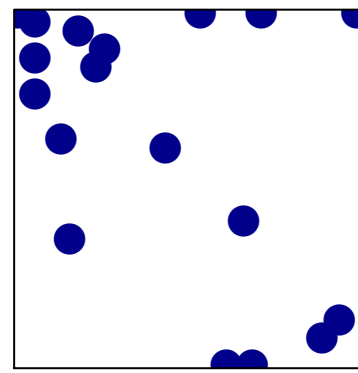
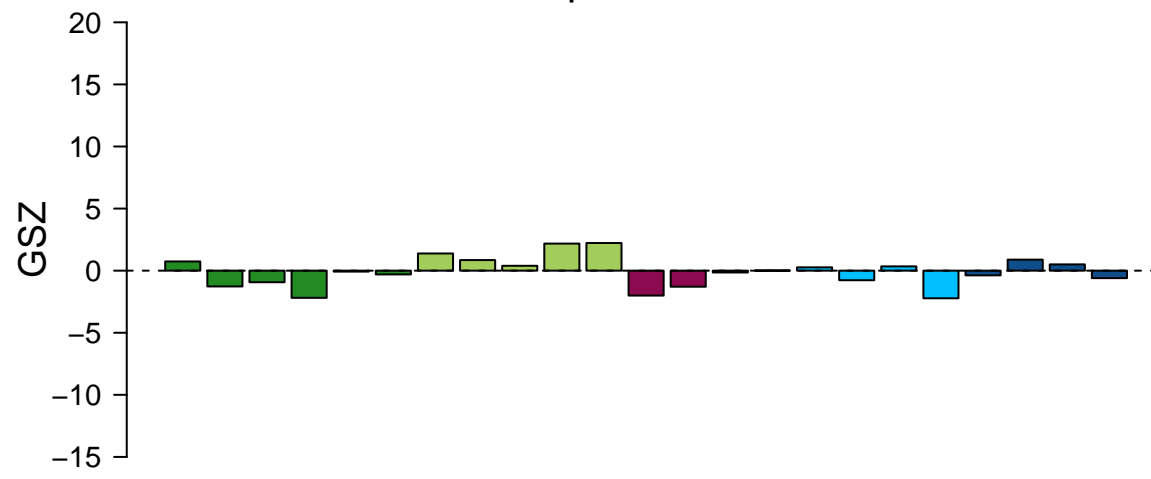
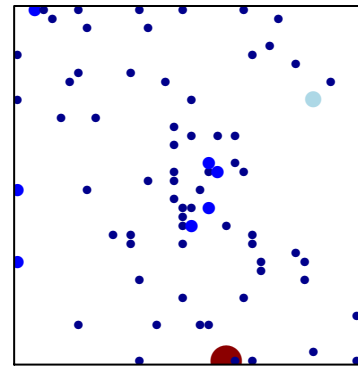
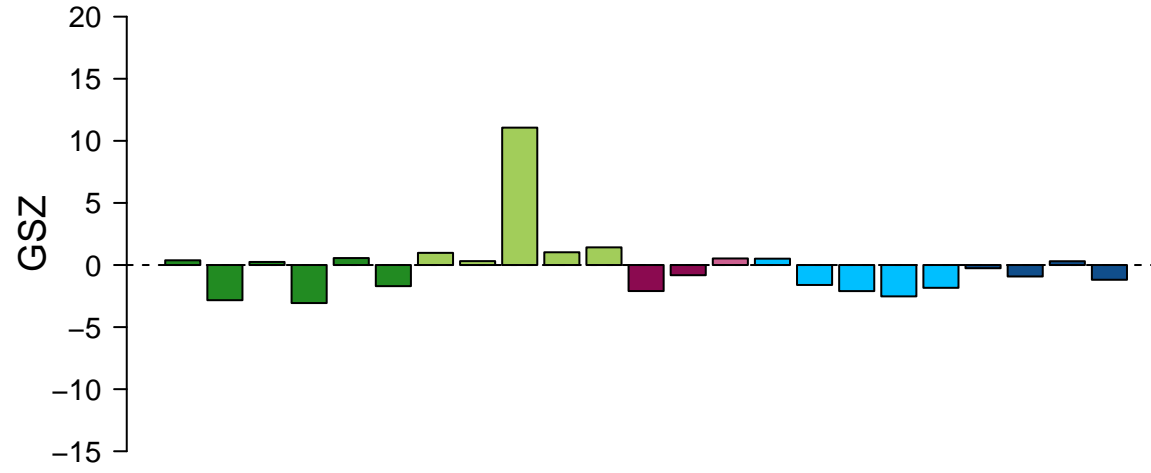


