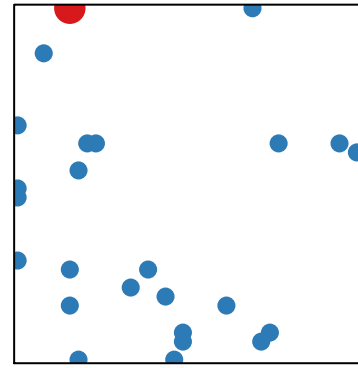
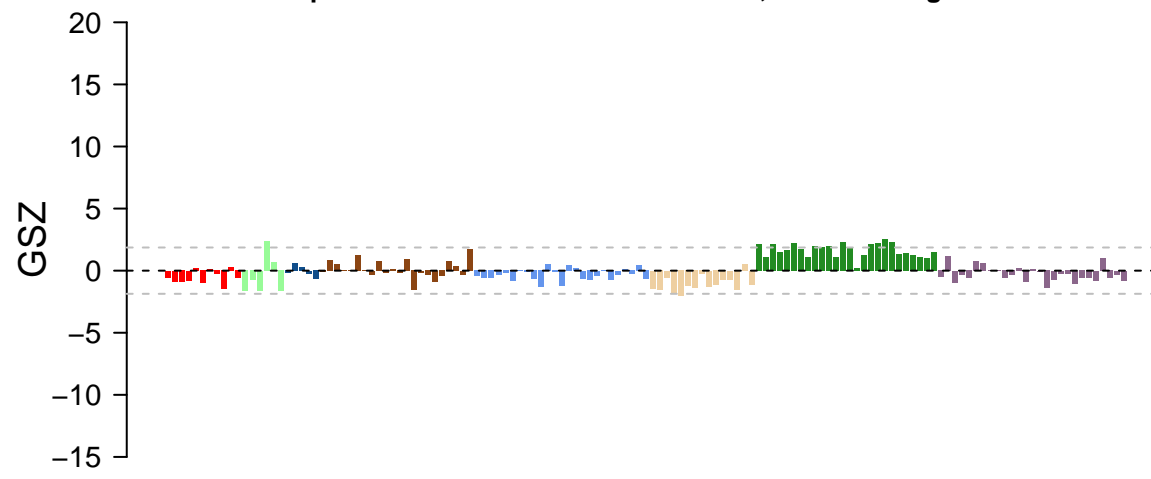
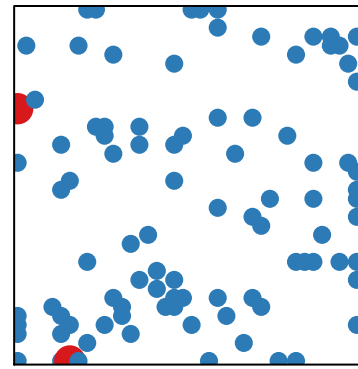
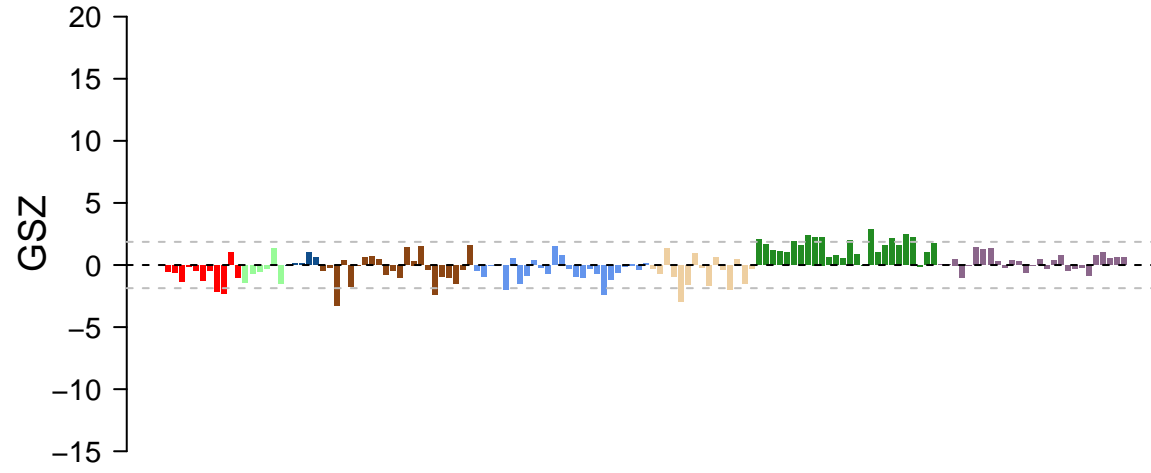


protein localization to chromosome, telomeric region



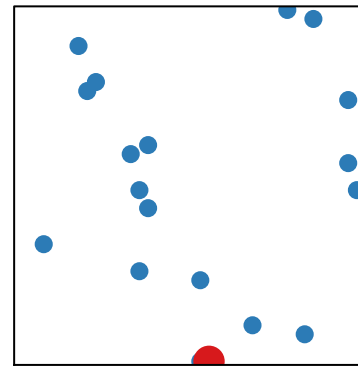
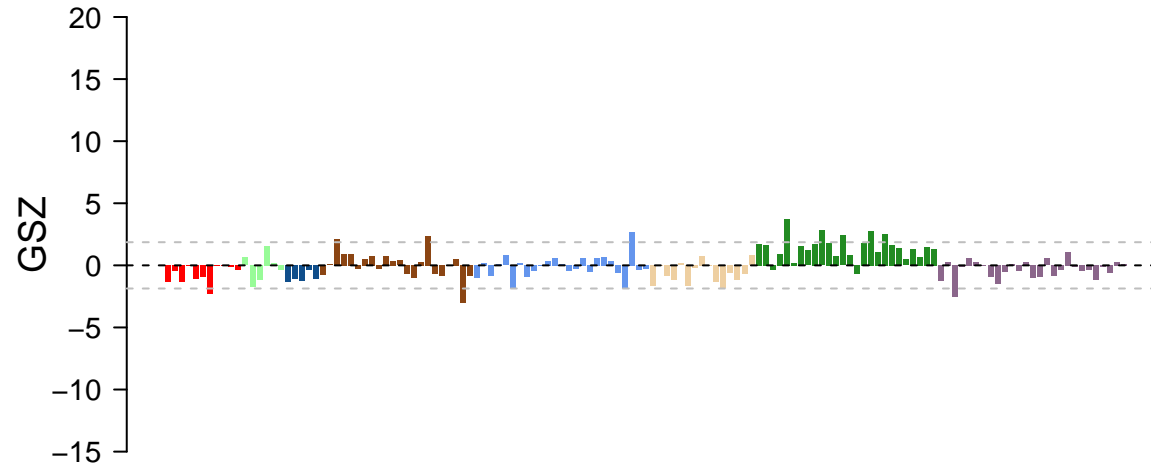
features = 10 , max = 2

