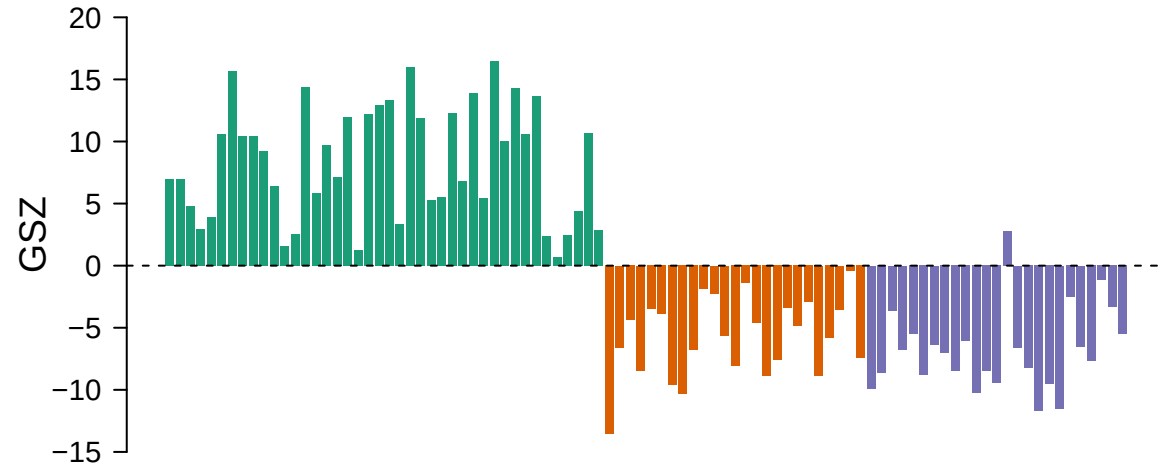
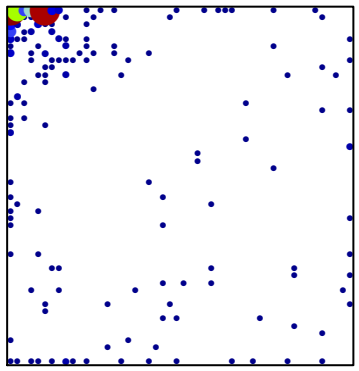
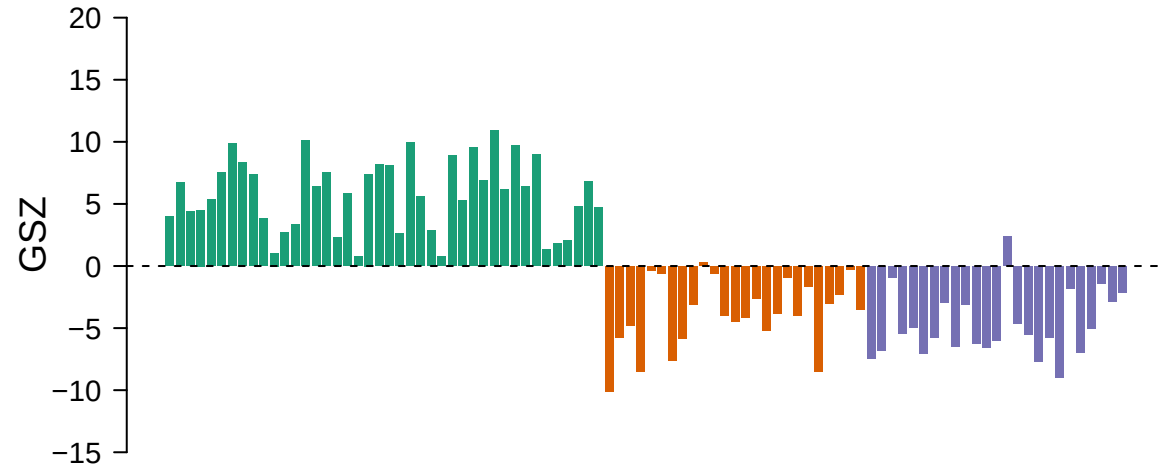
ZHANG_TLX_TARGETS_60HR_DN

