Selected genes from the study WALLACE_PROSTATE_CANCER_DN
REACTOME_REVERSIBLE_HYDRATION OF_CARBON_DIOXIDE
REACTOME_REVERSIBLE_HYDRATION OF_CARBON_DIOXIDE
REACTOME_REVERSIBLE_HYDRATION OF_CARBON_DIOXIDE
REACTOME_PROSTURAL_RIZ1_TARGETS_DN
KIM_MYCL1_AMPLIFICATION_TARGETS_DN
KIM_MYCL1_AMPLIFICATION_TARGETS_UP
GO_G_PROTEIN_COUPLED_NEUROTRANSMITTER_RECEPTOR_ACTIVITY
GO_G_PROTEIN_COUPLED_PURINERGIC_NUCLEOTIDE_RECEPTOR_SIGNALING_PATHWAY
GO_CARBONATE_DEHYDRATIASE_ACTIVITY
HASEGAWA_TUMORIGENESIS_BY_RET_C634R
GO_NEGATIVE_REGULATION_OF_ATPASE_ACTIVITY
KREPPEL_C1099_TARGETS_DN
GO_POSITIVE_REGULATION_OF_SYNAPTIC_TRANSMISSION_GABAERGIC
HOEGERKORP_C144_TARGETS_TEMPORAL_UP
NIELSEN_GIST_AND_SYNOVIAL_SARCOMA_UP
NIELSEN_GIST_AND_SYNOVIAL_SARCOMA_UP
NIELSEN_GIST_AND_SYNOVIAL_SARCOMA_UP
NIELSEN_GIST_AND_SYNOVIAL_SARCOMA_UP
NIELSEN_GIST_AND_SYNOVIAL_SARCOMA_UP
OOL_LIU_TOPP911_TARGETS
GO_HEMOGLOBIN_METABOLIC_PROCESS
GO_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE_BY_CIRCULATORY_RENIN_ANGIOTENSIN
WEBER_METHYLATED_IN_COLON_CANCER
GO_REGULATION_OF_PROTEIN_KINASE_C_SIGNALING
GO_REGULATION_OF_PROTEIN_KINASE_A_SIGNALING
SUZUKI_RESPONSE_TO_TSA_AND_DECITABINE_TB
GUTIERREZ_WALDENSTROEMS_MACROGLOBULINEMIA_1_DN
TUOMISTO_TUMOR_SUPPRESSION_BY_COL_13:a1_UP
GO_RESPONSE_TO_ANGIOTENSIN
NIELSEN_MALIGNAT_FIBROUS_FIISTIOCYTOMA_UP
GO_RESPONSE_TO_ANGIOTENSIN
NIELSEN_MALIGNAT_FIBROUS_FIISTIOCYTOMA_UP
GO_RESPONSE_TO_ANGIOTENSIN
NIELSEN_MALIGNAT_FIBROUS_FIISTIOCYTOMA_UP
GO_RESPONSE_TO_ANGIOTENSIN
NIELSEN_MALIGNAT_FIBROUS_FIISTIOCYTOMA_UP
GO_REGULATION_OF_CLE_PROJECTION_SIZE
GO_MATURE_B_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE
GO_CELLULAR_RESPONSE_TO_EXOSCENOUS_DISRNA GO_CELLULAR_RESPONSE_TO_EXOGENOUS_DSRNA GO_AMINO_SUGAR_CATABOLIC_PROCESS MODULE_354 MODULE_227 STEGER_ADIPOGENESIS_DN MODULE 227

