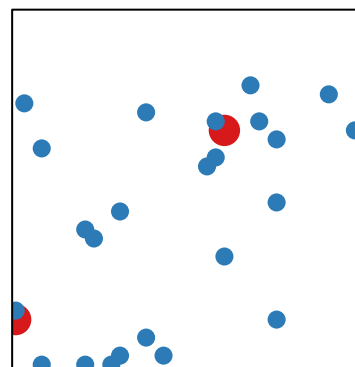
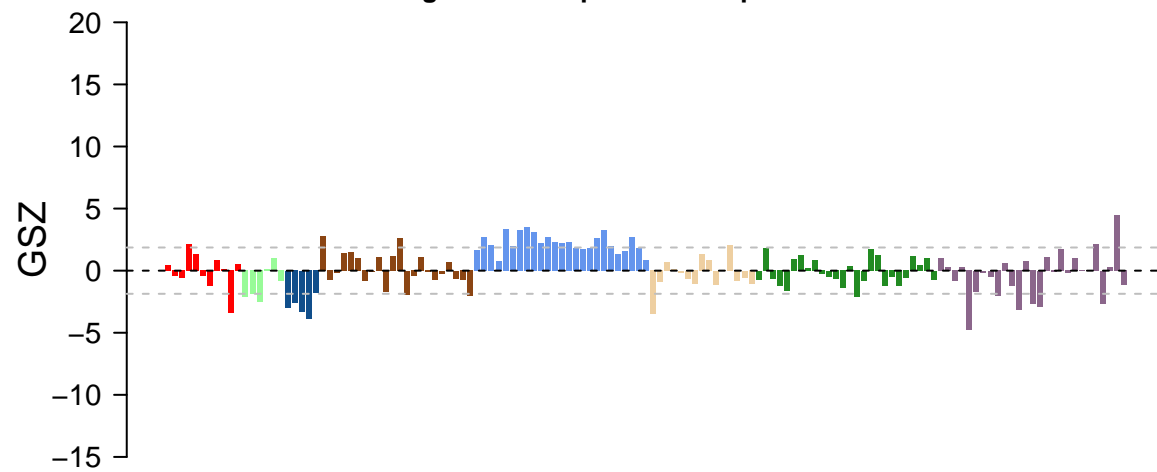
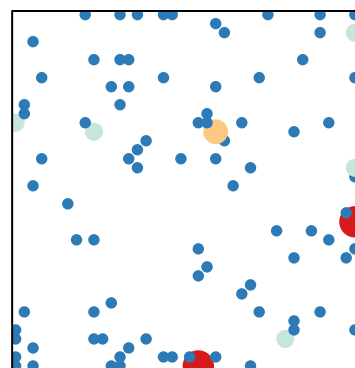
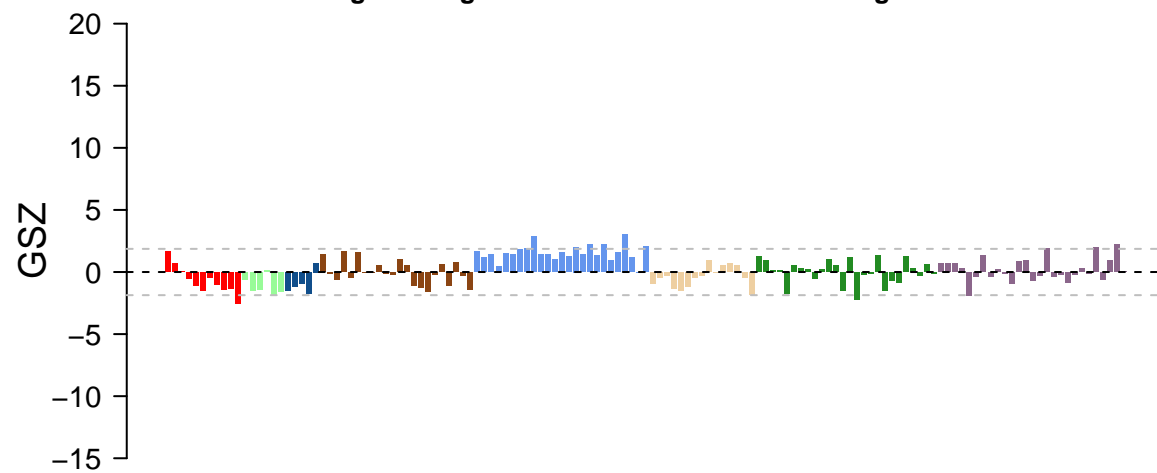


regulation of epithelial cell proliferation



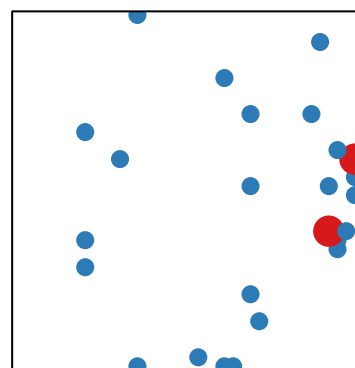
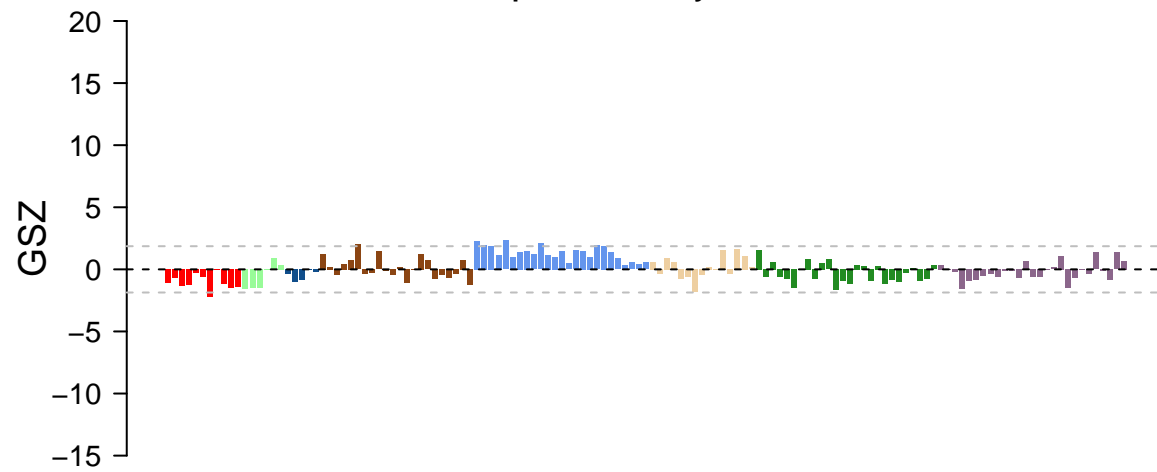
features = 13 , max = 2