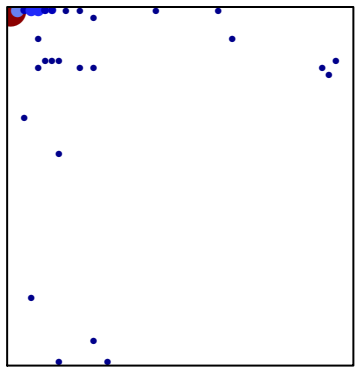




features = 267 , max = 24

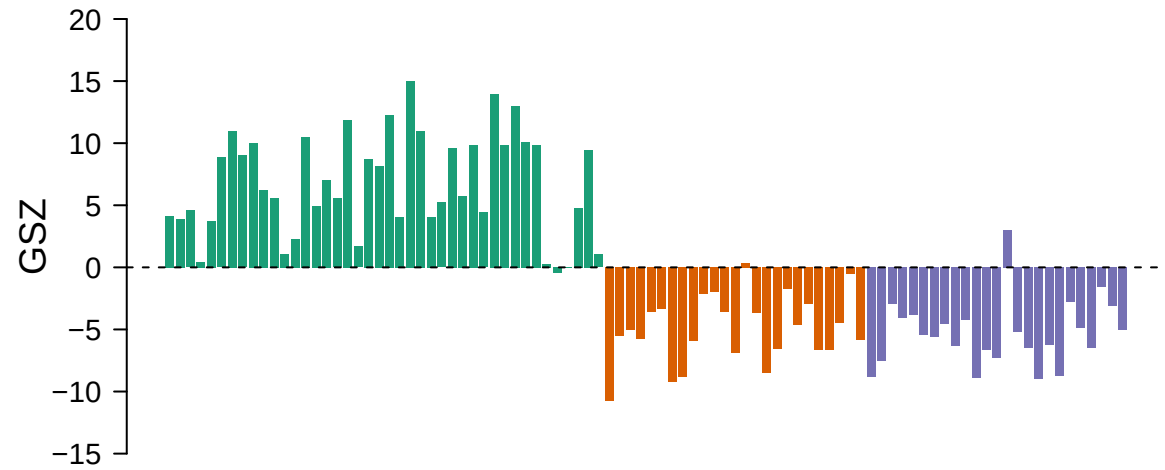
KAUFFMANN_MELANOMA_RELAPSE_UP

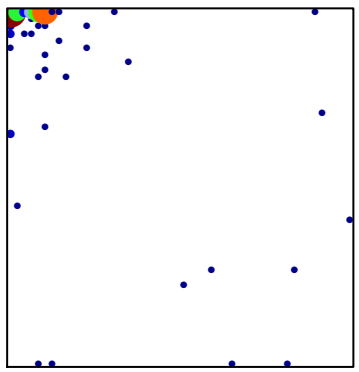




features = 59 , max = 16

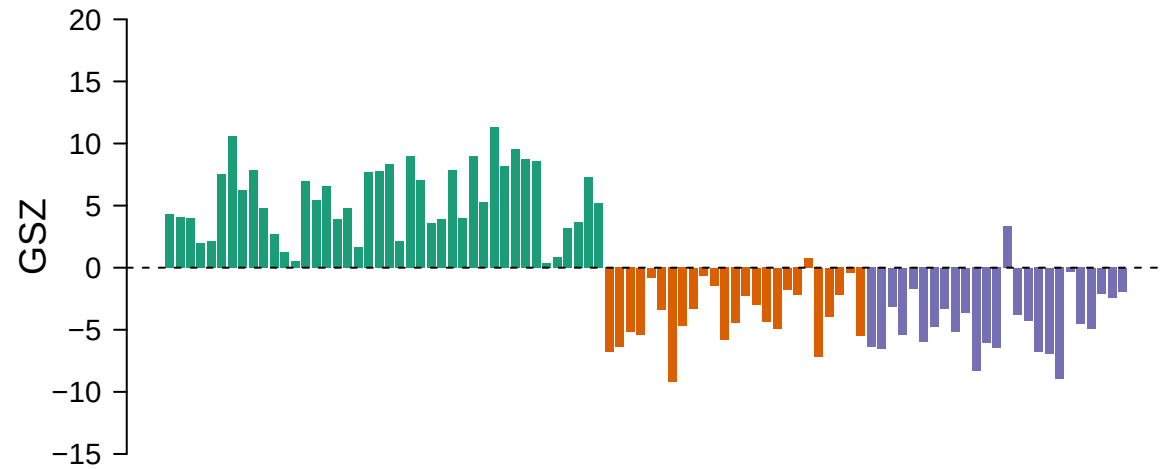
ZHANG_TLX_TARGETS_UP

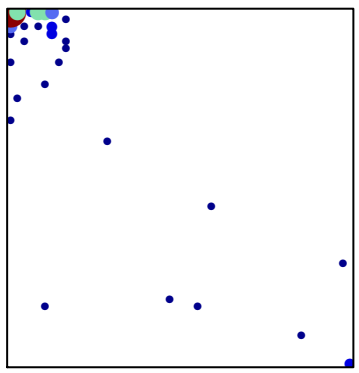




features = 87 , max = 14

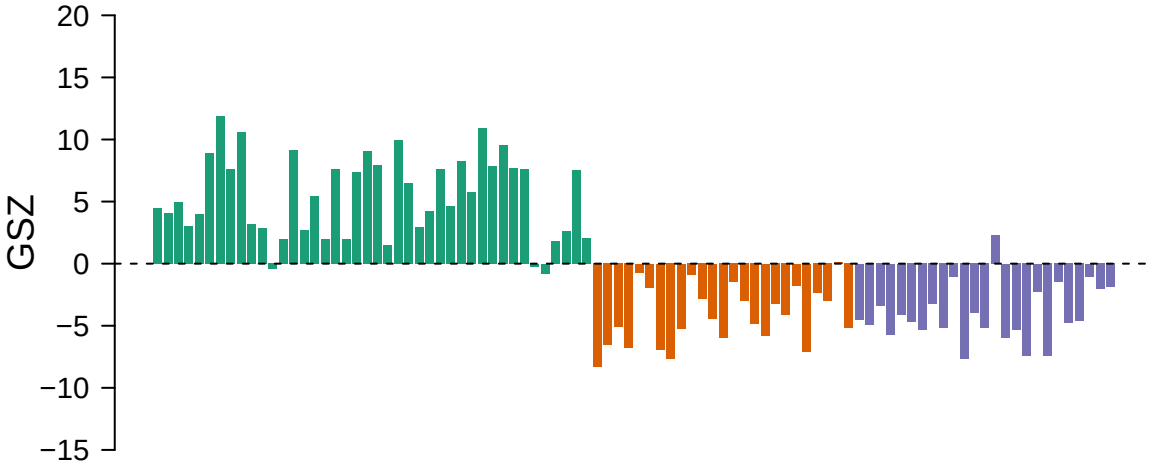