protein targeting to membrane



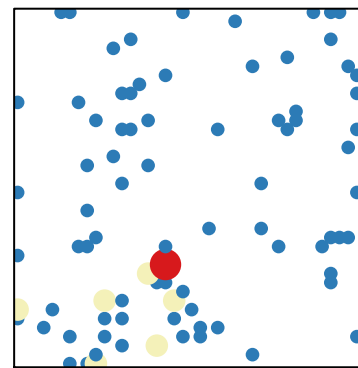
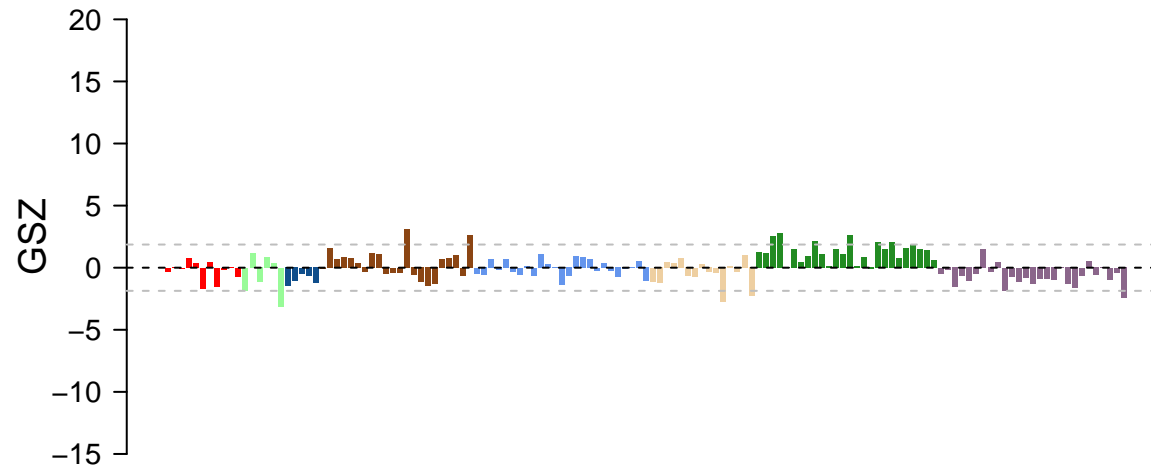
features = 45 , max = 2

neural precursor cell proliferation



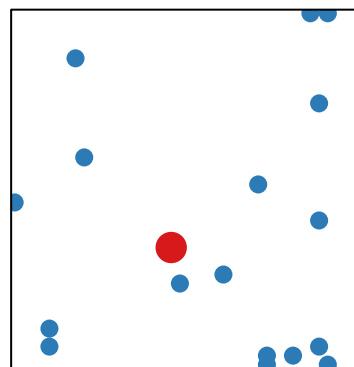
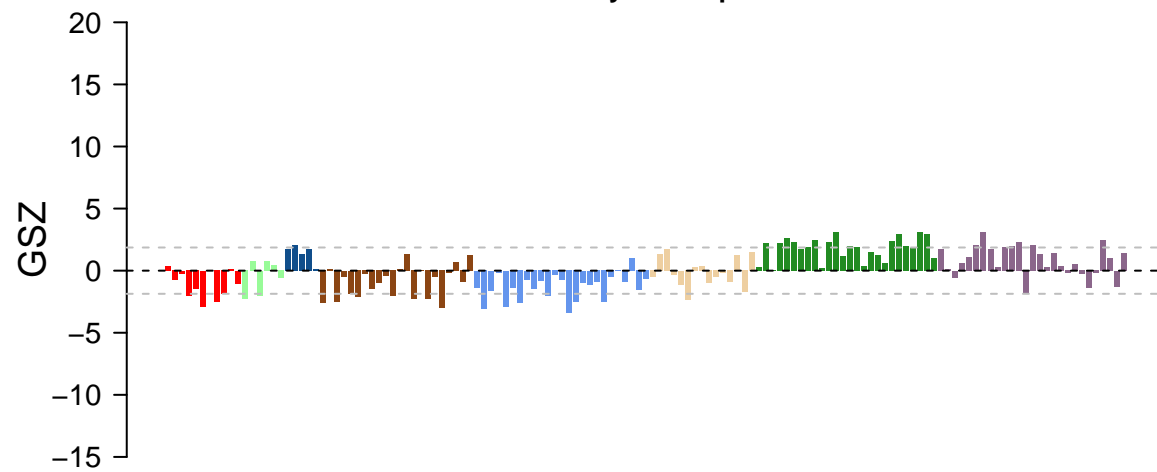
features = 12 , max = 2

