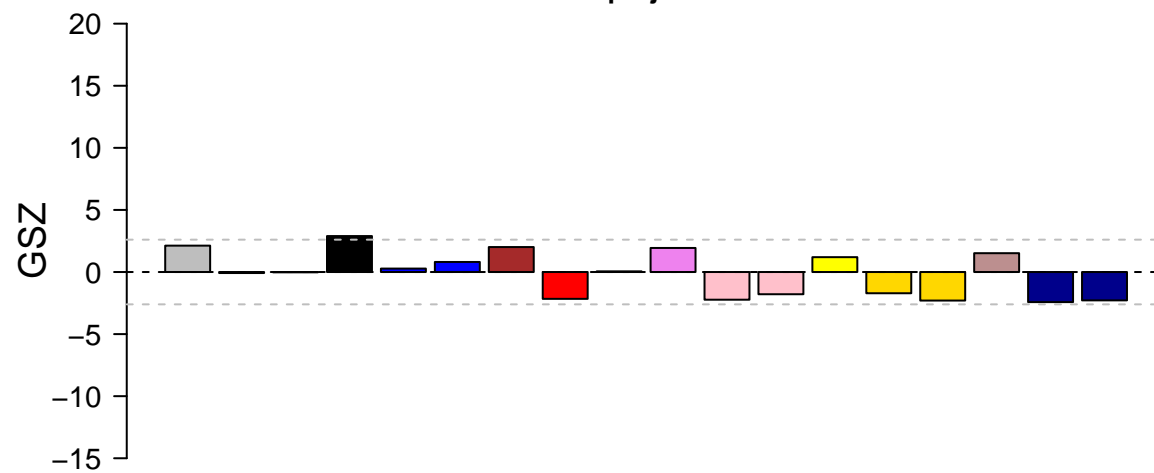
# features = 18 , max = 1

**cell differentiation**



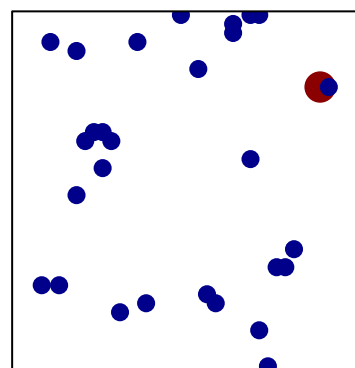
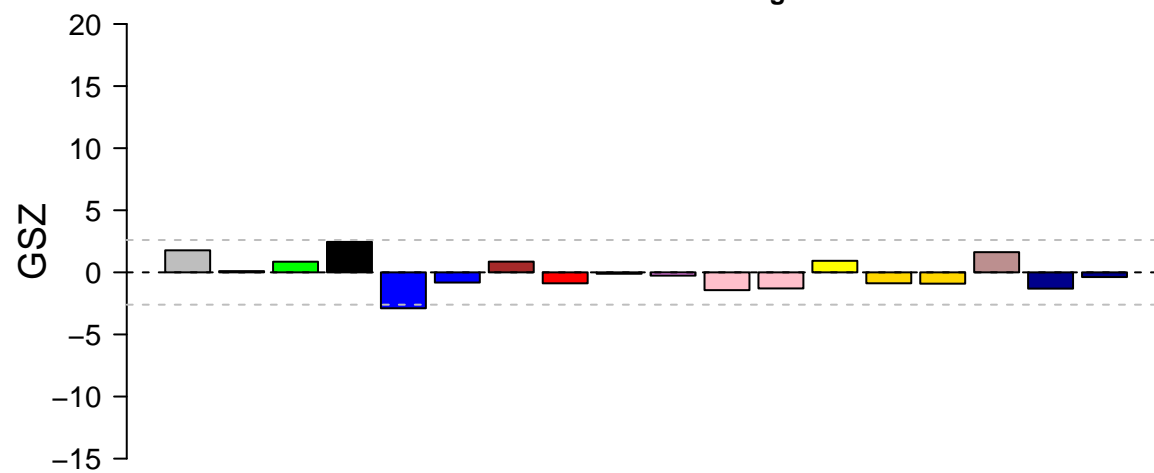
# features = 257 , max = 4