REN_BOUND_BY_E2F

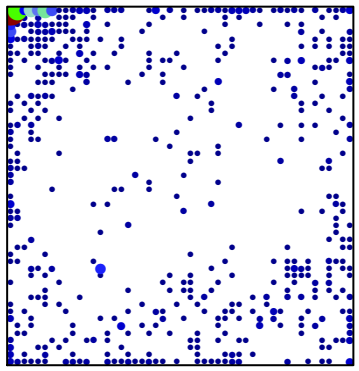




features = 57 , max = 9

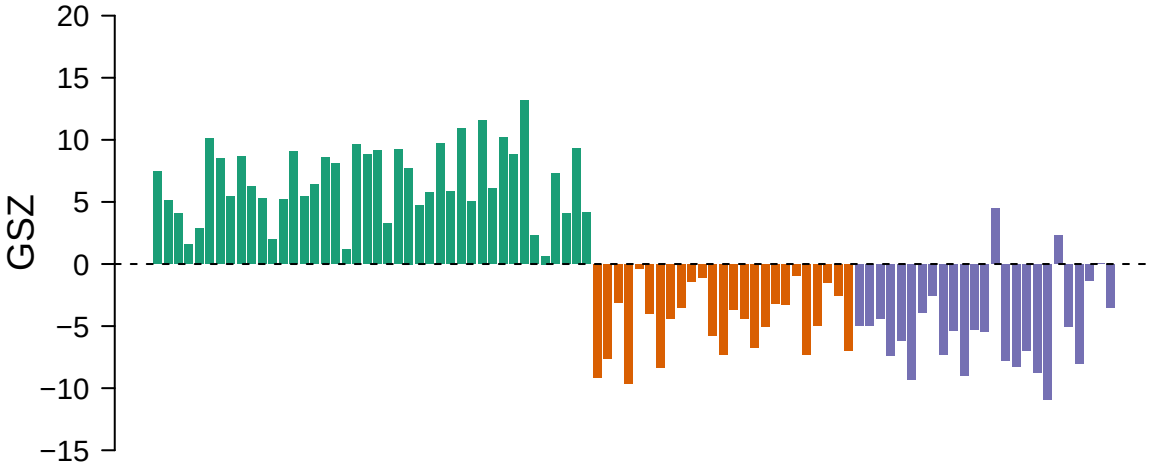
DNA metabolic process

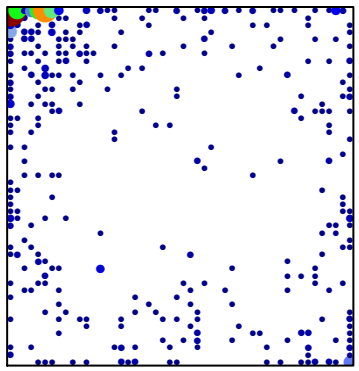




features = 768 , max = 32

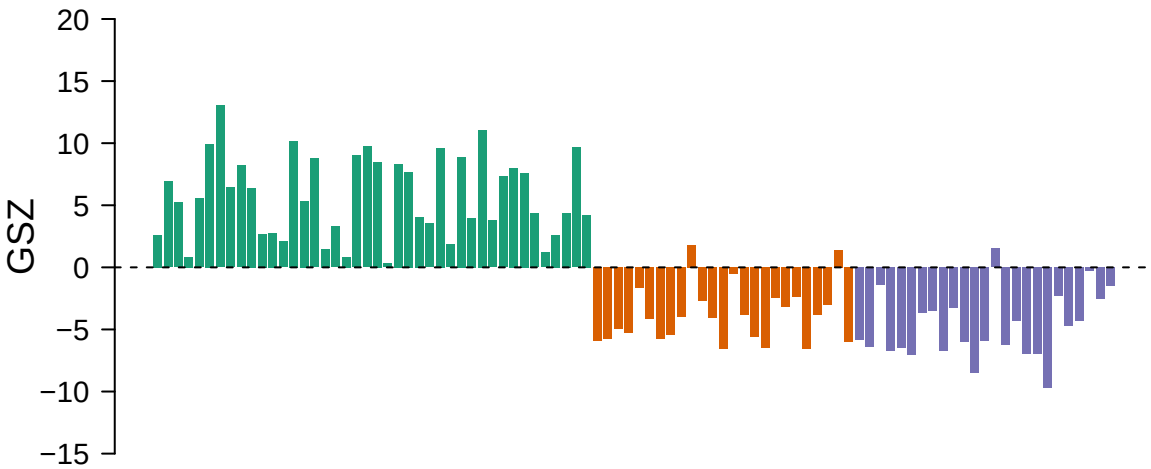
BERENJENO_TRANSFORMED_BY_RHOA_UP

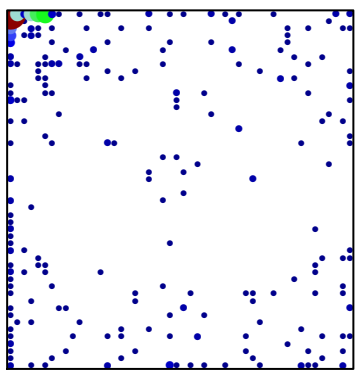




features = 505 , max = 28

