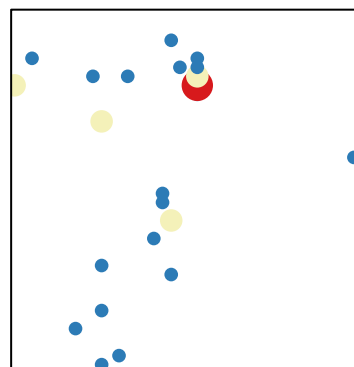
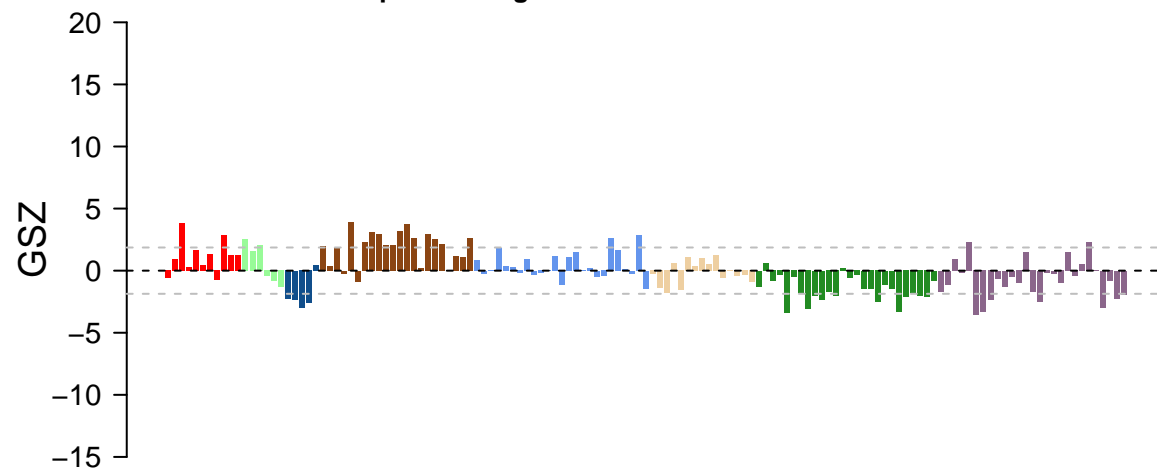
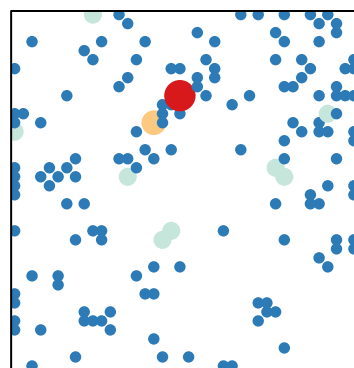
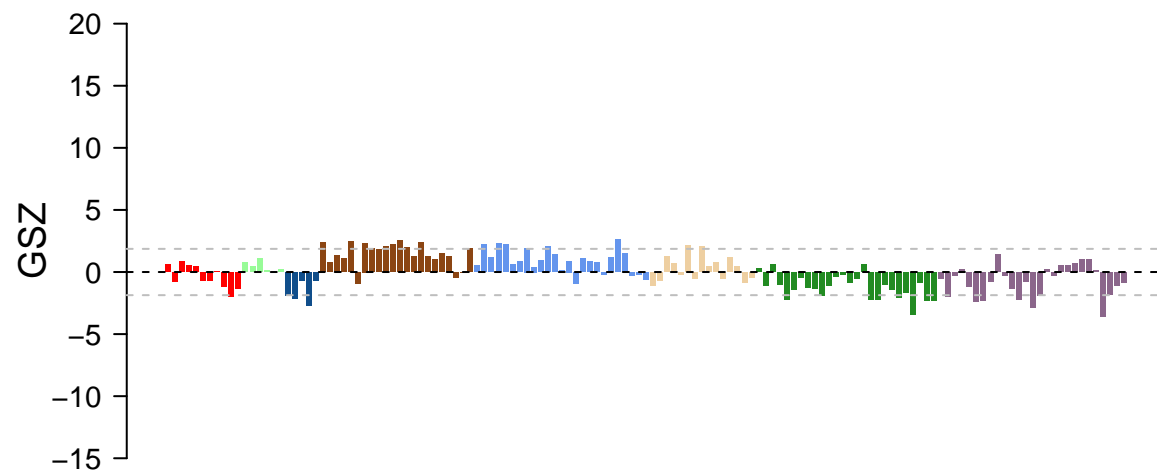


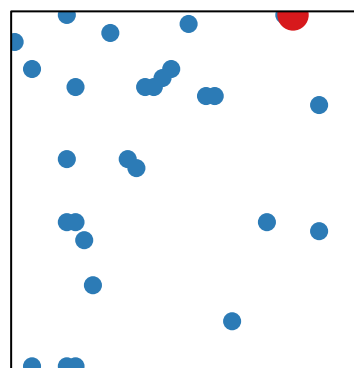
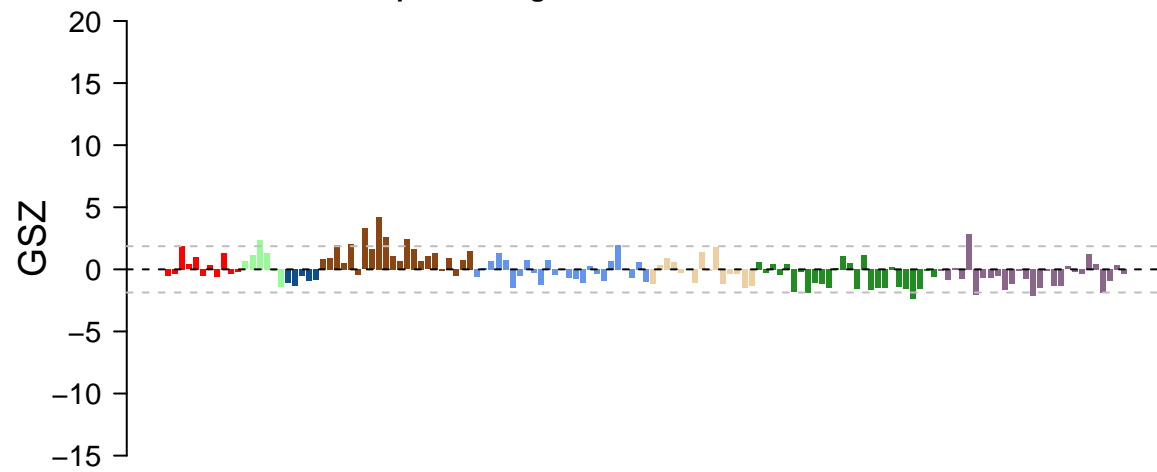
positive regulation of B cell differentiation



