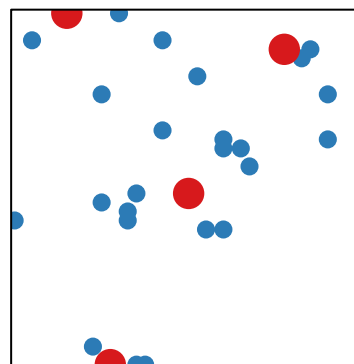
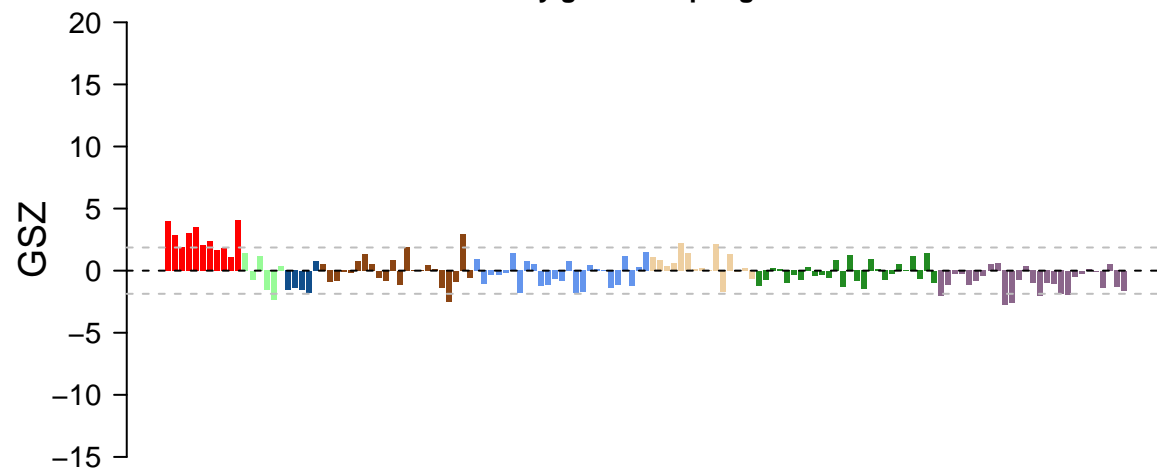
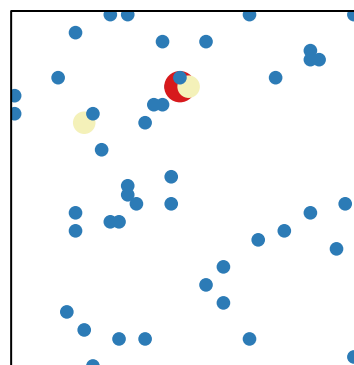
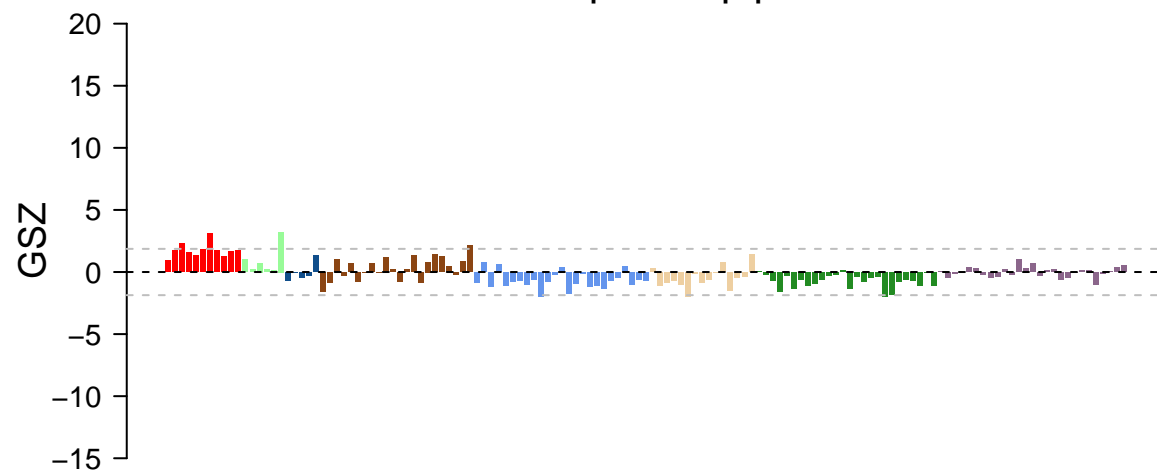


salivary gland morphogenesis



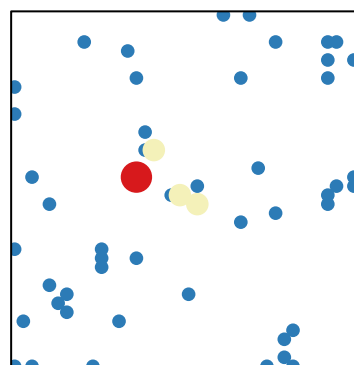
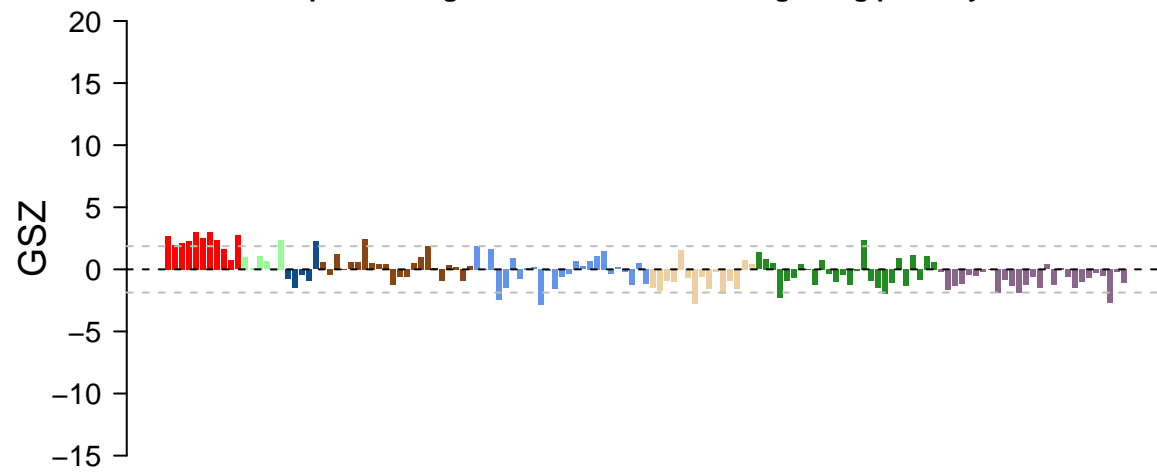
features = 10 , max = 2