negative regulation of cold-induced thermogenesis



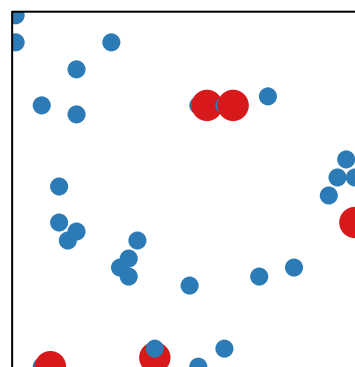
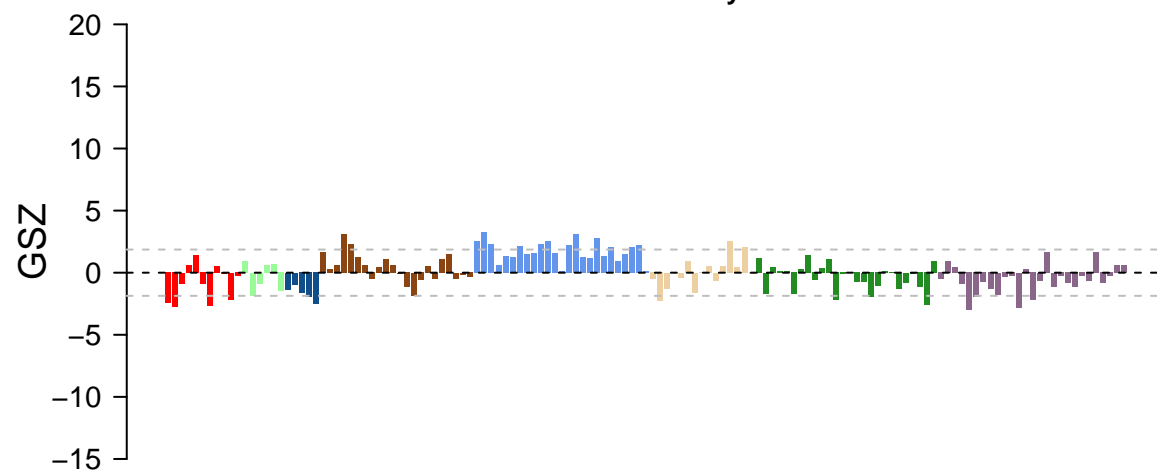
features = 46 , max = 4

protein deacetylation



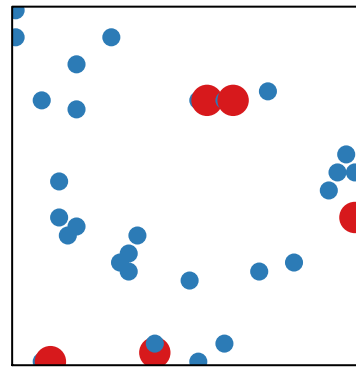
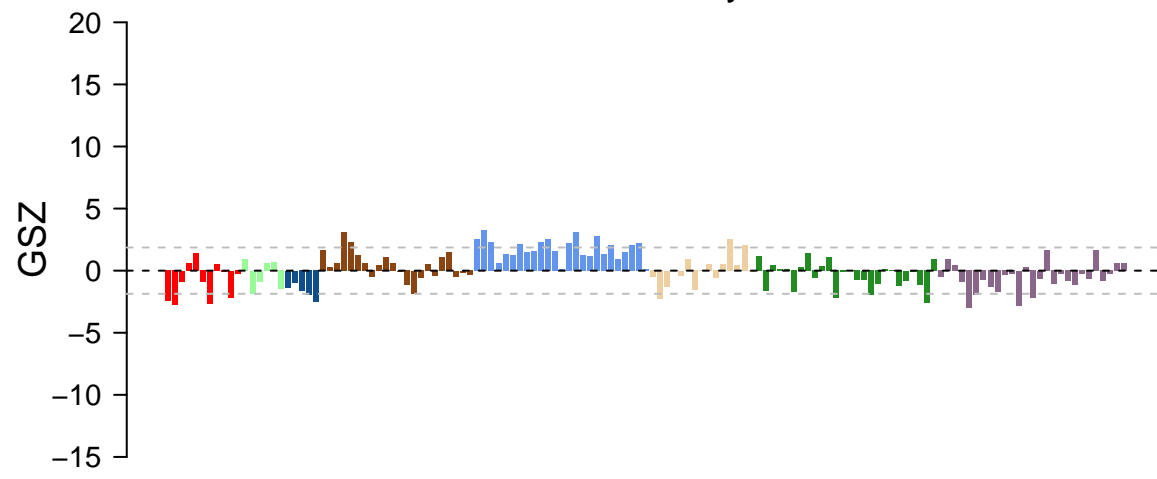
features = 13 , max = 2