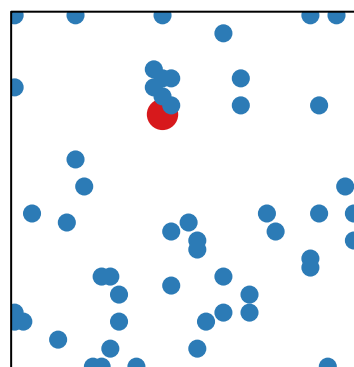
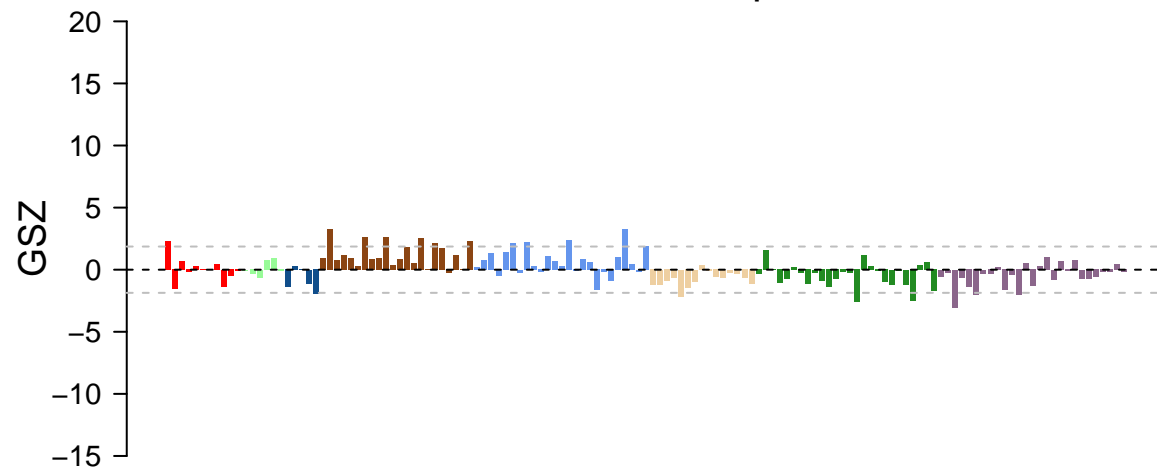
B cell differentiation



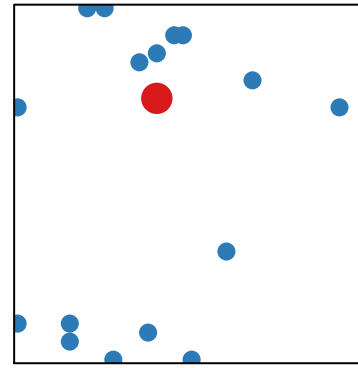
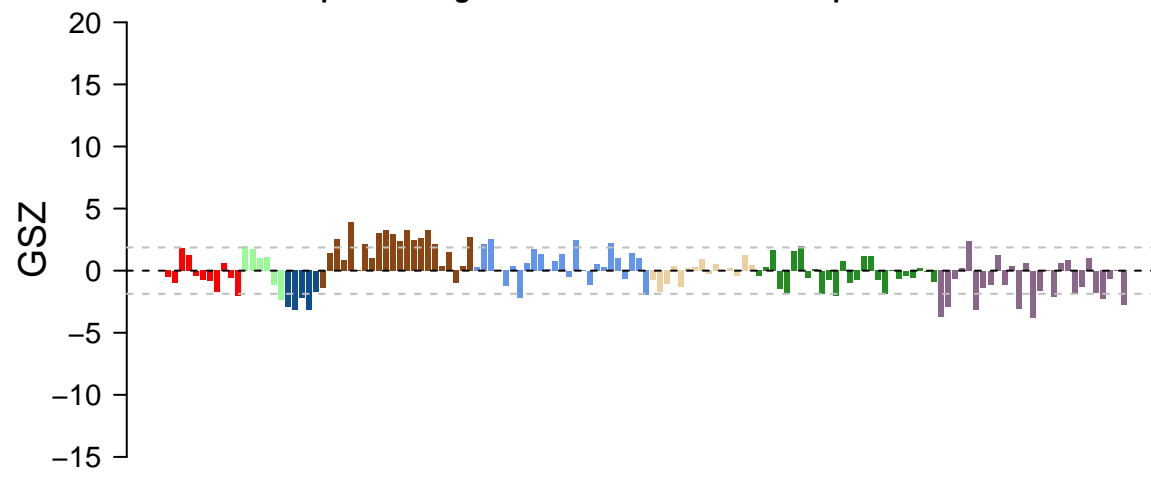
positive regulation of T cell chemotaxis



