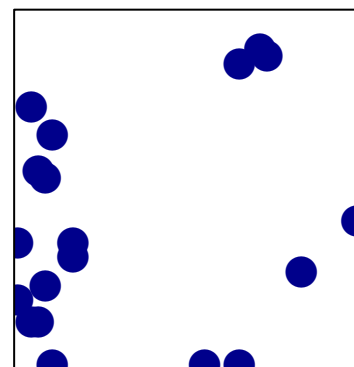
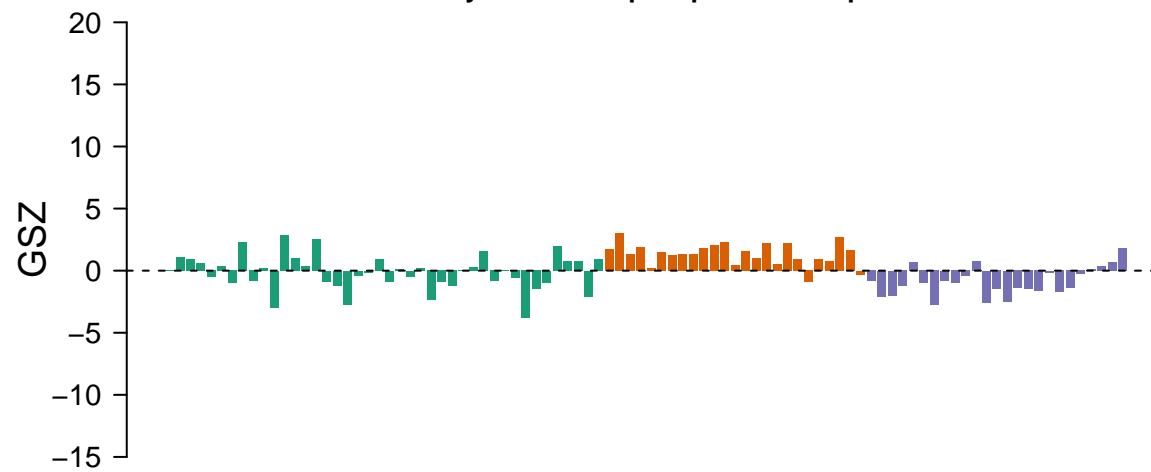
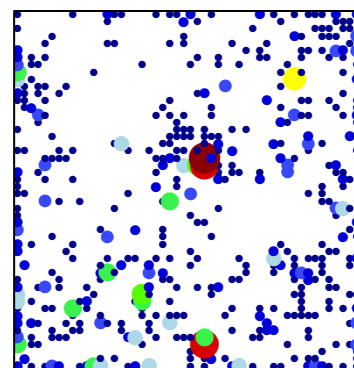
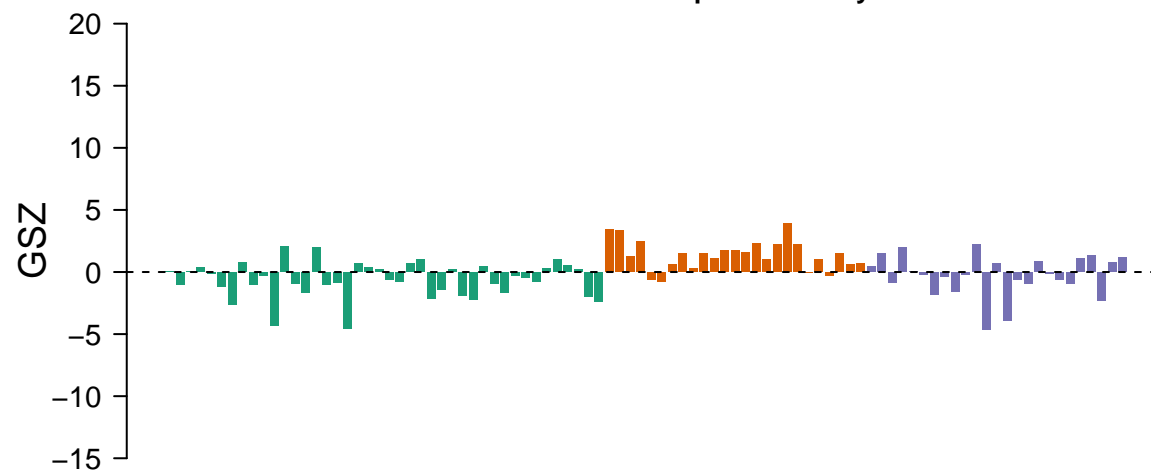