execution phase of apoptosis



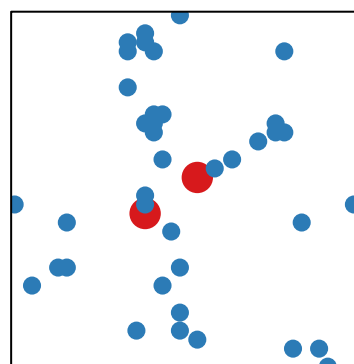
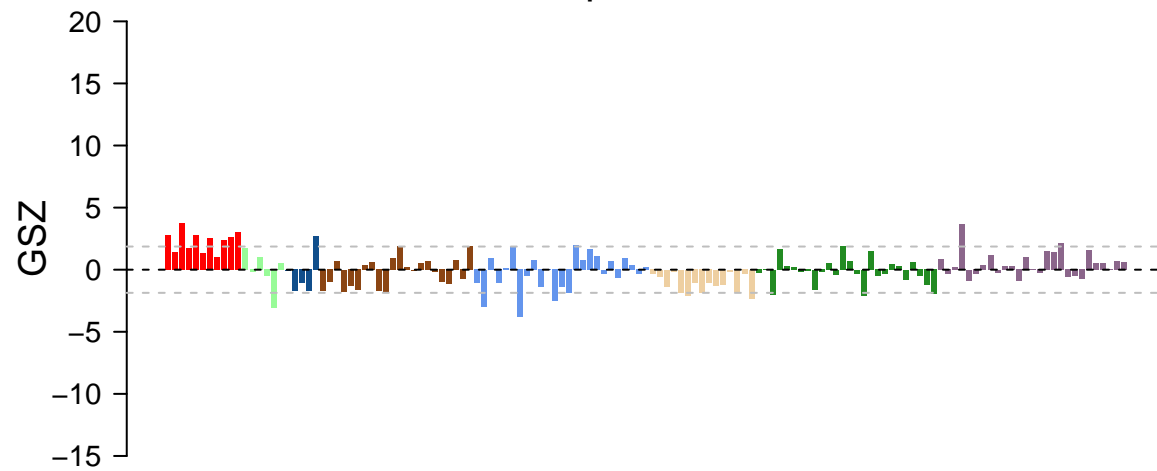
features = 20 , max = 3

positive regulation of smoothened signaling pathway



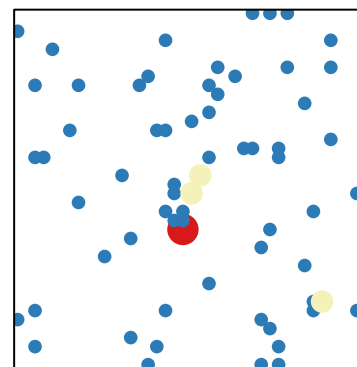
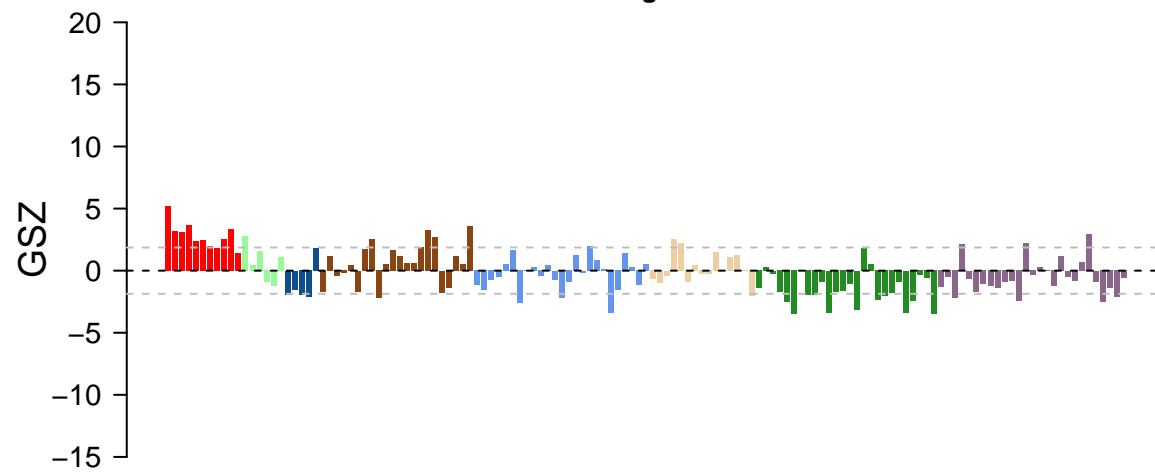
features = 30 , max = 3