endoplasmic reticulum unfolded protein response



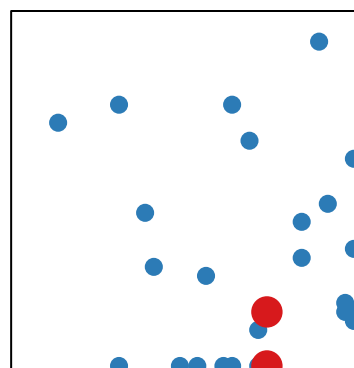
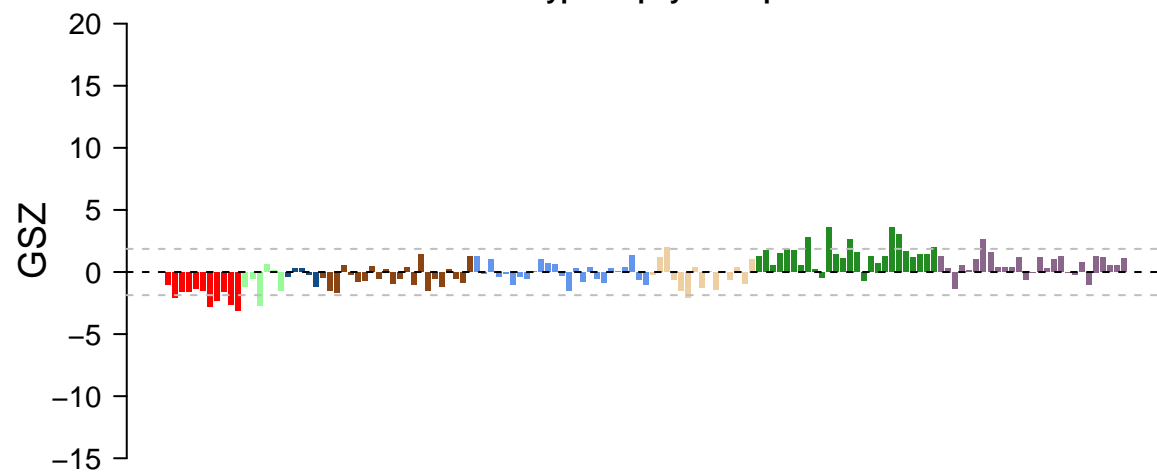
features = 43 , max = 3

cGMP biosynthetic process



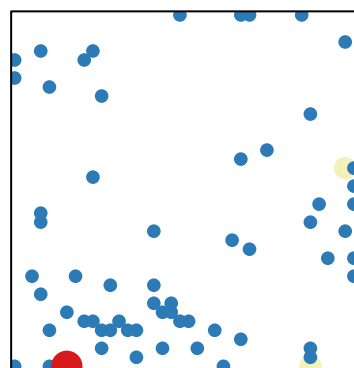
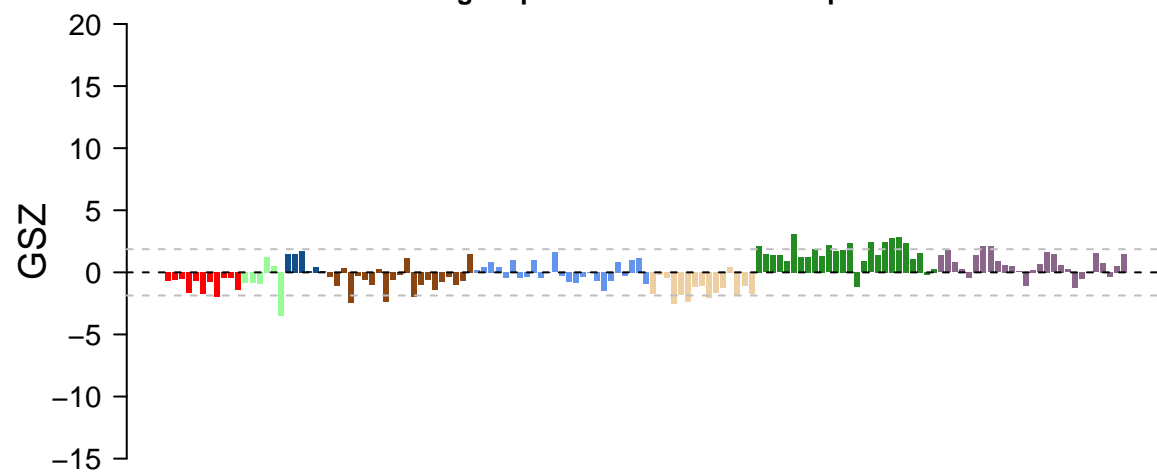
features = 11 , max = 2

cardiac muscle hypertrophy in response to stress



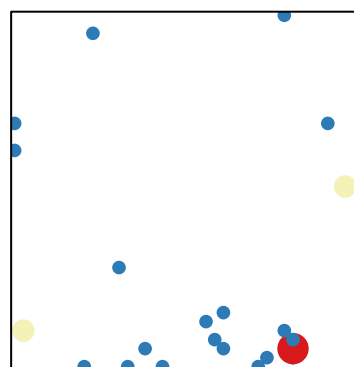
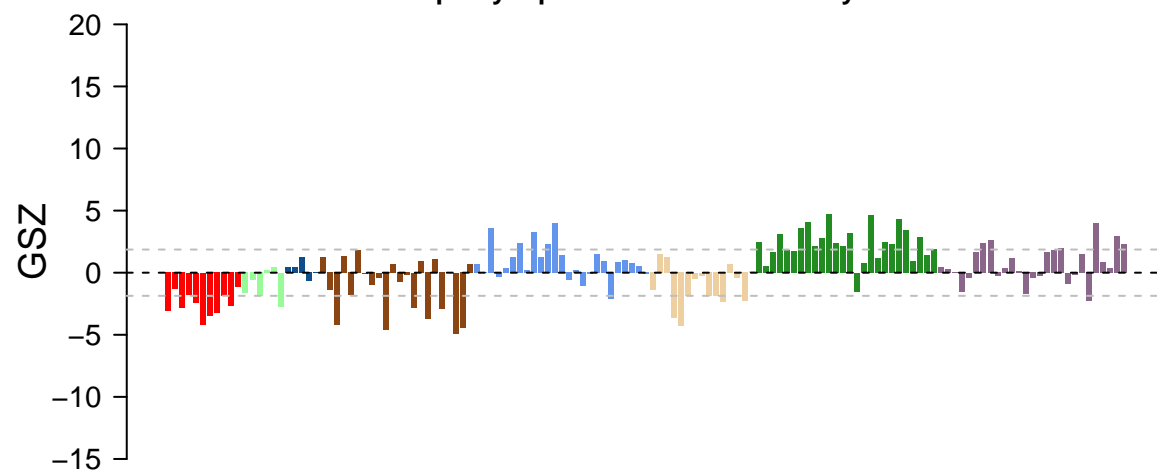
features = 14 , max = 2

