

	GSE10239_MEMORY_VS_KLRG1INT_EFF_CD8_TCELL_DN, GSE10239_MEMORY_VS_KLRG1INT_EFF_CD8_TCELL_DN HALLMARK_MITOTIC_SPINDLE, HALLMARK_MITOTIC_SPINDLE GSE10239_MEMORY_VS_KLRG1HIGH_EFF_CD8_TCELL_DN, GSE10239_MEMORY_VS_KLRG1HIGH_EFF_CD8_TCELL_DN GOCC_SPINDLE_POLE, GOCC_SPINDLE_POLE GSE7768_OVA_ALONE_VS_OVA_WITH_MPL_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP, GSE7768_OVA_ALONE_VS_OVA_WITH_MPL_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP GOBP_SPINDLE_ORGANIZATION, GOBP_SPINDLE_ORGANIZATION GSE37532_TREG_VS_TCONV_PPARG_KO_CD4_TCELL_FROM_LN_UP, GSE37532_TREG_VS_TCONV_PPARG_KO_CD4_TCELL_FROM_LN_UP GSE5589_WT_VS_IL6_KO_LPS_AND_IL10_STIM_MACROPHAGE_45MIN_UP, GSE5589_WT_VS_IL6_KO_LPS_AND_IL10_STIM_MACROPHAGE_45MIN_UP GSE5589_WT_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_UP, GSE5589_WT_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_UP GSE9650_EFFECTOR_VS_MEMORY_CD8_TCELL_UP, GSE9650_EFFECTOR_VS_MEMORY_CD8_TCELL_UP GSE19923_WT_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP, GSE19923_WT_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP GSE9601_UNTREATED_VS_P13K_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_UP, GSE9601_UNTREATED_VS_P13K_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_UP GSE7768_OVA_WITH_LPS_VS_OVA_WITH_MPL_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP, GSE7768_OVA_WITH_LPS_VS_OVA_WITH_MPL_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP GSE17186_CD21LOW_VS_CD21HIGH_TRANSITIONAL_BCELL_UP, GSE17186_CD21LOW_VS_CD21HIGH_TRANSITIONAL_BCELL_UP GOBP_MICROTUBULE_CYTOSKELETON_ORGANIZATION_INVOLVED_IN_MITOSIS, GOBP_MICROTUBULE_CYTOSKELETON_ORGANIZATION_INVOLVED_IN_MITOSIS GSE5503_MLN_DC_VS_PLN_DC_ACTIVATED_ALLOGENIC_TCELL_UP, GSE5503_MLN_DC_VS_PLN_DC_ACTIVATED_ALLOGENIC_TCELL_UP GSE37532_WT_VS_PPARG_KO_LN_TCONV_DN, GSE37532_WT_VS_PPARG_KO_LN_TCONV_DN GOCC_MIDBODY, GOCC_MIDBODY GSE6259_DEC205_POS_DC_VS_CD8_TCELL_UP, GSE6259_DEC205_POS_DC_VS_CD8_TCELL_UP GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_T425A_HEXON_INF_UP, GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_T425A_HEXON_INF_UP GSE7218_IGM_VS_IGG_SIGNAL_THROUGH_ANTIGEN_BCELL_UP, GSE7218_IGM_VS_IGG_SIGNAL_THROUGH_ANTIGEN_BCELL_UP GSE14415_ACT_TCONV_VS_ACT_NATURAL_TREG_DN, GSE14415_ACT_TCONV_VS_ACT_NATURAL_TREG_DN GSE8678_IL7R_LOW_VS_HIGH_EFF_CD8_TCELL_UP, GSE8678_IL7R_LOW_VS_HIGH_EFF_CD8_TCELL_UP GOBP_MITOTIC_SPINDLE_ORGANIZATION, GOBP_MITOTIC_SPINDLE_ORGANIZATION GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7_DN, GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7_DN GSE3720_UNSTIM_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_DN, GSE3720_UNSTIM_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_DN GSE13484_UNSTIM_VS_12H_YF17D_VACCINE_STIM_PBMC_UP, GSE13484_UNSTIM_VS_12H_YF17D_VACCINE_STIM_PBMC_UP