

GOBP_TRANSLATIONAL_INITIATION,GOBP_TRANSLATIONAL_INITIATION
GOCC_RIBOSOMAL_SUBUNIT,GOCC_RIBOSOMAL_SUBUNIT
RUBENSTEIN_SKELETAL_MUSCLE_T_CELL,S_RUBENSTEIN_SKELETAL_MUSCLE_T_CELL
GSE26301,THL_VS_TH17_RESTIMULATED_DAY15_POST_POLARIZATION_DN,GSE26301,THL_VS_TH17_RESTIMULATED_DAY15_POST_POLARIZATION_DN
GSE13547,CTRL_VS_ANTI_ILGM_STIM_BCELL_2H_DN,GSE13547,CTRL_VS_ANTI_ILGM_STIM_BCELL_2H_DN
GSE10239,NAIVE_VS_KLRG1GINT_EFF_CD8_TCELL_UP,GSE10239,NAIVE_VS_KLRG1GINT_EFF_CD8_TCELL_UP
GSE42088,UNINF_VS_LEISHMANIA_INF_DC_2H_DN,GSE42088,UNINF_VS_LEISHMANIA_INF_DC_2H_DN
GOMF_STRUCTURAL_CONSTITUENT_OF_RIBOSOME,GOMF_STRUCTURAL_CONSTITUENT_OF_RIBOSOME
GSE22886,NAIVE_TCELL_VS_DC_UP,GSE22886,NAIVE_TCELL_VS_DC_UP
GSE9650,EFFECTOR_VS_MEMORY_CD8_TCELL_DN,GSE9650,EFFECTOR_VS_MEMORY_CD8_TCELL_DN
GSE24574,BC1A_HRGT,THL_VS_TH17_RESTIMULATED_DAY15_POST_POLARIZATION_DN,GSE24574,BC1A_HRGT,THL_VS_TH17_RESTIMULATED_DAY15_POST_POLARIZATION_DN
GSE22886,IGC6A_MEMORY_BCELL_VS_BM_PLASMA_CELL_UP,GSE22886,IGC6A_MEMORY_BCELL_VS_BM_PLASMA_CELL_UP
GSE28726,ACT_CD4_TCELL_VS_ACT_VA2NEG_NKTCCELL_DN,GSE28726,ACT_CD4_TCELL_VS_ACT_VA2NEG_NKTCCELL_DN
GSE3720,UNSTIM_VS_PMA_STIM_VD1_GAMMADELTA_TCELL_UP,GSE3720,UNSTIM_VS_PMA_STIM_VD1_GAMMADELTA_TCELL_UP
FAN_OVARY_CLU_XBP1_SELK_HIGH_STROMAL_CELL,FAN_OVARY_CLU_XBP1_SELK_HIGH_STROMAL_CELL
GSE25088,WT_VS_STAT16_KO_MACROPHAGE_ROSIGLITAZONE_STIM_UP,GSE25088,WT_VS_STAT16_KO_MACROPHAGE_ROSIGLITAZONE_STIM_UP
REACTOME_REGULATION_OF_EXPRESSION_OF_SLITS_AND_ROBO4S,REACTOME_REGULATION_OF_EXPRESSION_OF_SLITS_AND_ROBO4S
GSE34156,NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_DN,GSE34156,NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_DN
GNP2_EIF356,GNP2_EIF356
GSE16385,IL4_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP,GSE16385,IL4_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP
GSE5099,DAY3_VS_DAY7_MCSF_TREATED_MACROPHAGE_UP,GSE5099,DAY3_VS_DAY7_MCSF_TREATED_MACROPHAGE_UP
GSE360,HIGH_DOSE_B_MALAYI_VS_M_TUBERCULOSIS_DC_UP,GSE360,HIGH_DOSE_B_MALAYI_VS_M_TUBERCULOSIS_DC_UP
GSE12366,PLASMA_CELL_VS_NAIVE_BCELL_DN,GSE12366,PLASMA_CELL_VS_NAIVE_BCELL_DN
GSE10239,NAIVE_VS_KLRG1GINT_EFF_CD8_TCELL_UP,GSE10239,NAIVE_VS_KLRG1GINT_EFF_CD8_TCELL_UP
HP_ABNORMAL_NEUTROPHIL_COUNT,HP_ABNORMAL_NEUTROPHIL_COUNT
GSE43955,10H_VS_60H_ACT_CD4_TCELL_WITH_TGFB_IL6_UP,GSE43955,10H_VS_60H_ACT_CD4_TCELL_WITH_TGFB_IL6_UP
