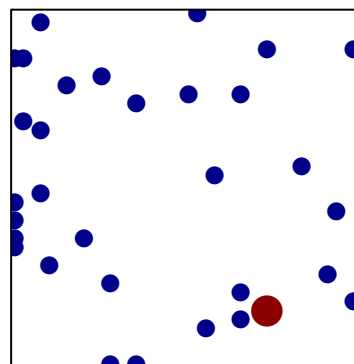
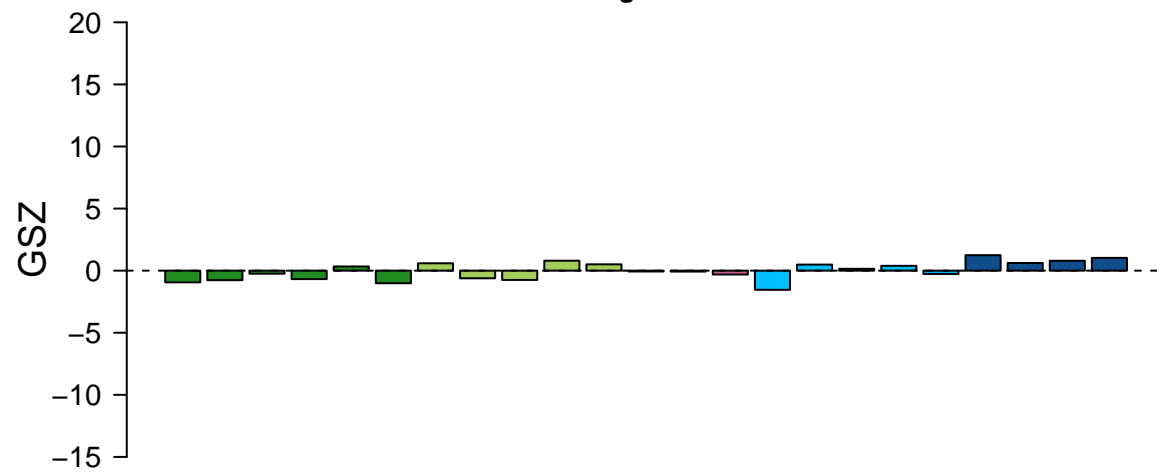
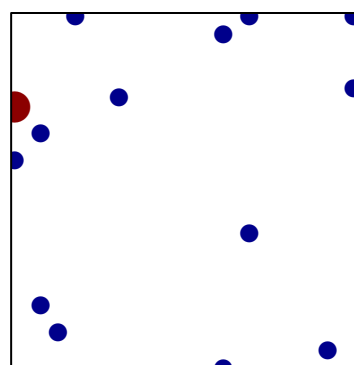
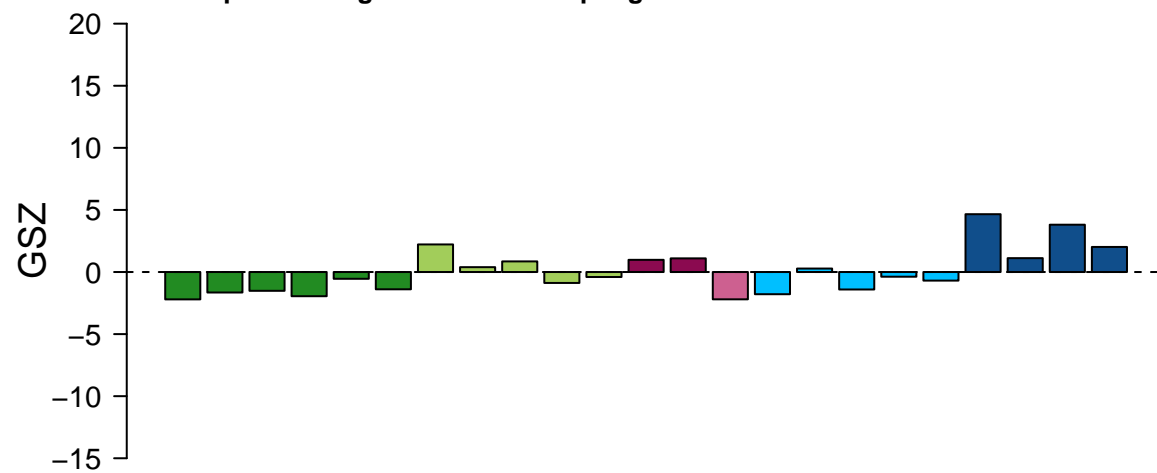