histone H4-K5 acetylation

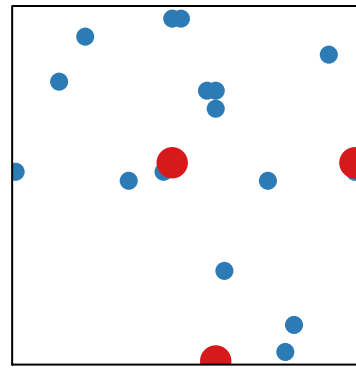
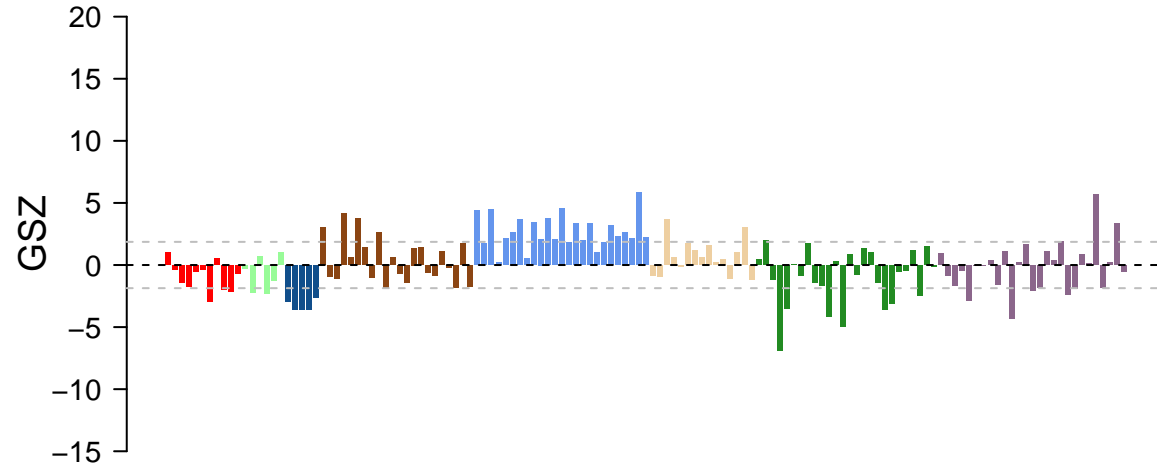


features = 16 , max = 2

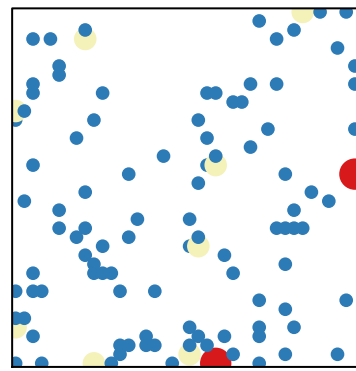
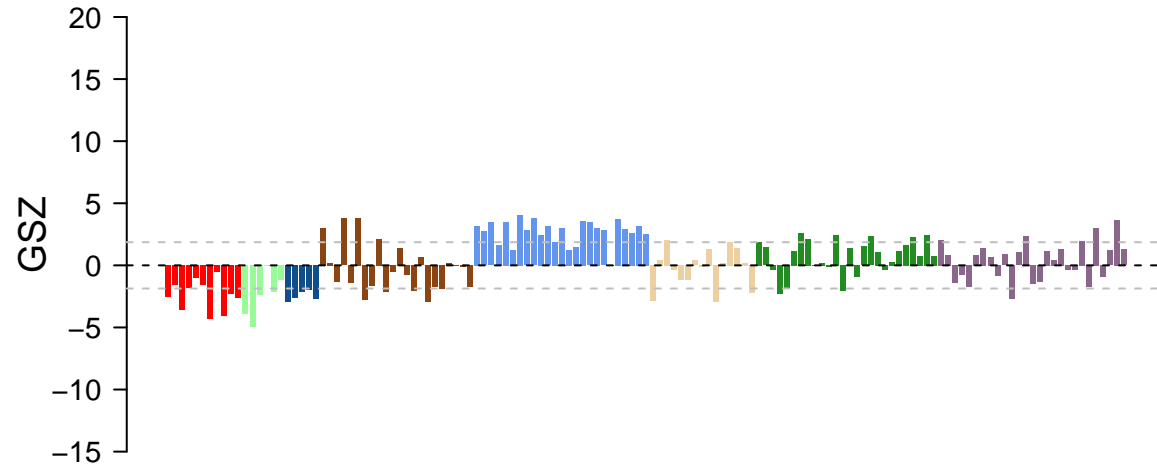
histone H4-K8 acetylation