cellular response to nitric oxide



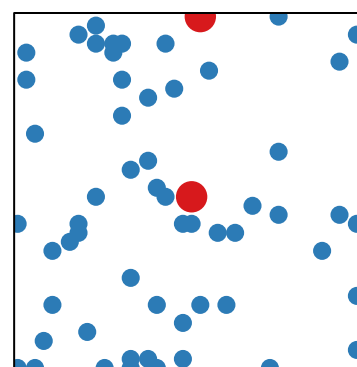
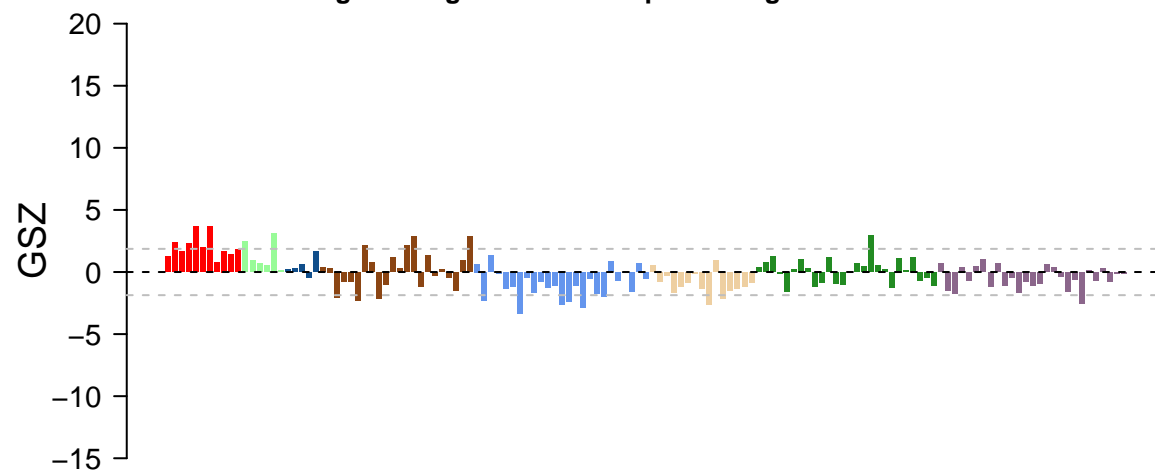
features = 15 , max = 2

odontogenesis



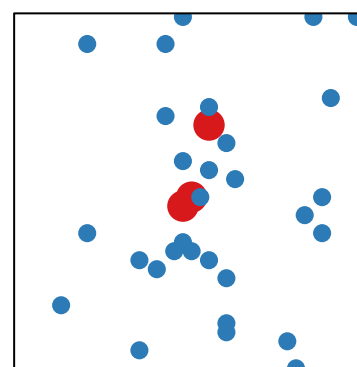
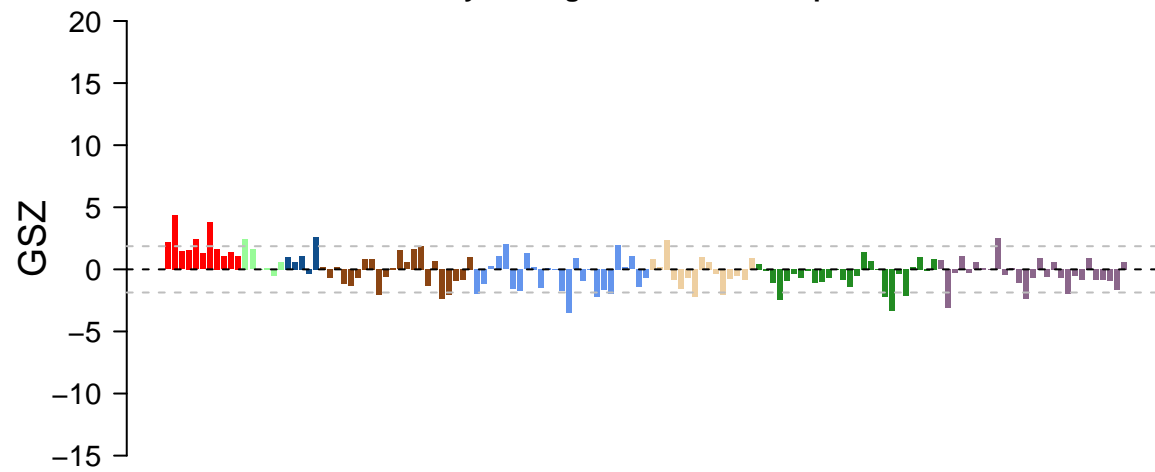
features = 34 , max = 3

negative regulation of Ras protein signal transduction



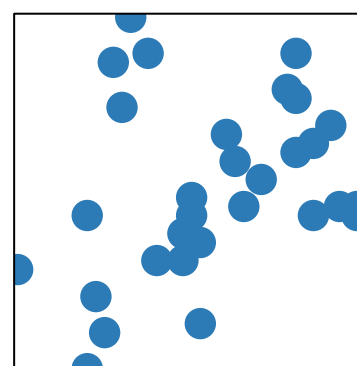
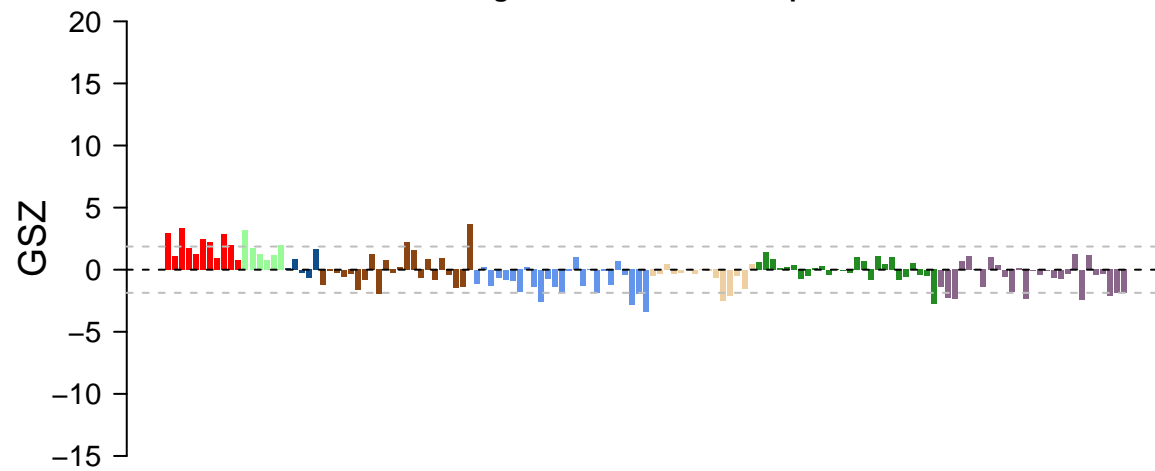
features = 25 , max = 2