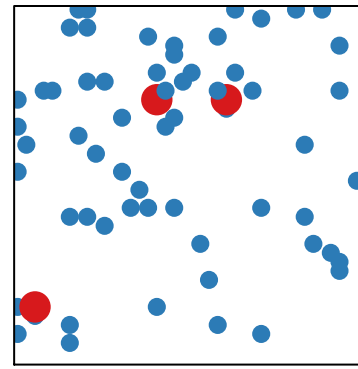
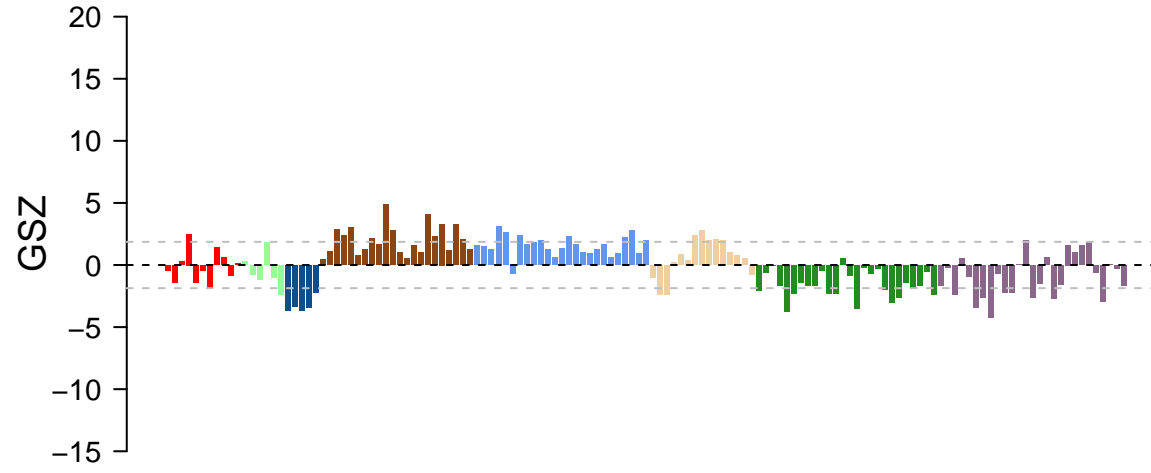
neural crest cell development



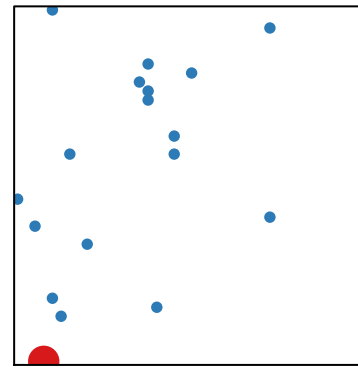
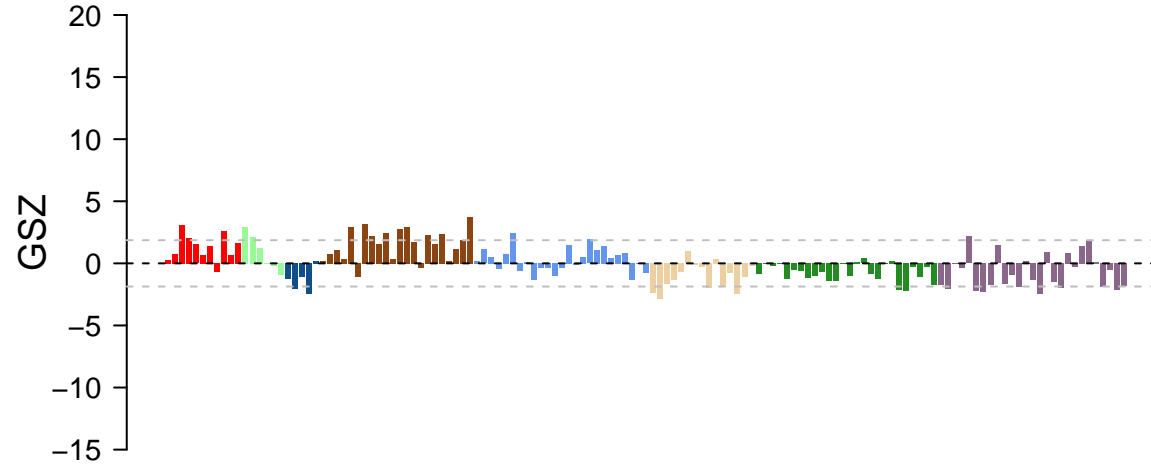
positive regulation of interleukin-1 beta production



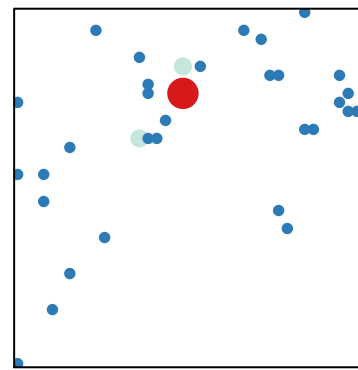
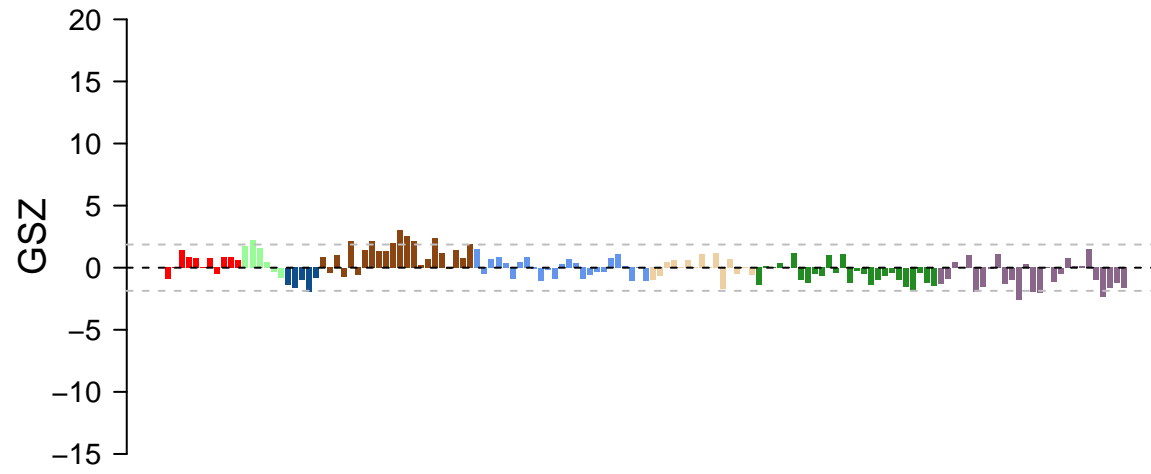
negative regulation of DNA binding