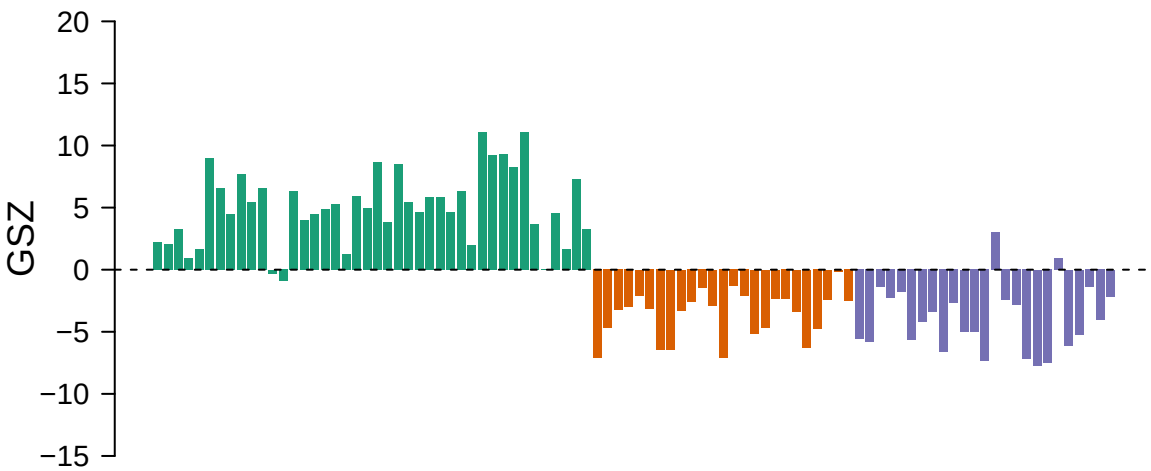
WANG_RESPONSE_TO_GSK3_INHIBITOR_SB216763_DN

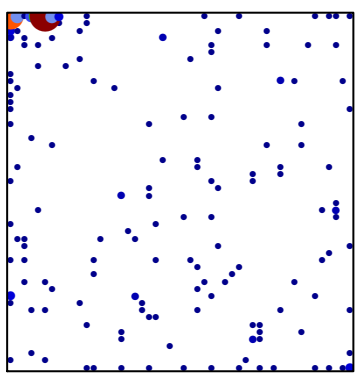




features = 335 , max = 24

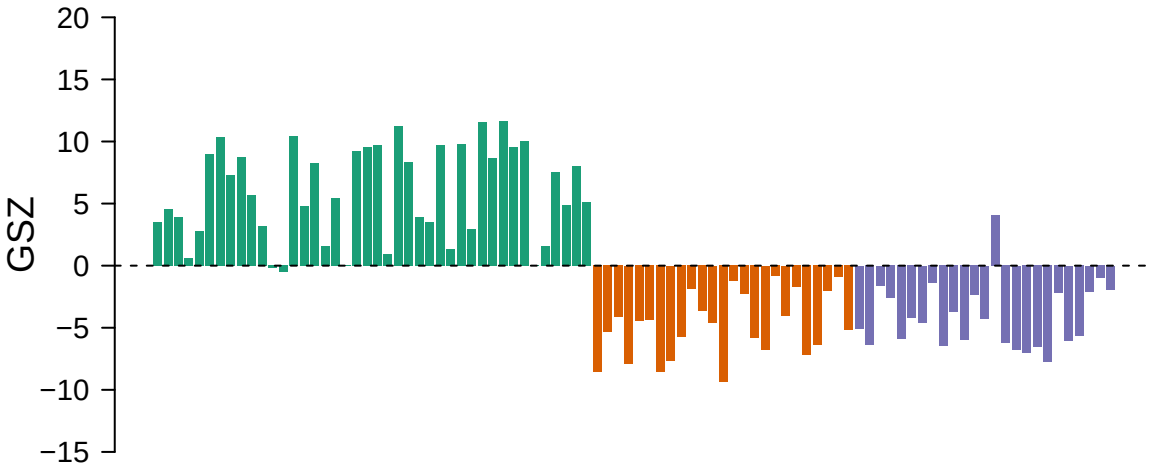
ZHENG_GLIOBLASTOMA_PLASTICITY_UP

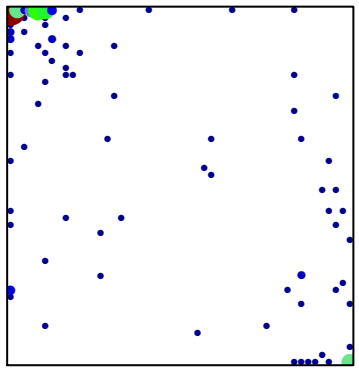




features = 224 , max = 19

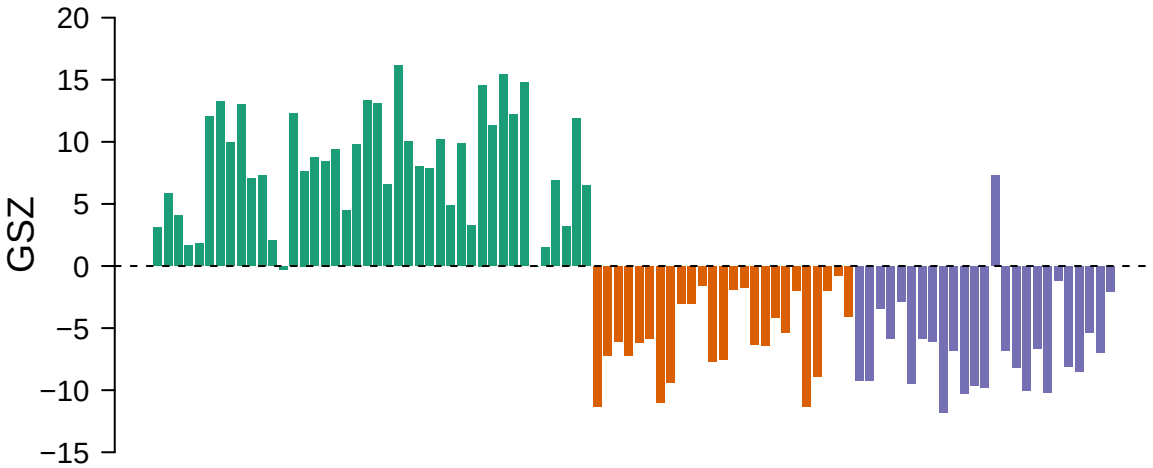
RUIZ_TNC_TARGETS_DN