lamellipodium membrane



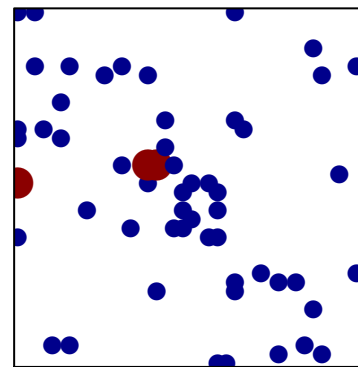
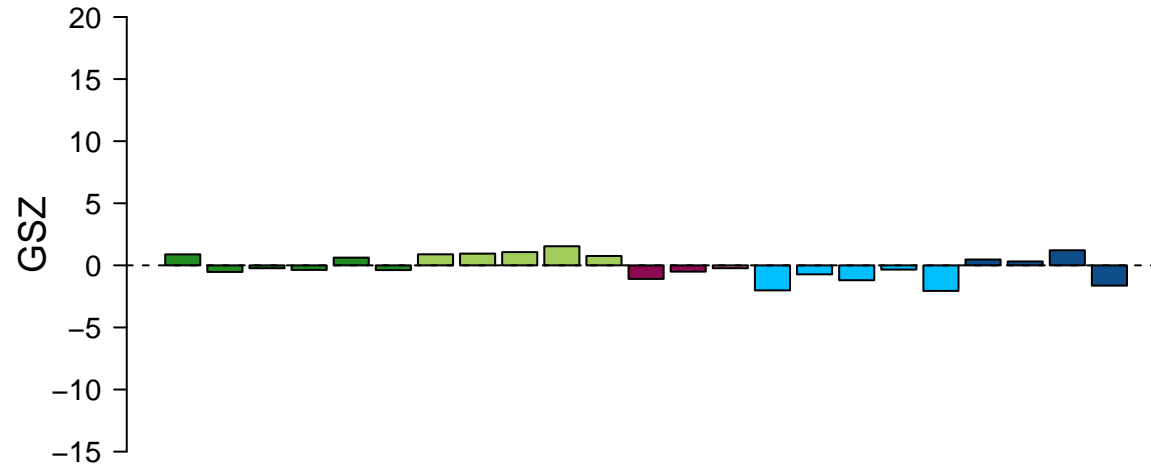
features = 18 , max = 1

phototransduction, visible light



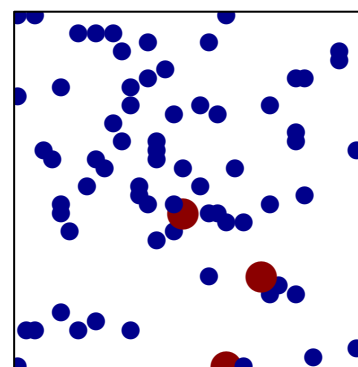
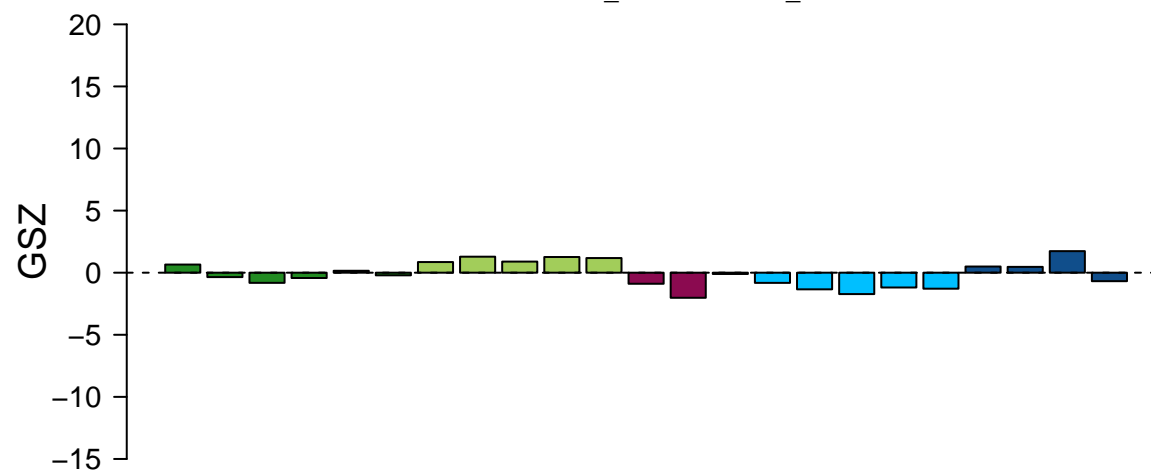
features = 89 , max = 7

sensory perception of pain



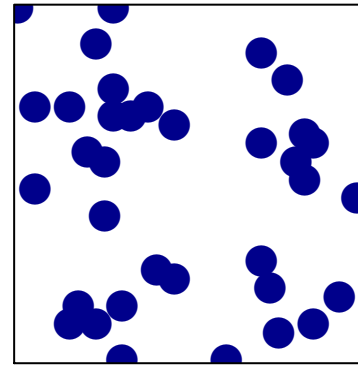
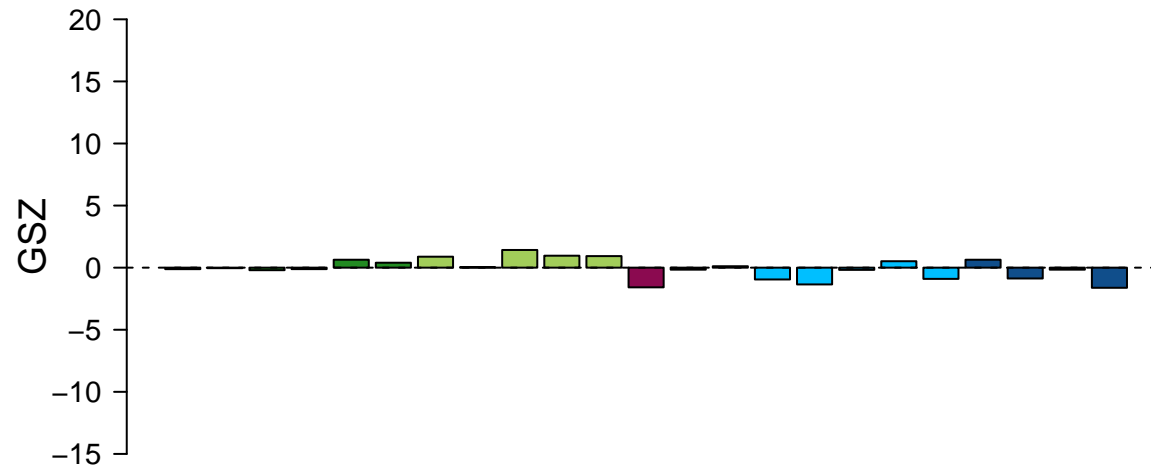
features = 59 , max = 2