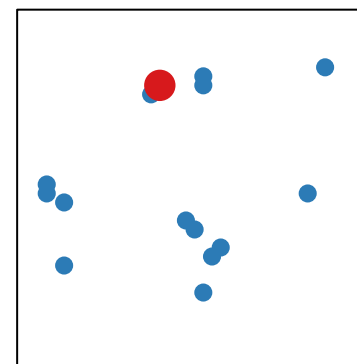
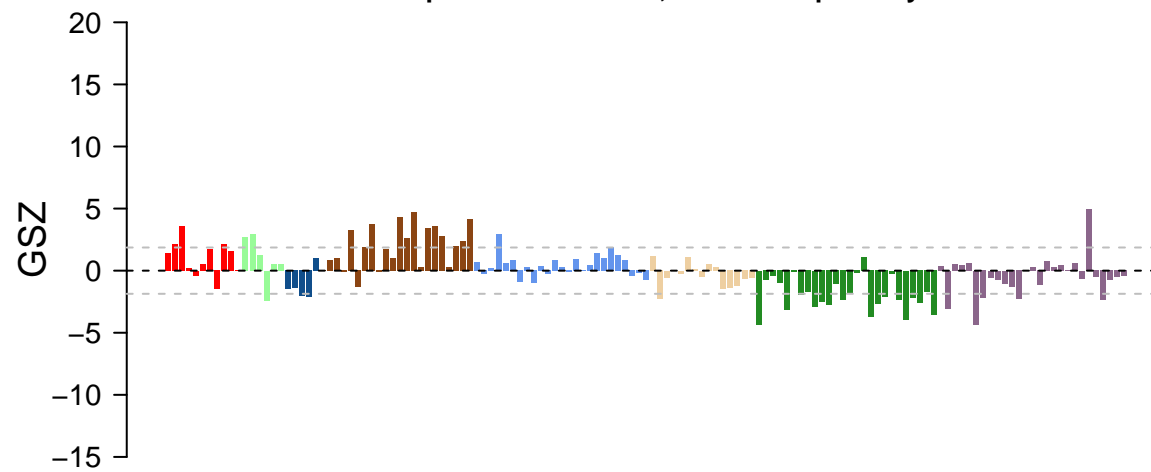
positive regulation of T cell cytokine production



regulation of B cell receptor signaling pathway

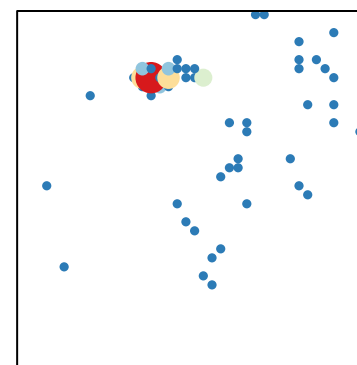
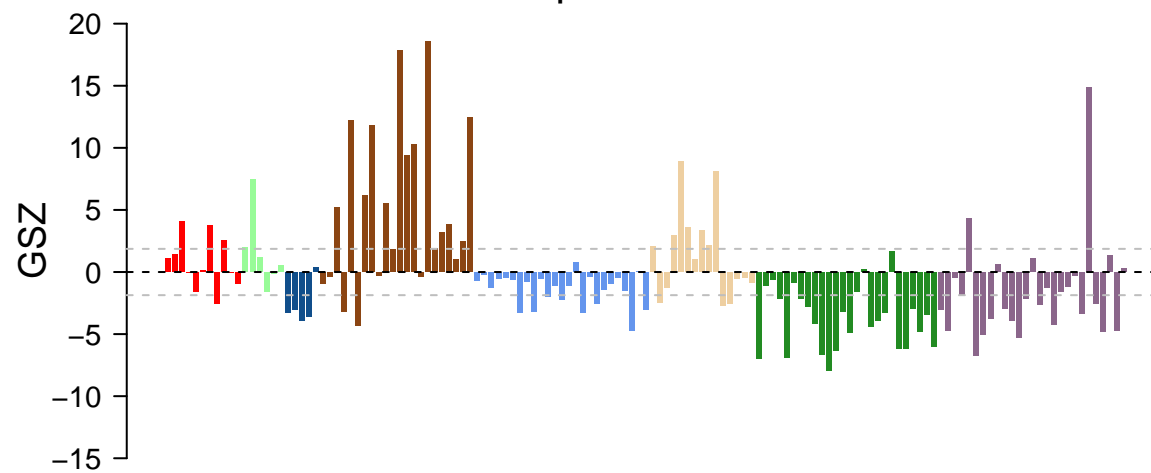