embryonic digestive tract development



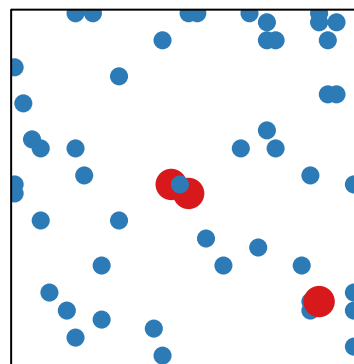
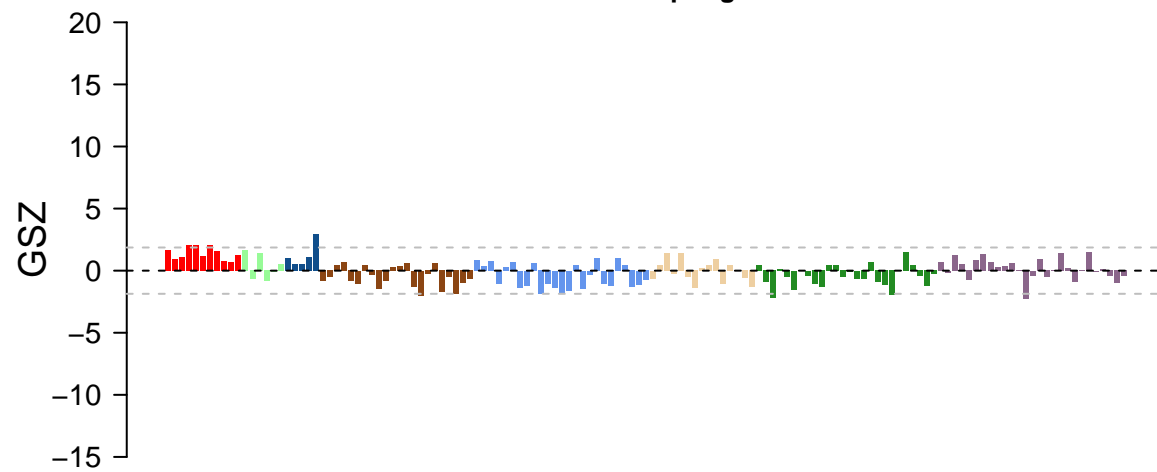
features = 18 , max = 2

drug transmembrane transport



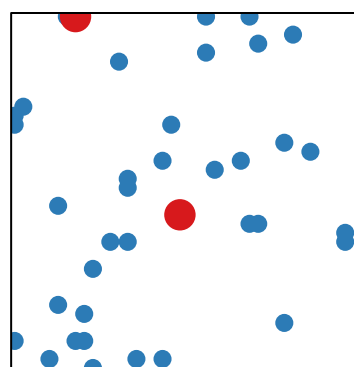
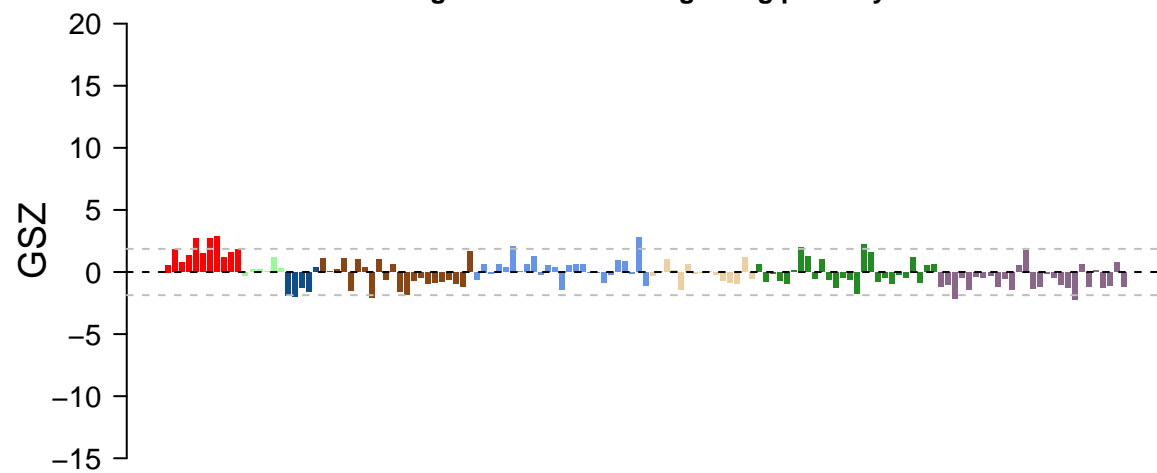
features = 19 , max = 1