### Golgi stack



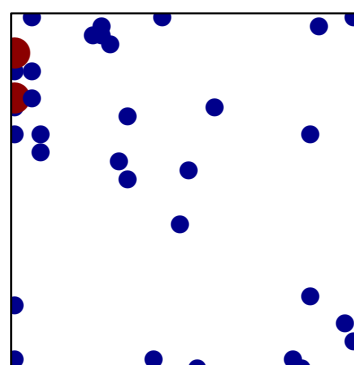
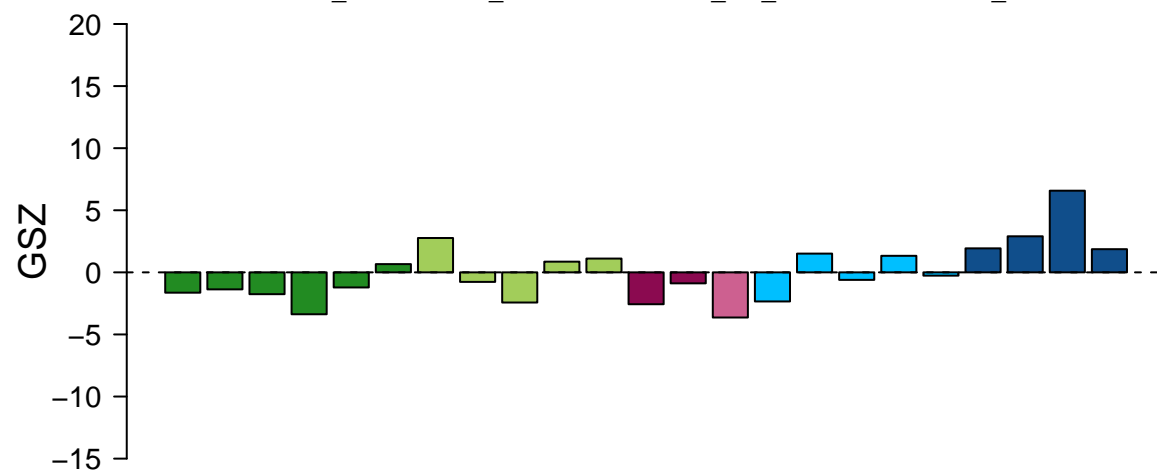
# features = 33 , max = 2

### positive regulation of macrophage derived foam cell differentiation



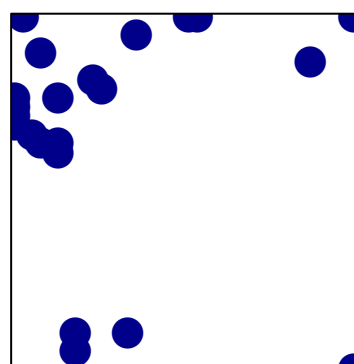
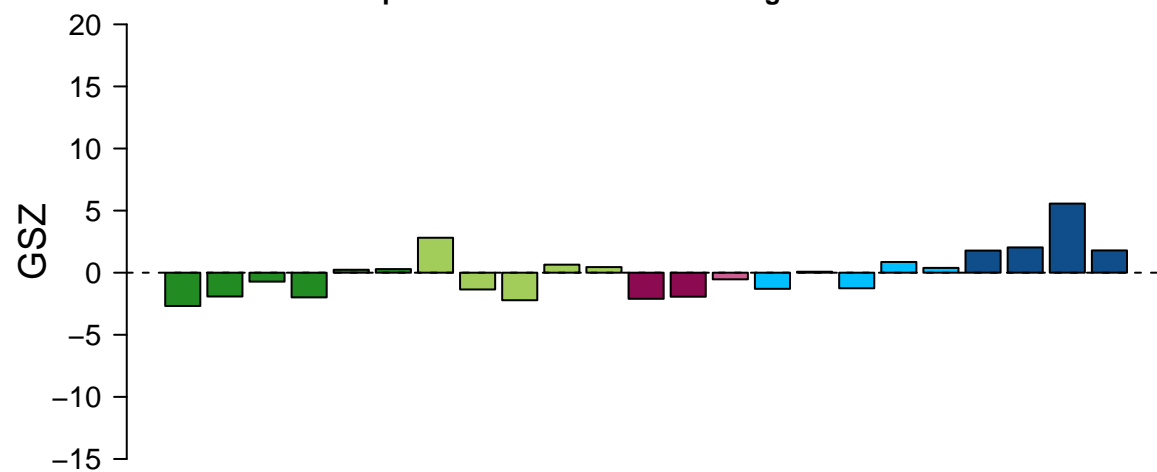
# features = 15 , max = 2