ATP synthesis coupled proton transport



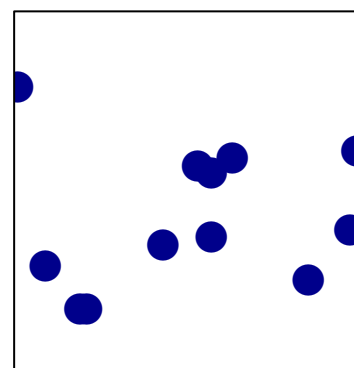
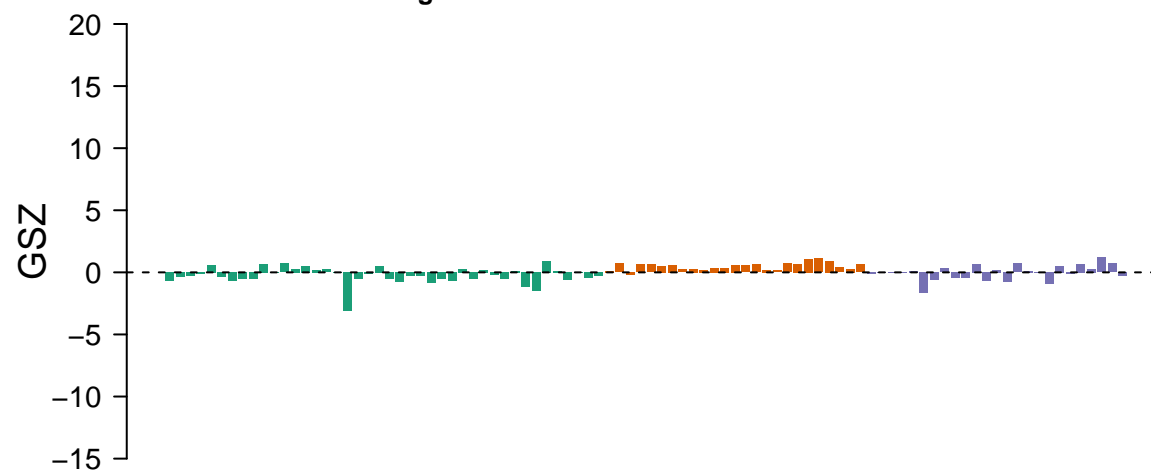
features = 19 , max = 1

transmembrane transporter activity



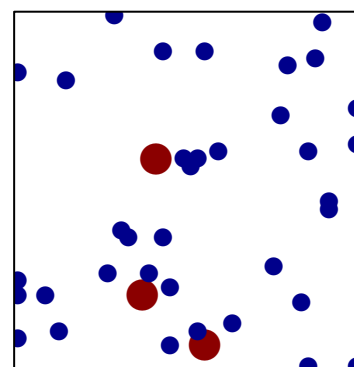
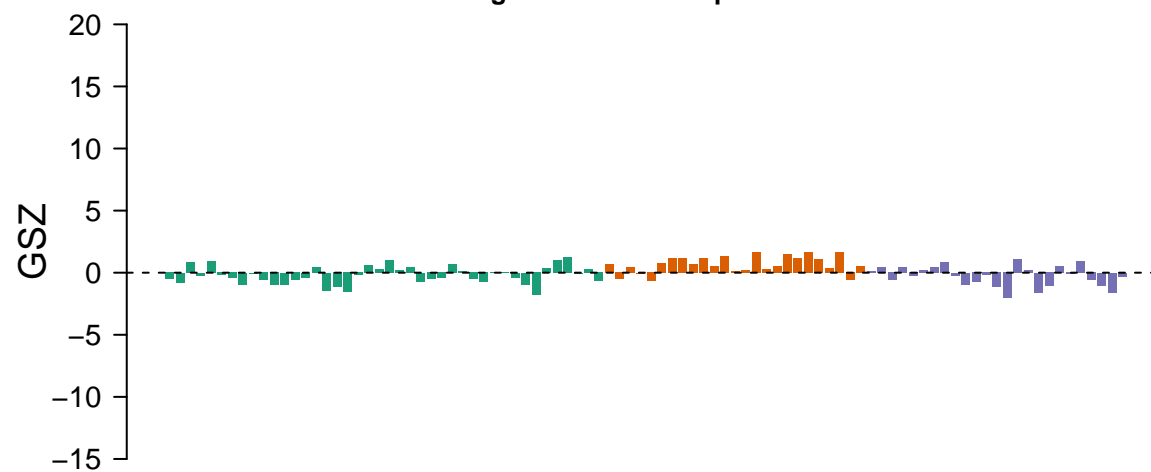
features = 695 , max = 10

regulation of neurotransmitter secretion



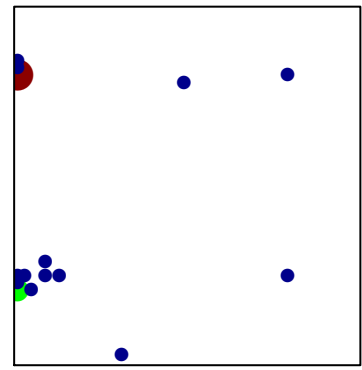
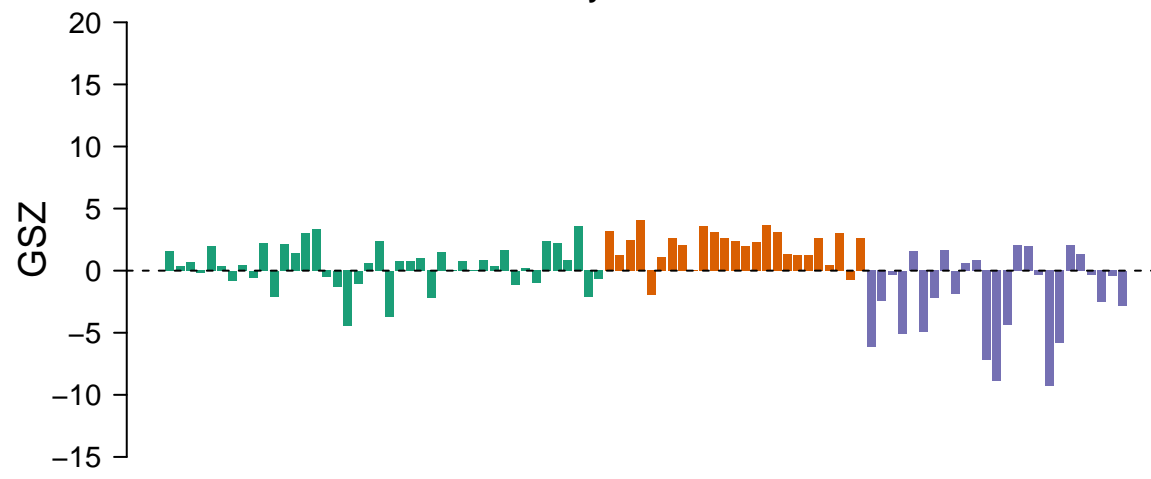
features = 12 , max = 1