GSE27241_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_TH17_POLARIZING_CONDITIONS_UP, GSE27241_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_TH17_POLARIZING_CONDITIONS_UP GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_LPS_STIM_DN, GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_LPS_STIM_DN GOBP_NUCLEOSOME_ORGANIZATION, GOBP_NUCLEOSOME_ORGANIZATION GSE21546_SAPIA_KO_VS_SAPIA_KO_AND_ELK1_KO_DP_THYMOCYTES_DN, GSE21546_SAPIA_KO_VS_SAPIA_KO_AND_ELK1_KO_DP_THYMOCYTES_DN GSE5542_IFNA_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_24H_DN, GSE5542_IFNA_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_24H_DN GSE25088_CTRL_VS_IL4_STIM_MACROPHAGE_UP, GSE25088_CTRL_VS_IL4_STIM_MACROPHAGE_UP GSE7509_UNSTIM_VS_IFNA_STIM_IMMATURE_DC_UP, GSE7509_UNSTIM_VS_IFNA_STIM_IMMATURE_DC_UP GSE3920_IFNB_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP, GSE3920_IFNB_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP GSE8685_IL2_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP, GSE8685_IL2_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP GSE2405_0H_VS_24H_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_DN, GSE2405_0H_VS_24H_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_DN GSE40274_CTRL_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_CTRL_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN GSE32901_NAIVE_VS_TH1_CD4_TCELL_DN, GSE32901_NAIVE_VS_TH1_CD4_TCELL_DN GSE21546_WT_VS_SAPIA_KO_AND_ELK1_KO_ANT1_CD3_STIM_DP_THYMOCYTES_DN, GSE21546_WT_VS_SAPIA_KO_AND_ELK1_KO_ANT1_CD3_STIM_DP_THYMOCYTES_DN GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_UP, GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_UP GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_UP, GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_UP GSE21379_WT_VS_SAP_KO_CD4_TCELL_DN, GSE21379_WT_VS_SAP_KO_CD4_TCELL_DN WP_INTEGRATED_BREAST_CANCER_PATHWAY, WP_INTEGRATED_BREAST_CANCER_PATHWAY GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_30MIN_DN, GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_30MIN_DN GSE45365_HEALTHY_VS_MCMV_INFECTION_CD11B_DC_IFNAR_KO_DN, GSE45365_HEALTHY_VS_MCMV_INFECTION_CD11B_DC_IFNAR_KO_DN GOBP_REGULATION_OF_MITOTIC_SISTER_CHROMATID_SEGREGATION, GOBP_REGULATION_OF_MITOTIC_SISTER_CHROMATID_SEGREGATION GSE37301_MULTIPOTENT_PROGENITOR_VS_PRO_BCELL_UP, GSE37301_MULTIPOTENT_PROGENITOR_VS_PRO_BCELL_UP GOBP_SISTER_CHROMATID_COHESION, GOBP_SISTER_CHROMATID_COHESION GOCC_CONDENSED_NUCLEAR_CHROMOSOME, GOCC_CONDENSED_NUCLEAR_CHROMOSOME GSE360_L_MAJOR_VS_M_TUBERCULOSIS_MAC_DN, GSE360_L_MAJOR_VS_M_TUBERCULOSIS_MAC_DN GSE7509_UNSTIM_VS_FCGRIIB_STIM_DC_UP, GSE7509_UNSTIM_VS_FCGRIIB_STIM_DC_UP GSE3982_DC_VS_BCELL_DN, GSE3982_DC_VS_BCELL_DN GOBP_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION MIR504_3P, MIR504_3P GOBP_CHROMOSOME_LOCALIZATION, GOBP_CHROMOSOME_LOCALIZATION GOBP_METAPHASE_PLATE_CONGRESSION, GOBP_METAPHASE_PLATE_CONGRESSION