complement activation, alternative pathway



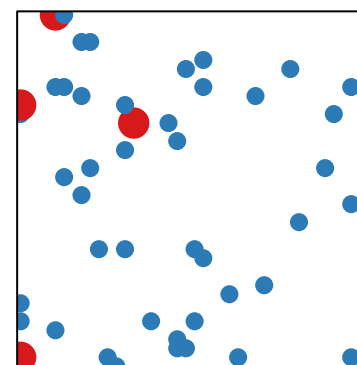
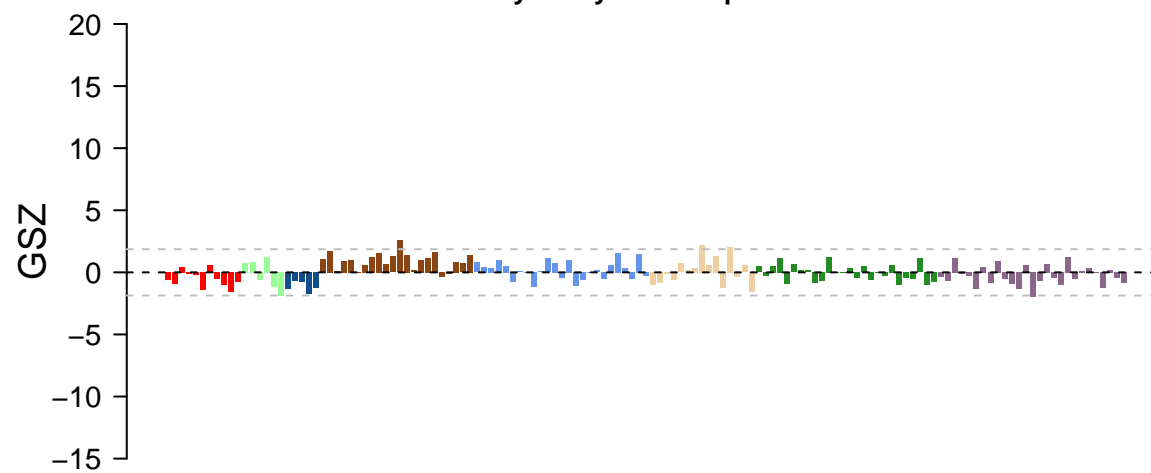
features = 13 , max = 2

complement activation



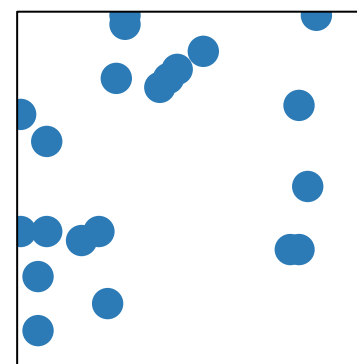
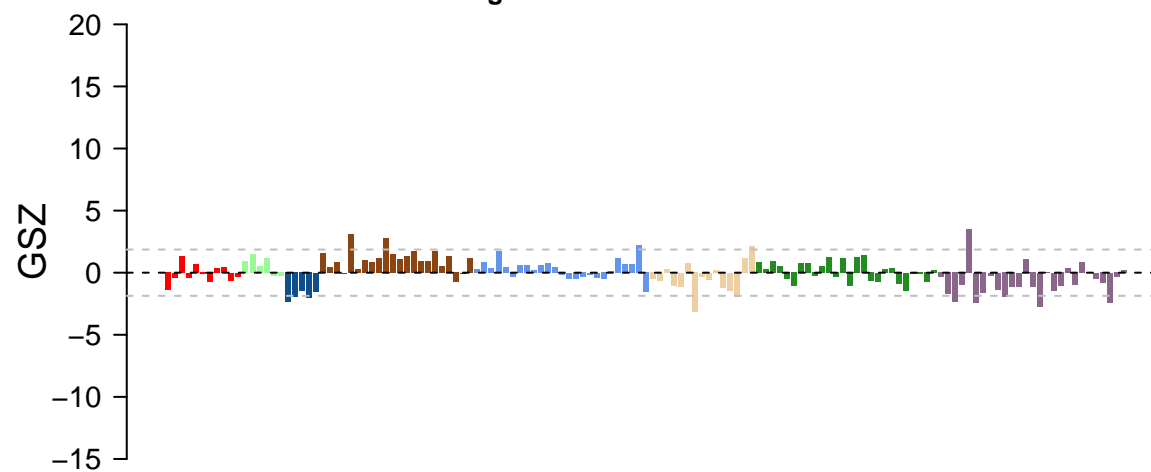
features = 47 , max = 6

erythrocyte development



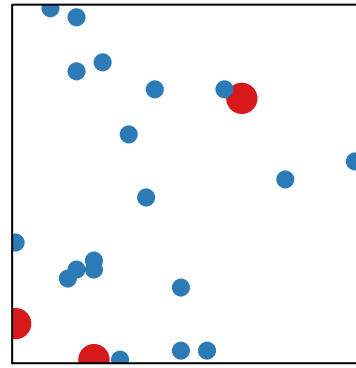
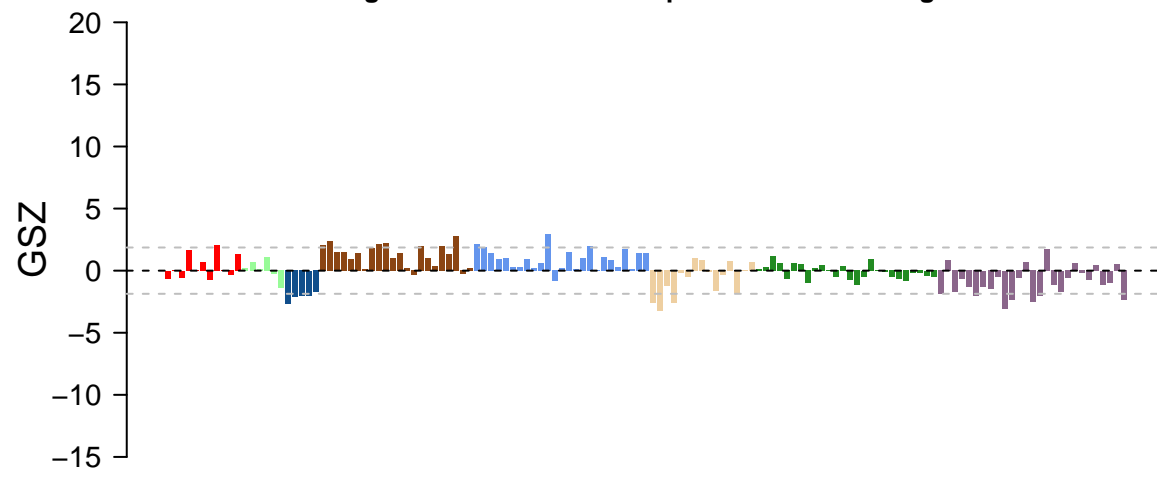
features = 19 , max = 2