regulation of blood pressure



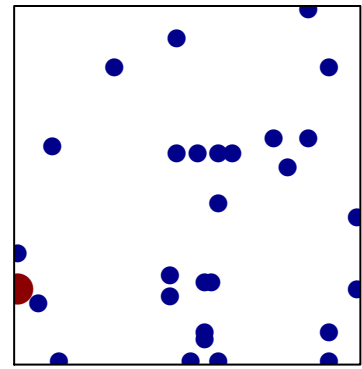
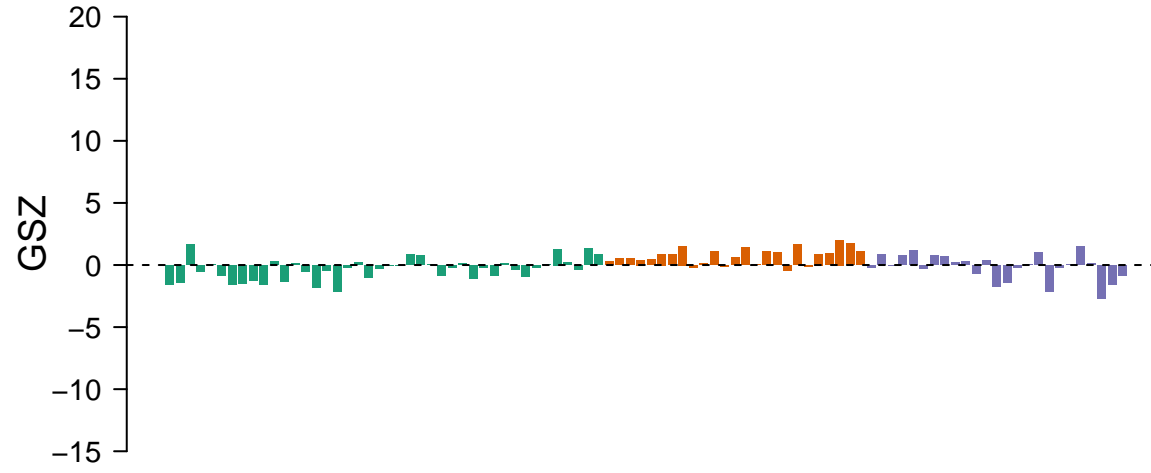
features = 42 , max = 2

melanocyte differentiation



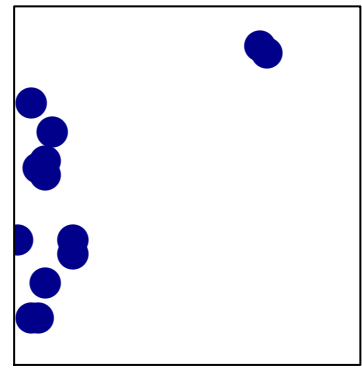
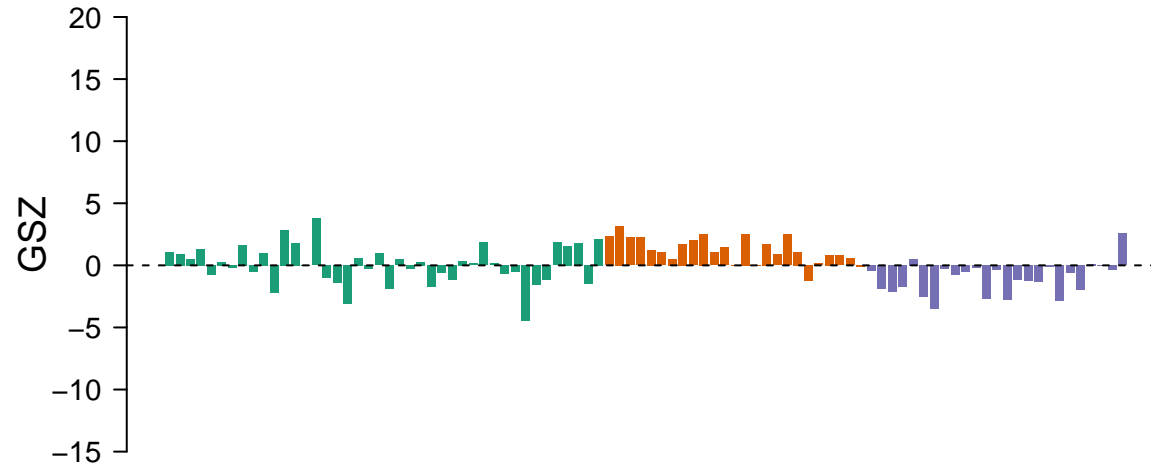
features = 18 , max = 3