GSE12366,PLASMA_CELL_VS_NAIVE_BCELL_DN,GSE12366,PLASMA_CELL_VS_NAIVE_BCELL_DN
GSE22886,IGM_MEMORY_BCELL_VS_BM_PLASMA_CELL_UP,GSE22886,IGM_MEMORY_BCELL_VS_BM_PLASMA_CELL_UP
GSE25123,WT_VS_PPARG_KO_MACROPHAGE_DN,GSE25123,WT_VS_PPARG_KO_MACROPHAGE_DN
GSE1184,UNTREATED_VS_CSF1_IN_MAC_UP,GSE1184,UNTREATED_VS_CSF1_IN_MAC_UP
GSE10239,NAIVE_VS_DAY15_TREG_CD8_TCELL_UP,GSE10239,NAIVE_VS_DAY15_TREG_CD8_TCELL_UP
GOCC_LARGE_RIBOSOMAL_SUBUNIT,GOCC_LARGE_RIBOSOMAL_SUBUNIT
GSE9650,NAIVE_VS_EFF_CD8_TCELL_UP,GSE9650,NAIVE_VS_EFF_CD8_TCELL_UP
GSE34156,NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_UP,GSE34156,NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_UP
GSE22886,NAIVE_CD4_TCELL_VS_48H_ACT_TH1_UP,GSE22886,NAIVE_CD4_TCELL_VS_48H_ACT_TH1_UP
GSE36392,EOSINOPHIL_VS_NEUTROPHIL_IL25_TREATED_LUNG_DN,GSE36392,EOSINOPHIL_VS_NEUTROPHIL_IL25_TREATED_LUNG_DN
RUVBL1_TARGET_GENES,RUVBL1_TARGET_GENES
GSE3720,UNSTIM_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP,GSE3720,UNSTIM_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP
KAECH_DAYS_EFF_VS_MEMORY_CD8_TCELL_DN,KAECH_DAYS_EFF_VS_MEMORY_CD8_TCELL_DN
GSE2196,HEALTHY_VS_OBESE_MOUSE_SKIN_GAMMADELTA_TCELL_DN,GSE2196,HEALTHY_VS_OBESE_MOUSE_SKIN_GAMMADELTA_TCELL_DN
GSE37301,HEMATOPOIETIC_STEM_CELL_VS_PRO_BCELL_DN,GSE37301,HEMATOPOIETIC_STEM_CELL_VS_PRO_BCELL_DN
GSE1925,CTRL_VS_IFNG_PRIMED_MACROPHAGE_3H_IFNG_STIM_UP,GSE1925,CTRL_VS_IFNG_PRIMED_MACROPHAGE_3H_IFNG_STIM_UP
GSE25088,WT_VS_STAT16_KO_MACROPHAGE_UP,GSE25088,WT_VS_STAT16_KO_MACROPHAGE_UP
GOLDRATH_EFF_VS_MEMORY_CD8_TCELL_UP,GOLDRATH_EFF_VS_MEMORY_CD8_TCELL_UP
QLPBMC_ZOSTAVAX_AGE_50_75YO_CORRELATED_WITH_CONTRACTION_OF_VZV_SPECIFIC_T_CELLS_PEAK_TO_28DAY_AT_IDV_NEGATIVE,QLPBMC_ZOSTAVAX_AGE_50_75YO_CORRELATED_WITH_CONTRACTION_OF_VZV_SPECIFIC_T_CELLS_PEAK_TO_28DAY_AT_IDV_NEGATIVE
GSE360,DC_VS_MAC1_DONOVANI_DN,GSE360,DC_VS_MAC1_DONOVANI_DN
GSE5099,MONOCYTE_VS_CLASSICAL_M1_MACROPHAGE_DN,GSE5099,MONOCYTE_VS_CLASSICAL_M1_MACROPHAGE_DN
KAECH_NAIVE_VS_DAYS_EFF_CD8_TCELL_UP,KAECH_NAIVE_VS_DAYS_EFF_CD8_TCELL_UP
GSE17721,LPS_VS_PAMCSK4_16H_BMDC_DN,GSE17721,LPS_VS_PAMCSK4_16H_BMDC_DN
GSE9960,HEALTHY_VS_CRAM_POS_SIPSIS_PBMIC_UP,GSE9960,HEALTHY_VS_CRAM_POS_SIPSIS_PBMIC_UP
GSE22886,NAIVE_CD4_TCELL_VS_48H_ACT_TH2_UP,GSE22886,NAIVE_CD4_TCELL_VS_48H_ACT_TH2_UP