regulation of T cell activation

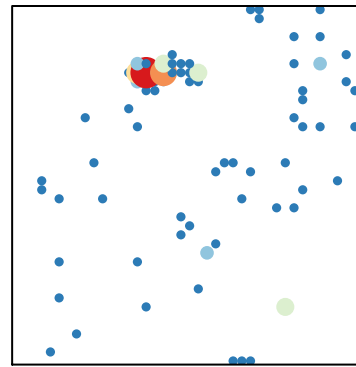
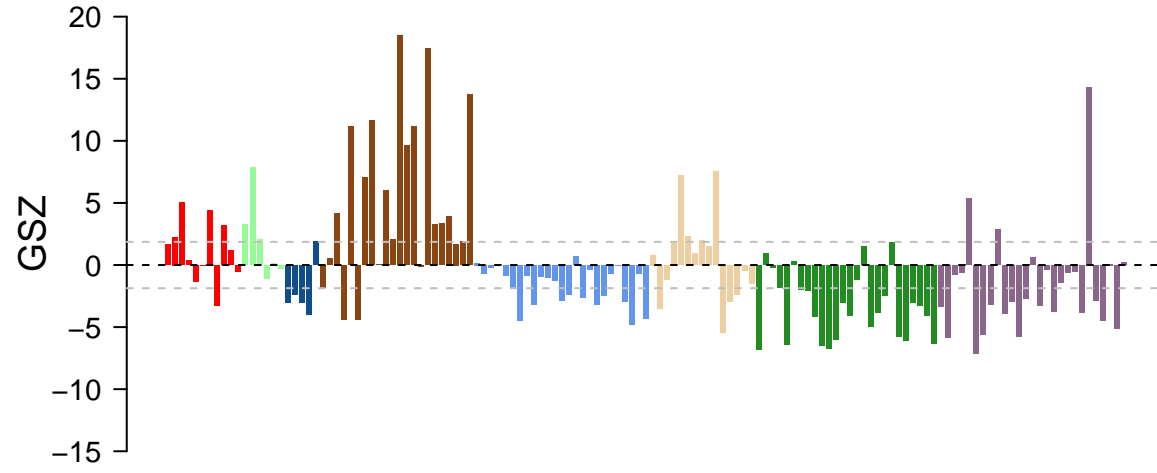


features = 10 , max = 1

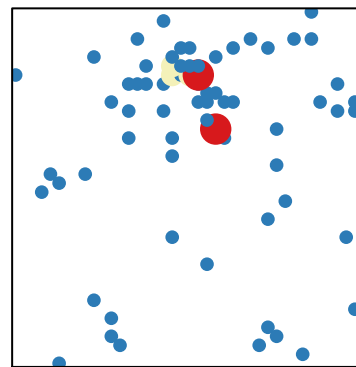
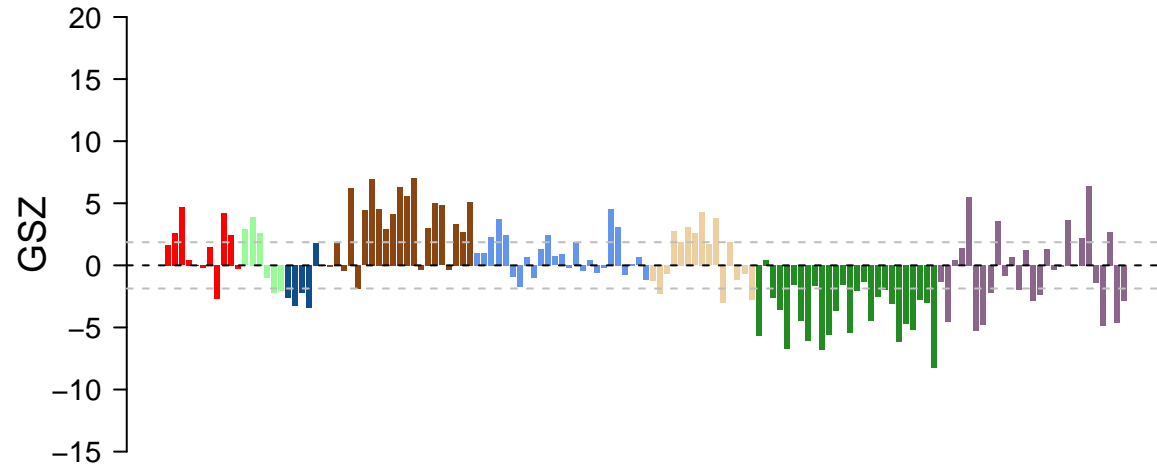
signal transduction in response to DNA damage