Golgi to plasma membrane transport



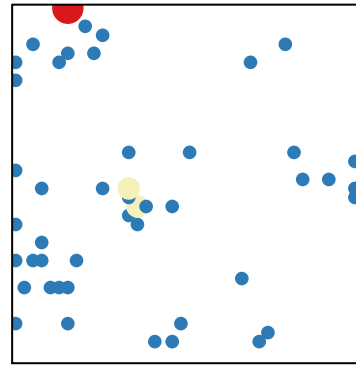
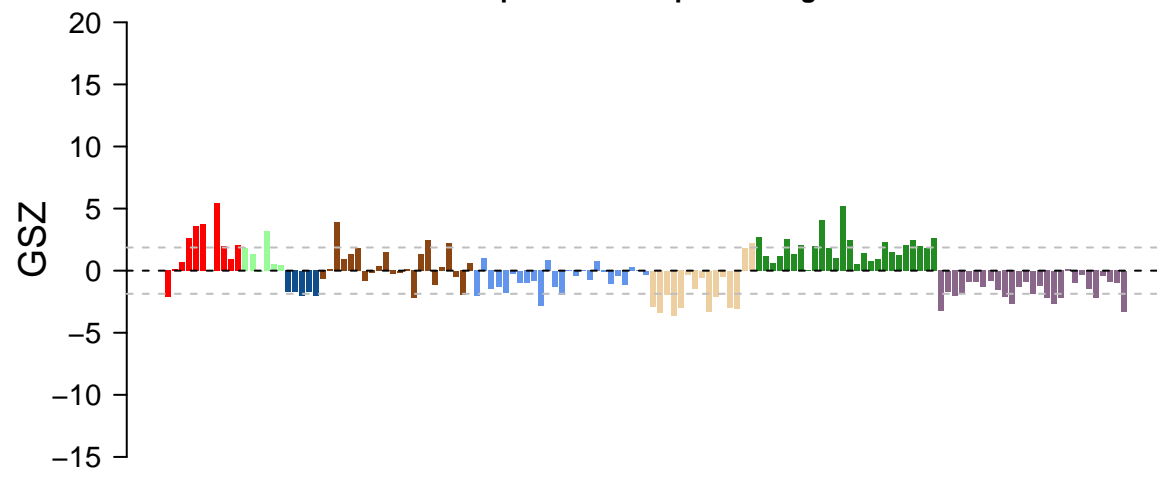
features = 26 , max = 3

presynaptic membrane assembly



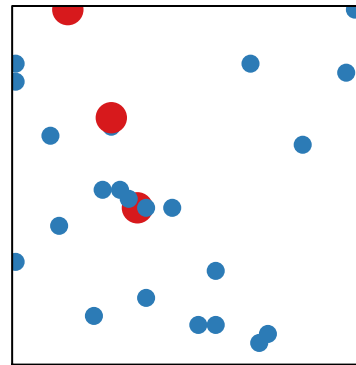
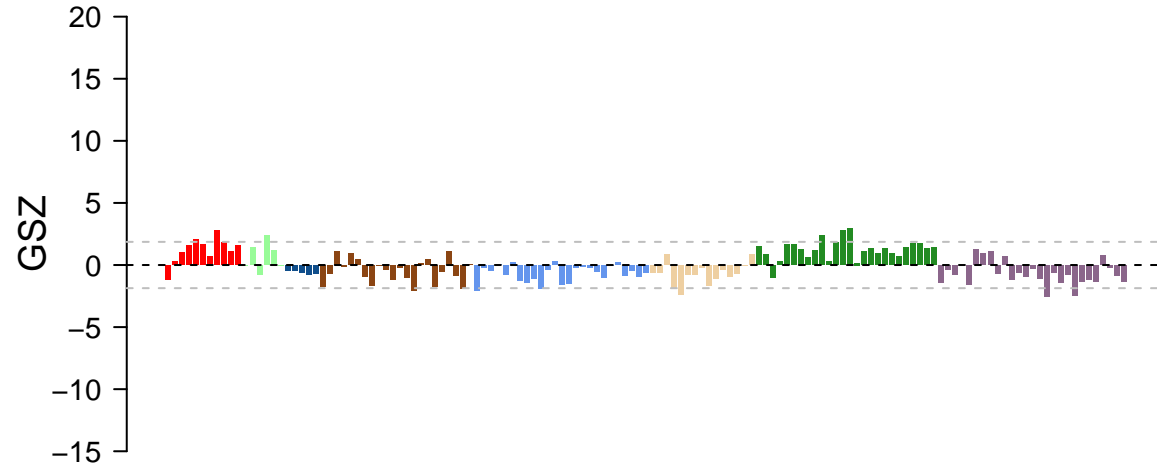
features = 10 , max = 3