GSE26156,DOUBLE_POSITIVE_VS_CD4_SINGLE_POSITIVE_THYMOCYTE_UP,GSE26156,DOUBLE_POSITIVE_VS_CD4_SINGLE_POSITIVE_THYMOCYTE_UP
GSE27786,LSK_VS_CD8_TCELL_DN,GSE27786,LSK_VS_CD8_TCELL_DN
GSE20727,H2O2_VS_ROS_INHIBITOR_TREATED_DC_DN,GSE20727,H2O2_VS_ROS_INHIBITOR_TREATED_DC_DN
GSE17721,CTRL_VS_PAMCSK4_24H_BMDC_UP,GSE17721,CTRL_VS_PAMCSK4_24H_BMDC_UP
GSE39,CDPPOS_VS_CDPPOS_DC_CULTURE_DN,GSE39,CDPPOS_VS_CDPPOS_DC_CULTURE_DN
GSE1400,TRANSLATED_PNA_VS_MRNA_4H_LPS_DC_DN,GSE1400,TRANSLATED_PNA_VS_MRNA_4H_LPS_DC_DN
GSE3720,LPS_VS_PMA_STIM_VD1_GAMMADELTA_TCELL_DN,GSE3720,LPS_VS_PMA_STIM_VD1_GAMMADELTA_TCELL_DN
GOBP_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM,GOBP_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM
GAVIN_FOXP3_TARGETS_CLUSTER_T7,GAVIN_FOXP3_TARGETS_CLUSTER_T7
GSE5099,CLASSICAL_M1_VS_ALTERNATIVE_M2_MACROPHAGE_UP,GSE5099,CLASSICAL_M1_VS_ALTERNATIVE_M2_MACROPHAGE_UP
QLPBMC_ZOSTAVAX_AGE_50_75YO_CORRELATED_WITH_EXPANSION_OF_VZV_SPECIFIC_T_CELLS_TO_PEAK_AT_IDV_NEGATIVE,QLPBMC_ZOSTAVAX_AGE_50_75YO_CORRELATED_WITH_EXPANSION_OF_VZV_SPECIFIC_T_CELLS_TO_PEAK_AT_IDV_NEGATIVE
ACEVEDO_NORMAL_TSSUE_ADJACENT_TO_LIVER_TUMOR_UP,ACEVEDO_NORMAL_TSSUE_ADJACENT_TO_LIVER_TUMOR_UP
GOBP_PROTEIN_STABILIZATION,GOBP_PROTEIN_STABILIZATION
GSE26488,WT_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN,GSE26488,WT_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN
GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM,GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM
REACTOME_SELENOAMINO_ACID_METABOLISM,REACTOME_SELENOAMINO_ACID_METABOLISM
REACTOME_RESPONSE_OF_HIF2A_K1_GCN2_TO_AMINO_ACID_DEFICIENCY,REACTOME_RESPONSE_OF_HIF2A_K1_GCN2_TO_AMINO_ACID_DEFICIENCY
GSE4685,NAIVE_CD4_TCELL_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN,GSE4685,NAIVE_CD4_TCELL_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GOBP_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS,GOBP_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_CELLULAR_AMIDE_METABOLIC_PROCESS,GOBP_POSITIVE_REGULATION_OF_CELLULAR_AMIDE_METABOLIC_PROCESS
GSE34156,UNTREATED_VS_24H_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_DN,GSE34156,UNTREATED_VS_24H_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_DN