ODONNELL_METASTASIS_UP



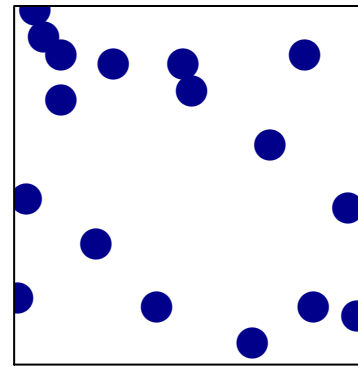
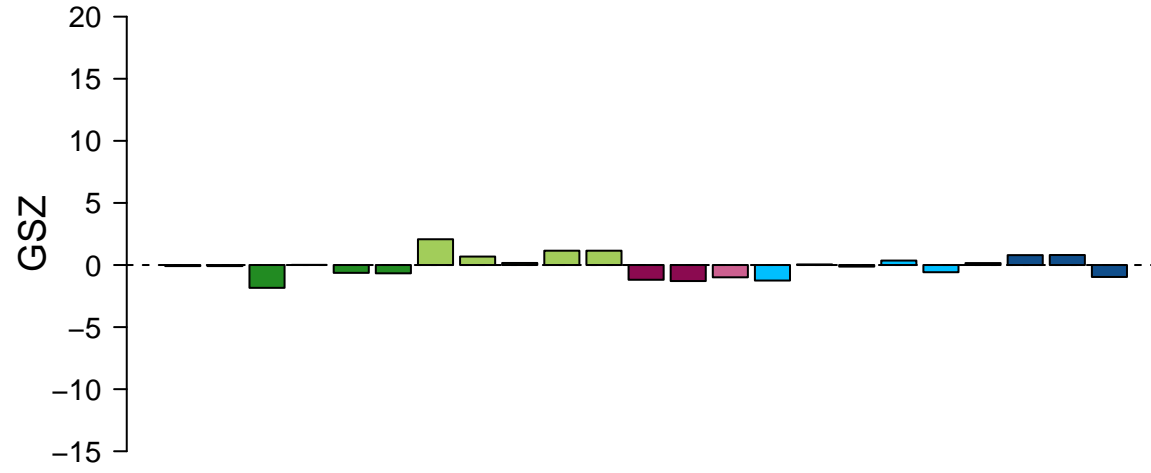
features = 74 , max = 2

GAUSSMANN_MLL_AF4_FUSION_TARGETS_G_DN



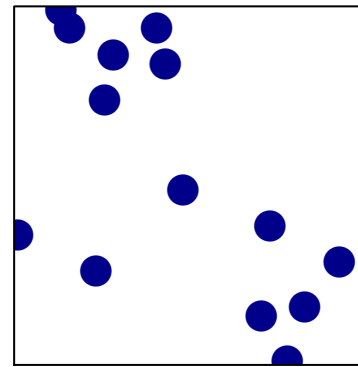
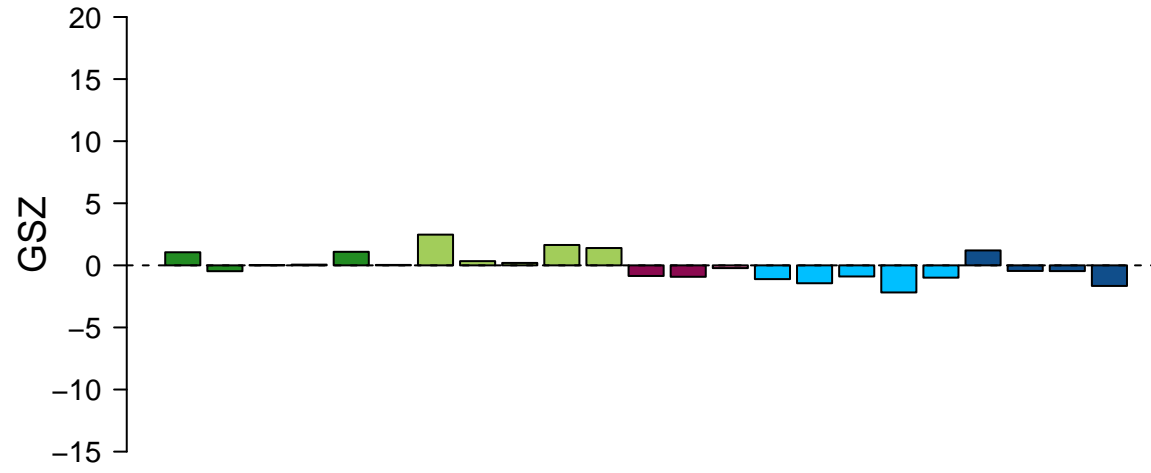
features = 35 , max = 1