hair follicle morphogenesis



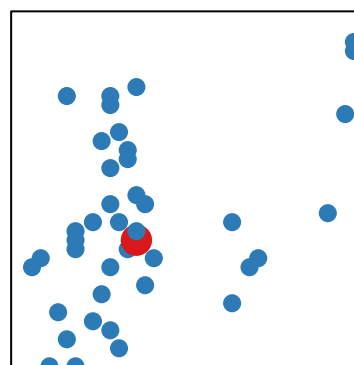
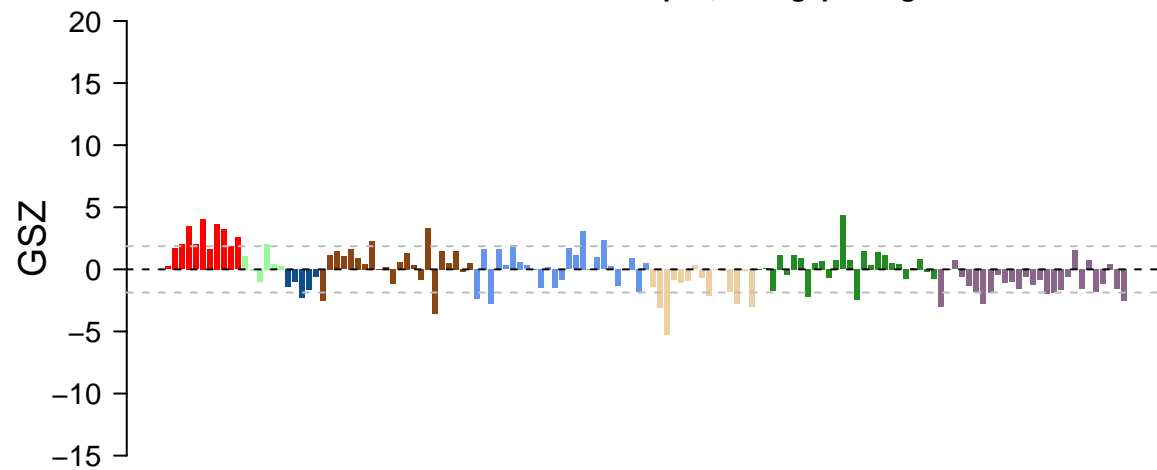
features = 25 , max = 2

regulation of Notch signaling pathway



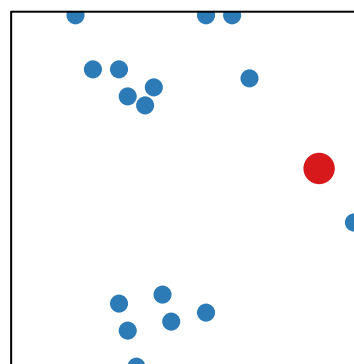
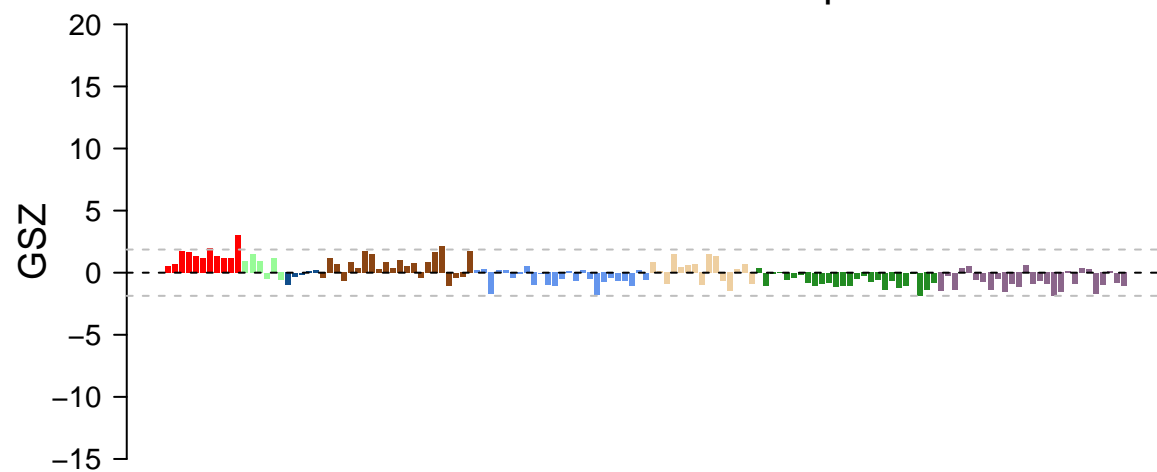
features = 17 , max = 2

nucleotide–excision repair, DNA gap filling



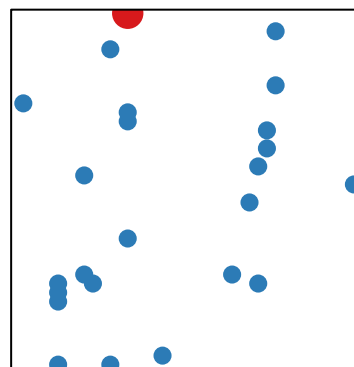
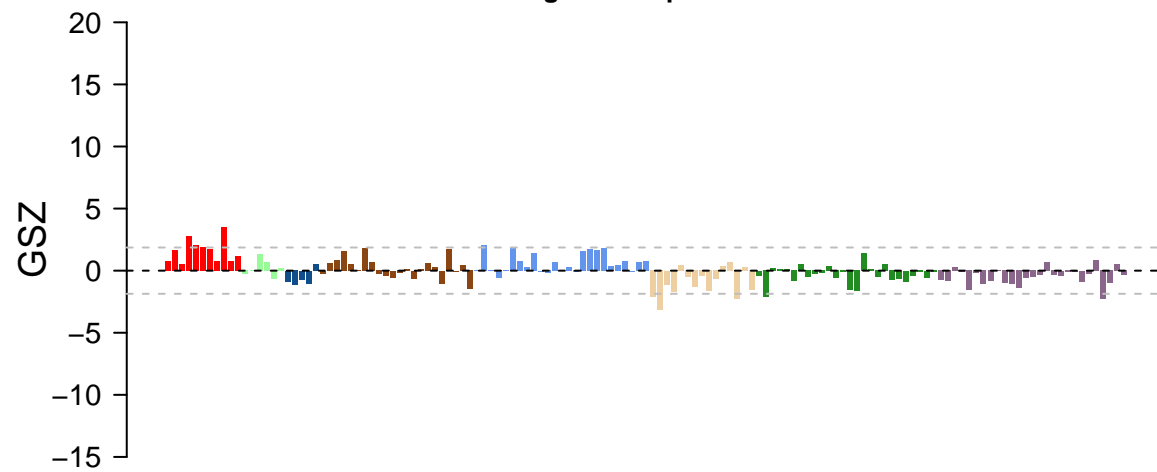
features = 22 , max = 2