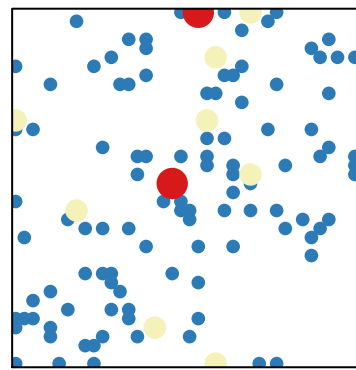
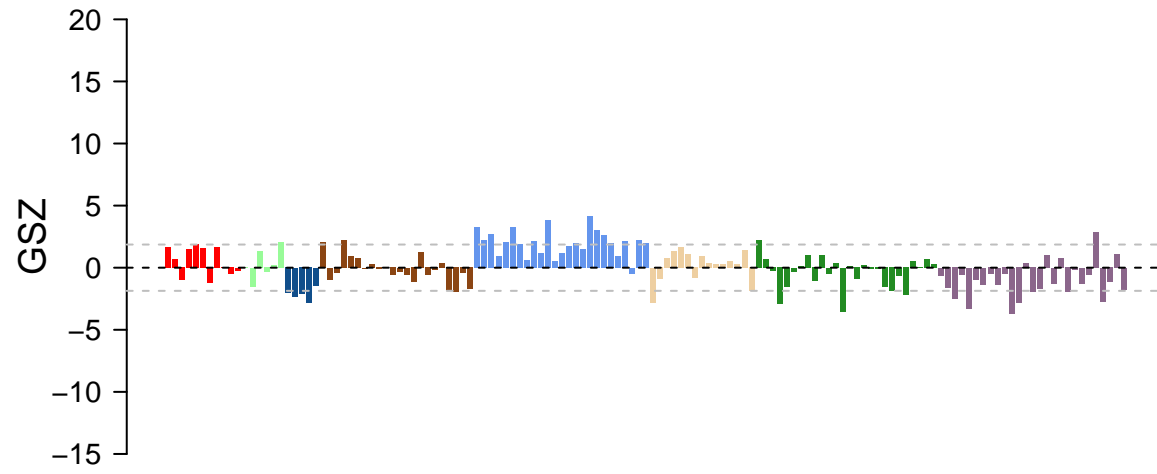
planar cell polarity pathway involved in neural tube closure



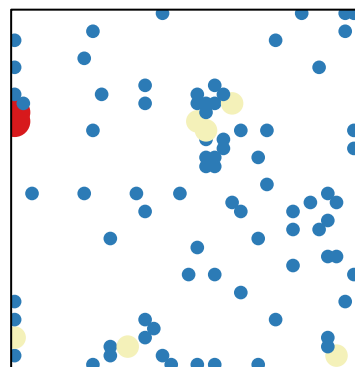
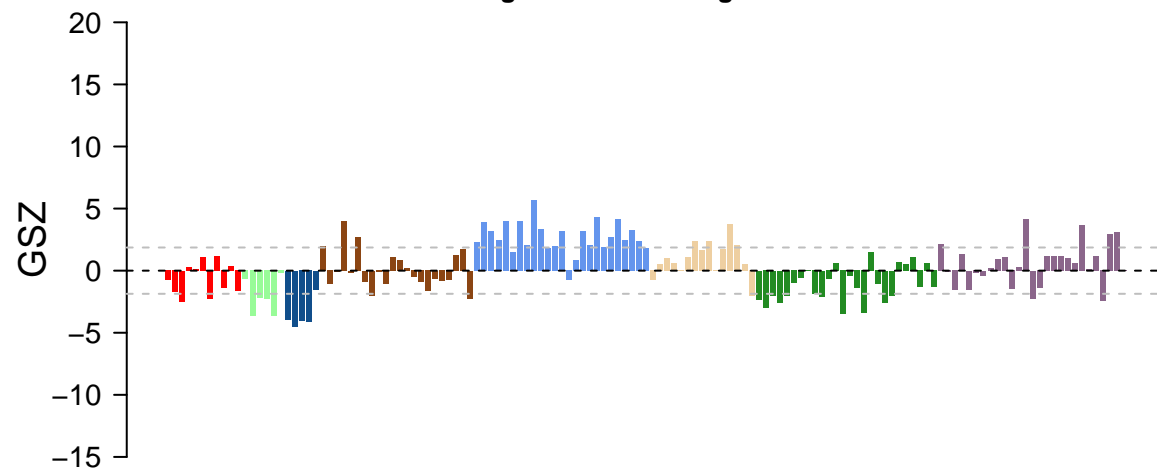
somitogenesis



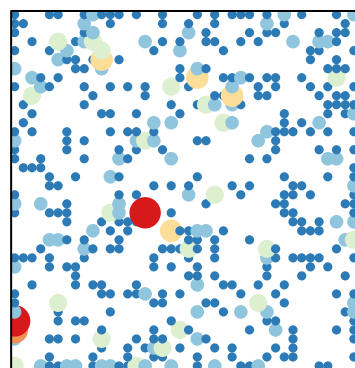
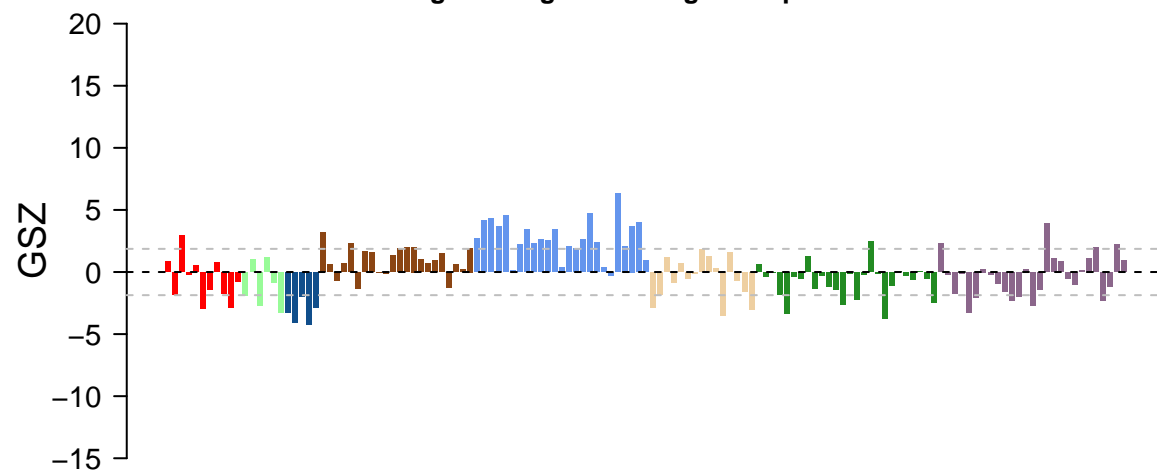
embryonic digit morphogenesis