KYNG_NORMAL_AGING_UP



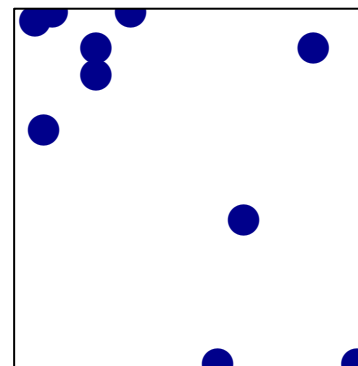
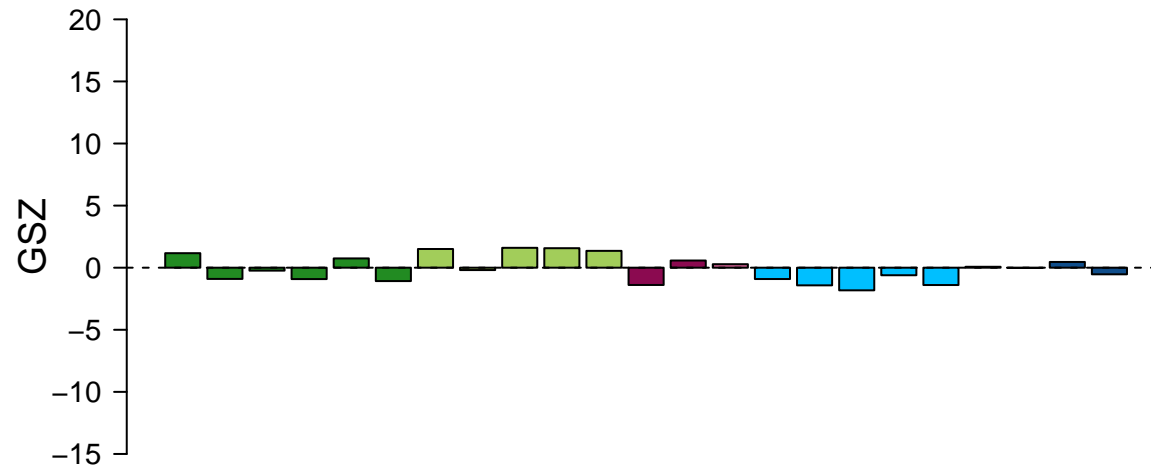
features = 17 , max = 1

REACTOME_REGULATION_OF_AMPK_ACTIVITY_VIA_LKB1



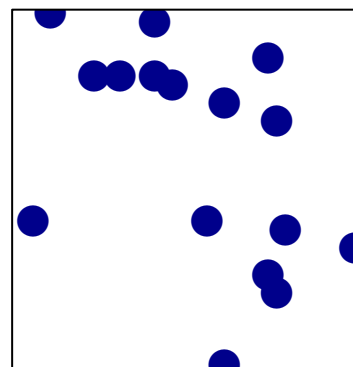
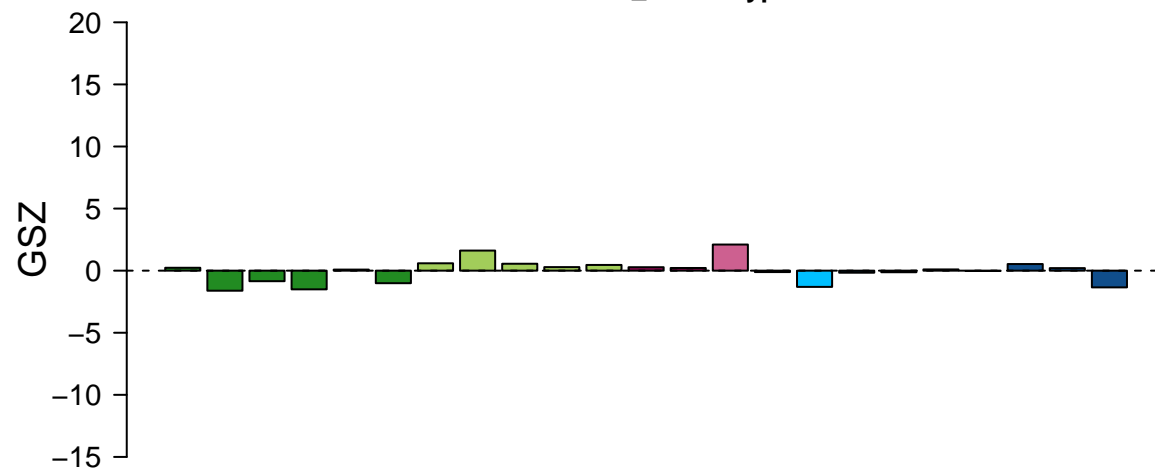
features = 14 , max = 1

negative regulation of catenin import into nucleus



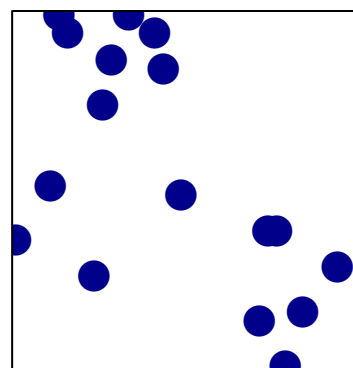
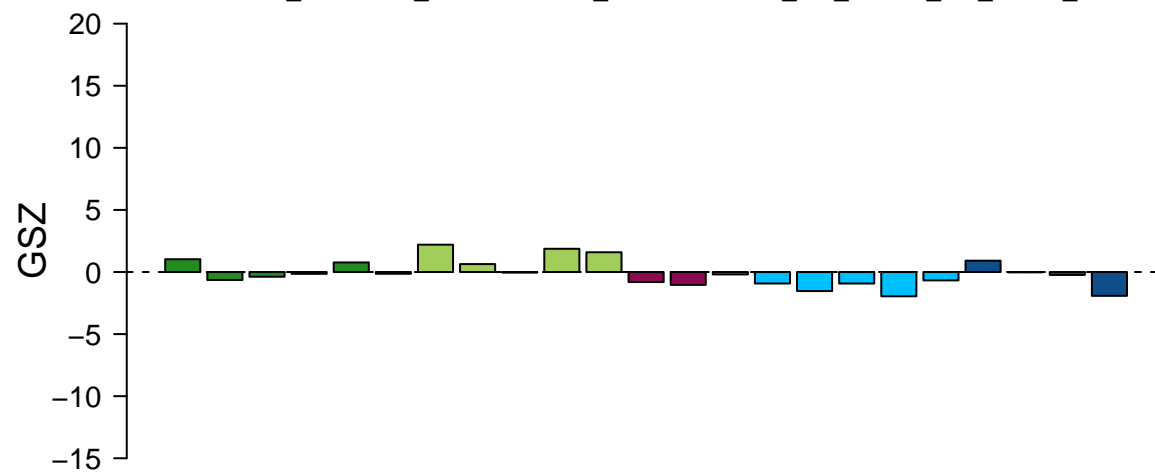
features = 10 , max = 1