STEGER_ADIPOGENESIS_DN

GO_MULTIVESICULAR_BODY_SORTING_PATHWAY
MODULE 335
GO_REGULATION_BY_VIRUS_OF_VIRAL_PROTEIN_LEVELS_IN_HOST_CELL
BIOCARTA_AHSP_PATHWAY
GCGCCTT_MIR525_MIR524
RORIE_TARGETS_OF_EWSR1_FLI1_FUSION_DN
ROETH_TERT_TARGETS_UD
GO_POSITIVE_REGULATION_OF_BONE_REMODELING GO_POSITIVE_REGULATION_OF_BONE_REMODELING GO C2H2 ZINC FINGER DOMAIN ZEMBUTSU SENSITIVITY TO CYCLOPHOS NIELSEN GIST VS SYNOVIAL SARC HO LIVER CANCER VASCULAR I GO_PURINERGIC_NUCLEOTIDE_RECEPTOR_SIGNALING GO BETA AMYLOID METABOLIC PROC GO_ADENYLATE_CYCLASE_ACTIVATING_DOPAMINE RECEPTOR SIGNALING PATH BUCKANOVICH T LYMPHOCYTE HOMING ON TUMOR KEGG PROXIMAL TUBULE BICARBONATE RECLAMAT GO_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE_BY_RENIN_ANGIOTEN GO_REGULATION_OF_SYSTEMIC_ARTÉRIAL_BLOOD_PRESSURE_BY_RENIN_ANGIOTENSIN MODULE 294

GO_POSITIVE_REGULATION_OF_INSULIN_SECRETION_INVOLVED_IN_CELLULAR_RESPONSE_TO_GLUCOSE_STIMULUS
GO_REGULATION_OF_INSULIN_SECRETION_INVOLVED_IN_CELLULAR_RESPONSE_TO_GLUCOSE_STIMULUS
GO_REGULATION_OF_INSULIN_SECRETION_INVOLVED_IN_CELLULAR_RESPONSE_TO_GLUCOSE_STIMULUS
GO_REGULATION_OF_INSULIN_SECRETION_INVOLVED_IN_CELLULAR_RESPONSE_TO_GLUCOSE_STIMULUS
GO_REGULATION_OF_INSULIN_SECRETION_INVOLVED_INVOLVED_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVOLVEN_INVO WURSC...
GÖ CELLULAIN
GÖ CELLULAIN
KEGG_GLYCINE_SERINE_AINE_

GO_NEGATIVE_REGULATION_OF_NOTCH_SIGNALING_PAIL....
chr13q32

YAO TEMPORAL_RESPONSE TO PROGESTERONE CLUSTER 3
GO_POSITIVE_REGULATION OF OSTEOCLAST_DIFFERENTIATION DORN_ADENOVIRUS_INFECTION 12HR_UP_
TRAYNOR_RETT_SYNDROM_UP_
BANDRES_RESPONSE TO CARMUSTIN MGMT 48HR_UP_
RIZ_ERYTHROID_DIFFERENTIATION 6HR
WANG_IMMORTALIZED_BY_HOXA9_AND_MEIST_DN
REACTOME_ACTIVATED_NOTCH1_TRANSMITS_SIGNAL_TO_THE_NUCLEUS_
TSENG_ADIPOGENIC_POTENTIAL_UP_
GO_NEGATIVE_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_MARIADASON_RESPONSE_TO_CURCUMIN_SULINDAC_5_MARSON_FOXP3_TARGETS_STIMULATED_UP_
FERRANDO_HOX11_NEIGHBORS_GO_BICARBONATE_TRANSPORT_GO_RESPONSE_TO_PID_HNF3B_PATHWAY_COMES_ACENTS_UITARGET_TARGET_SULINDAL_UP_
TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TARGET_TAR PID HNF3B PATHWAY
BACOLOD_RESISTANCE_TO_ALKYLATING AGENTS UP
WELCH GATA1 TARGETS
MARCHINI TRABECTEDIN RESISTANCE UP
AMIT EGF RESPONSE 240 MCF10A
BOGNI_TREATMENT_RELATED_MYELOID_LEUKEMIA_UP NAKAYAMA_SOFT_TISSUE_TUMORS_PACE_TO_
DASU_ILE_SIGNALING_SOFT_DISCRIPTION