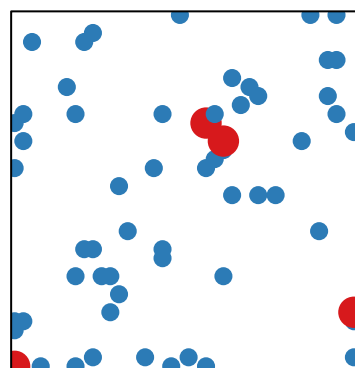
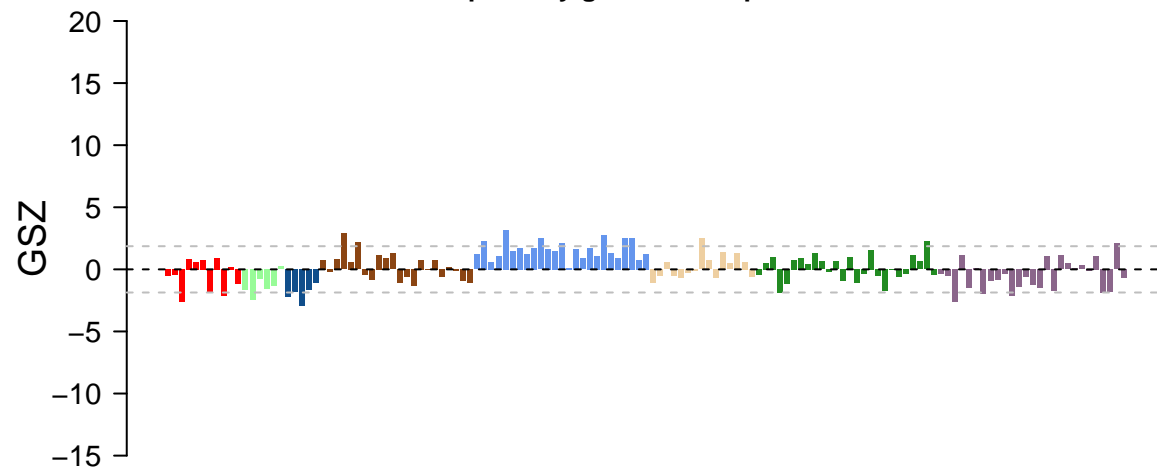
regulation of neurogenesis



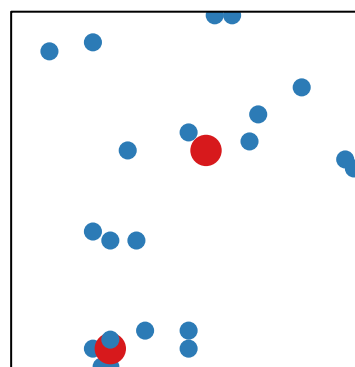
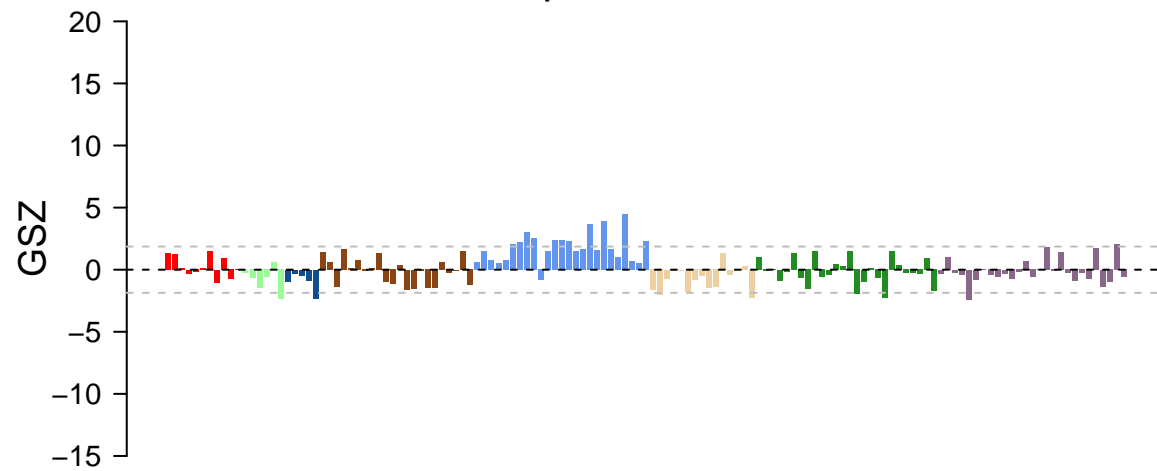
negative regulation of gene expression



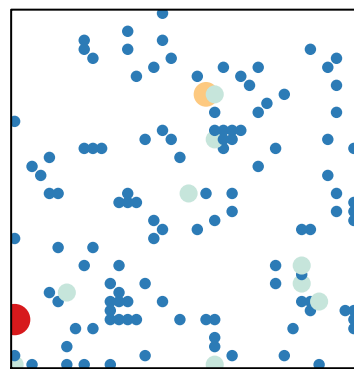
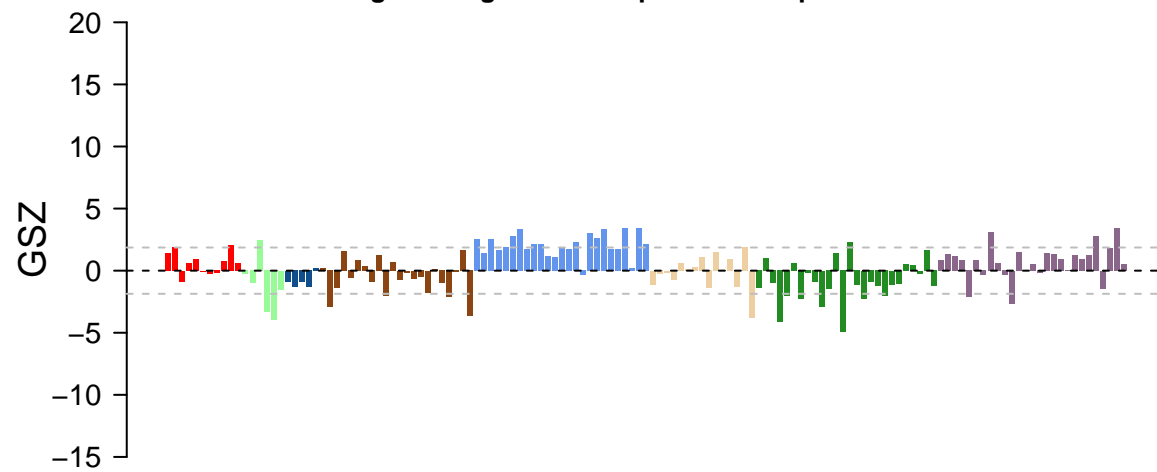
pituitary gland development