GSE21670_IL6_VS_TGFB_AND_IL6_TREATED_CD4_TCELL_DN, GSE21670_IL6_VS_TGFB_AND_IL6_TREATED_CD4_TCELL_DN GSE3982_EFF_MEMORY_VS_CENT_MEMORY_CD4_TCELL_UP, GSE3982_EFF_MEMORY_VS_CENT_MEMORY_CD4_TCELL_UP KEGG_PROGESTERONE_MEDIATED_OOCYTE_MATURATION, KEGG_PROGESTERONE_MEDIATED_OOCYTE_MATURATION GOBP_PROTEIN_DEPOLYMERIZATION, GOBP_PROTEIN_DEPOLYMERIZATION MIR5089_5P, MIR5089_5P KIM_WT1_TARGETS_8HR_DN, KIM_WT1_TARGETS_8HR_DN GOMF_MICROTUBULE_MOTOR_ACTIVITY, GOMF_MICROTUBULE_MOTOR_ACTIVITY MIR320A_5P, MIR320A_5P GSE29164_UNTREATED_VS_CD8_TCELL_TREATED_MELANOMA_DAY7_DN, GSE29164_UNTREATED_VS_CD8_TCELL_TREATED_MELANOMA_DAY7_DN GOBP_MITOTIC_METAPHASE_PLATE_CONGRESSION, GOBP_MITOTIC_METAPHASE_PLATE_CONGRESSION GSE27786_NKTCCELL_VS_ERYTHROBLAST_DN, GSE27786_NKTCCELL_VS_ERYTHROBLAST_DN GSE3720_LPS_VS_PMA_STIM_VD1_GAMMADELTA_TCELL_UP, GSE3720_LPS_VS_PMA_STIM_VD1_GAMMADELTA_TCELL_UP GOBP_NUCLEAR_ENVELOPE_ORGANIZATION, GOBP_NUCLEAR_ENVELOPE_ORGANIZATION GOBP_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION, GOBP_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION GSE10211_UV_INACT_SENDAI_VS_LIVE_SENDAI_VIRUS_TRACHEAL_EPITHELIAL_CELLS_DN, GSE10211_UV_INACT_SENDAI_VS_LIVE_SENDAI_VIRUS_TRACHEAL_EPITHELIAL_CELLS_DN GSE6259_FLT3L_INDUCED_VS_WT_SPLENIC_DC_33D1_POS_DN, GSE6259_FLT3L_INDUCED_VS_WT_SPLENIC_DC_33D1_POS_DN REACTOME_FACTORS_INVOLVED_IN_MEGAKARYOCYTE_DEVELOPMENT_AND_PLATELET_PRODUCTION, REACTOME_FACTORS_INVOLVED_IN_MEGAKARYOCYTE_DEVELOPMENT_AND_PLATELET_PRODUCTION GOBP_POSITIVE_REGULATION_OF_INTERFERON_GAMMA_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERFERON_GAMMA_PRODUCTION HP_DEPRESSED_NASAL_RIDGE, HP_DEPRESSED_NASAL_RIDGE HP_GASTROINTESTINAL_CARCINOMA, HP_GASTROINTESTINAL_CARCINOMA GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP, GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP GOBP_CHROMATIN_SILENCING, GOBP_CHROMATIN_SILENCING GSE19888_ADENOSINE_A3R_INH_VS_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_VS_TCELL_MEMBRANES_ACT_MAST_CELL_DN LEE_LIVER_CANCER_ACOX1_UP, LEE_LIVER_CANCER_ACOX1_UP MODULE_189, MODULE_189 GOCC_REPLISOME, GOCC_REPLISOME GOBP_MITOTIC_SISTER_CHROMATID_COHESION, GOBP_MITOTIC_SISTER_CHROMATID_COHESION GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_DN, GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_DN MODULE_127, MODULE_127 DESCARTES_FETAL_MUSCLE_ERYTHROBLASTS, DESCARTES_FETAL_MUSCLE_ERYTHROBLASTS GOMF_DNA_POLYMERASE_ACTIVITY, GOMF_DNA_POLYMERASE_ACTIVITY YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_15, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_15 REACTOME_MITOTIC_TELOPHASE_CYTOKINESIS, REACTOME_MITOTIC_TELOPHASE_CYTOKINESIS HP_ADENOCARCINOMA_OF_THE_LARGE_INTESTINE, HP_ADENOCARCINOMA_OF_THE_LARGE_INTESTINE WP_PRIMARY_FOCAL_SEGMENTAL_GLOMERULOSCLEROSIS_FSGS, WP_PRIMARY_FOCAL_SEGMENTAL_GLOMERULOSCLEROSIS_FSGS HUI_MAPK14_TARGETS_UP, HUI_MAPK14_TARGETS_UP