endosome to melanosome transport



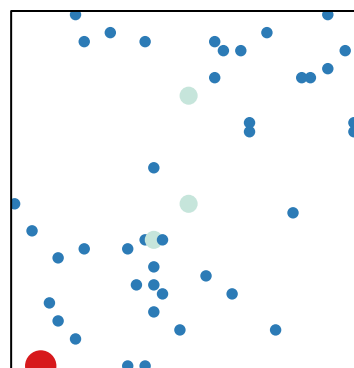
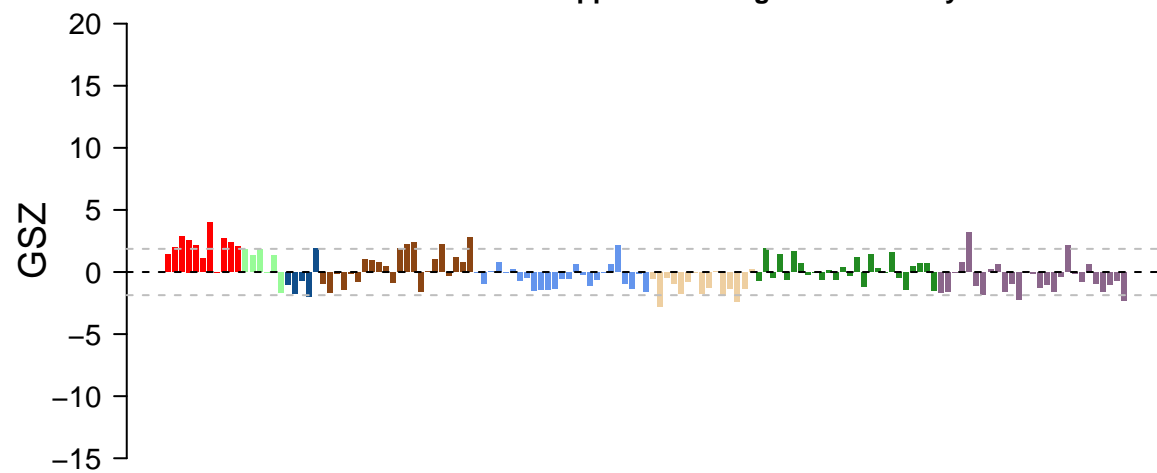
features = 10 , max = 2

left/right axis specification



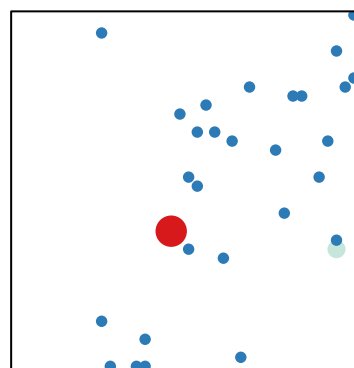
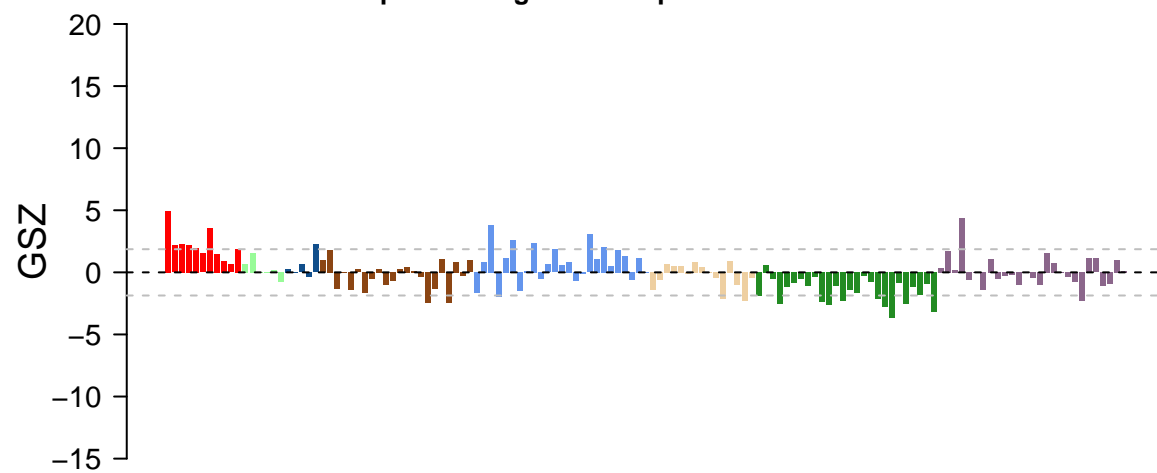
features = 12 , max = 2