### COLIN\_PILOCYTIC\_ASTROCYTOMA\_VS\_GLIOMASTOMA\_UP



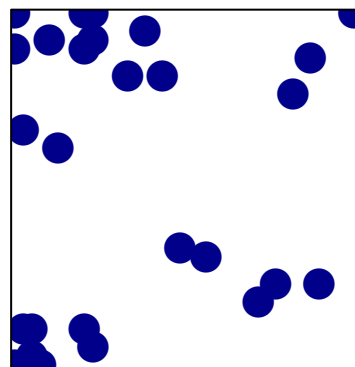
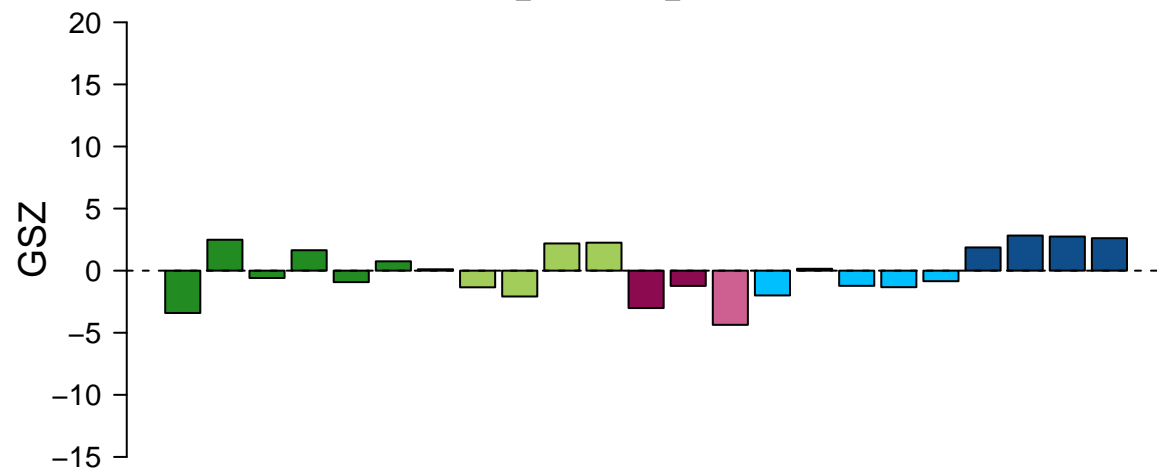
# features = 35 , max = 2

### cellular response to vascular endothelial growth factor stimulus



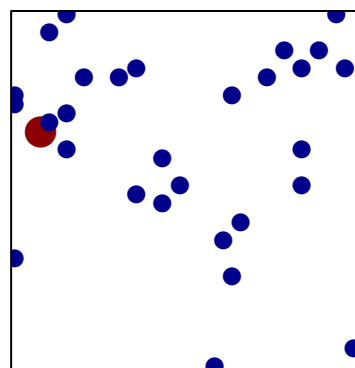
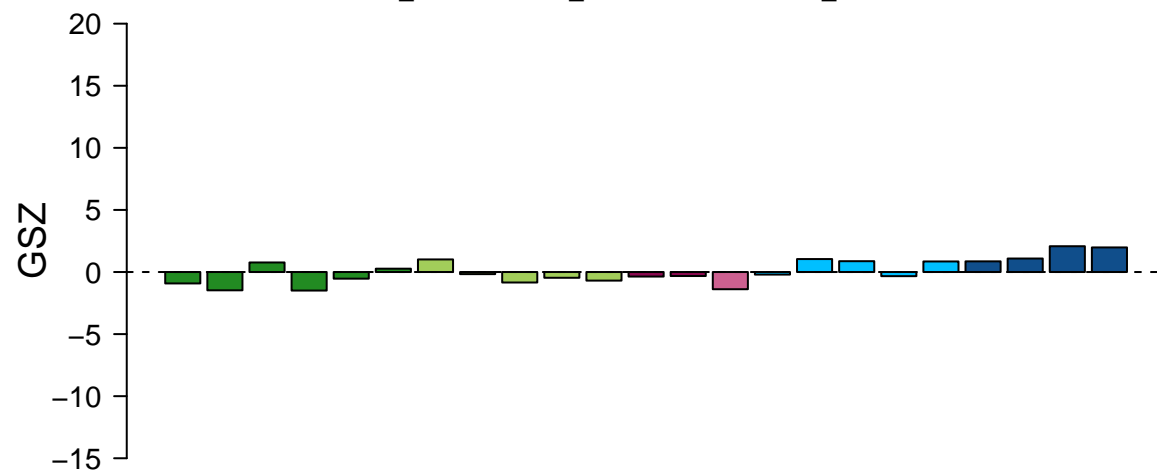
# features = 22 , max = 1