**cell projection**



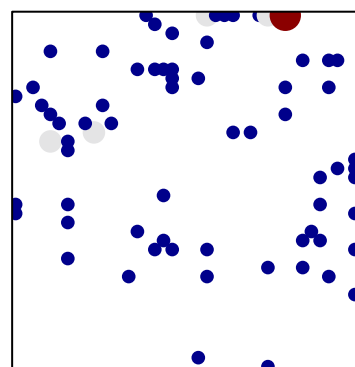
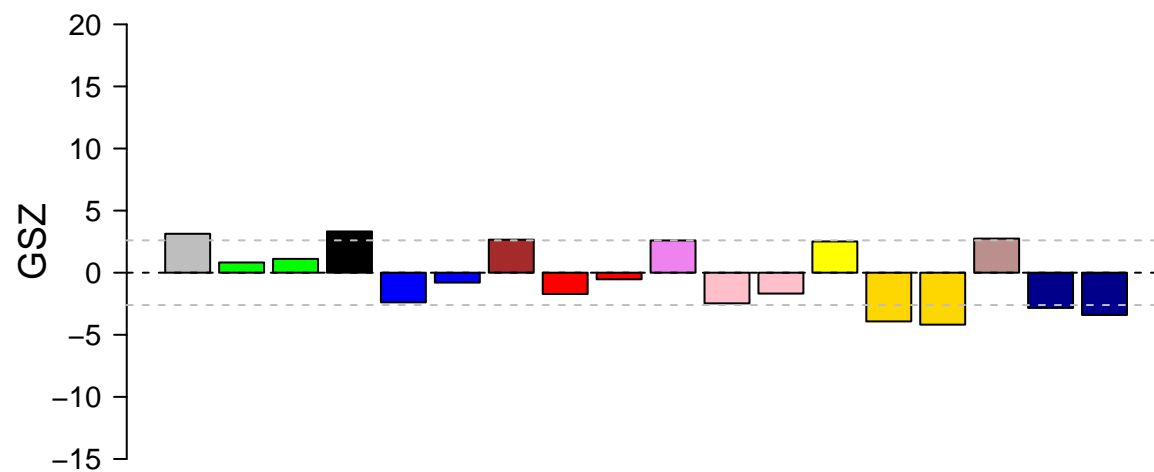
# features = 70 , max = 2

**chromatin remodeling**



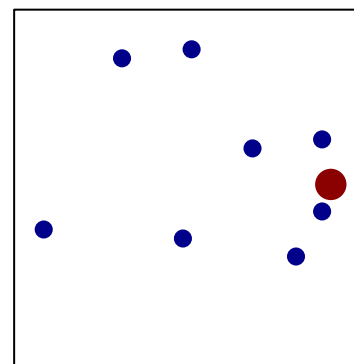
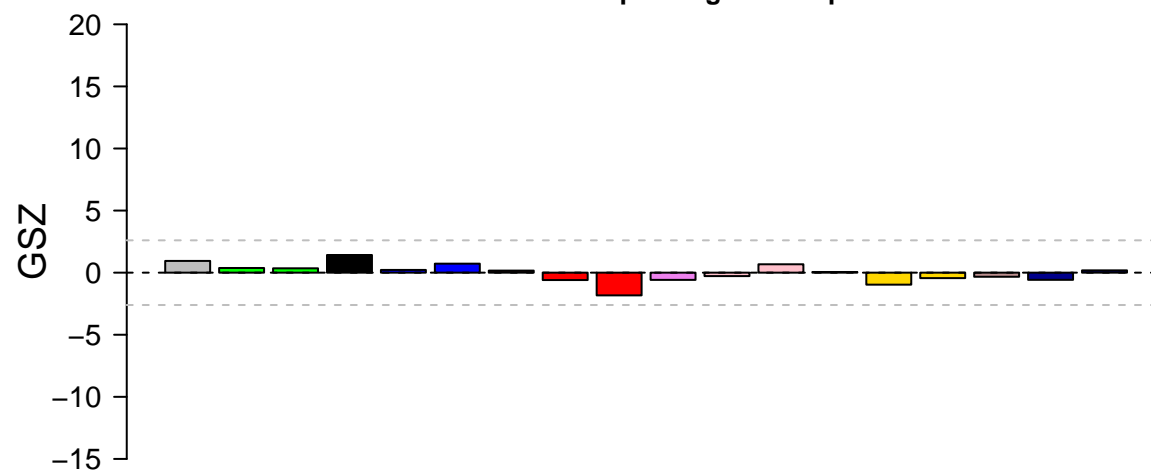
# features = 30 , max = 2