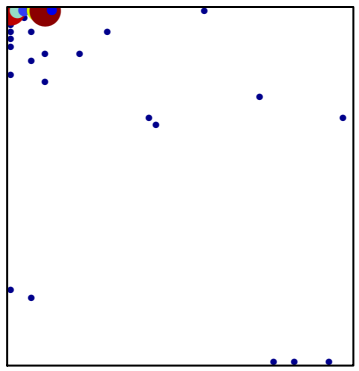




features = 136 , max = 16

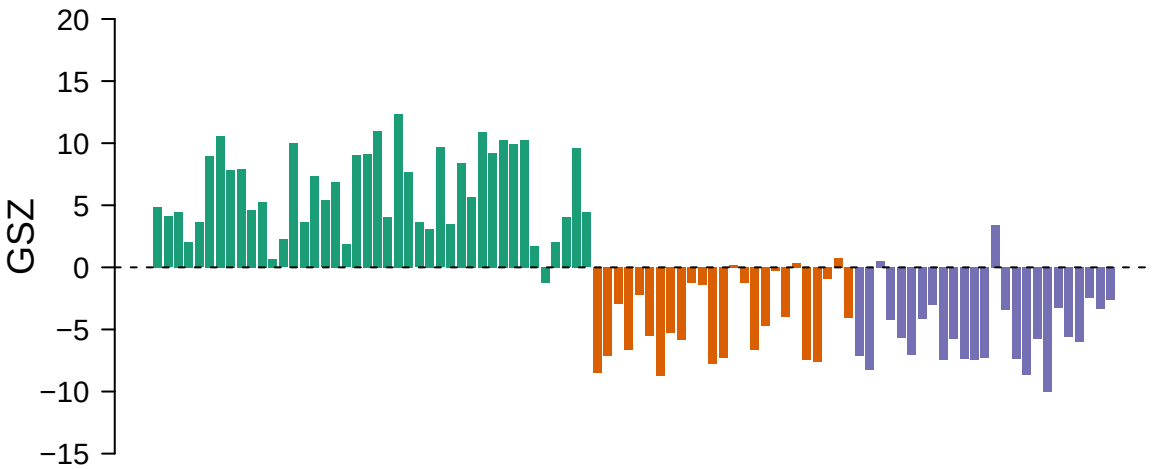
GRAHAM_NORMAL_QUIESCENT_VS_NORMAL_DIVIDING_DN

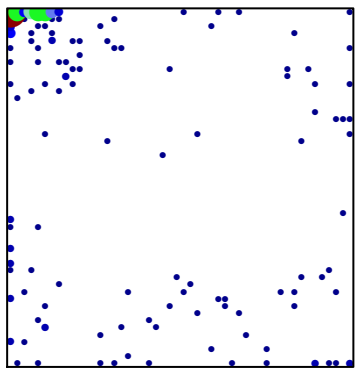




features = 81 , max = 15

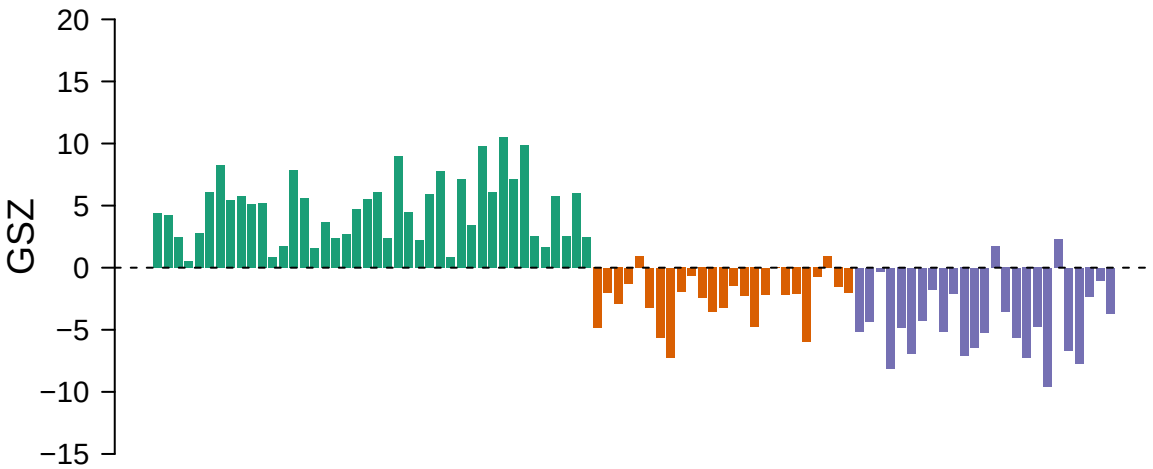
FUJII_YBX1_TARGETS_DN

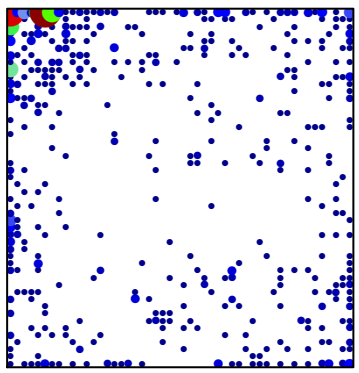




features = 198 , max = 20

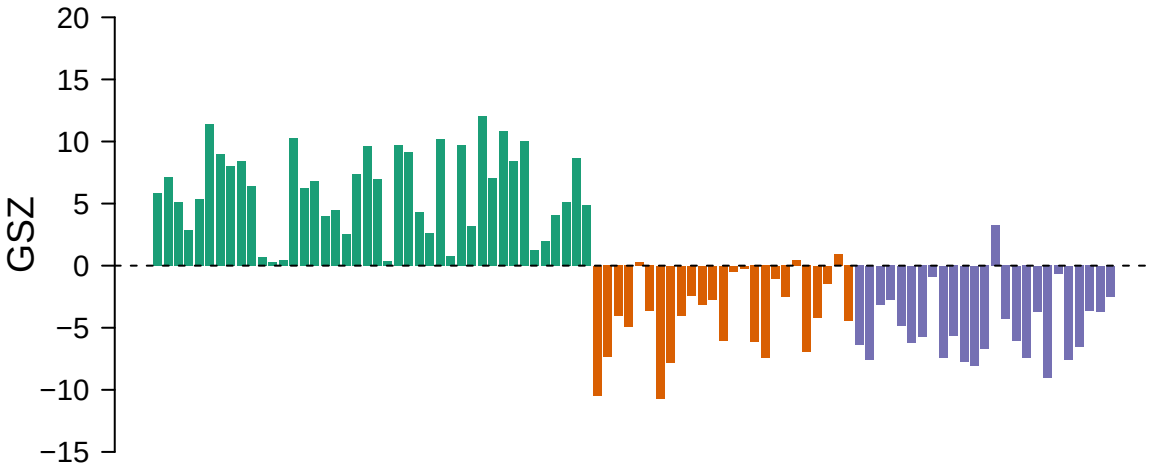