PID\_GLYPICAN\_1PATHWAY



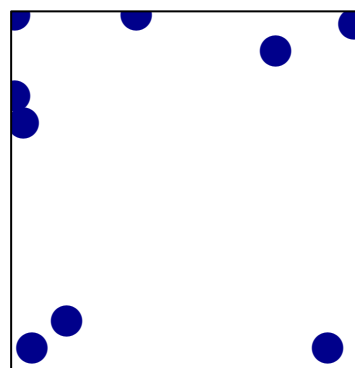
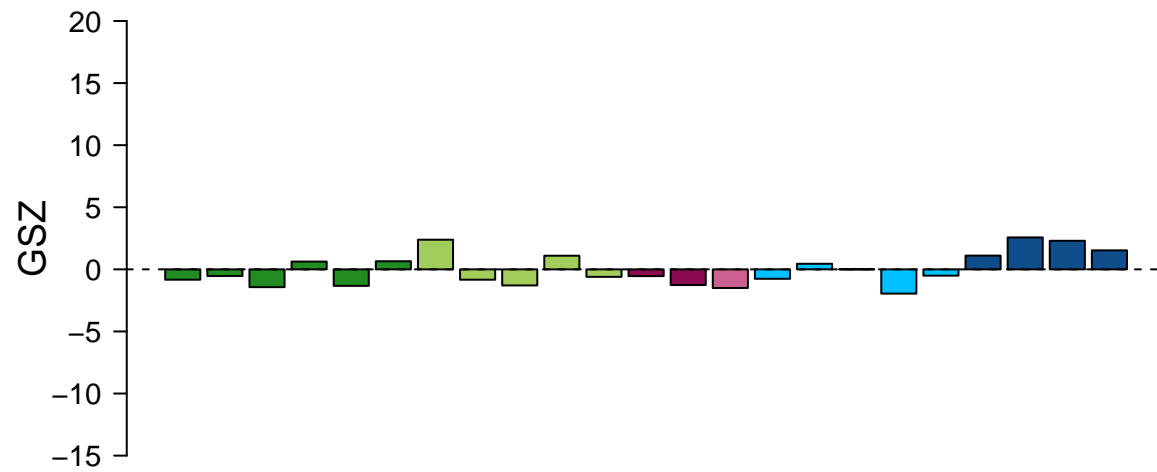
# features = 27 , max = 1

RIZ\_ERYTHROID\_DIFFERENTIATION\_HEMGN



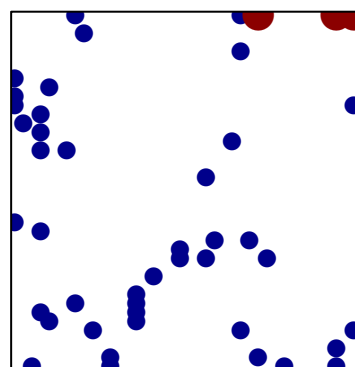
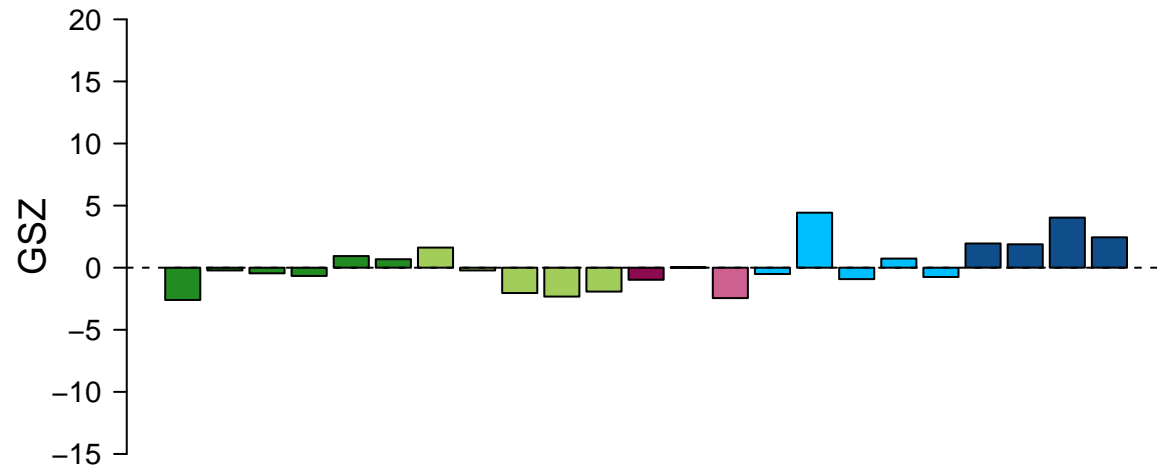
# features = 31 , max = 2