DASU_ILE_SIGNALING_SOFT_DISCRIPTION

GO_POSITIVE_REGULATION_OF_G_PROTEIN_COUPLED_RECEPTOR_PROTEIN_SIGNALING_PROTEIN_BISSUE_
GO_PYSTEDVAL_PROSPHATE_BISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSUE_TISSU DASU_IL6_SIGNALING_SCAR DAVICIONI_PAX_FOXO1_SIGNATURE_IN_ĀRMS_UP
RELA_DN.V1_DN
HENDRICKS_SMARCA4_TARGĒTS_DN
NKX25_01
GNF2_BNIP3L
GNF2_RAD23A
REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION
GO_REGULATION_OF_PH
CERVERA_SDHB_TARGĒTS_S
GSE41176_UNSTIM_VS_ANTI_IGM_STIM_TAK1_KO_BCELL_3H_DN
GNF2_TAL1
GNF2_SPTB
GNF2_TAL1
GNF2_SPTB
GNF2_TAL1
GNF2_SPTB
GNF2_ANK1
REACTOME_SIGNALING_BY_NOTCH1 REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCATE_DEFENDING

GSE41176_UNSTIM_VS_ANTLIGM_STIM_VS_CROWNING

GSE4136_DVY_VS_DVY_VS_CROWNING

GSE4136_DVY_VS_DVY_VS_CROWNING

GSE4136_DVY_VS_DVY_VS_CROWNING

GSE4136_DVY_VS_DVY_VS_CROWNING

GSE4136_DVY_VS_DVY_VS_CROWNING

GSE4136_DVY_VS_DVY_VS_CROWNING

GSE42601_MMATURE_CD4_SINGLE_PGSTIVE_CG4_DVY_DVY_VS_CROWNING

GSE42601_MMATURE_CD4_SINGLE_PGSTIVE_CG4_DVY_DVY_VS_CROWNING

GSE42601_MMATURE_CD4_SINGLE_PGSTIVE_CG4_DVY_DVY_US_CROWNING

MIXECESEALUE_LC5_WITH_FLX_MIRE

GSE4360_NEXTW_VS_CROWNING

GSE4360_DVS_MIRE_DVX_US_CG5_DVY_DVY_US_CROWNING

WS_GSE4360_DVS_MIRE_DVS_CROWNING

GO_PGSTIVE_REGULATION_OF_BISS_CROWNING

GO_PGSTIVE_REGULATION_OF_BISS_CROWNING

GO_PGSTIVE_REGULATION_OF_BISS_CROWNING

GO_PGSTIVE_REGULATION_OF_BISS_CROWNING

GSE4360_DVS_MIRE_DVS_CROWNING

GSE4360_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MIRE_DVS_MI GSE22886_IGG_IGA_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_I GAUSSMANN_MLL_AF4_FUSION_TARGETS_ATI GSE15330_GRANULOCYTE_MONOCYTE_PROGENITOR_VS_PRO_BCELL_I GSE26030_UNSTIM_VS_RESTIM_TH17_DAY15_POST_POLARIZATION_I GSE30971_2H_VS_4H_LPS_STIM_MACROPHAGE_WBP7_KOTI GSE21379_WT_VS_SAP_KO_TFH_CD4_TCELL_I FOXM1_ OCT1_Q5_ GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_10MIN_T GSE25087_TREG_VS_TCONV_FETUS_I GSE32986_UNSTIM_VS_GMCSF_STIM_DC_T GSE2000/_INLO_VO_TGATTTRY_GFIT_01

GSE32986_UNSTIM_VS_GMCSF_STIM_DC_UP

MEISSNER_BRAIN_HCP_WITH_H3K4ME3_AND_H3K27ME3

GSE46606_IRF4HIGH_VS_IRF4MID_CD40L_TL2_IE5_DAY1_STIMULATED_BCELL_DN

GSE39820_IL1B_IL6_VS_IL1B_IL6_US_A_TREATED_CD4_TCELL_DN

GSE2128_C57BL6_VS_NOD_THYMOCYTE_DN

PASQUALUCCI_LYMPHOMA_BY_GC_STAGE_DN

GSE28237_FOLLICULAR_VS_EARLY_GC_BCELL_UP

GO_TRANSCRIPTIONAL_ACTIVATOR_ACTIVITY_RNA_POLYMERASE_II_CORE_PROMOTER_PROXIMAL_REGION_SEQUENCE_SPECIFIC_BINDING
GSE4748_CYANOBACTERIUM_LPSLIKE_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP

VDR_Q3

CSE31062_CTPL_VS_ANT_IGM_STIM_BCELL_16H_UP GSE21063 CTRL VS ANTI IGM STIM BCELL 16H UP
GSE21546_WT_VS_SAP1A_KO_AND_ELK1_KO_ANTI_CD3_STIM_DP_THYMOCYTES_DN
OCT_06
OCT_06
GSE20715 0H VS_24H OZONE_TLR4_KO_LUNG_UP
GSE3039_ALPHAALPHA_VS_ALPHABETA_CD8_TCELL_DN
GSE21796_LSK_VS_NKTCELL_DN
GSE19941_IL10_KO_VS_IL10_KO_AND_NFKBP50_KO_UNSTIM_MACROPHAGE_DN
GSE45382_UNTREATED_VS_TGFB_TREATED_MACROPHAGES_DN
GSE40274_CTRL_VS_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
E2F1_UP_VT_DN
STY_02
RREB1_01
GSE40274_CTRL_VS_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE40274_CTRL_VS_SAP_KO_CD4_TCELL_DN
GSE40274_CTRL_VS_SAP_KO_CD4_TCELL_DN
GSE40274_CTRL_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP_MSSARWEN_TSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_NSMN_UP_TC_ GSE21063 CTRL_VS_ANTI_IGM_STIM_BCELL_16H GSE21546_WT_VS_SAP1A_KO_AND_ELK1_KO_ANTI_CD3_STIM_DP_THYMOCYTES GSE13493 CD4INTCD8POS VS CD8POS THYMOCYTE UP
GSE32164_RESTING_DIFFERENTIATED VS_CMYC_INFIBITED_MACROPHAGE_UP
GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_DN
GSE5960_TH1_VS_ANERGIC_TH1_UP
GSE45739_NRAS_KO_VS_WT_UNSTIM_CD4_TCELL_DN
GSE39820_CTRL_VS_LTB_ILB_IL6_IL23A_CD4_TCELL_DN
GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD8_TCELL_DN
PAX4_04
SOX5_01 SOX5 01
GSE17721_0.5H_VS_24H_LPS_BMDC_DN
SPZ1_01
GSE46606_UNSTIM_VS_CD40L_IL2_IL5 1DAY STIMULATED IRF4 KO BCELL_DN
GSE30083 SP1 VS SP2 THYMOCYTE_DN
GSE5542_IFNG_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_24H_UP
AREB6_03
TITF1_Q3 AREB6 03
TITF1 03
GSE3982 BCELL VS TH1 UP
GSE20366_CD103_KLRG1_DP_VS_DN_TREG_DN
VTATTTNR MEF2_02
GSE3039_ALPHABETA_CD8_TCELL_VS_B2_BCELL_UP
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_52H_DN
GSE20754_WT_VS_TCF1_KO_MEMORY_CD8_TCELL_DN
GSE30754_WT_VS_TCF1_KO_MEMORY_CD8_TCELL_DN
GSE30754_WT_VS_TCF1_KO_MEMORY_CD8_TCELL_DN
GSE30754_WT_VS_TCF1_KO_MEMORY_CD8_TCELL_DN
GSE30754_WT_VS_TCF1_KO_MEMORY_CD8_TCELL_DN
UP WANG_MLL_TARGETS
GSE3203_HEALTHY_VS_INFLUENZA_INFECTED_LN_BCELL_UP
GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_UP
GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_UP
GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_UP
GSE374179_THPOK_KO_VS_WT_VA14I_NKTCELL_DN
GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GFP_TREG_DN
GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GFP_TREG_DN
GSE40274_CTRL_VS_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE20366_CD103_POS_VS_CD103_KLRG1_DP_TREG_DN
GSE20366_CD103_POS_VS_CD103_KLRG1_DP_TREG_DN
GSE20366_CD103_POS_VS_CD103_KLRG1_DP_TREG_DN
GSE40143_CTRL_VS_LMP2A_TRANSDUCED_CD10_POS_GC_BCELL_DN
SENESE_HDAC1_AND_HDAC2_TARGETS_DN
SENESE_HDAC1_AND_HDAC2_TARGETS_DN SENESE_HDAC1_AND_HDAC2_TARGETS_ SOX9 GSE3982 MAST CELL VS THI
GARGALOVIC RESPONSE TO OXIDIZED PHOSPHOLIPIDS GREY
GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCEIL_60H GO_COFACTOR_BIOSYNTHETIC_PROCES GO_COFACTOR_BIOSYNTHETIC_PROCES GSE34515_CD16_POS_MONOCYTE_VS_DC_L GSE12198_NK_VS_NK_ACT_EXPANSION_SYSTEM_DERIVED_NK_CELL_D BENPORATH_ES_WITH_H3K27ME MODITION GSE20715_0H_VS_6H_OZONE LUNG_UP MODULE 19 ACROPHAGE 12H MBOVIS BCG STIM DN 424_CD161_HIGH_VS_INT_CD8_TCELL_UP MODULE 220 23_WT_VS_PPARG_KO_MACROPHAGE_UP GSE22935_WT_VS_MYD88_KO_MACROPHAGE_12H_MBOVIS_BCGSE33424_CD161_HIGH_VS_INT_CD8 GSE25123 WT_VS_PPARG_KO_MACRO GSE17721_0.5H_VS_12H_GARDIQUIMOD GSE41867_NAIVE_VS_DAY15_LCMV_CONE13_EFFECTOR_CD8 GSE43863_DAY6_EFF_VS_DAY150_MEM_TH1_CD4_TCELL_T KAECH_DAY15_EFF_VS_MEMORY_CD8_TCELL_T LINDGREN_BLADDER_CANCER_CLUSTER_; GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_T GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_24H_T GO_MYELOID_CELL_HOMEOSTAS GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_T MODULE_ SERVITJA_LIVER_HNF1A_TARGETS_ CREB_ CMITTIGT_UNKNOT SMTTTTGT_UŇ GSE23568_ID3_KO_VS_WT_CD8_TC GREGORY_SYNTHETIC_LETHAL_WITH_I GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS GO_TRANSCRIPTIONAL_ACTIVATOR_ACTIVITY_RNA_POLYMERASE_II_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_T GREĞÖRY SYNTHETIC LETHAL WITH IMATINIE
GSE5542 UNTREATED UN IPING TREATED EPITHELIAL CELLS 24H UP
TRANSCRIPTIONAL ACTIVATOR ACTIVITY_RNA_POLYMERASE_II_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_BINDING