replication fork processing



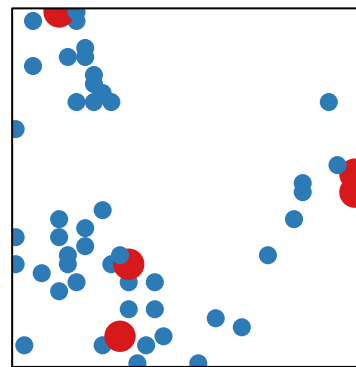
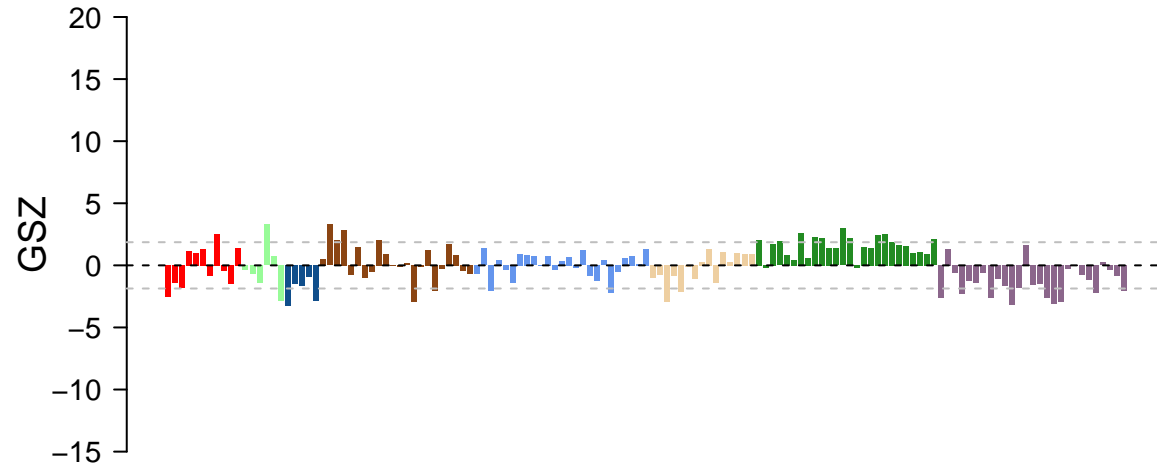
features = 24 , max = 3

cellular response to hydroxyurea



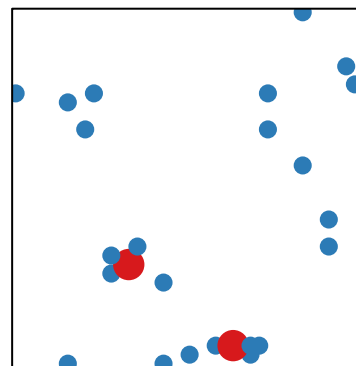
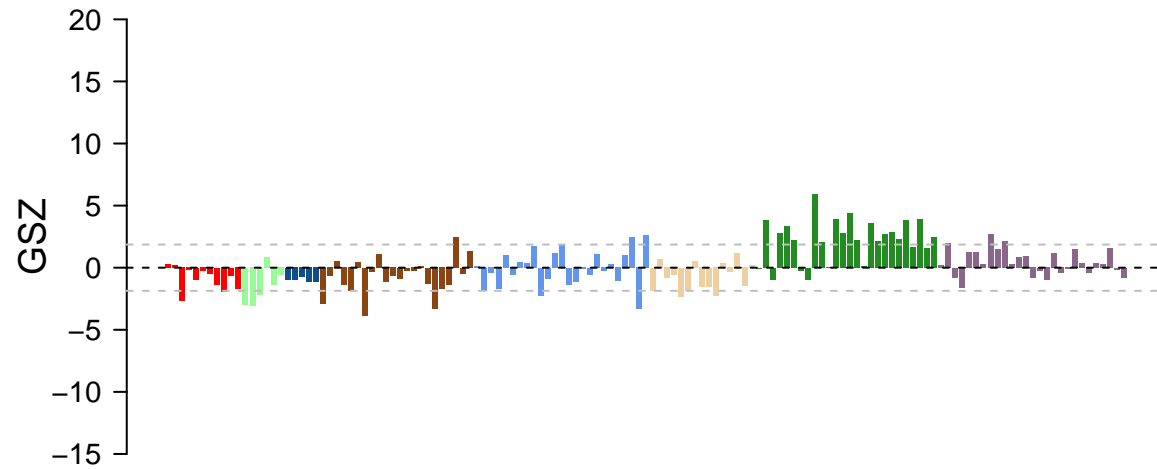
features = 11 , max = 2

spliceosomal complex assembly



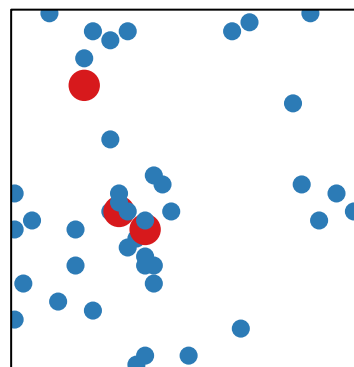
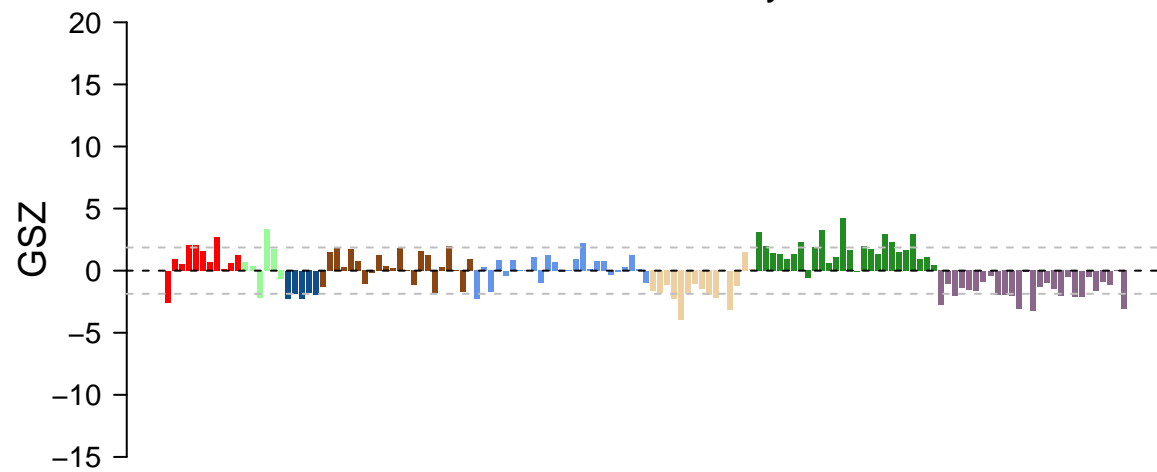
features = 25 , max = 2