VERHAAK_NL subtype



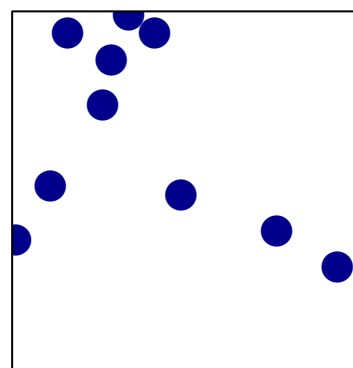
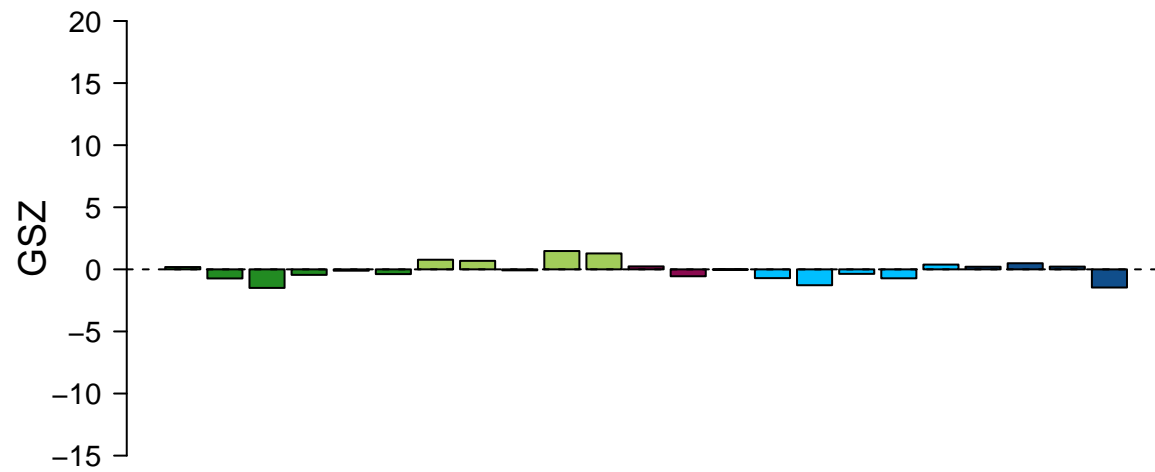
features = 16 , max = 1

REACTOME_ENERGY_DEPENDENT_REGULATION_OF_MTOR_BY_LKB1_AMPK



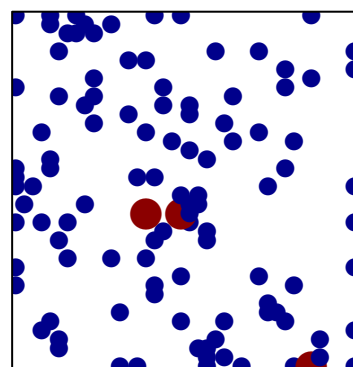
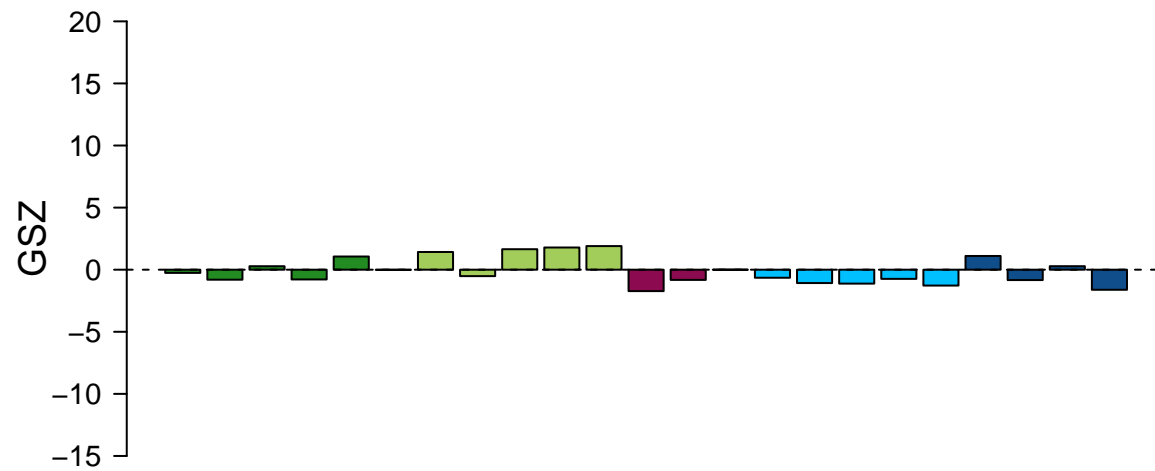
features = 17 , max = 1