BENDPORATH EFT STATE
CAIRO HEPATOBLASTOMA_UP

GSE9509_LPS_VS_LPS_AND_L10_STMI_L10_KO MARCOPHAGE_BONG
GSE9509_LPS_VS_LPS_AND_L10_STMI_L10_KO MARCOPHAGE_BONG
GSE9509_LPS_VS_LPS_AND_L10_STMI_L10_KO MARCOPHAGE_BONG
GSE47274_MEMORY_US_HIGH_L12_STM_NK_CELL_UP

GO_TRANSCRIPTION_FACTOR_ACTIVITY_RNA_POLYMERASE_II_CORE_PROMOTER_PROXIMAL_REGION_SEQUENCE_SPECIFIC_BINDING
GSE47274_MEMORY_GSELL_VS_FLASMABING_UP

GO_TRANSCRIPTION_FACTOR_ACTIVITY_RNA_POLYMERASE_II_CORE_PROMOTER_PROXIMAL_REGION_SEQUENCE_SPECIFIC_BINDING
GSE47174_MEMORY_GSELL_VS_FLASMABING_UP

GSE47174_TARGETS_OF_EWSR_FLI_FUSION_SEQUENCE_SPECIFIC_BINDING
GSE41176_UNSTIM_US_ANTI_LGROWNED_BE_FLASMABING_US_
KINSEY_TARGETS_OF_EWSR_FLI_FUSION_SEQUENCE_SPECIFIC_BINDING
GSE41176_UNSTIM_US_ANTI_LGROWNED_BE_FLASMABING_US_
GSE3403_LT_US_LASMABING_US_FLASMABING_US_FLASMABING_US_
GSE3403_LT_US_LASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_FLASMABING_US_ pval < 0.05 **FALSE** TRUE GSE3920_IFNA_VS_IFNB_TREATED_ENDOTHELIAL_CELL_ MODULE_ GSE24972_WT_VS_IRF8_KO_MARGINAL_ZONE_SPLEEN_BCELL_ GO_NEGATIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION_OF_CHROMOSOME_ORGANIZATION_CELL_ MODULE_ GSE17721_LPS_VS_PAM3CSK4_1H_BMDC_ GSE19941_IL10_KO_VS_IL10_KO_AND_NFKBP50_KO_LPS_STIM_MACROPHAGE_ TBK1.DF_ REACTONE ANTORS, STATE MESS TO STATE OF SECTION OF SECT GSE25677_MPL_VS_R848_STIM_BCELL PXR JUBAN_TARGETS_OF_SPI1_AND_FLI1 SREBP MARY_BRONCHIAL_EPITHELIAL_CELLS GSE2025 UNTREATED VS PROGESTERONE, TREATED, COJ. TCELL DN
GSE566 STRONG, VS. WEAK, DC. STIMULATED, COJ. TOTAL DISTRICT
GSE566 STRONG, VS. WEAK, DC. STIMULATED, COJ. TCELL DN
GSE37301 HEMATOPOLETIC, STEM, CELL, VS. PRO BC. CHIEF AND GSE39301 UNTREATED VS. IPNA TREATED FINANCIA COLD BY
GSE3920 IPNA VS. IPNA TREATED FINANCIA COLD BY
GSE3930 PRE VS. POST ILG INJECTION IPNA VY LIVER UP
GSE3442 CTRL, VS. HERT TRANSDICED ON GSE37788 INVESTIGATION OF COLD BY
GSE3442 CTRL, VS. HERT TRANSDICED ON GSE34411, NAIVE VS. IGM MEMOSYT COLD BY
GSE34411, NAIVE VS. IGM MEMOSYT COLD BY
GSE3440 MEMORY SCELL DAYY. VS. GERMINAL CENTER SCELL DAYY. DN
GSE34411, NAIVE VS. IGM MEMOSYT COLD BY
GSE3646 CTRL, VS. TSST ACT. 161 MEMORY COLD BY COLD BY
GSE36476 CTRL, VS. TSST ACT. 161 MEMORY COLD BY COLD BY
GSE36476 CTRL, VS. TSST ACT. 161 MEMORY COLD BY COLD BY
GSE36476 CTRL, VS. TSST ACT. 161 MEMORY COLD BY COLD BY
GSE340274 LEFT VS. FOXP3. AND ASSESS BY COLD BY
GSE3640 DOUBLE NEGASSE 4066 NAIVE VS. EFFECTOR COST COLD BY