LI_CISPLATIN_RESISTANCE_DN



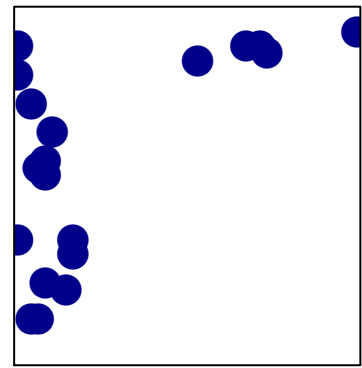
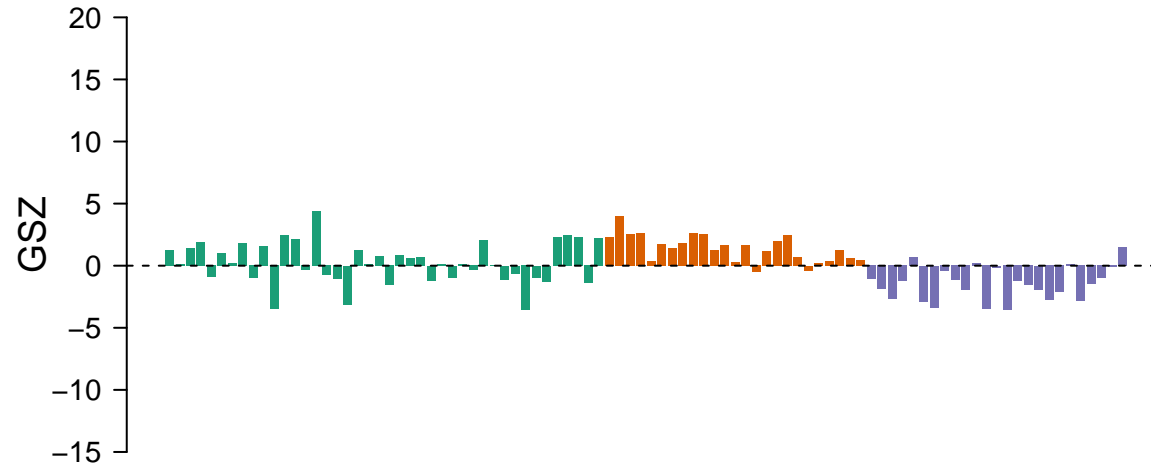
features = 30 , max = 2

REACTOME_FORMATION_OF_ATP_BY_CHEMIOSMOTIC_COUPLING



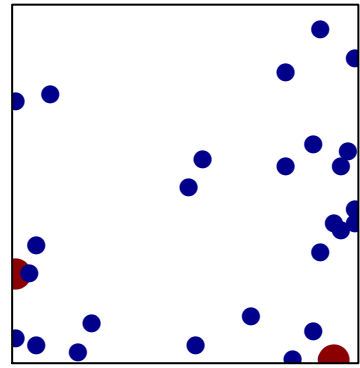
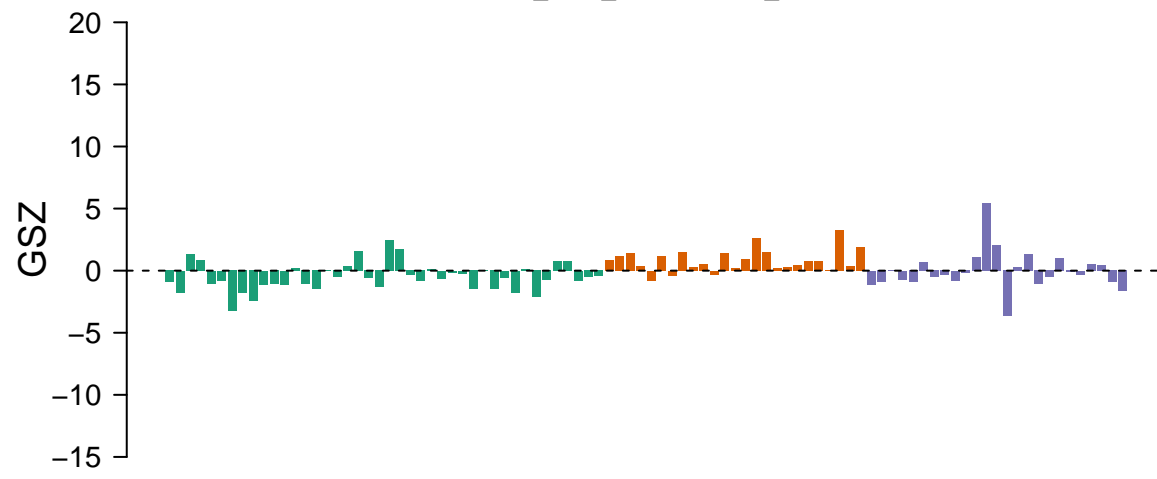
features = 13 , max = 1