cellular response to leucine starvation

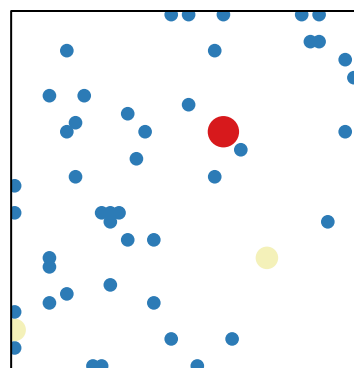
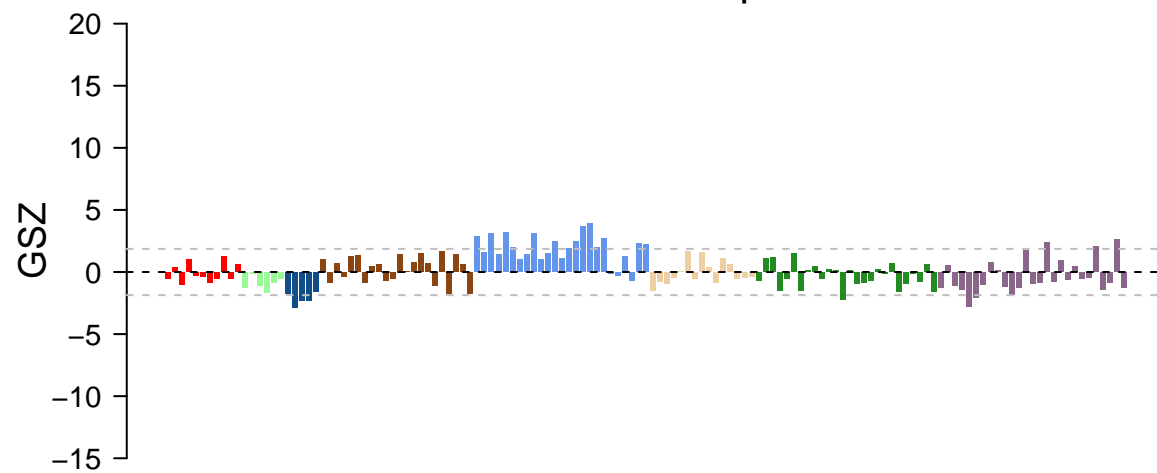


negative regulation of epithelial cell proliferation



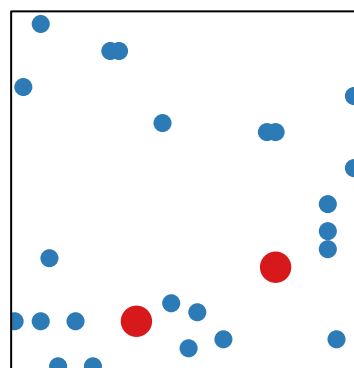
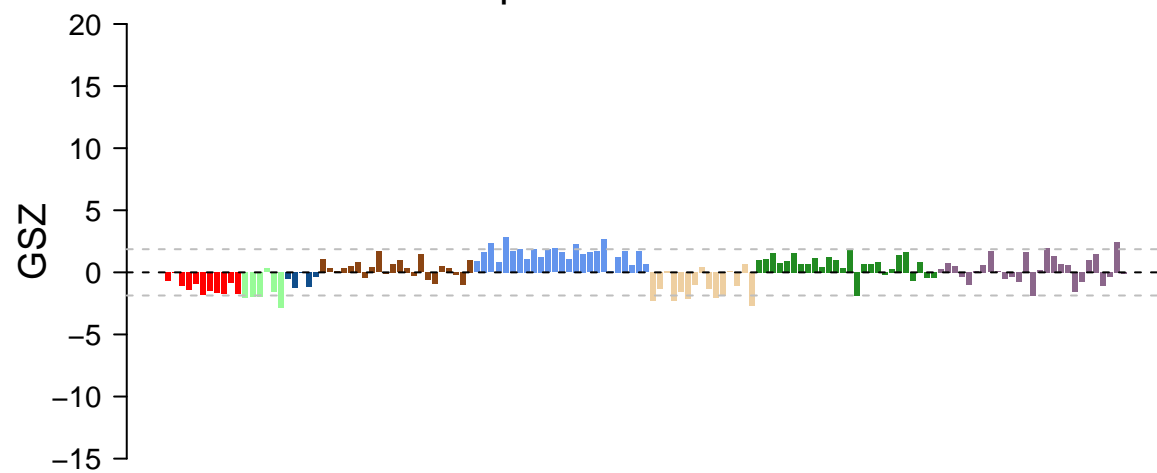
features = 62 , max = 4