CASORELLI_ACUTE_PROMYELOCYTIC_LEUKEMIA_DN

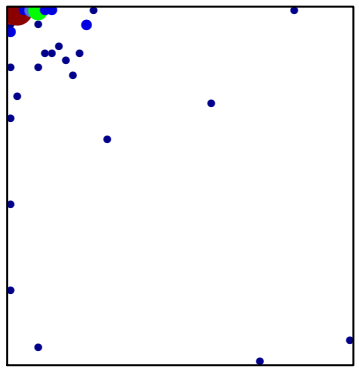




features = 615 , max = 19

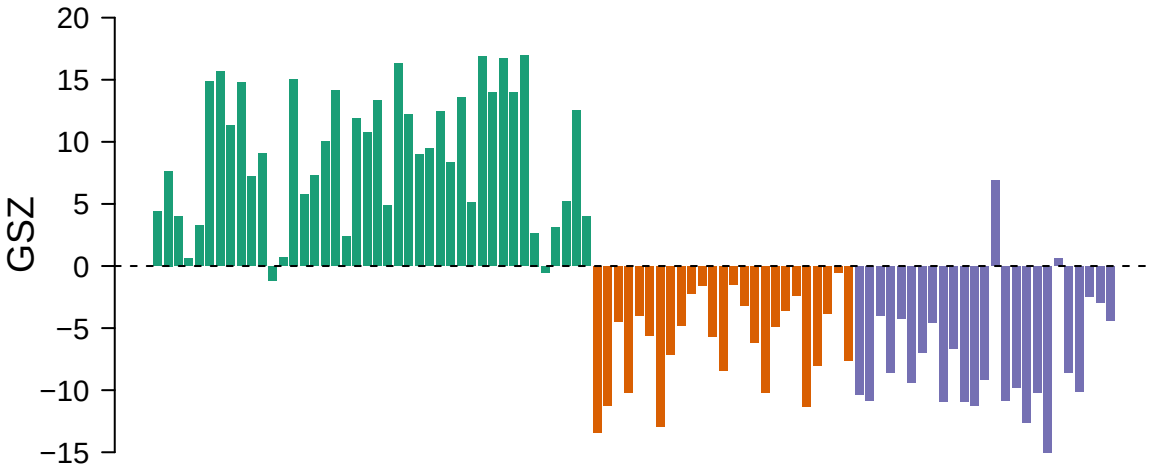
SONG_TARGETS_OF_IE86_CMV_PROTEIN

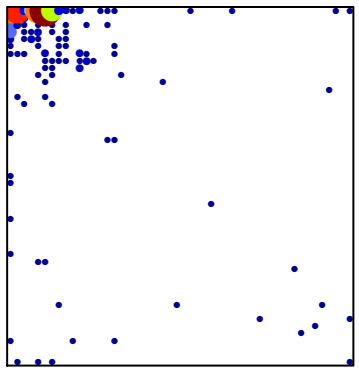




features = 58 , max = 9

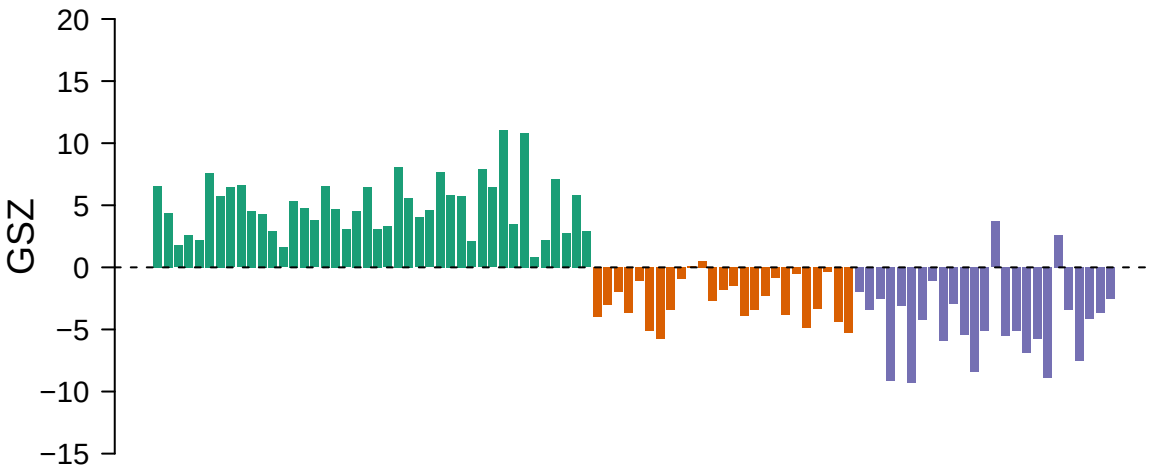
HALLMARK_E2F_TARGETS

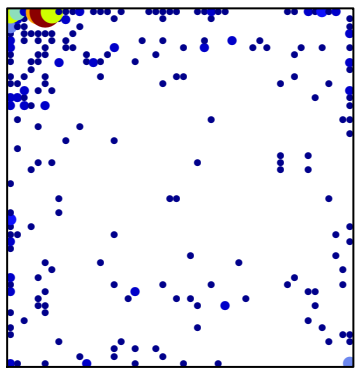