isoprenoid biosynthetic process



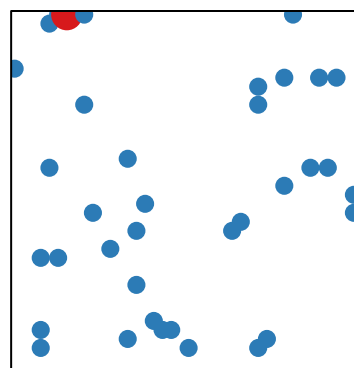
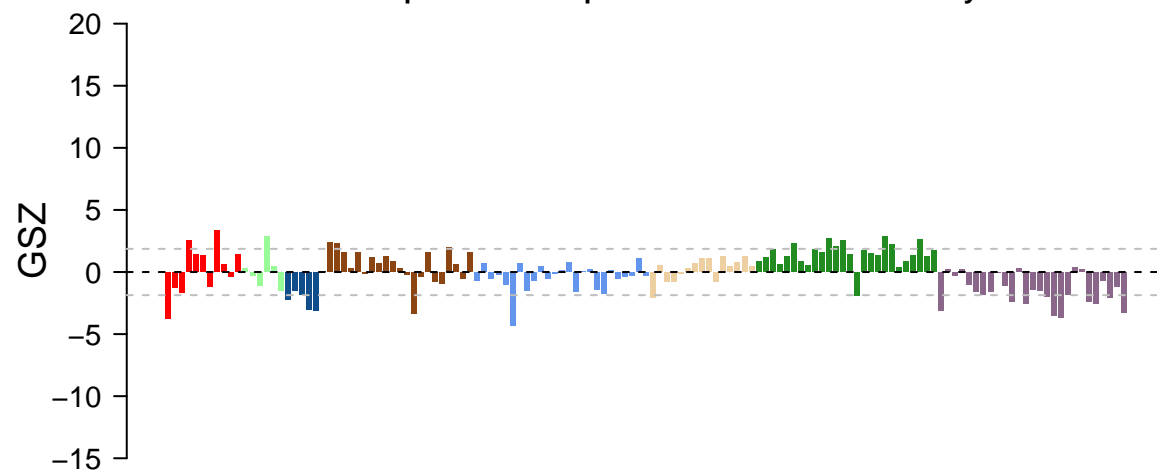
features = 15 , max = 2

DNA helicase activity



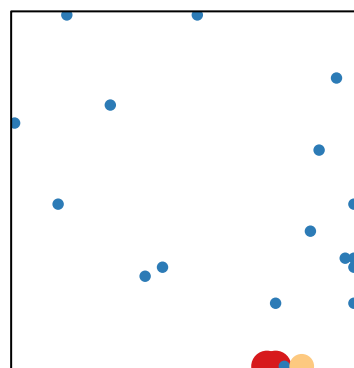
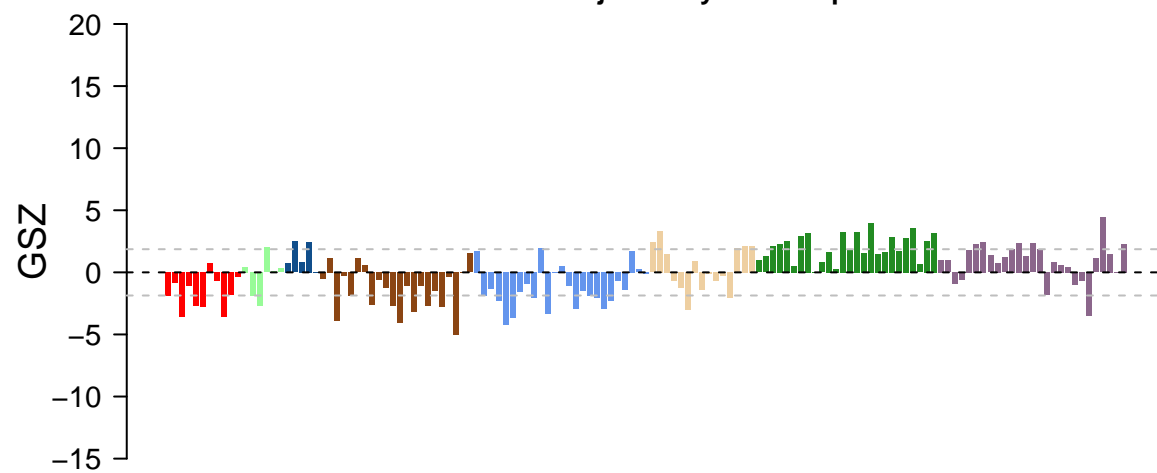
features = 20 , max = 2