**cilium**



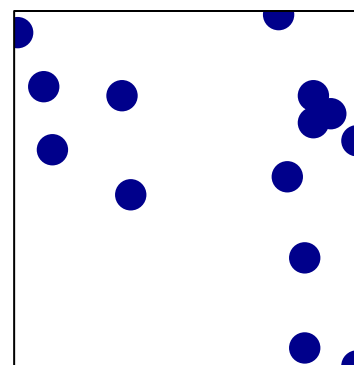
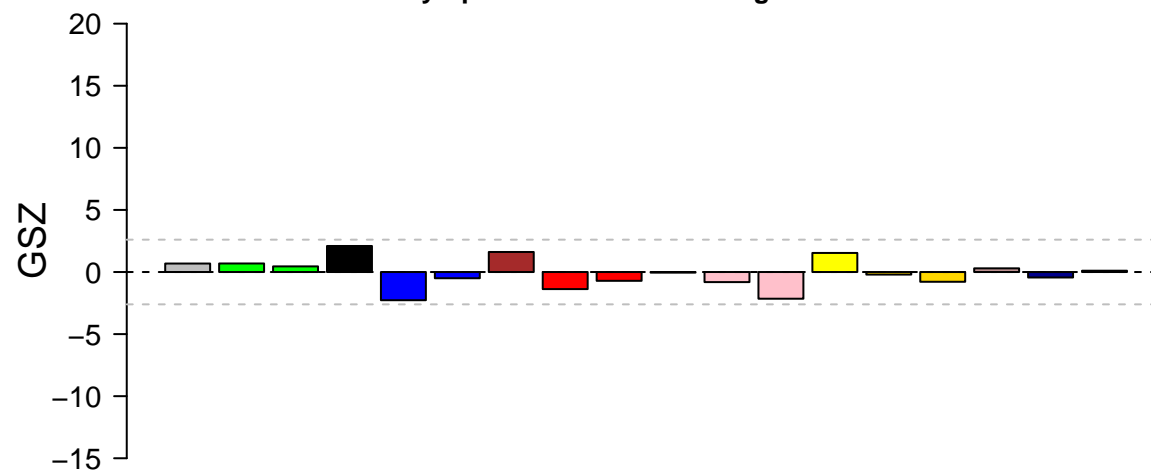
# features = 76 , max = 3

**cullin-RING ubiquitin ligase complex**



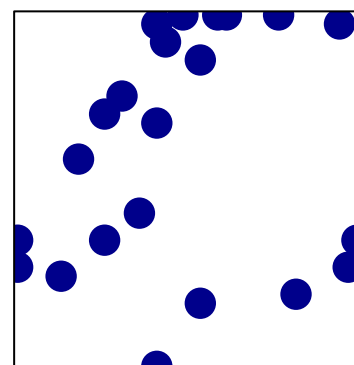
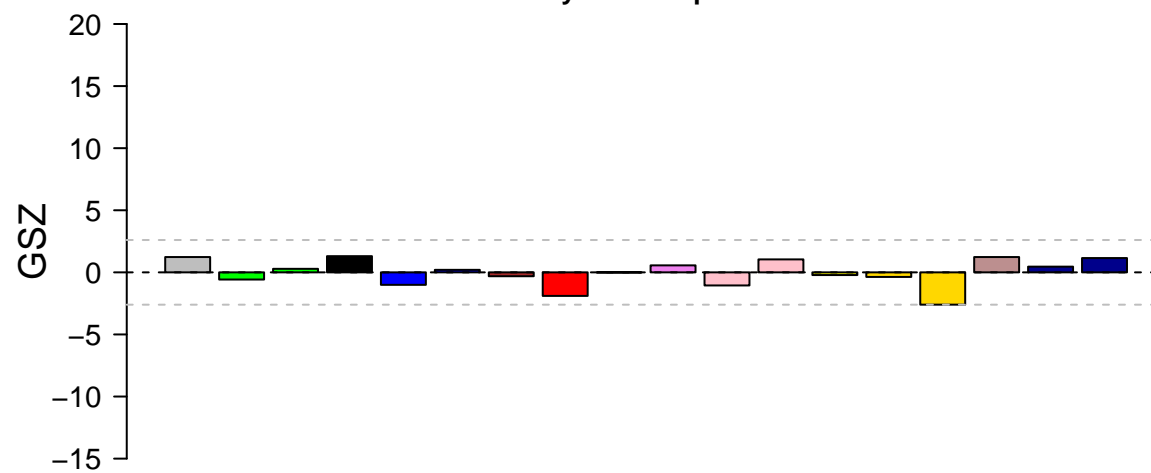
# features = 10 , max = 2