mitochondrial proton-transporting ATP synthase complex

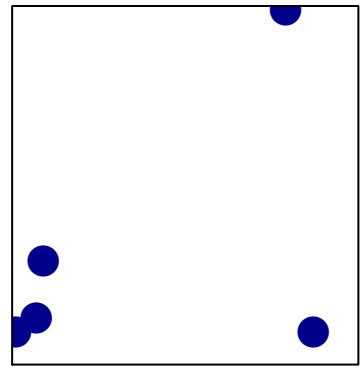
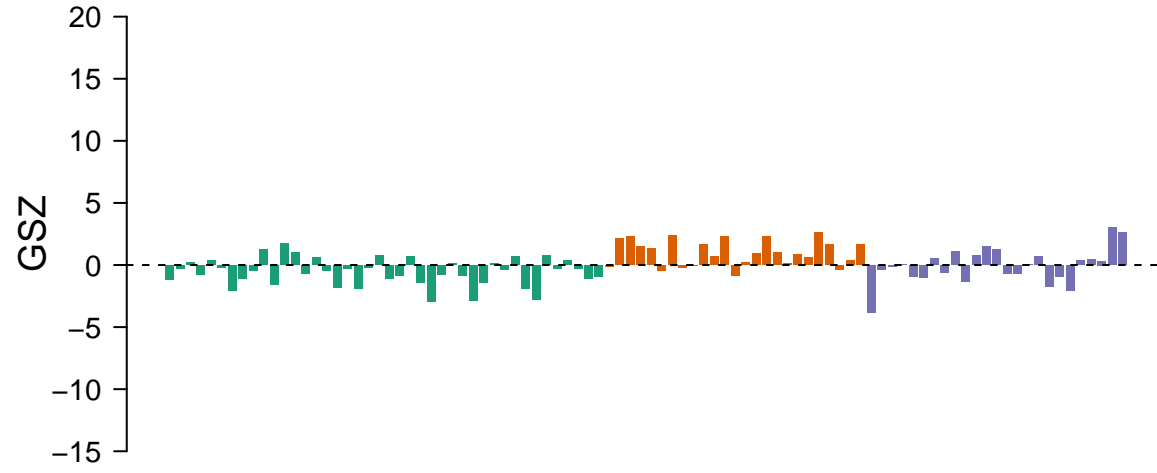


features = 19 , max = 1

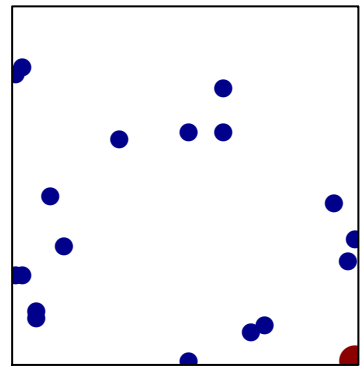
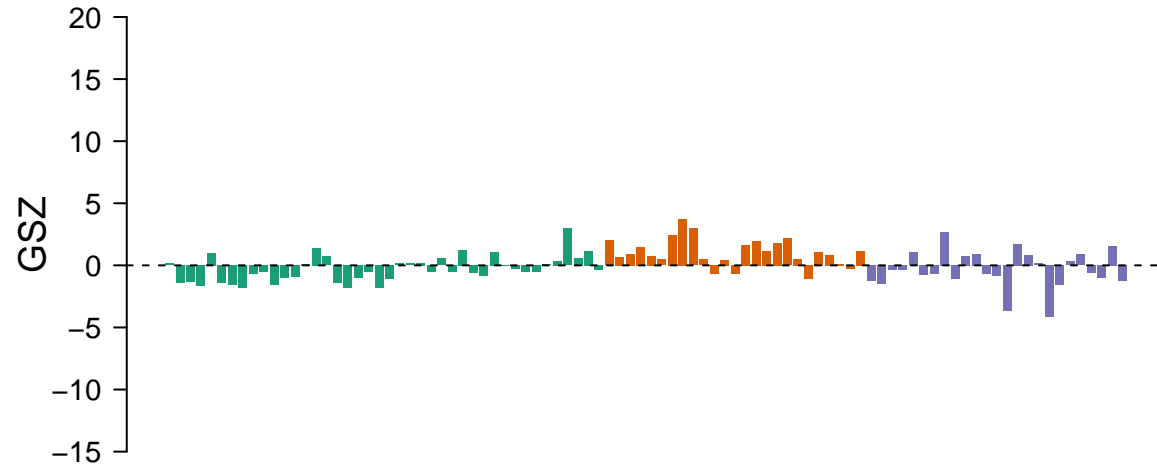
HAN_JNK_SINGALING_UP



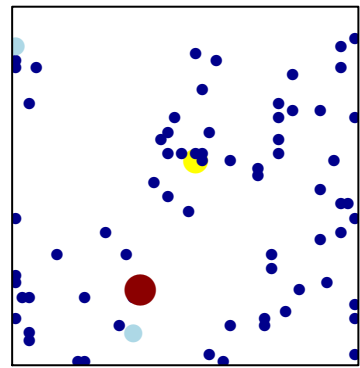
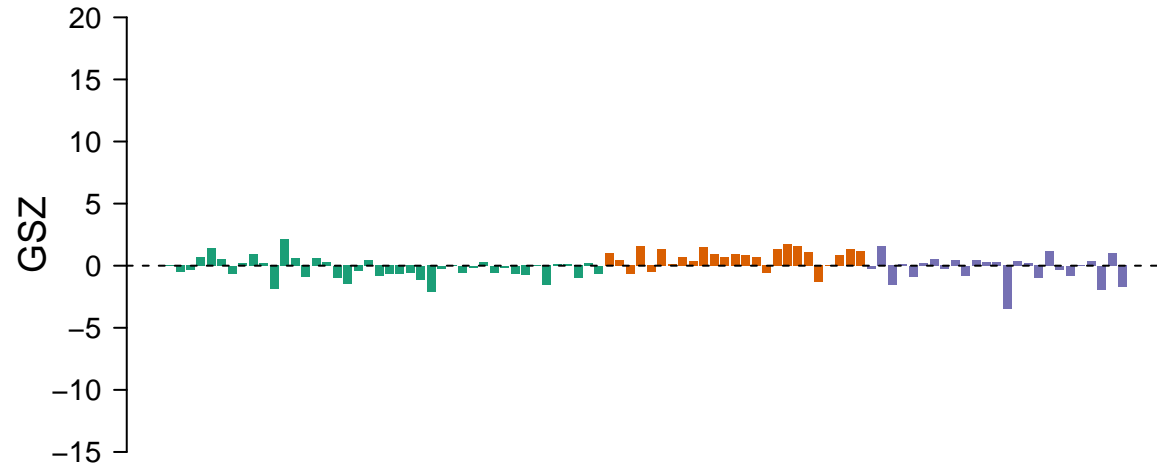
TURJANSKI_MAPK11_TARGETS