KUMAMOTO\_RESPONSE\_TO\_NUTLIN\_3A\_UP



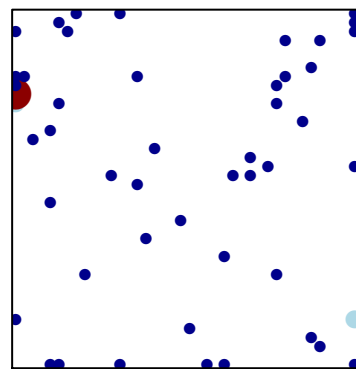
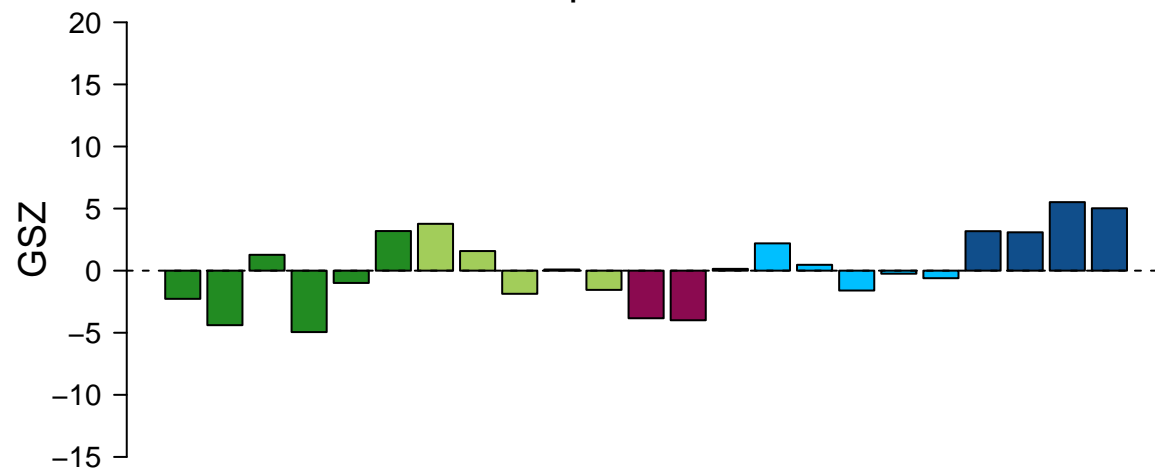
# features = 9 , max = 1

positive regulation of inflammatory response



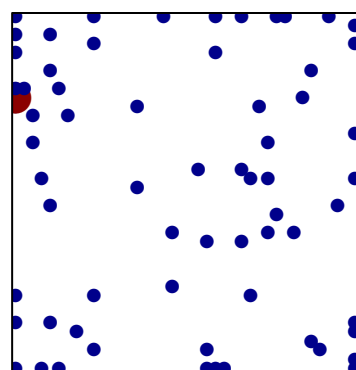
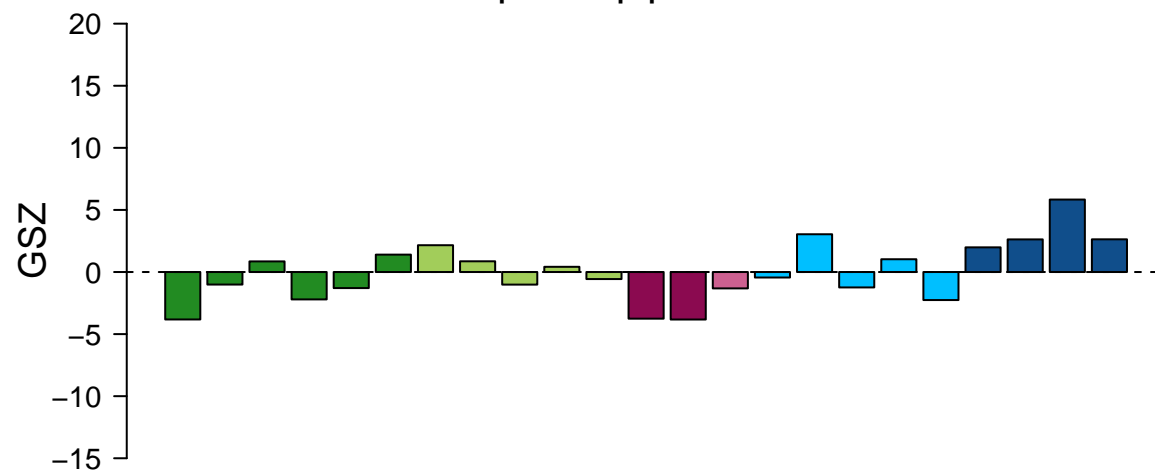
# features = 48 , max = 2