GSE37416,12H_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN,GSE37416,12H_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GSE32034,UNTREATED_VS_ROSIGLITAZONE_TREATED_1Y6C_HIGH_MONOCYTE_DN,GSE32034,UNTREATED_VS_ROSIGLITAZONE_TREATED_1Y6C_HIGH_MONOCYTE_DN
MYBL1_TARGET_GENES,MYBL1_TARGET_GENES
GSE1033,1H_VS_24H_POLYIC_STIM_DC_DN,GSE1033,1H_VS_24H_POLYIC_STIM_DC_DN
GSE34205,HEALTHY_VS_FLU_INF_INFANT_PBMIC_UP,GSE34205,HEALTHY_VS_FLU_INF_INFANT_PBMIC_UP
HP_ANEMIA_OF_INADEQUATE_PRODUCTION,HP_ANEMIA_OF_INADEQUATE_PRODUCTION
GSE29949,CD8_POS_CD8_SPLEEN_VS_MONOCYTE_BONE_MARROW_DN,GSE29949,CD8_POS_CD8_SPLEEN_VS_MONOCYTE_BONE_MARROW_DN
GSE1460,DP_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_DN,GSE1460,DP_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_DN
GOMB_TRANSLATION_REGULATOR_ACTIVITY,GOMB_TRANSLATION_REGULATOR_ACTIVITY
GSE6269,HEALTHY_VS_FLU_INF_PBMIC_UP,GSE6269,HEALTHY_VS_FLU_INF_PBMIC_UP
REACTOME_EUKARYOTIC_TRANSLATION_ELONGATION,REACTOME_EUKARYOTIC_TRANSLATION_ELONGATION
GSE1925,3H_VS_24H_IFNG_STIM_MACROPHAGE_DN,GSE1925,3H_VS_24H_IFNG_STIM_MACROPHAGE_DN
KEGG_RIBOSOME,KEGG_RIBOSOME
GOMB_TRANSLATION_FACTOR_ACTIVITY_RNA_BINDING,GOMB_TRANSLATION_FACTOR_ACTIVITY_RNA_BINDING
MORF_JUND,MORF_JUND
GSE25087,TREG_VS_TCONV_ADULT_DN,GSE25087,TREG_VS_TCONV_ADULT_DN
GSE365,DUSP1_VS_WT_SPLENOCYTES_DN,GSE365,DUSP1_VS_WT_SPLENOCYTES_DN
GSE365,CTRL_VS_LPS_INJECTED_SPLENOCYTES_DN,GSE365,CTRL_VS_LPS_INJECTED_SPLENOCYTES_DN
ZZZ3_TARGET_GENES,ZZZ3_TARGET_GENES
GOBP_RIBOSOMAL_SMALL_SUBUNIT_BIOGENESIS,GOBP_RIBOSOMAL_SMALL_SUBUNIT_BIOGENESIS
GOBP_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE,GOBP_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE
JOHNSTONE_PARV1_TARGETS_2_UP,JOHNSTONE_PARV1_TARGETS_2_UP
GOBP_REGULATION_OF_TRANSLATIONAL_INITIATION,GOBP_REGULATION_OF_TRANSLATIONAL_INITIATION
HP_ABNORMALITY_OF_THE_FRONTAL_HAIRLINE,HP_ABNORMALITY_OF_THE_FRONTAL_HAIRLINE
GOBP_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS,GOBP_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
GOMB_TRANSLATION_REGULATOR_ACTIVITY_NUCLEIC_ACID_BINDING,GOMB_TRANSLATION_REGULATOR_ACTIVITY_NUCLEIC_ACID_BINDING
GSE36392,TYPE_2_MYELOID_VS_EOSINOPHIL_IL25_TREATED_LUNG_UP,GSE36392,TYPE_2_MYELOID_VS_EOSINOPHIL_IL25_TREATED_LUNG_UP
GSE17721,PAMCSK4_VS_GADQIUMOD_24H_BMDC_UP,GSE17721,PAMCSK4_VS_GADQIUMOD_24H_BMDC_UP
LAKE_ADULT_KIDNEY_C4_ENDOTHELIAL_CELLS_AEA_AND_DVR,LAKE_ADULT_KIDNEY_C4_ENDOTHELIAL_CELLS_AEA_AND_DVR