HOFMANN_MYELODYSPLASTIC_SYNDROM_LOW_RISK_DN, HOFMANN_MYELODYSPLASTIC_SYNDROM_LOW_RISK_DN GOBP_POSITIVE_REGULATION_OF_MITOTIC_NUCLEAR_DIVISION, GOBP_POSITIVE_REGULATION_OF_MITOTIC_NUCLEAR_DIVISION GOBP_NUCLEOLAR_LARGE_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GOBP_NUCLEOLAR_LARGE_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_I REACTOME_MYOGENESIS, REACTOME_MYOGENESIS GOMF_DNA_DIRECTED_DNA_POLYMERASE_ACTIVITY, GOMF_DNA_DIRECTED_DNA_POLYMERASE_ACTIVITY GOBP_ENDOCRINE_SYSTEM_DEVELOPMENT, GOBP_ENDOCRINE_SYSTEM_DEVELOPMENT MODULE_203, MODULE_203 GOBP_SYNAPSE_MATURATION, GOBP_SYNAPSE_MATURATION MODULE_89, MODULE_89 HP_SUPERNUMERARY_NIPPLE, HP_SUPERNUMERARY_NIPPLE GOBP_POSITIVE_REGULATION_OF_NUCLEAR_DIVISION, GOBP_POSITIVE_REGULATION_OF_NUCLEAR_DIVISION MSX1_01, MSX1_01 SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX1_UP, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX1_UP GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_OF_NUCLEOLAR_LARGE_RRNA_BY_RNA_POLYMERASE_I, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_OF_NUCLEOLAR_LARGE_RRNA_BY_RNA_POLYMERASE_I GOCC_CELL_CORTX_REGION, GOCC_CELL_CORTX_REGION WP_ARYL_HYDROCARBON_RECEPTOR_NETPATH, WP_ARYL_HYDROCARBON_RECEPTOR_NETPATH GSE9988_LPS_VS_LOW_LPS_MONOCYTE_UP, GSE9988_LPS_VS_LOW_LPS_MONOCYTE_UP GOCC_CHROMOCENTER, GOCC_CHROMOCENTER PRDM5_TARGET_GENES, PRDM5_TARGET_GENES MODULE_90, MODULE_90 GSE10147_IL3_AND_HIVP17_VS_IL3_AND_CPG_STIM_PDC_UP, GSE10147_IL3_AND_HIVP17_VS_IL3_AND_CPG_STIM_PDC_UP GOBP_SPINDLE_MIDZONE_ASSEMBLY, GOBP_SPINDLE_MIDZONE_ASSEMBLY GSE13522_CTRL_VS_T_CRUZY_Y_STRAIN_INF_SKIN_BALBC_MOUSE_DN, GSE13522_CTRL_VS_T_CRUZY_Y_STRAIN_INF_SKIN_BALBC_MOUSE_DN TURASHVILI_BREAST_CARCINOMA_DUCTAL_VS_LOBULAR_UP, TURASHVILI_BREAST_CARCINOMA_DUCTAL_VS_LOBULAR_UP GOCC_SYNAPTONEMAL_STRUCTURE, GOCC_SYNAPTONEMAL_STRUCTURE GOBP_REGULATION_OF_TRANSCRIPTION_OF_NUCLEOLAR_LARGE_RRNA_BY_RNA_POLYMERASE_I, GOBP_REGULATION_OF_TRANSCRIPTION_OF_NUCLEOLAR_LARGE_RRNA_BY_RNA_POLYMERASE_I JACKSON_DNMT1_TARGETS_DN, JACKSON_DNMT1_TARGETS_DN GOMF_NADPLUS_KINASE_ACTIVITY, GOMF_NADPLUS_KINASE_ACTIVITY GOBP_NEGATIVE_REGULATION_OF_REPRODUCTIVE_PROCESS, GOBP_NEGATIVE_REGULATION_OF_REPRODUCTIVE_PROCESS HP_CONICAL_TOOTH, HP_CONICAL_TOOTH RIZ_ERYTHROID_DIFFERENTIATION_HEMGN, RIZ_ERYTHROID_DIFFERENTIATION_HEMGN GOBP_REGULATION_OF_SYNAPSE_MATURATION, GOBP_REGULATION_OF_SYNAPSE_MATURATION GOBP_NATURAL_KILLER_CELL_CYTOKINE_PRODUCTION, GOBP_NATURAL_KILLER_CELL_CYTOKINE_PRODUCTION BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_24HR_DN, BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_24HR_DN SANSOM_WNT_PATHWAY_REQUIRE_MYC, SANSOM_WNT_PATHWAY_REQUIRE_MYC GOBP_POSITIVE_REGULATION_OF_UBIQUITIN_PROTEIN_LIGASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_UBIQUITIN_PROTEIN_LIGASE_ACTIVITY GOMF_SPHINGOMYELIN_PHOSPHODIESTERASE_ACTIVITY, GOMF_SPHINGOMYELIN_PHOSPHODIESTERASE_ACTIVITY