response to cAMP



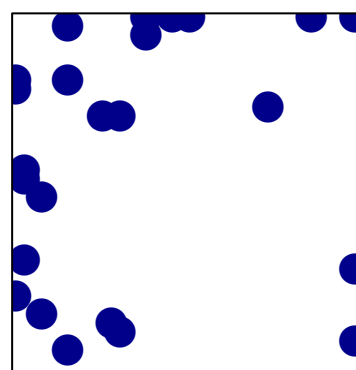
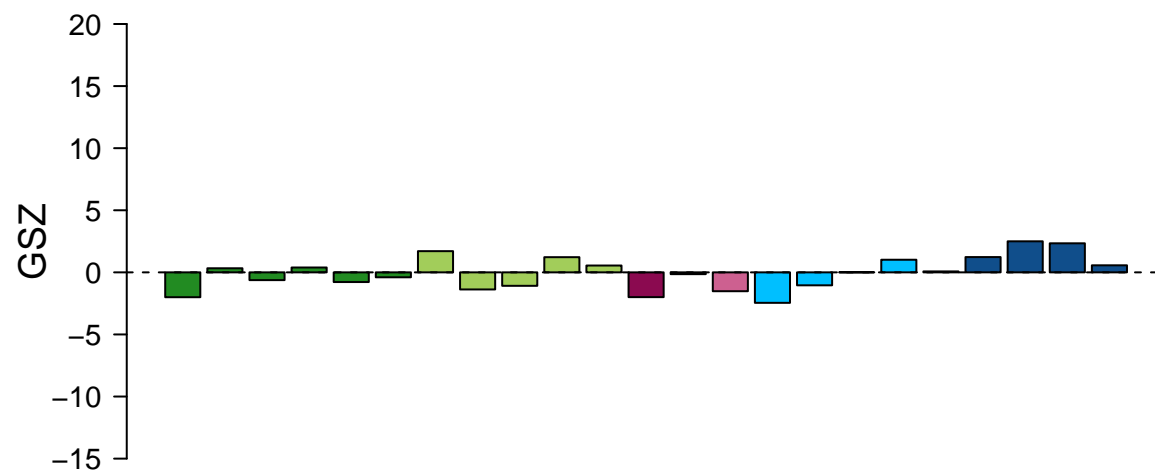
# features = 54 , max = 4