REACTOME_REGULATION_OF_RHEB_GTPASE_ACTIVITY_BY_AMPK



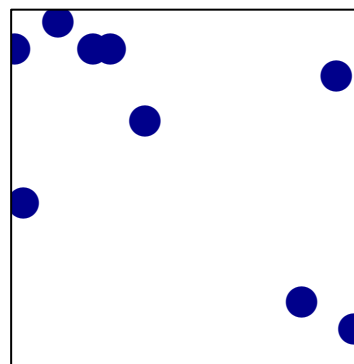
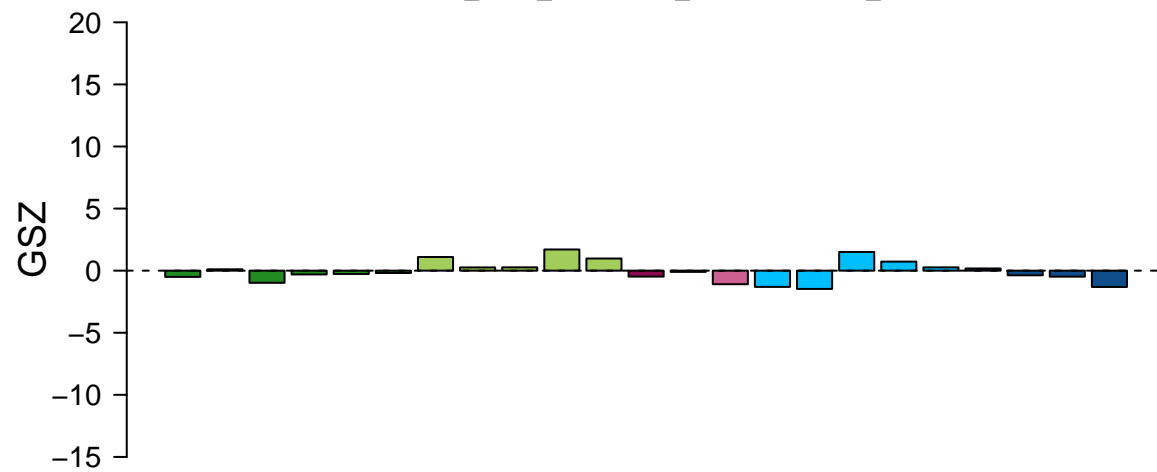
features = 10 , max = 1

XU_GH1_EXOGENOUS_TARGETS_DN



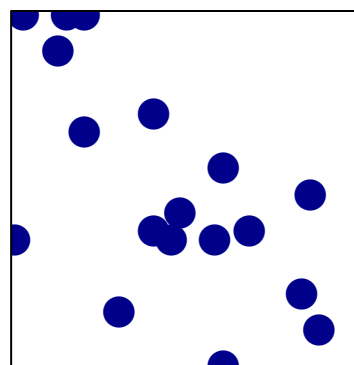
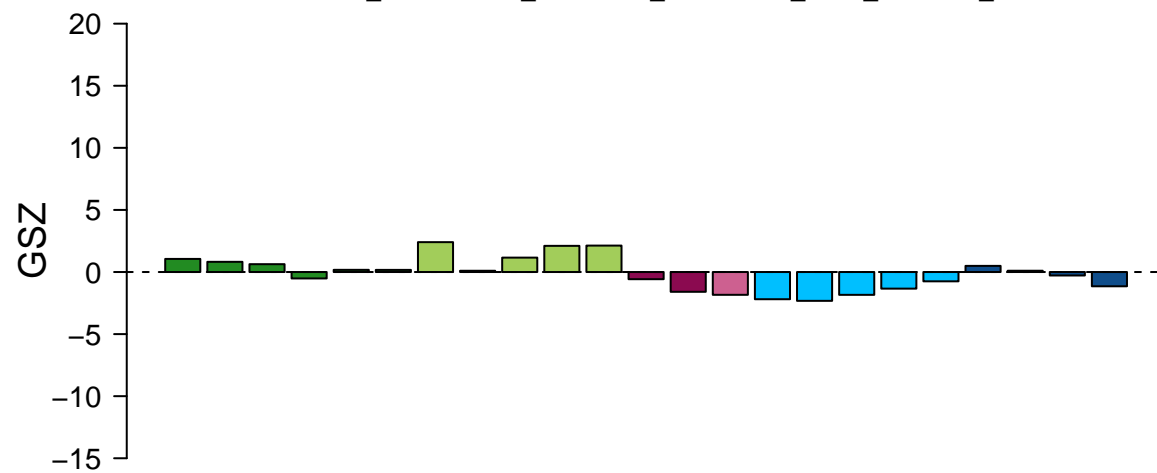
features = 113 , max = 2