features = 197 , max = 17

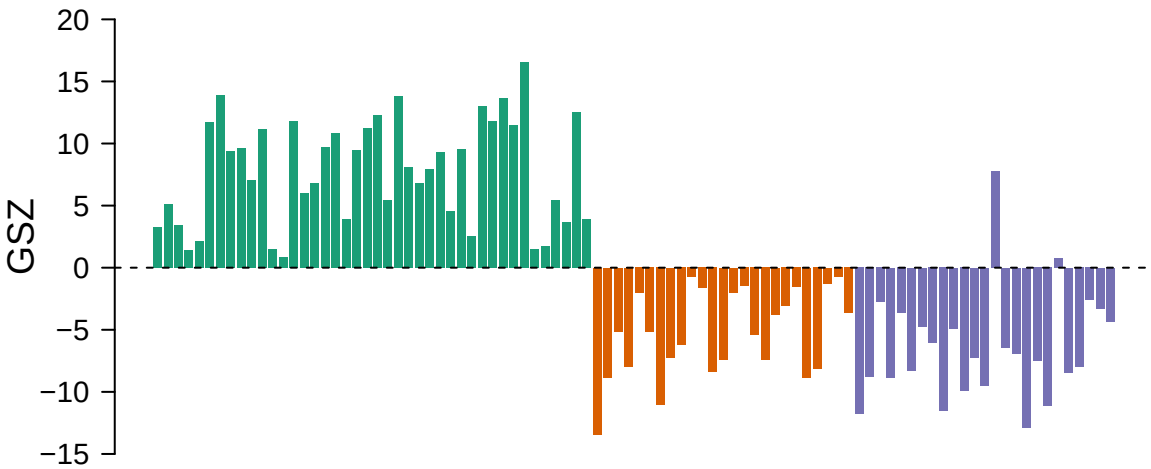
TARTE_Plasmablast signature

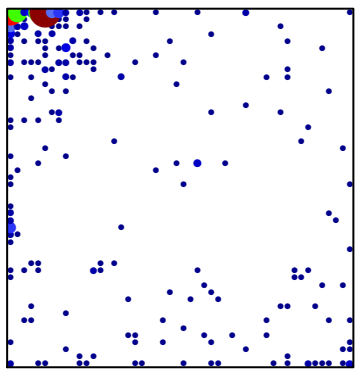




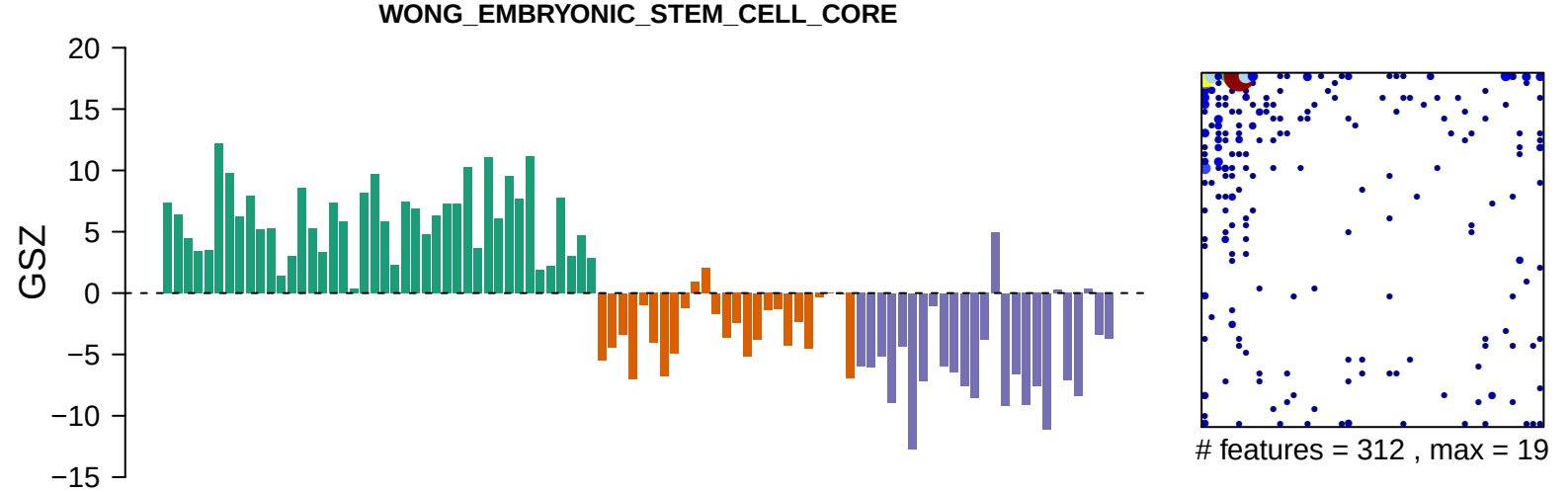
features = 294 , max = 12

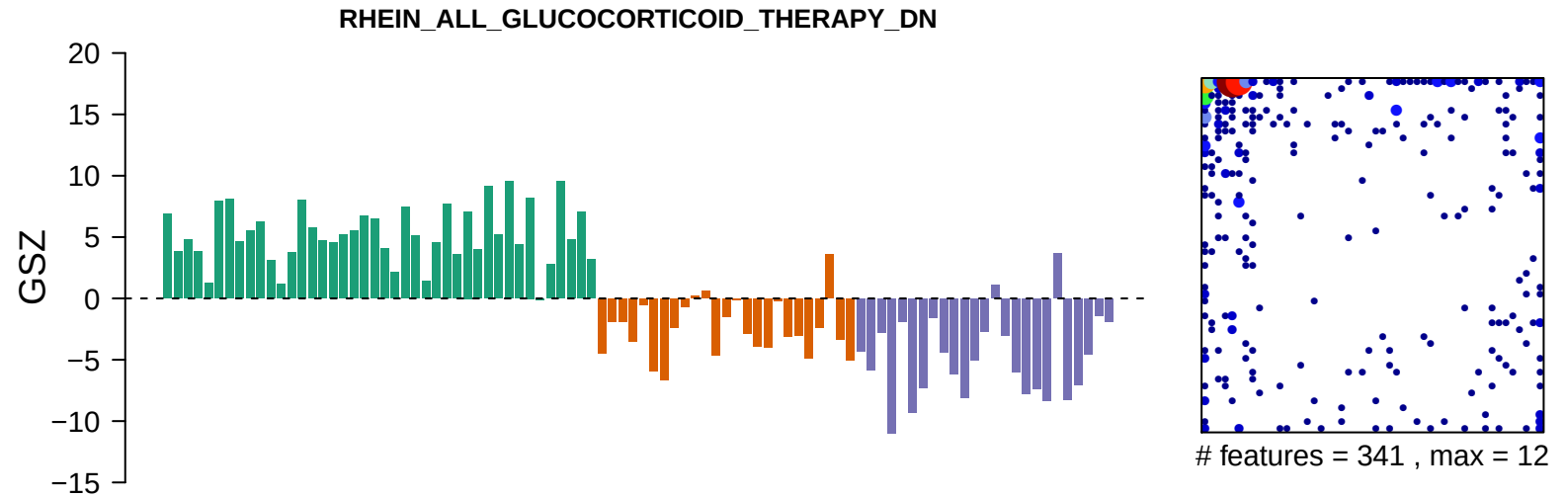
BLUM_RESPONSE_TO_SALIRASIB_DN

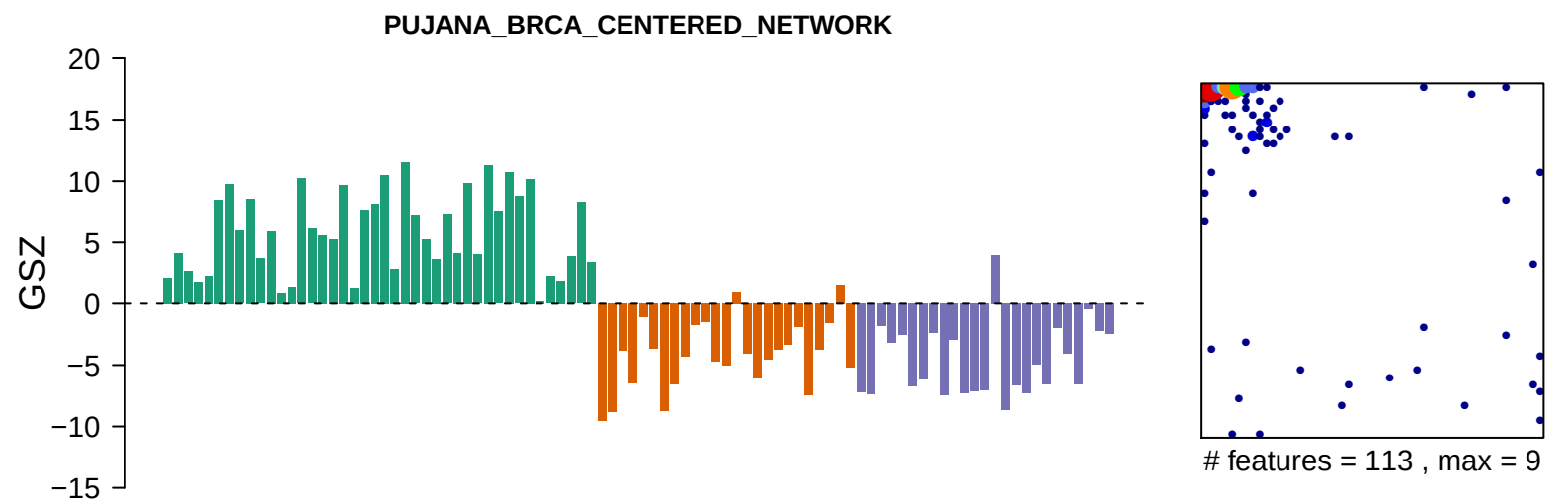


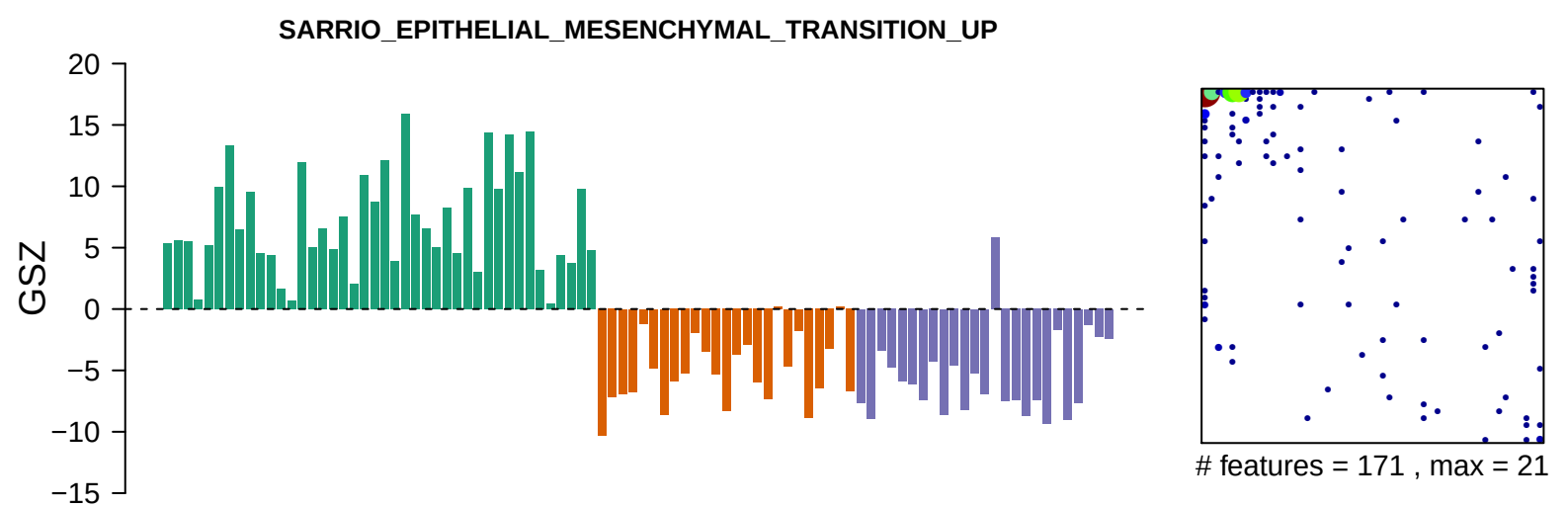


features = 327 , max = 25


WONG_EMBRYONIC_STEM_CELL_CORE
GSZ
20
15
10
5
0
−5
−10
−15
features = 312 , max = 19


RHEIN_ALL_GLUCOCORTICOID_THERAPY_DN
GSZ
20
15
10
5
0
−5
−10
−15
features = 341 , max = 12


PUJANA_BRCA_CENTERED_NETWORK
GSZ
20
15
10
5
0
−5
−10
−15
features = 113 , max = 9


SARRIO_EPITHELIAL_MESENCHYMAL_TRANSITION_UP
GSZ
20
15
10
5
0
−5
−10
−15
features = 171 , max = 21