GSE370 DOUBLE NEGASSE 4066 NAIVE VS. EFFECTOR COST TCELL UN
GSE37186 NAIVE VS. COST IM BILOCK MIS. COST TCELL UN
GSE37186 NAIVE VS. COST IM BILOCK MIS. COST TCELL UN
GSE3746 CTRL, VS. FOXP3. TRANSDIVE COST TCELL UN
GSE3750 DOUBLE NEGASSE 40666 NAIVE VS. FOXP3. TRANSDIVE COST TCELL UN
GSE376 DOUBLE NEGASSE 40666 NAIVE VS. FOXP3. TRANSDIVE COST TO BY
GSE376 DOUBLE NEGASSE 40666 NAIVE VS. FOXP3. TRANSDIVE COST TO BY
GSE376 DOUBLE NEGASSE 40666 NAIVE VS. FOXP3. TRANSDIVE COST TO BY
GSE376 DOUBLE NEGASSE 40666 NAIVE VS. TSC TSC TSC TSC TSC TSC TSC TSC TSC GSE10329_CD4_TOLLE_WANG_CLIM2_TARGETS_DN
KIM_WTT_TARGETS_DN
KIM_WTT_TARGETS_DN
KIM_WTT_TARGETS_DN
GSE4111 WT VS_SOC33_KO_MOROPPHAGE_LOS_TARGETS_DN
GSE41567_NATIVE_VS_DAY15_CMV_EFFECTOR_CD8_TOELL_OP_
GSE36476_CTRL_VS_TSST_ACT_40H_MOROY_CD4_TOELL_VOLING_UP_
GSE36476_CTRL_VS_TSST_ACT_40H_MOROY_CD4_TOELL_VOLING_UP_
KIMOVYS_KA_MIR.OBB_TARGETS_DN
GSE41350_TREG_VS_TEFF_IN_LIZRB_KO_UP_
GO_GLUCOSAMINE_CONTAINING_COMPOUND METABOLIC_PROCESS_GSE11057_NATIVE_VS_CENT_MEMORY_CD4_TOELL_VOLING_UP_
GSE31057_NATIVE_VS_CENT_MEMORY_CD4_TOELL_DN
GSE413867_NATIVE_VS_CENT_MEMORY_CD4_TOELL_DN
GSE31687_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TEFF_CTOR_CD8_TOELL_DN
GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TEFF_UP_
GSE37302_UNSTIM_VS_LPS_STIM_VD2_GAMMADELTA_TOELL_UP_
GSE363888_STAT5_AB_KNOCKIN_VS_WTGSE41350_TREG_VS_TEFF_UP_
GSE3736888_STAT5_AB_KNOCKIN_VS_WTGSE41350_TREG_VS_TEFF_UP_
GSE3736_NATIVE_VS_CD210W_TRANSITION_ETER_LIC_WS_ARMS_KNONC_UP_
HALLMARK_UNPCLOBED_PROTEIN_RESPONSE_
GSE3736_NATIVE_VS_CD210W_TRANSITION_ETER_LIC_WS_TOEN_BLOOD_UP_
HALLMARK_UNPCLOBED_PROTEIN_RESPONSE_
GSE37486_NATIVE_VS_CD210W_TRANSITION_ETER_LIC_UP_ARMS_KNONC_UP_
GSE3747_CPG_DNA_VS_LUNTREATED_IN_DC_DOT_
HALLMARK_UNPCLOBED_PROTEIN_RESPONSE_
GSE37486_NATIVE_VS_CD310W_TRANSITION_ETER_LIC_UP_ARMS_KNONC_UP_
GSE37407_CPG_DNA_VS_LUNTREATED_IN_DC_DOT_
GSE4110N_TRANSITION_LIC_LIC_UP_ARMS_KNONC_UP_
GSE37568_IL4_VS_IL4_AND_DEXAMETHASONE_TREATED_MOROPHAGE_DN_
GSE4110N_TRANSITION_LIC_UP_ARMS_KNONC_UP_
GSE37568_UNSTEM_ARMS_KNONC_UP_ARMS_KNONC_UP_
GSE37533_UNTREATED_VS_PIOGALIZATONE_TREATED_CD4_TOELL_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNONC_UP_ARMS_KNON GSE32901 TH17 EMRICHED VS TH17 NEG CD4 TCELL UP GSE11961_MEMORY_BCELL DAY7 VS PLASMA CELL DAY7 DN GSE11961_MEMORY_BCELL DAY7 VS PLASMA CELL DAY7 DN GSE12366 GC VS NAIVE BCELL DN GSE17721 LPS VS POLYIC 1H BMDC DN GSE36476 CTRL VS TSST ACT 72H MEMORY CD4 