CLAUS_PGR_POSITIVE_MENINGIOMA_UP



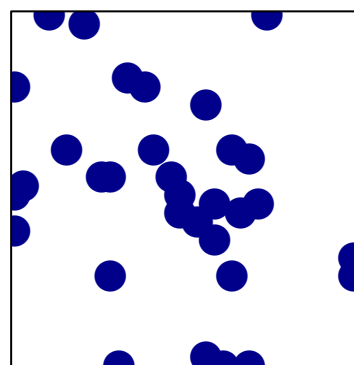
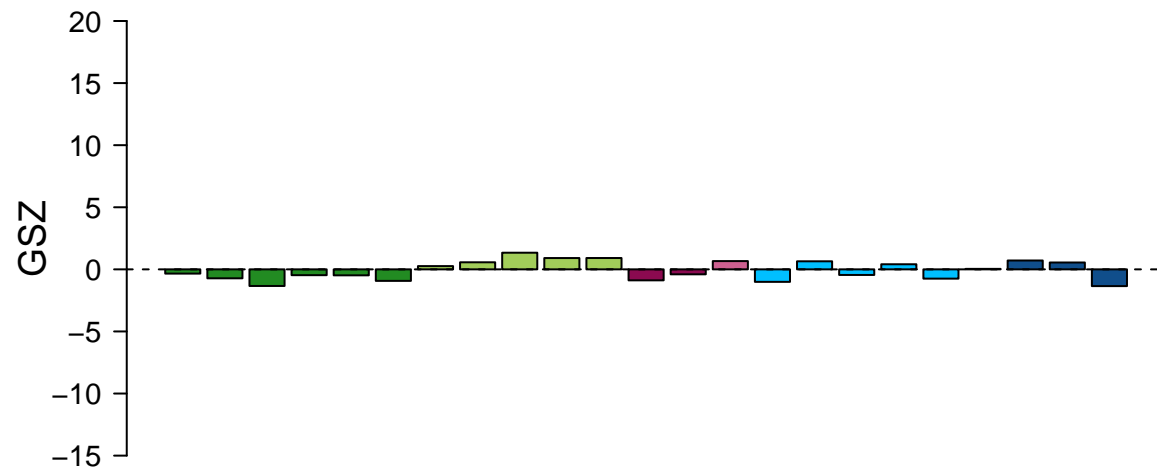
features = 9 , max = 1

SETLUR_PROSTATE_CANCER_TMPRSS2_ERG_FUSION_DN



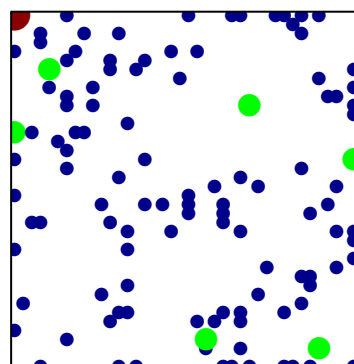
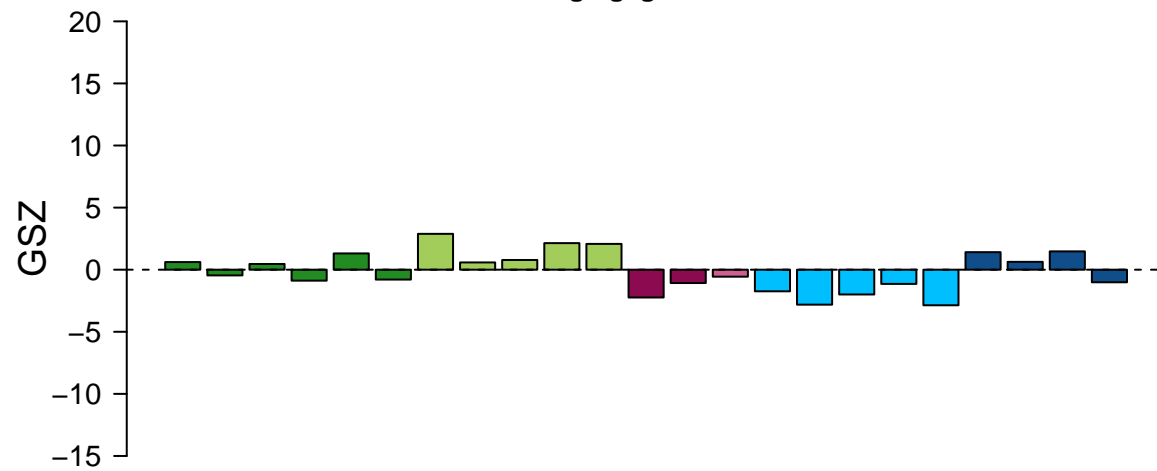
features = 18 , max = 1

insulin secretion



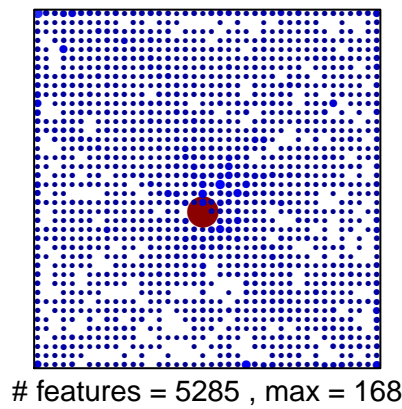
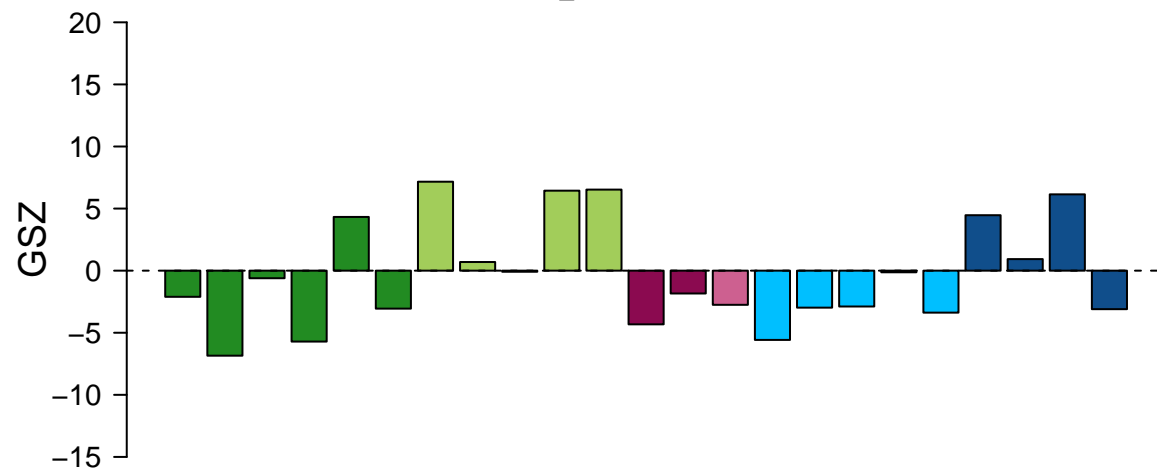
features = 32 , max = 1