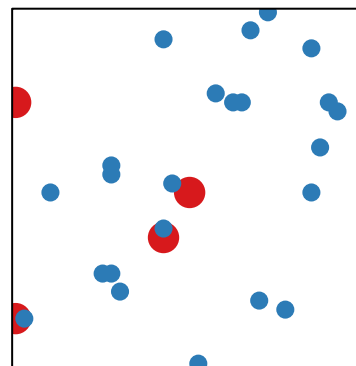
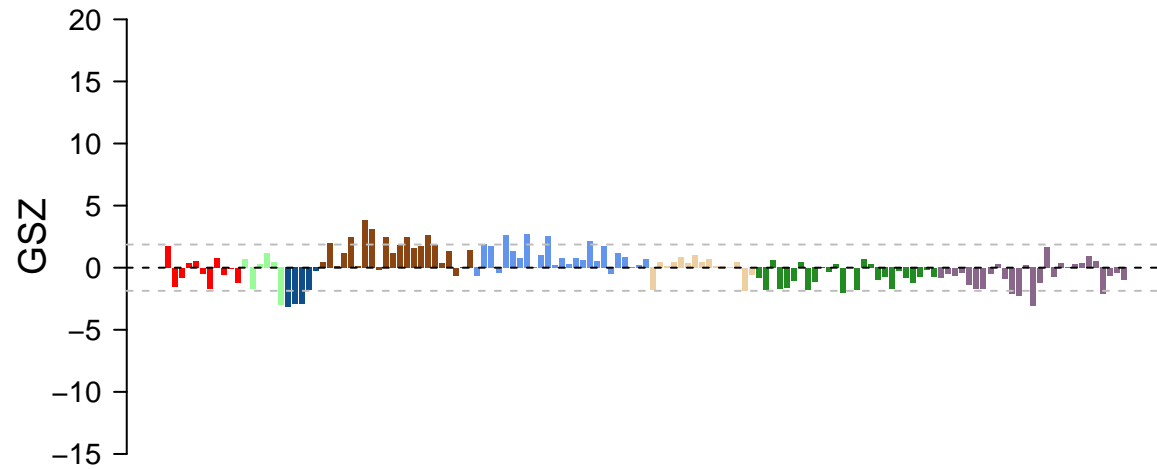
regulation of complement activation



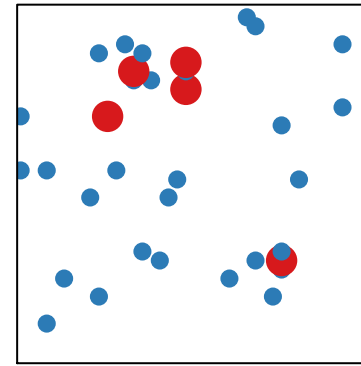
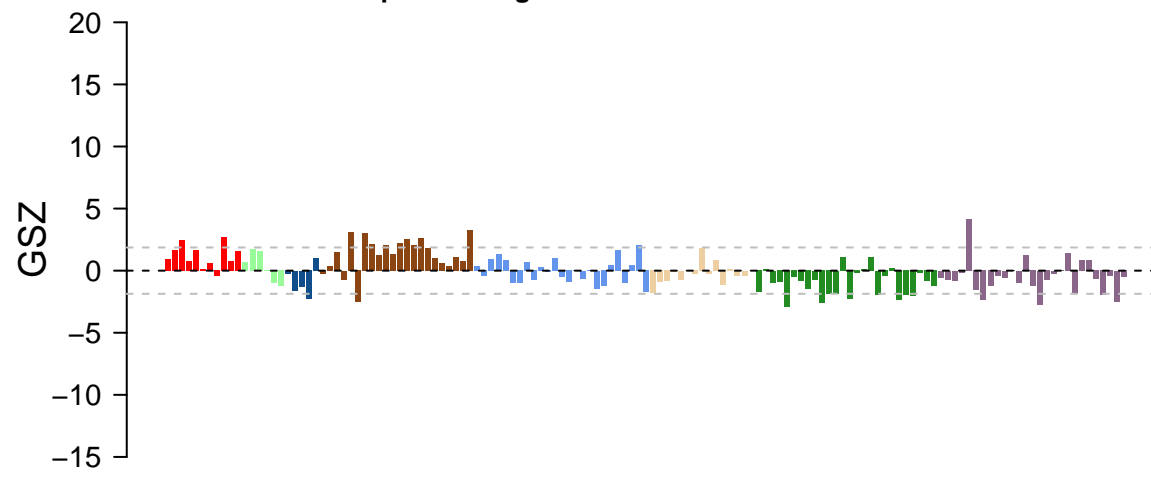
positive regulation of interferon-gamma production



cardiac right ventricle morphogenesis

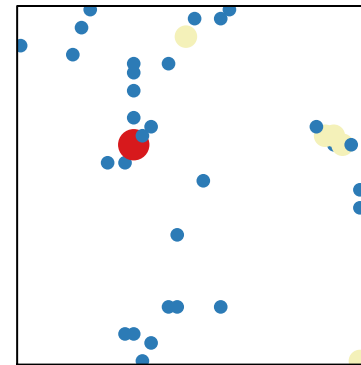
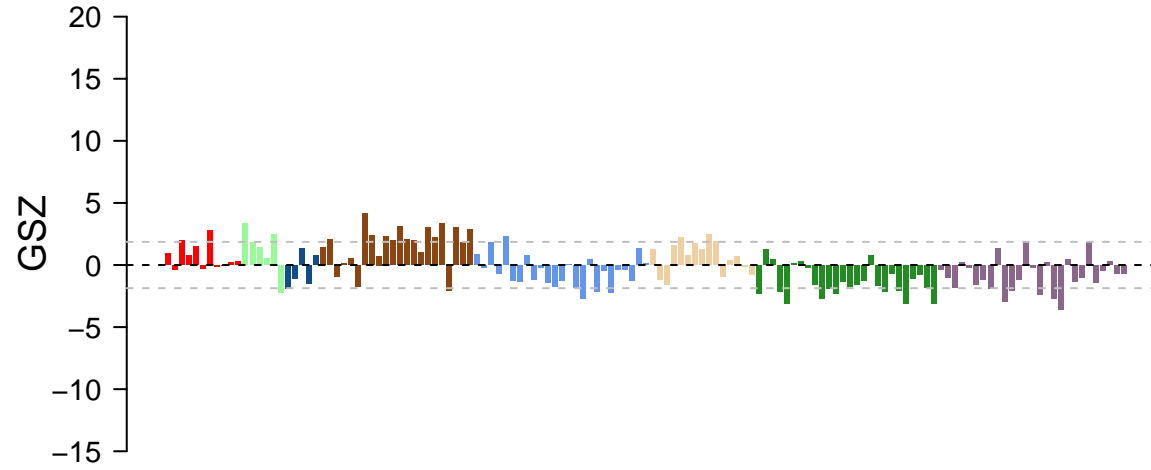