DNA replication-independent nucleosome assembly



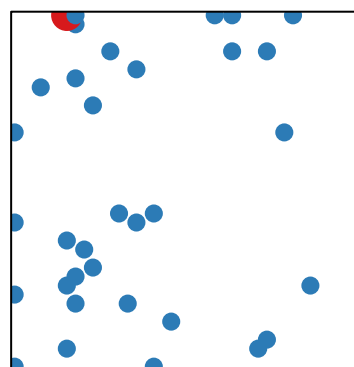
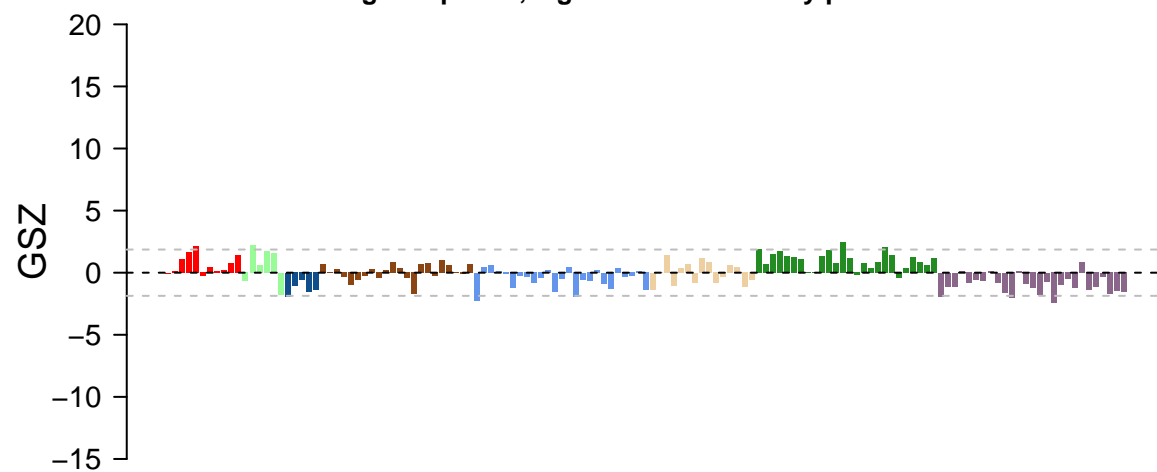
features = 20 , max = 2

cerebellar Purkinje cell layer development



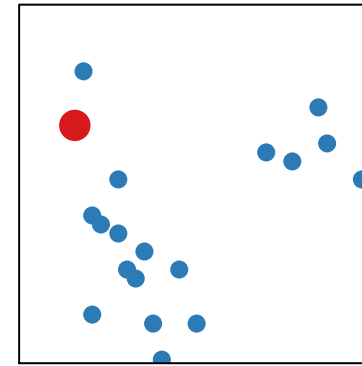
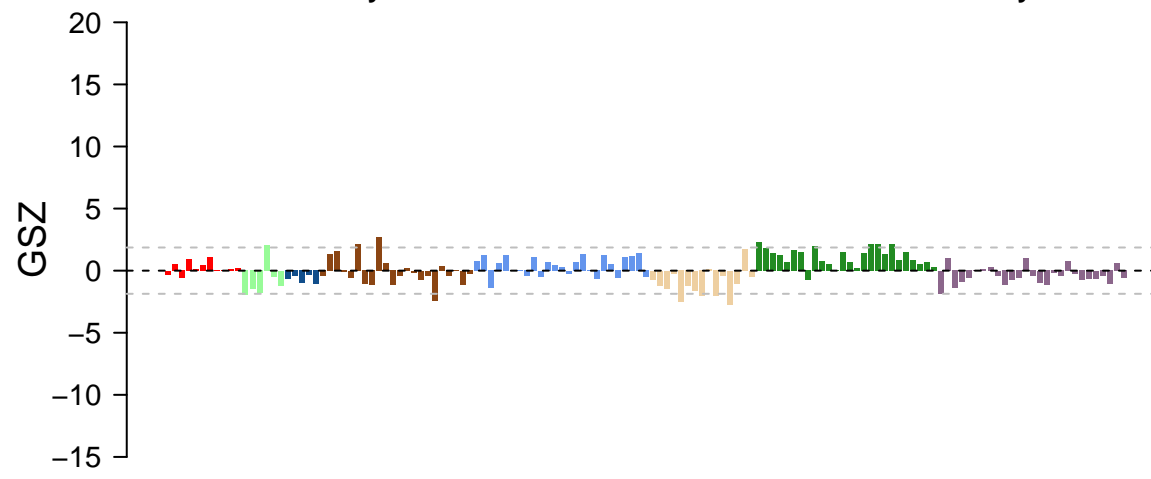
features = 13 , max = 4

DNA damage response, signal transduction by p53 class mediator



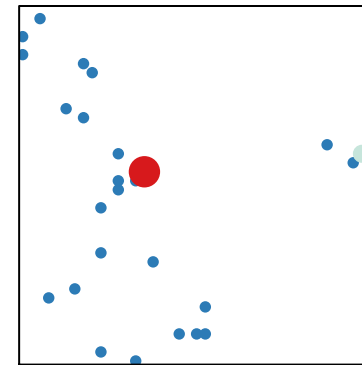
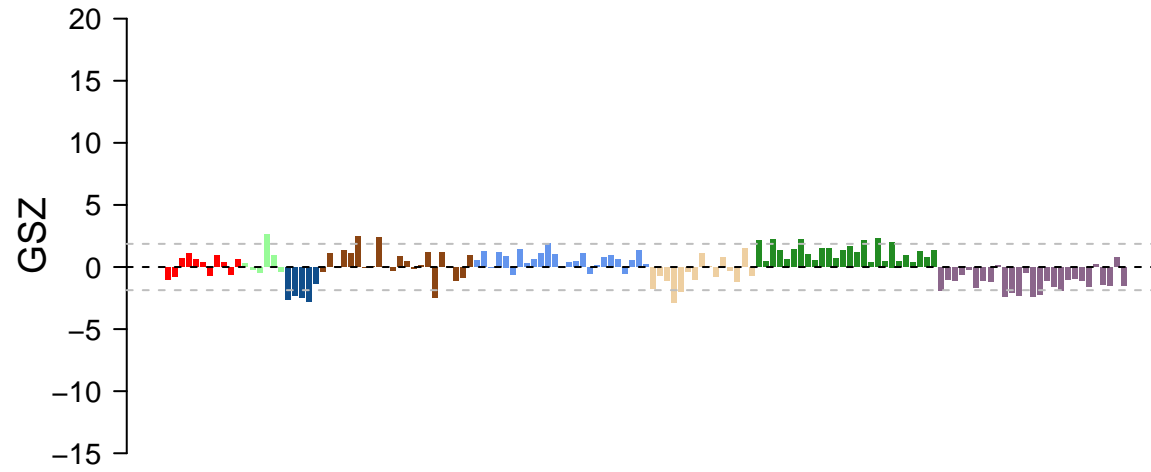
features = 15 , max = 2