**cytoplasmic microtubule organization**



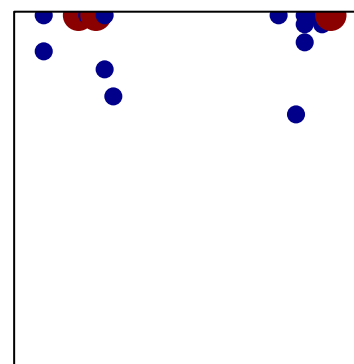
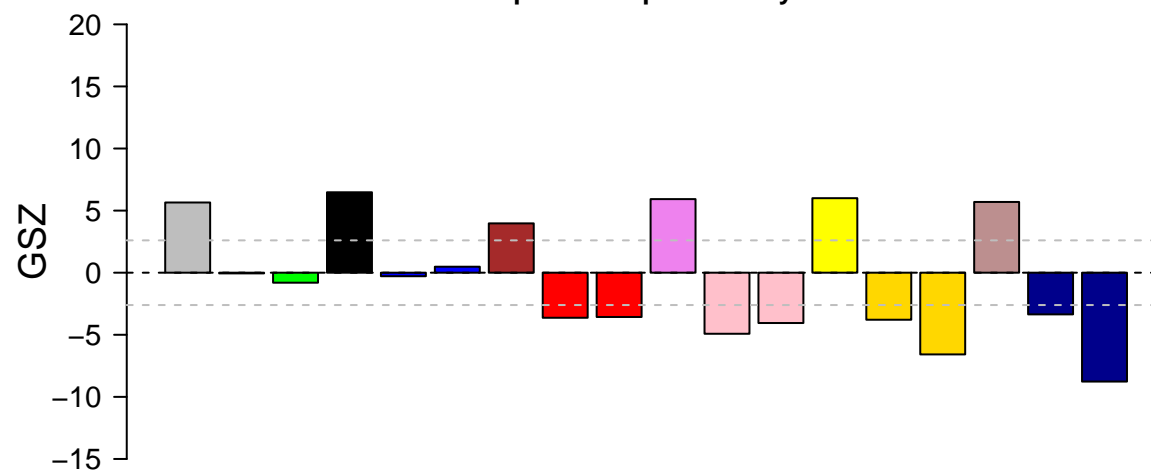
# features = 14 , max = 1