HORVATH_aging_genes meth DOWN

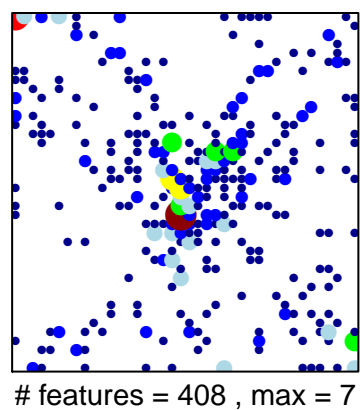
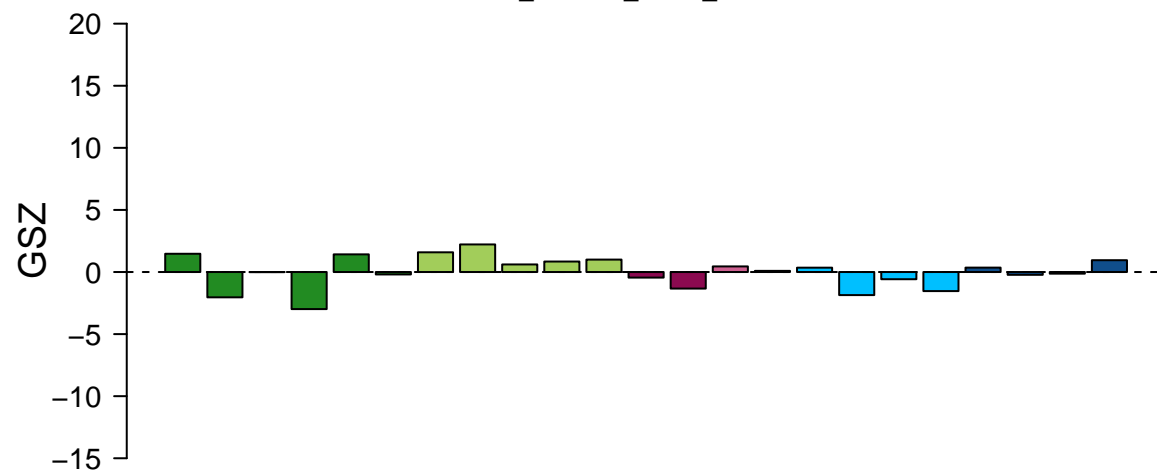


features = 123 , max = 3

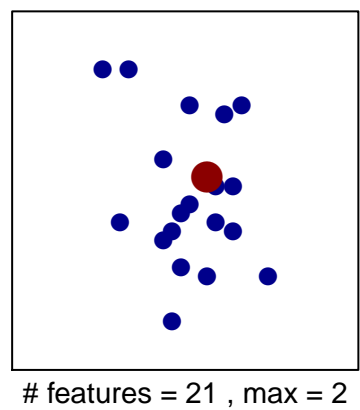
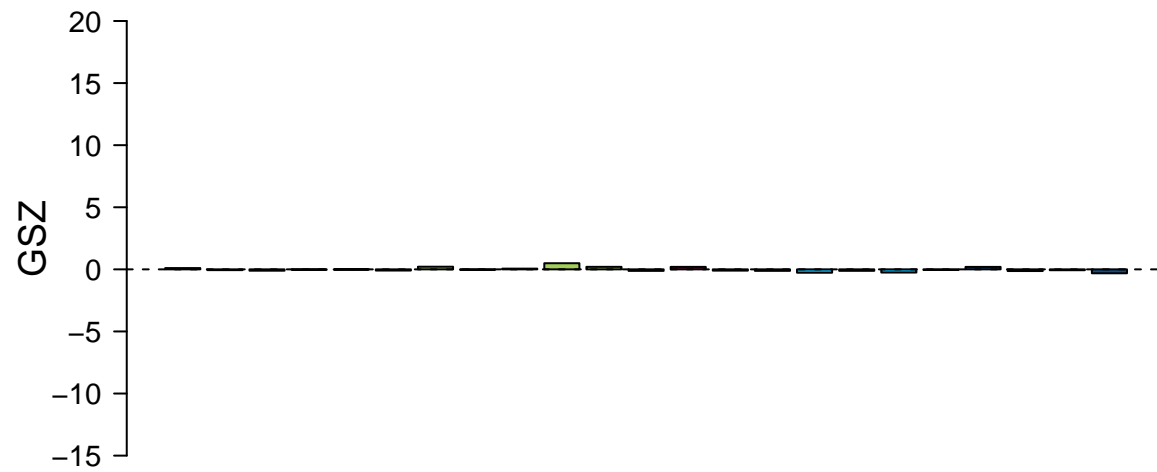
LU_BPDE 0.005 DN



Mid_Frontal_Lobe_TxTrans



REACTOME_LIGAND_GATED_ION_CHANNEL_TRANSPORT



YAUCH_HEDGEHOG_SIGNALING_PARACRINE_DN