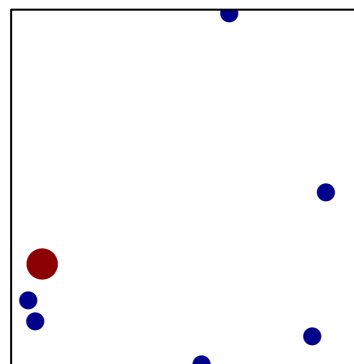
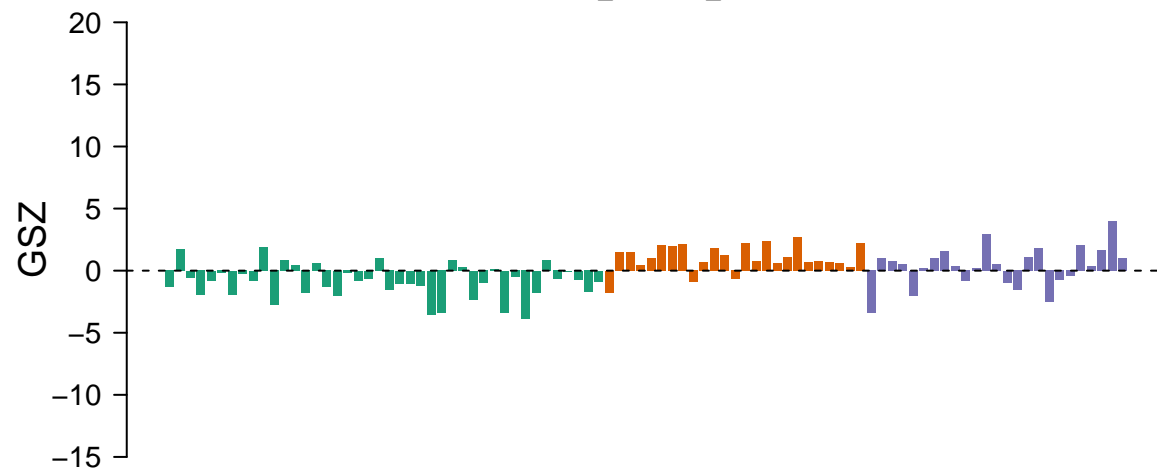
HERNANDEZ_ABERRANT_MITOSIS_BY_DOCETACEL_4NM_UP



sodium ion transport

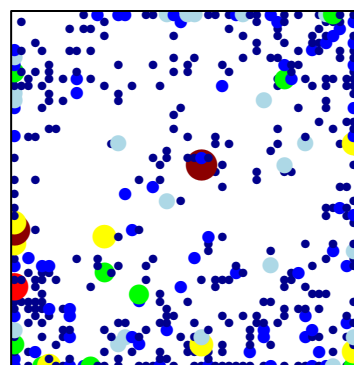
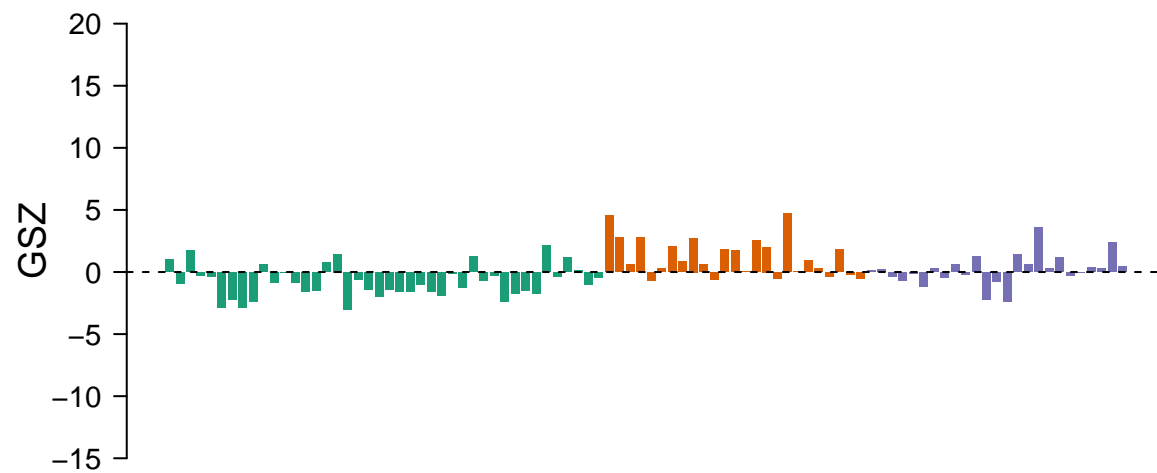