activation of NF-kappaB-inducing kinase activity



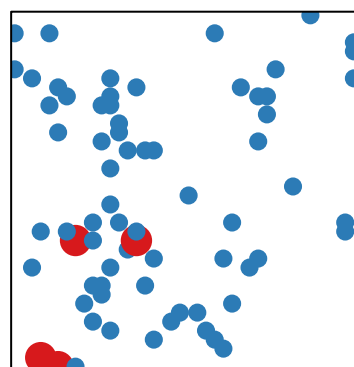
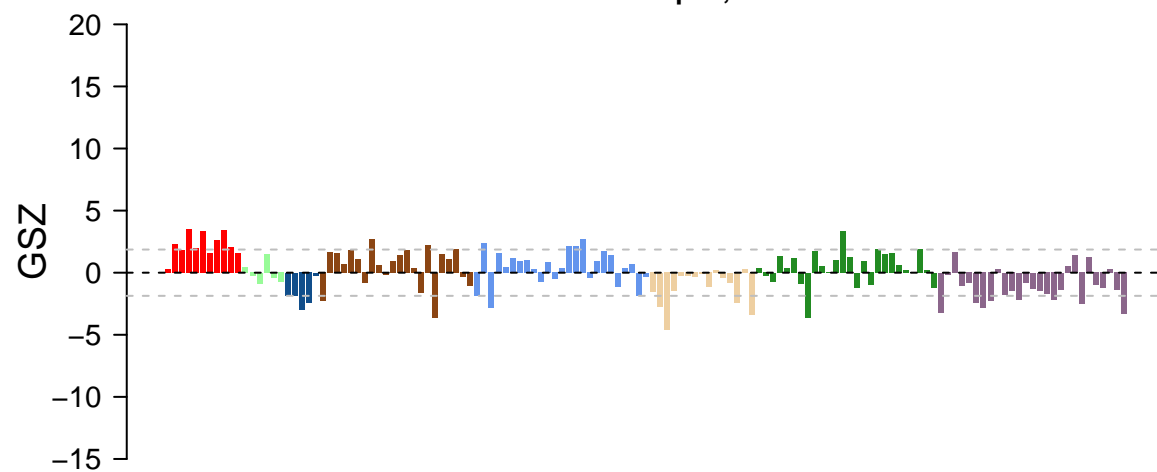
features = 18 , max = 4

positive regulation of positive chemotaxis



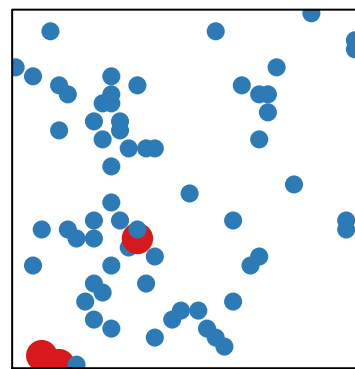
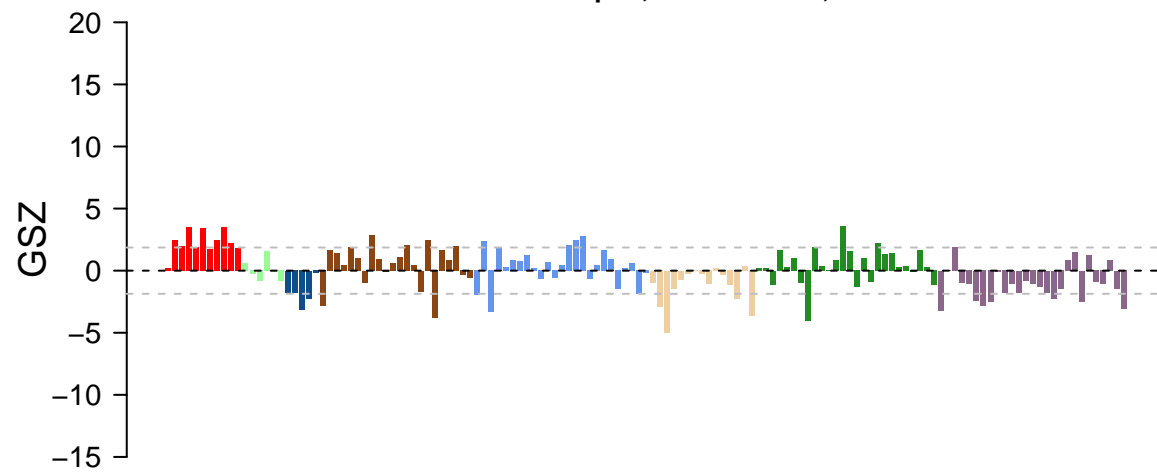
features = 11 , max = 4

nucleotide-excision repair, DNA incision



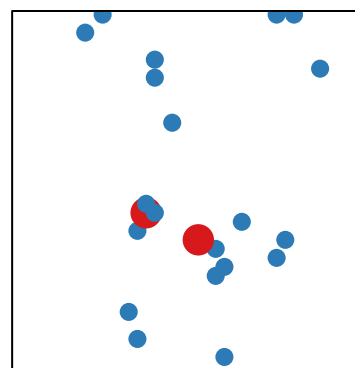
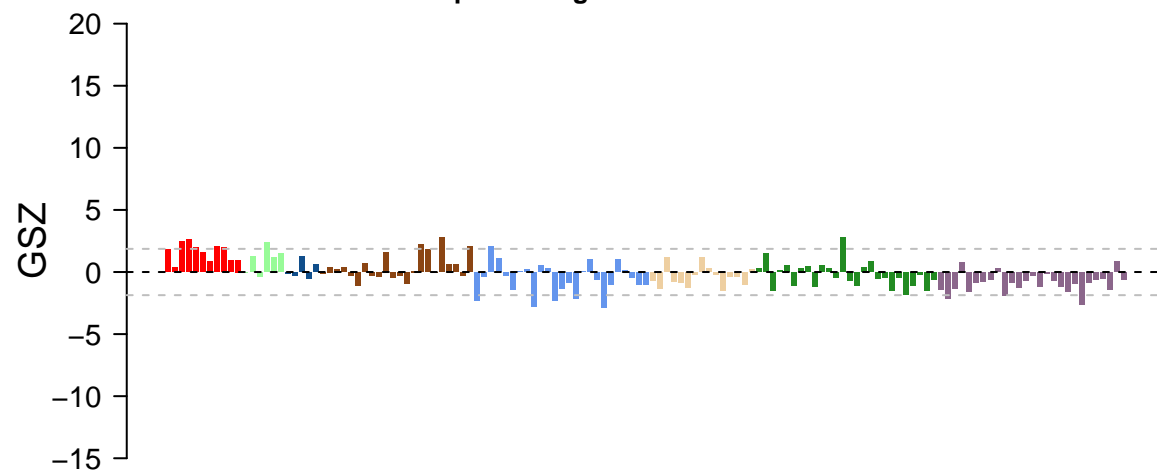
features = 36 , max = 2