**dynein complex**



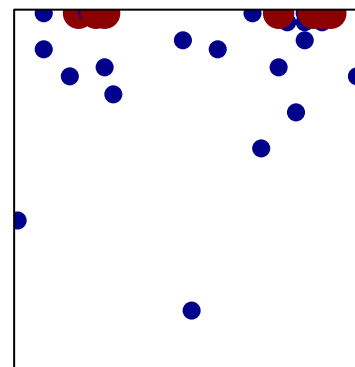
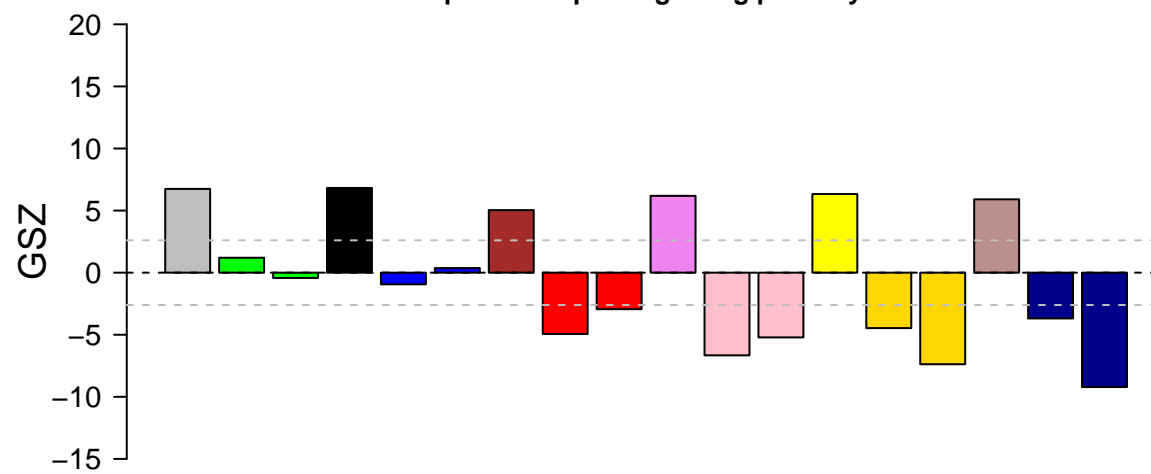
# features = 22 , max = 1

**ephrin receptor activity**



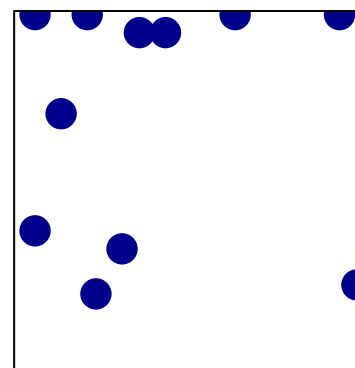
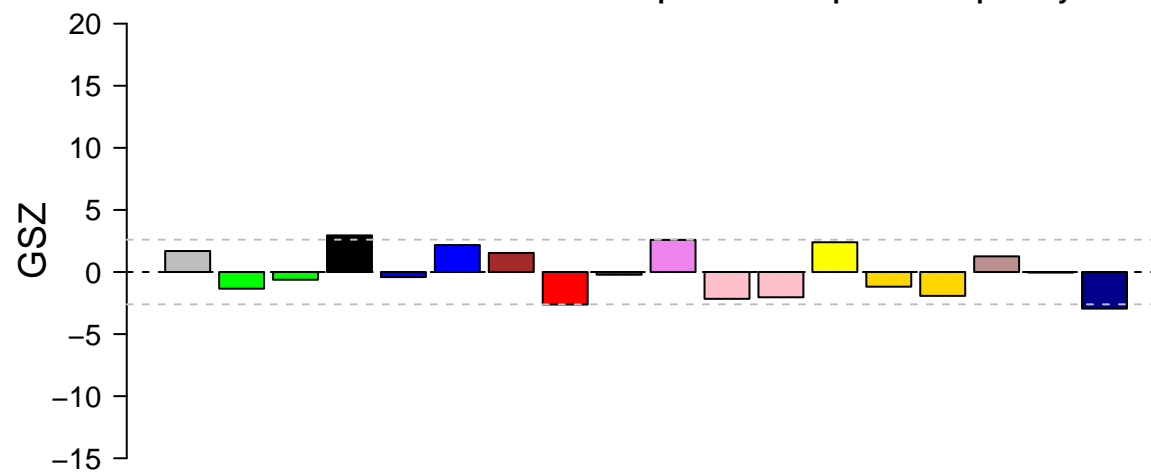
# features = 19 , max = 2