TURJANSKI_MAPK7_TARGETS



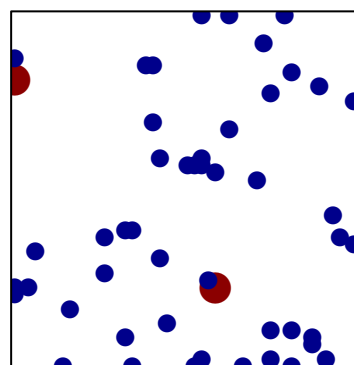
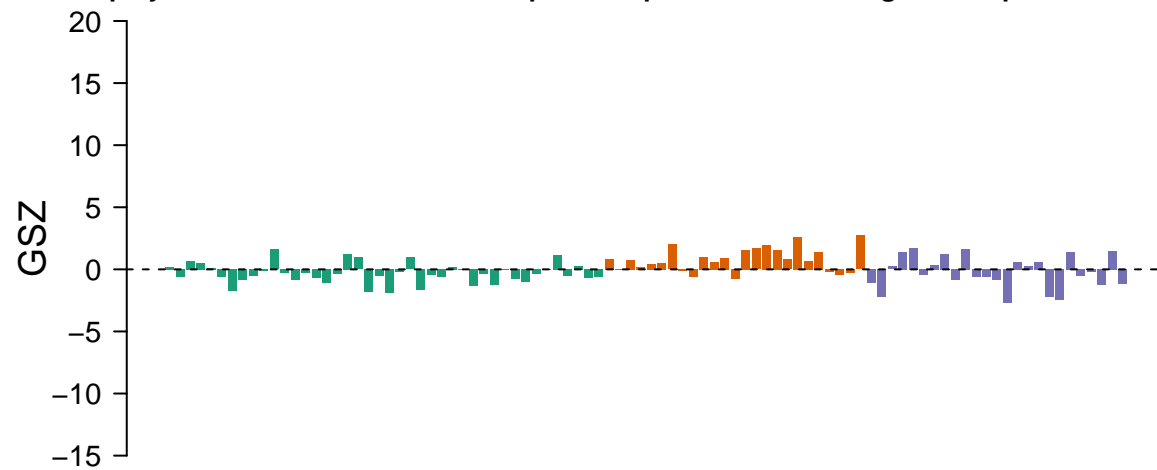
features = 8 , max = 2

endosome



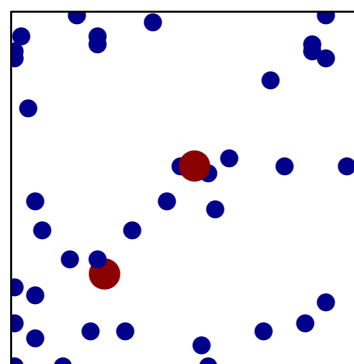
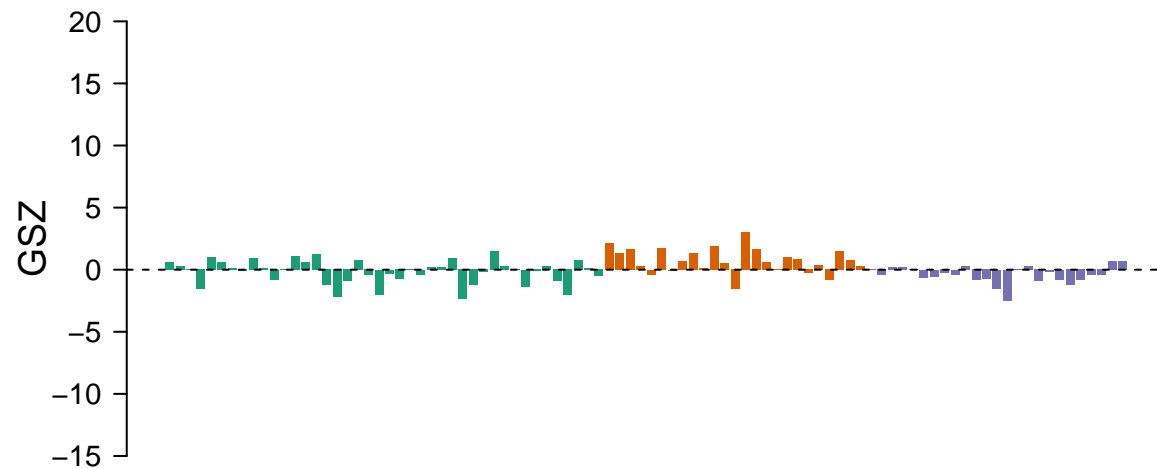
features = 616 , max = 7

RNA polymerase II distal enhancer sequence-specific DNA binding transcription factor act



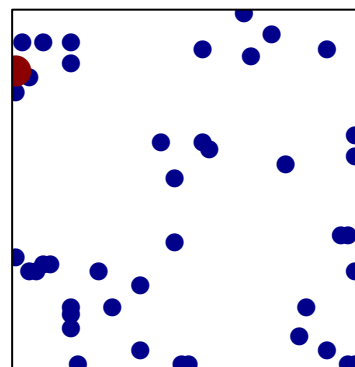
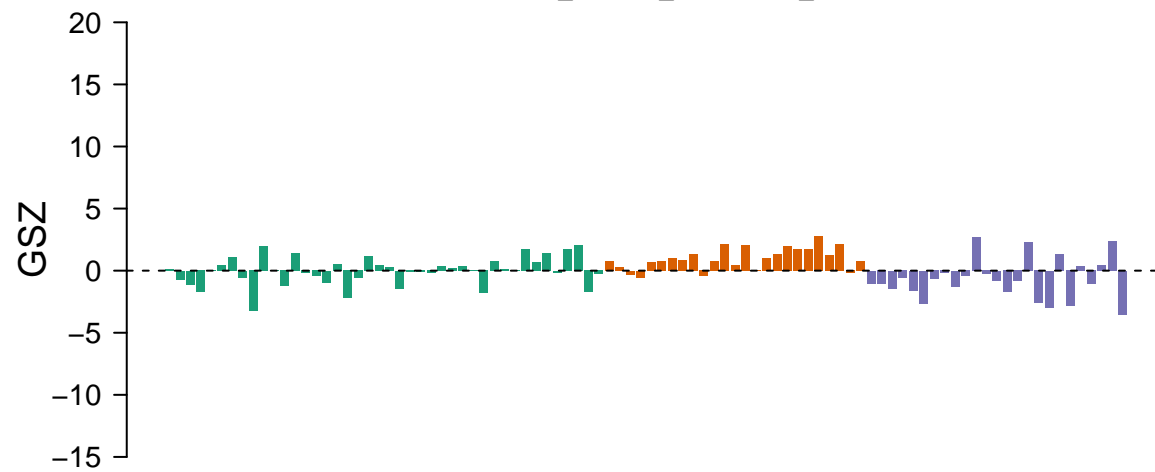
features = 52 , max = 2