response to peptide hormone



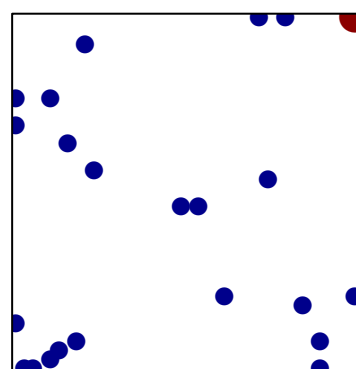
# features = 68 , max = 3

hsa-miR-1250



# features = 24 , max = 1

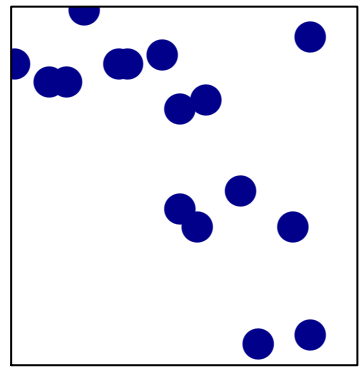
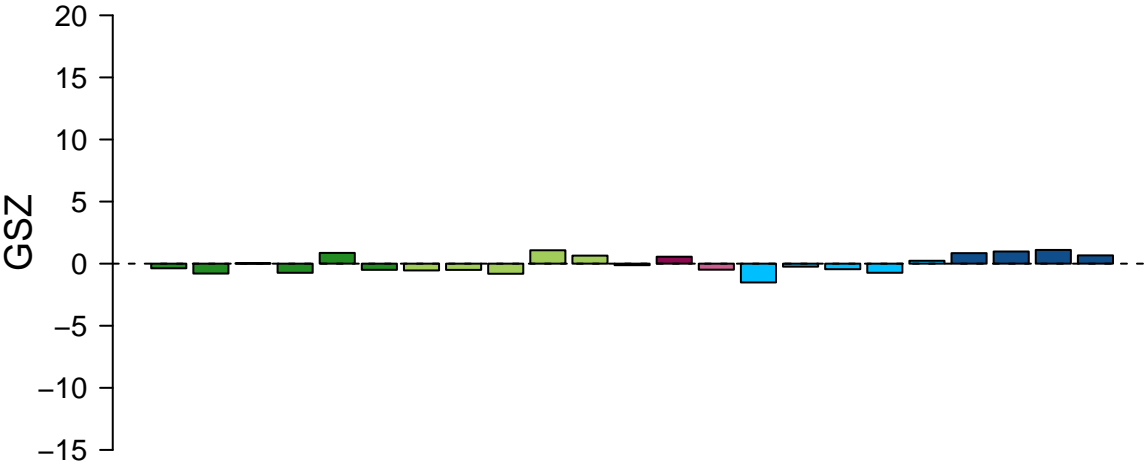
CHIANG\_LIVER\_CANCER\_SUBCLASS\_INTERFERON\_UP



# features = 24 , max = 2

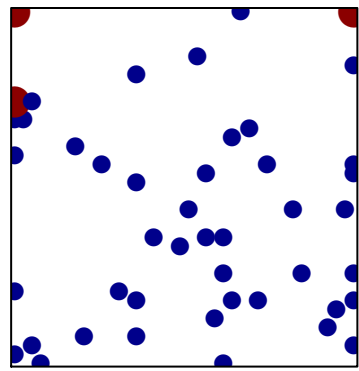
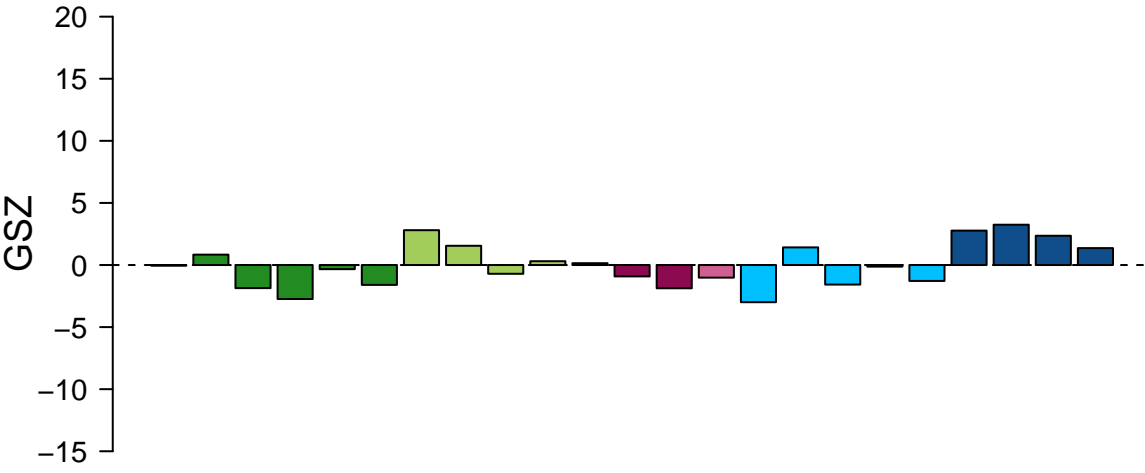