positive regulation of T cell differentiation



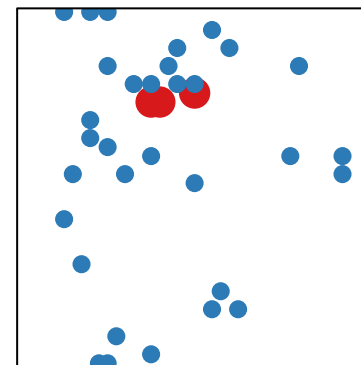
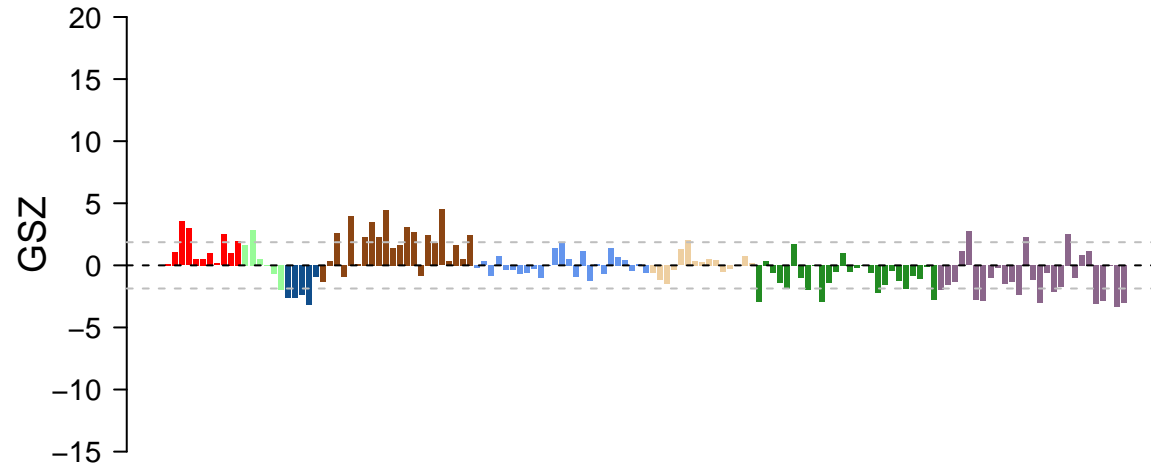
features = 18 , max = 2

cell junction assembly



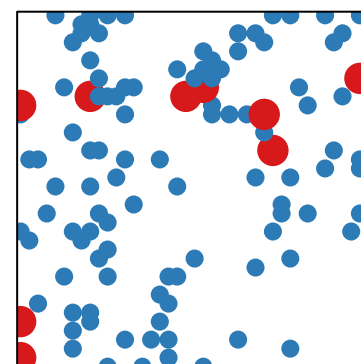
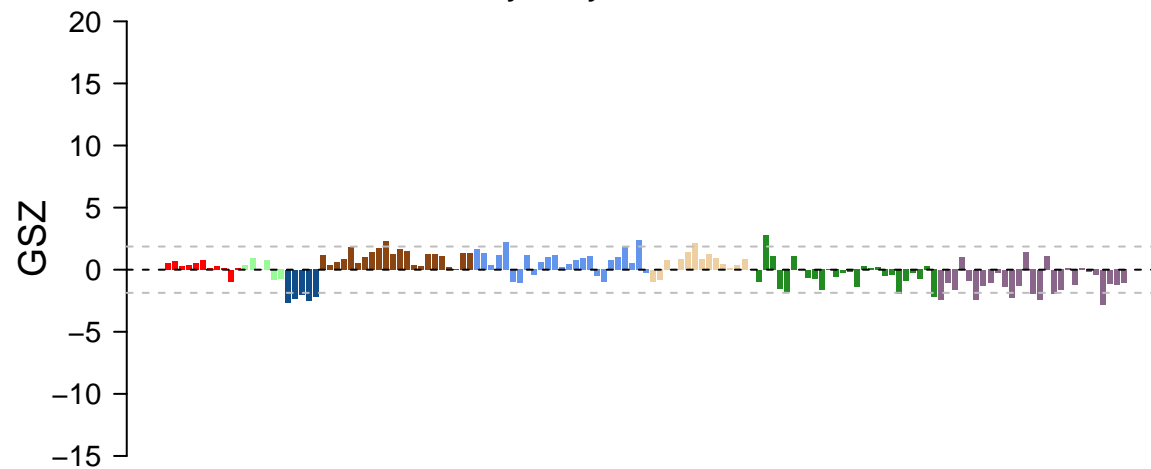
features = 17 , max = 3

cellular response to interferon-beta



features = 19 , max = 2

erythrocyte differentiation



features = 46 , max = 2