CHIANG_LIVER_CANCER_SUBCLASS_INTERFERON_DN

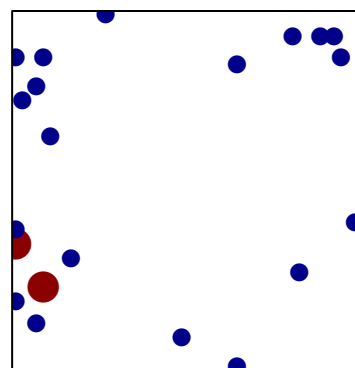
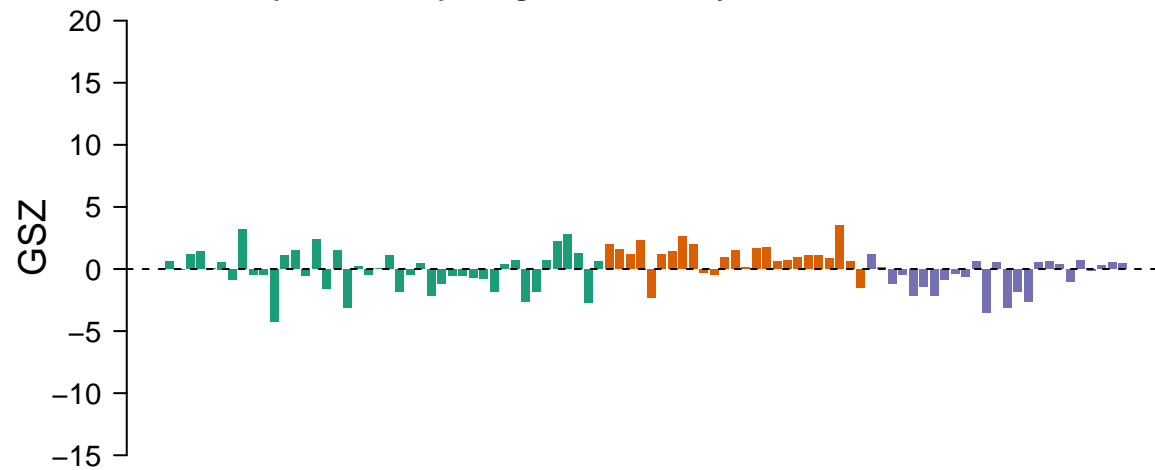


features = 42 , max = 2

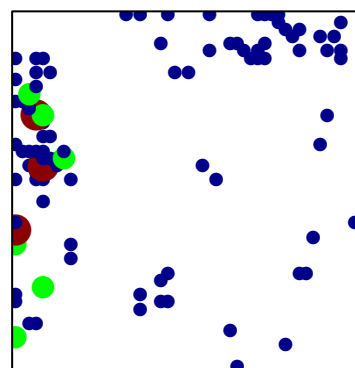
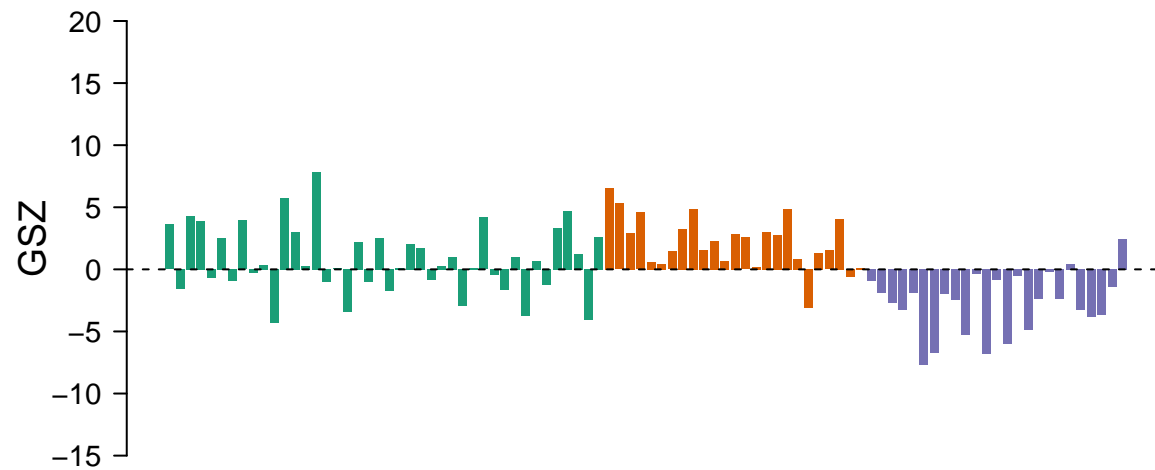
MARSON_FOXP3_TARGETS_DN



proton-transporting ATPase activity, rotational mechanism



KEGG_OXIDATIVE_PHOSPHORYLATION



hydrogen ion transmembrane transporter activity