transmembrane receptor protein serine/threonine kinase signaling pathway



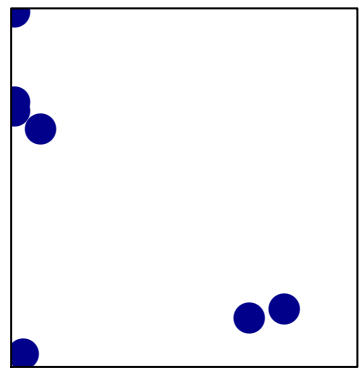
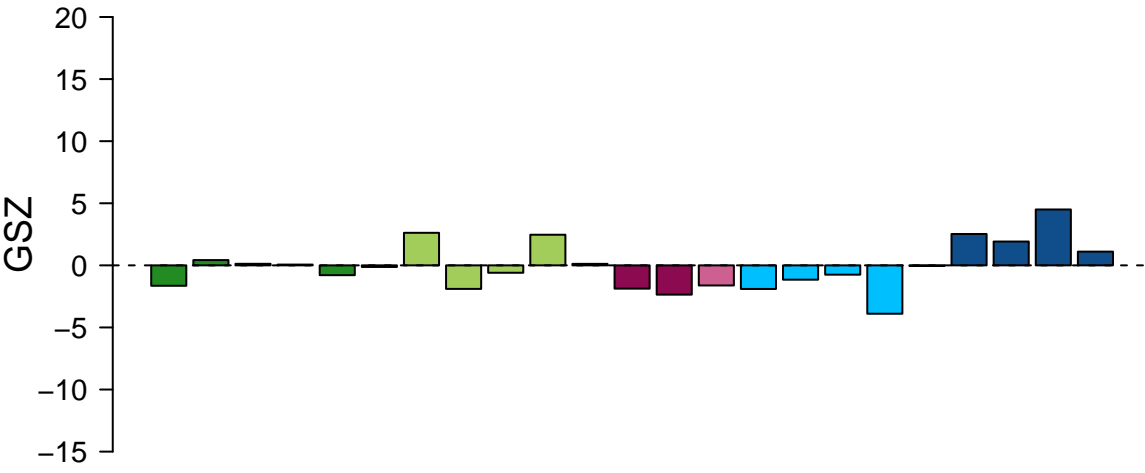
# features = 16 , max = 1

Subero\_DLBCCL\_hypo\_meth



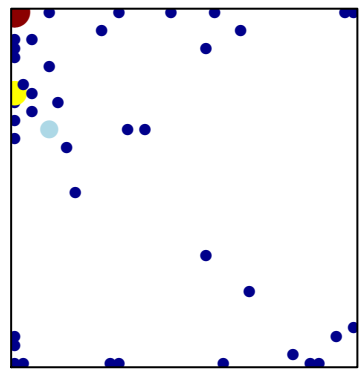
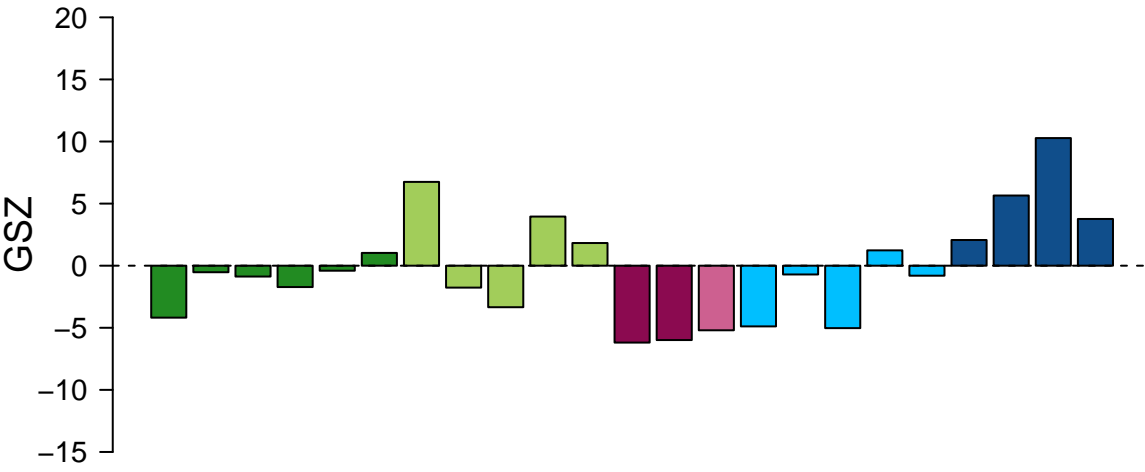
# features = 49 , max = 2

LOPEZ\_MESOTELIOMA\_SURVIVAL\_TIME\_DN



# features = 7 , max = 1

CHIARADONNA\_NEOPLASTIC\_TRANSFORMATION\_KRAS\_CDC25\_DN



# features = 49 , max = 4