aminoacyl-tRNA metabolism involved in translational fidelity



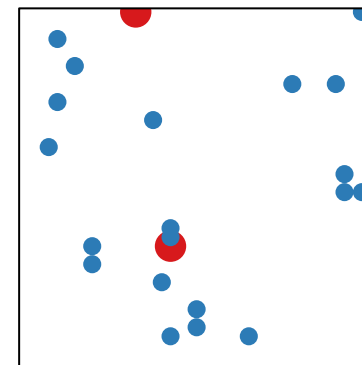
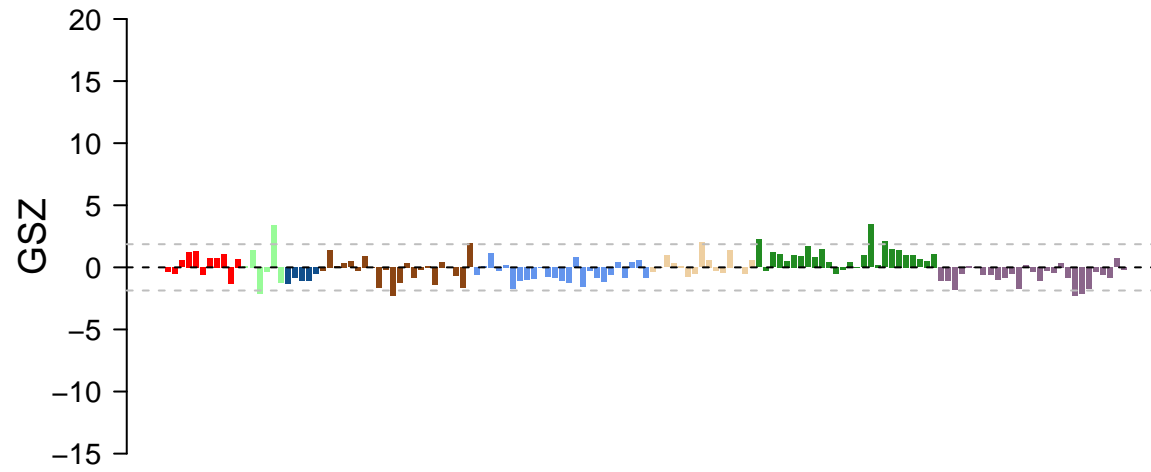
features = 11 , max = 2

IRES-dependent viral translational initiation



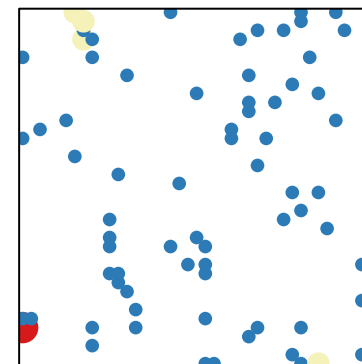
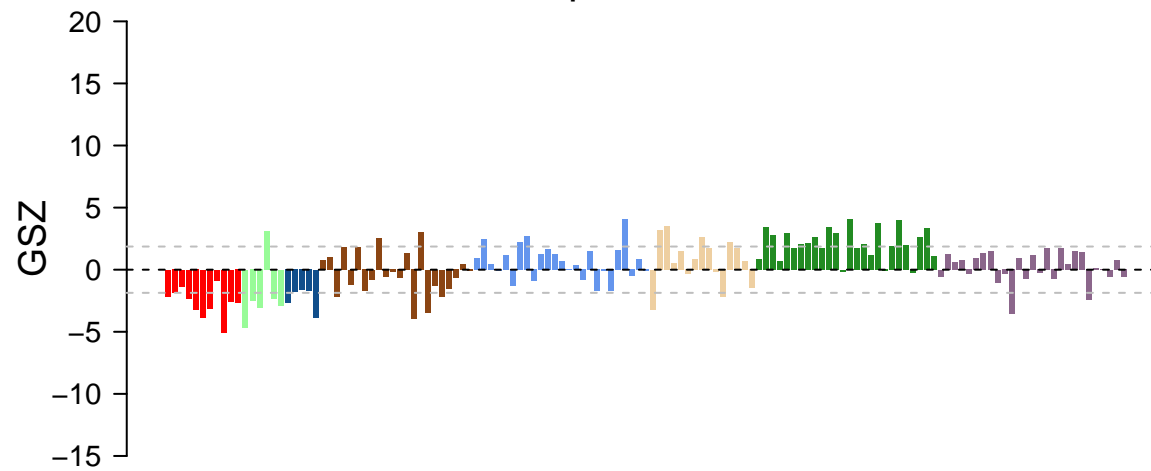
features = 10 , max = 4

regulation of MAP kinase activity



features = 10 , max = 2

cellular response to BMP stimulus



features = 36 , max = 3