GOMF_BMP_RECEPTOR_BINDING, GOMF_BMP_RECEPTOR_BINDING GOBP_RESPONSE_TO_LEAD_ION, GOBP_RESPONSE_TO_LEAD_ION chr15q23, chr15q23 GOBP_NEGATIVE_REGULATION_OF_CELL_DIVISION, GOBP_NEGATIVE_REGULATION_OF_CELL_DIVISION GOBP_REGULATION_OF_CELL_AGING, GOBP_REGULATION_OF_CELL_AGING GOBP_OOGENESIS, GOBP_OOGENESIS HP_AGAMMAGLOBULINEMIA, HP_AGAMMAGLOBULINEMIA GOBP_POSITIVE_REGULATION_OF_RECEPTOR_BINDING, GOBP_POSITIVE_REGULATION_OF_RECEPTOR_BINDING GOBP_RESPONSE_TO_L_ASCORBIC_ACID, GOBP_RESPONSE_TO_L_ASCORBIC_ACID GOBP_LIPOXIN_METABOLIC_PROCESS, GOBP_LIPOXIN_METABOLIC_PROCESS GOBP_POSITIVE_REGULATION_OF_SYNAPSE_MATURATION, GOBP_POSITIVE_REGULATION_OF_SYNAPSE_MATURATION REACTOME_SYNTHESIS_OF_LIPOXINS_LX, REACTOME_SYNTHESIS_OF_LIPOXINS_LX GOBP_LEUKOTRIENE_B4_METABOLIC_PROCESS, GOBP_LEUKOTRIENE_B4_METABOLIC_PROCESS LU_TUMOR_VASCULATURE_DN, LU_TUMOR_VASCULATURE_DN GOBP_GLOMERULAR_EPITHELIAL_CELL_DEVELOPMENT, GOBP_GLOMERULAR_EPITHELIAL_CELL_DEVELOPMENT GOBP_ALPHA_LINOLENIC_ACID_METABOLIC_PROCESS, GOBP_ALPHA_LINOLENIC_ACID_METABOLIC_PROCESS REACTOME_ALPHA_LINOLENIC_OMEGA3_AND_LINOLEIC_OMEGA6_ACID_METABOLISM, REACTOME_ALPHA_LINOLENIC_OMEGA3_AND_LINOLEIC_OMEGA6_ACID_METABOLISM BUSSLINGER_GASTRIC_TUFT_CELLS, BUSSLINGER_GASTRIC_TUFT_CELLS TORCHIA_TARGETS_OF_EWSR1_FL11_FUSION_TOP20_DN, TORCHIA_TARGETS_OF_EWSR1_FL11_FUSION_TOP20_DN GOBP_NATURAL_KILLER_CELL_MEDIATED_IMMUNE_RESPONSE_TO_TUMOR_CELL, GOBP_NATURAL_KILLER_CELL_MEDIATED_IMMUNE_RESPONSE_TO_TUMOR_CELL HP_ABNORMALITY_OF_EPIDERMAL_MORPHOLOGY, HP_ABNORMALITY_OF_EPIDERMAL_MORPHOLOGY GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_MEDIATED_IMMUNE_RESPONSE_TO_TUMOR_CELL, GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_MEDIATED_IMMUNE_RESPONSE_TO_TUMOR_CELL REACTOME_PKA_MEDIATED_PHOSPHORYLATION_OF_CREB, REACTOME_PKA_MEDIATED_PHOSPHORYLATION_OF_CREB LMNB2_TARGET_GENES, LMNB2_TARGET_GENES GOMF_EXTRACELLULAR_MATRIX_CONSTITUENT_CONFERRING_ELASTICITY, GOMF_EXTRACELLULAR_MATRIX_CONSTITUENT_CONFERRING_ELASTICITY GOCC_LAMININ_COMPLEX, GOCC_LAMININ_COMPLEX GOBP_ASSEMBLY_OF_ACTOMYOSIN_APPARATUS_INVOLVED_IN_CYTOKINESIS, GOBP_ASSEMBLY_OF_ACTOMYOSIN_APPARATUS_INVOLVED_IN_CYTOKINESIS GOBP_TONGUE_DEVELOPMENT, GOBP_TONGUE_DEVELOPMENT DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_RESPIRATORY_HORIZONTA Basal_Cells, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_RESPIRATORY_HORIZONTA Basal_Cells GOBP_POSITIVE_REGULATION_OF_BLOOD_CIRCULATION, GOBP_POSITIVE_REGULATION_OF_BLOOD_CIRCULATION GOMF_CAMP_BINDING, GOMF_CAMP_BINDING MIR517A_3P_MIR517B_3P_MIR517C_3P, MIR517A_3P_MIR517B_3P_MIR517C_3P HP_ABNORMAL_URINE_POTASSIUM_CONCENTRATION, HP_ABNORMAL_URINE_POTASSIUM_CONCENTRATION KEGG_ALPHA_LINOLENIC_ACID_METABOLISM, KEGG_ALPHA_LINOLENIC_ACID_METABOLISM MNX1_TARGET_GENES, MNX1_TARGET_GENES HP_ABSENT_NIPPLE, HP_ABSENT_NIPPLE	
ND_IL10_STIM_IL10_KO_MACROPHAGE_UP, GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_UP		