one-carbon metabolic process



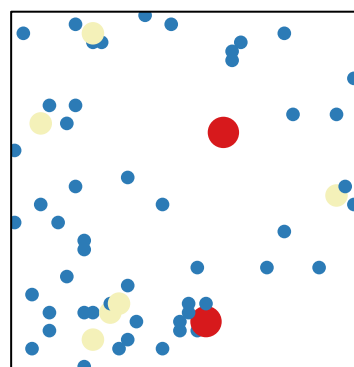
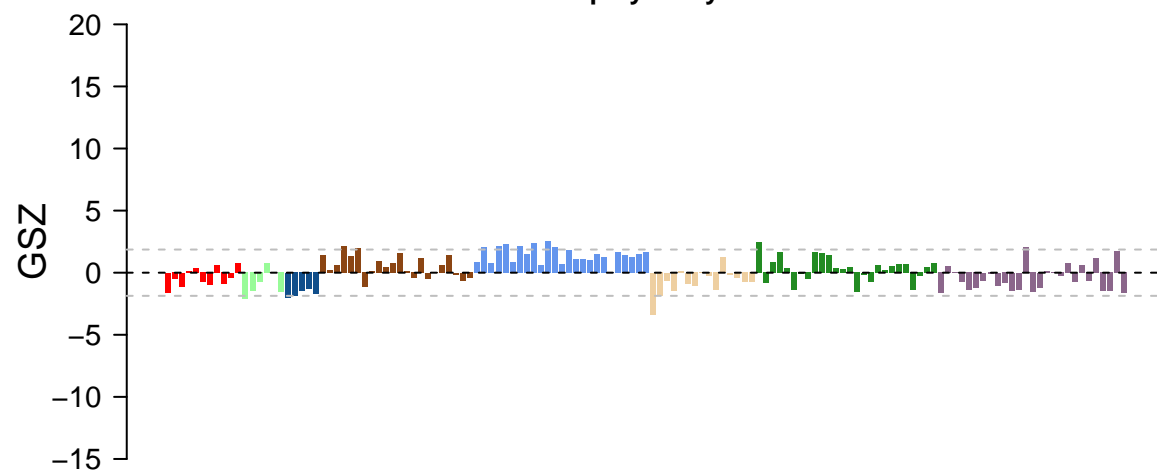
features = 22 , max = 3

protein hexamerization



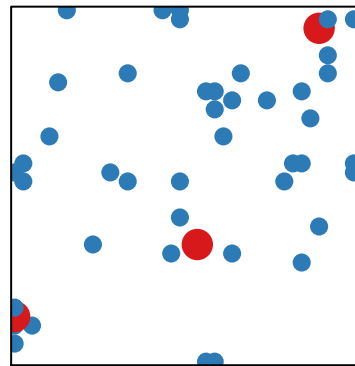
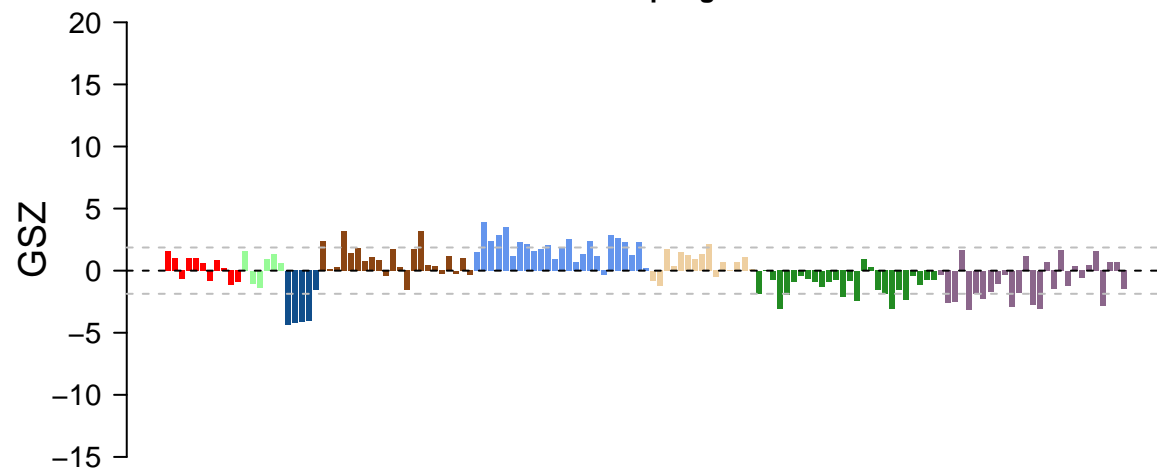
features = 13 , max = 2

mRNA polyadenylation



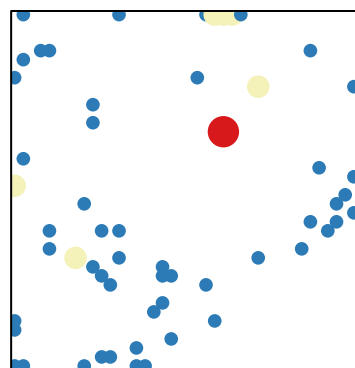
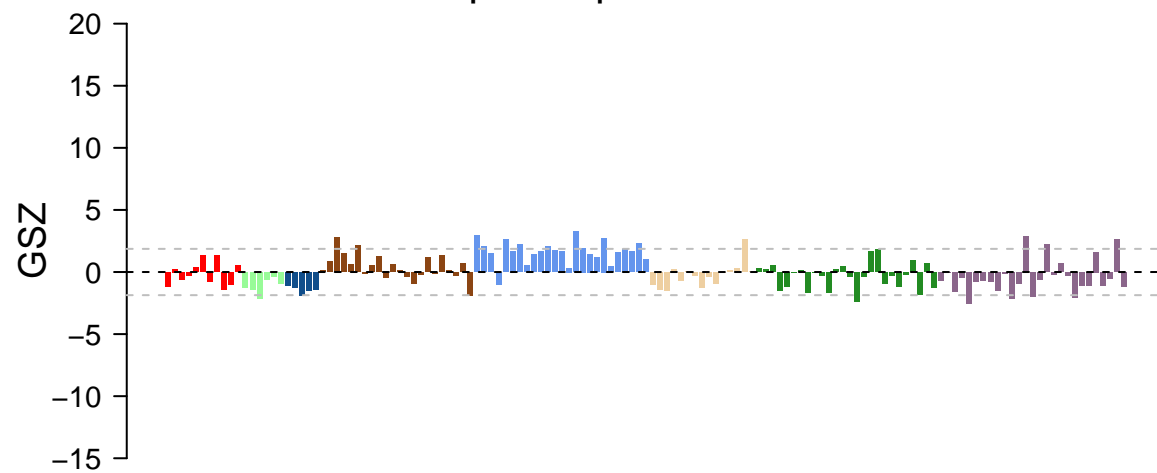
features = 28 , max = 3