TCELL YOUNG UP GSE15735 CTRL VS HDAC INHIBITOR TREATED CD4 TCELL 12H DN AMUNDSON_POOR_SURVIVAL_AFTER GAMMA RADIATION 8G GO_AMINO_ACID_TRANSPORT GSE6259 DEC205 POS DC VS BCELL DN GSE11961_UNSTIM_VS_ANTI_IGM_AND_CD40_STIM_6H_FOLLICULAR_BCELL_DN GO_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE GNF2_RAP1B GO_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSU
GNF2_RAF
GSE17721_PAM3CSK4_VS_GADIQUIMOD_2H_BMDC
GSE11961_MEMORY_BCELL_DAY7_VS_PLASMA_CELL_DAY7
GO_PIGMENT_GRANULE_LOCALIZATI
GSE5679_CTRL_VS_RARA_AGONIST_AM580_TREATED_DC
GSE19888_ADENOSINE_A3R_INH_VS_ACT_WITH_INHIBITOR_PRETREATMENT_IN_MAST_CELL_
DOUGLAS_BMI1_TARGETS_
GSE27859_DC_VS_CD11C_INT_F480_HI_MACROPHAGE_
ACACTTC_MIR409
GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_CORTEX_THYMUS_
GSE14308_TH1_VS_NATURAL_TREGG_
GSE22886_NEUTROPHIL_VS_DC_
GO_BRUSH_BORD_
GSE28408_LY6G_POS_VS_NEG_DC_
GSE3920_UNTREATED_VS_IFNG_TREATED_FIBROBLAST_
GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_
GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_
E2F1_
RASHI_RESPONSE_TO_IONIZING_RADIATION RASHI RESPONSE TO IONIZING RADIATION 3
MULLIGHAN NPM1 MUTATED SIGNATURE 2 UP
GSE29164_DAY3_VS_DAY7_CD8_TCELL_TREATED_MELANOMA_UP
GSE37532_WT_VS_PPARG_KO_LN_TCONV_UP
GSE17301_CTRL_VS_48H_IFNA2_STIM_CD8_TCELL_UP
GSE17721_LPS_VS_GARDIQUIMOD_1H_BMDC_DN
RB_P107_DN_V1_UP
GO_CLUSTER_OF_ACTIN_BASED_CELL_PROJECTIONS
GSE24634_NAIVE_CD4_TCELL_VS_DAY10_IL4_CONV_TREG_DN
GSE22103_LPS_VS_GMCSF_AND_IFNG_STIM_NEUTROPHIL_UP
MAX_01 GSE27241_WT_VS_RORGT_KO_TH17_POLARIZED_CD4_TCELL_T GSE17721_LPS_VS_PAM3CSK4_16H_BMDC_T MORI_IMMATURE_B_LYMPHOCYTE_I MORI_IMMATURE_B_LYMPHOCYTE_I MORI_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_T OISHI_CHOLANGIOMA_STEM_CELL_LIKE_I ARNI 01
TCCCRNNRTGC_UNKNOWN
GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP
MORF_TNFRSF6
HOELZEL_NF1_TARGETS_UP
ACATTCC_MIR1_MIR206
MORF_MAP2K7 GSE3394_CTRL_VS_B_ABORTUS_4H_MC_GTE_L_VS_NTCELL_UP

GSE3384_CTRL_VS_B_ABORTUS_4H_MC_GTE_L_UP

GSE19888_CTRL_VS_T_CELL_WISPARCELL_VS_NTCELL_UP

GSE19888_CTRL_VS_T_CELL_WISPARCELL_VS_NTCELL_UP

GSE19888_CTRL_VS_T_CELL_WISPARCELL_VS_NTCELL_UP

GSE19888_CTRL_VS_T_CELL_WISPARCELL_VS_NTCELL_UP

GSE19888_CTRL_VS_T_CELL_WISPARCELL_VS_NTCELL_UP

GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCYTE_MONOCOMANS_GRANULOCAMAS_GRANULOCYTE_MONOCOMANS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS_GRANULOCAMAS 0.25 0.50 0.00 0.75 1.00 1.25 Normalized Enrichment Score