nucleotide-excision repair, DNA incision, 5'-to lesion



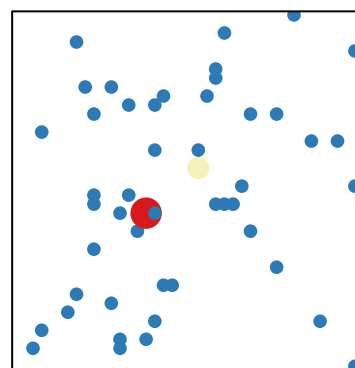
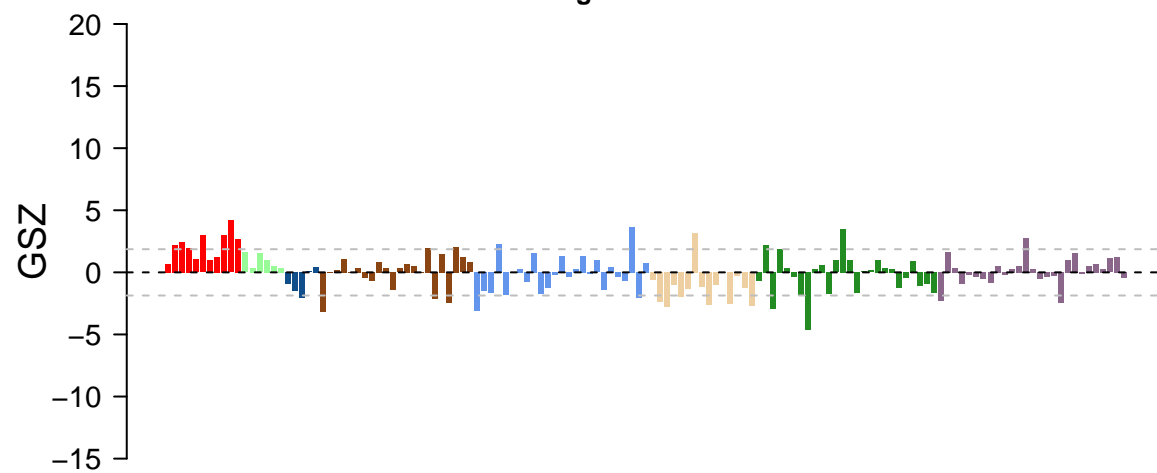
features = 35 , max = 2

trophoblast giant cell differentiation



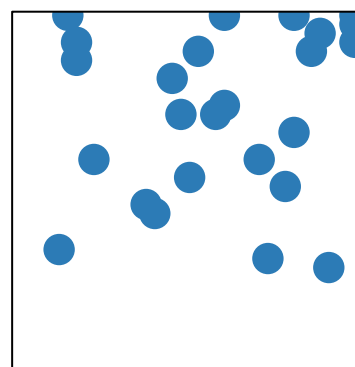
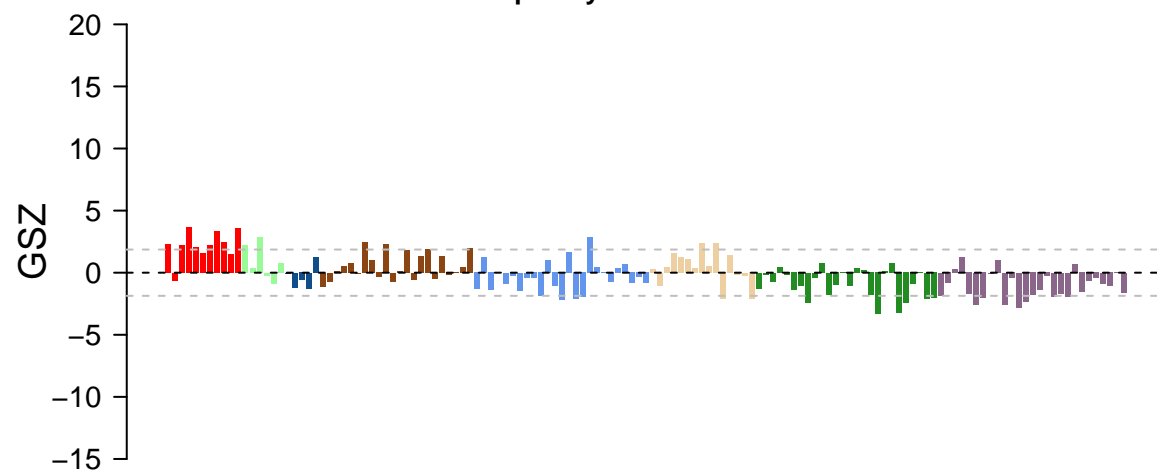
features = 12 , max = 2

nucleobase-containing small molecule interconversion



features = 25 , max = 3

hepatocyte differentiation



features = 13 , max = 1