cochlea morphogenesis



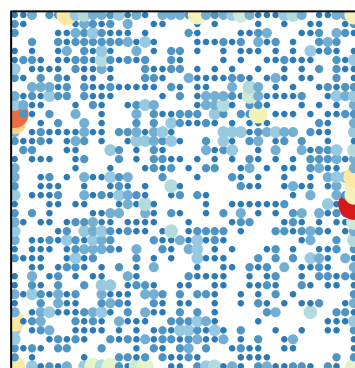
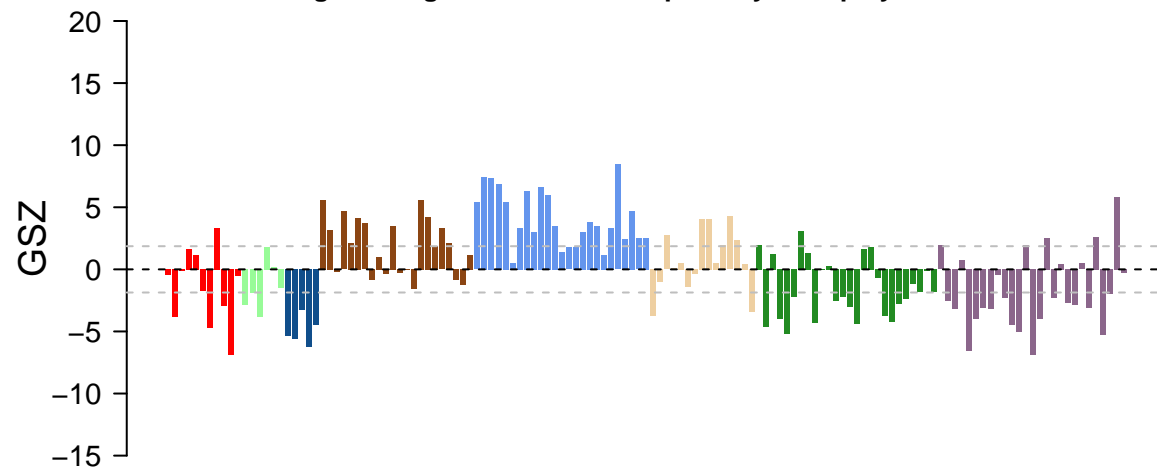
features = 25 , max = 2

protein export from nucleus



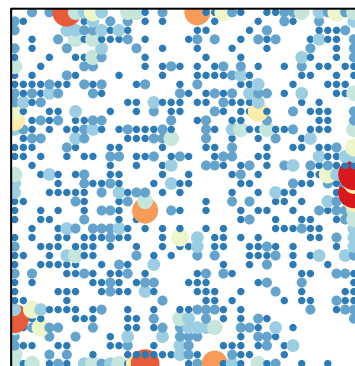
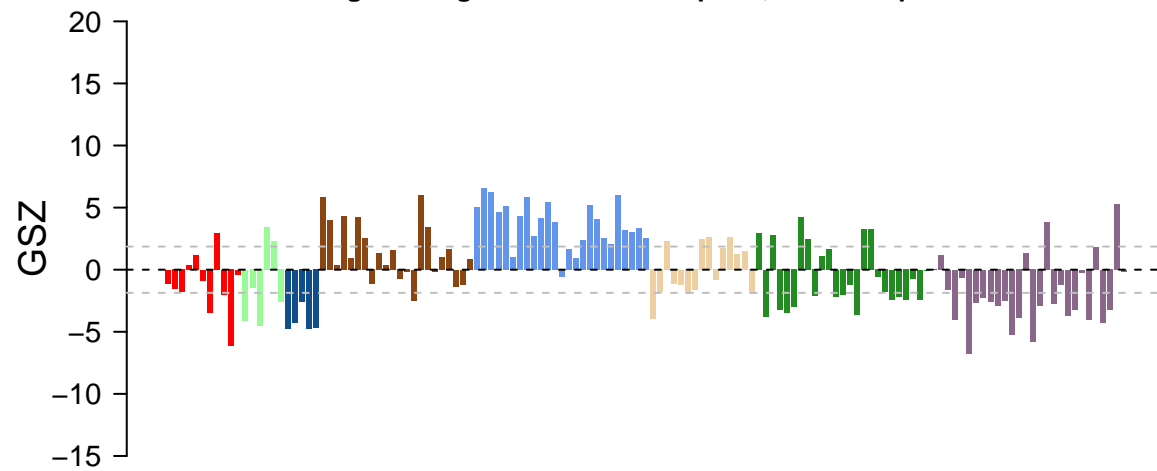
features = 26 , max = 3

negative regulation of transcription by RNA polymerase II



features = 783 , max = 15

negative regulation of transcription, DNA-templated



features = 541 , max = 10