### ephrin receptor signaling pathway



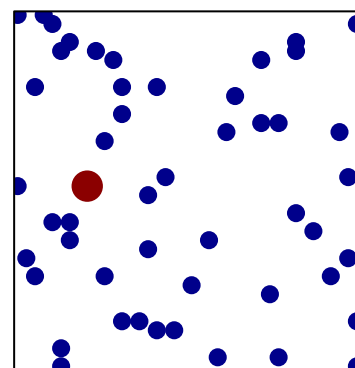
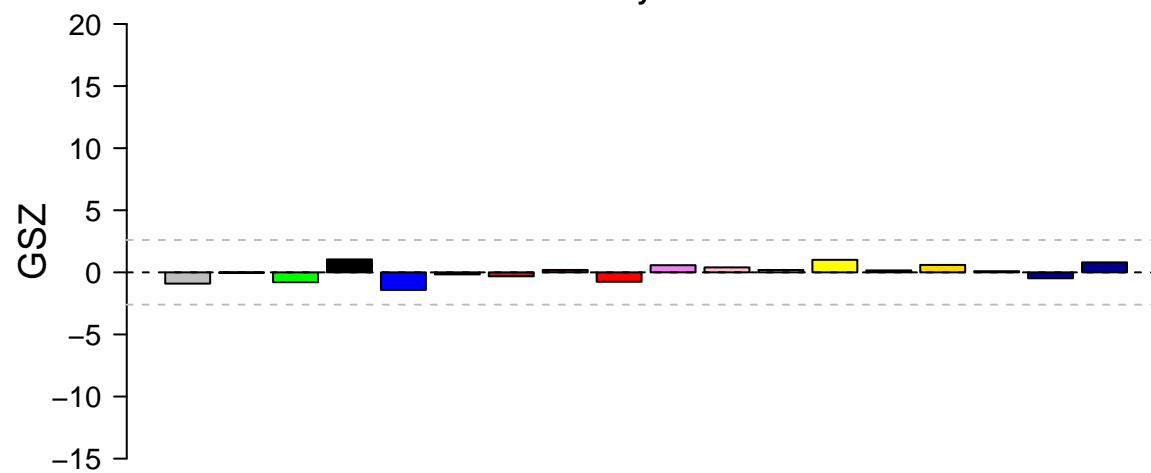
# features = 34 , max = 2

### establishment or maintenance of epithelial cell apical/basal polarity



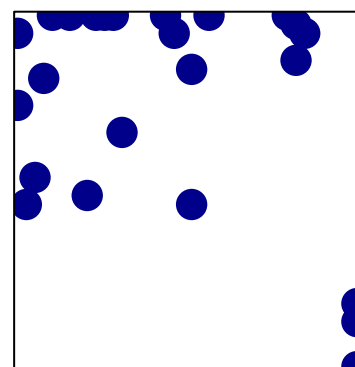
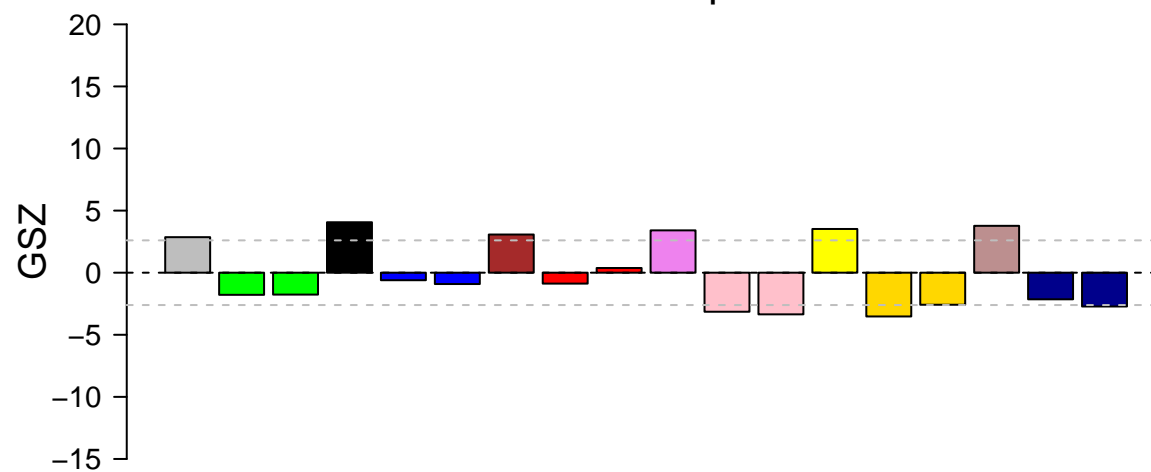
# features = 11 , max = 1

### exocytosis



# features = 51 , max = 2

### forebrain development



# features = 24 , max = 1