ZHENG_CORD_BLOOD_C7_PUTATIVE_LYMPHOID_PRIMED_MULTIPOTENT_PROGENTEROR_1,ZHENG_CORD_BLOOD_C7_PUTATIVE_LYMPHOID_PRIMED_MULTIPOTENT_PROGENTEROR_1
LAKE_ADULT_KIDNEY_C7_PROXIMAL_TUBULE_EPITHELIAL_CELLS_S3,LAKE_ADULT_KIDNEY_C7_PROXIMAL_TUBULE_EPITHELIAL_CELLS_S3
LAKE_ADULT_KIDNEY_C6_PROXIMAL_TUBULE_EPITHELIAL_CELLS_FIBRINOGEN_POS_S3,LAKE_ADULT_KIDNEY_C6_PROXIMAL_TUBULE_EPITHELIAL_CELLS_FIBRINOGEN_POS_S3
HP_RENAL_AGENESIS,HP_RENAL_AGENESIS
GSE13485,DAY3_VS_DAY7_YF17D_VACCINE_PBMIC_UP,GSE13485,DAY3_VS_DAY7_YF17D_VACCINE_PBMIC_UP
GSE22886,NAIVE_BCELL_VS_DC_UP,GSE22886,NAIVE_BCELL_VS_DC_UP
REACTOME_SIGNALING_BY_HEDGEHOG,REACTOME_SIGNALING_BY_HEDGEHOG
ZMYM2_TARGET_GENES,ZMYM2_TARGET_GENES
GSE3982,MAC_VS_CENT_MEMORY_CD4_TCELL_DN,GSE3982,MAC_VS_CENT_MEMORY_CD4_TCELL_DN
GSE25087,TREG_VS_TCONV_FETUS_DN,GSE25087,TREG_VS_TCONV_FETUS_DN
BILANGES_SERUM_AND_RAPAMYCIN_SENSITIVE_GENES,BILANGES_SERUM_AND_RAPAMYCIN_SENSITIVE_GENES
REACTOME_SIGNALING_BY_TGF_BETA_RECEPTOR_COMPLEX,REACTOME_SIGNALING_BY_TGF_BETA_RECEPTOR_COMPLEX
THEILGAARD_NEUTROPHIL_AT_SKIN_WOUND_UP,THEILGAARD_NEUTROPHIL_AT_SKIN_WOUND_UP
GSE21670,IL6_VS_TGFB_AND_IL4_TREATED_CD4_CONV_UP,GSE21670,IL6_VS_TGFB_AND_IL4_TREATED_CD4_CONV_UP
GSE24634,NAIVE_CD4_TCELL_VS_DAY10_IL4_CONV_TREG_UP,GSE24634,NAIVE_CD4_TCELL_VS_DAY10_IL4_CONV_TREG_UP
MULLIGHAN_NPM1_SIGNATURE_3_DN,MULLIGHAN_NPM1_SIGNATURE_3_DN
GSE2770,TGFB_AND_IL4_IL12_TREATED_ACT_CD4_TCELL_2H_UP,GSE2770,TGFB_AND_IL4_IL12_TREATED_ACT_CD4_TCELL_2H_UP
GSE6269,HEALTHY_VS_STAPH_AUREUS_INF_PBMIC_DN,GSE6269,HEALTHY_VS_STAPH_AUREUS_INF_PBMIC_DN
REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_DIFFERENTIATION_OF_HSCS,REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_DIFFERENTIATION_OF_HSCS
GSE1445,INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_UP,GSE1445,INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_UP
GSE9006,TYPE_1_DIABETES_AT_DX_VS_1MONTH_POST_DX_PBMIC_DN,GSE9006,TYPE_1_DIABETES_AT_DX_VS_1MONTH_POST_DX_PBMIC_DN
GCCNKCCTATNK_UNKNOWN,GCCNKCCTATNK_UNKNOWN
RBM15_TARGET_GENES,RBM15_TARGET_GENES
CHNG_MULTIPLE_MYELOMA_HYPERPLOID_UP,CHNG_MULTIPLE_MYELOMA_HYPERPLOID_UP
GSE1460,CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_DN,GSE1460,CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_DN
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_CONJUGATION_OR_REMOVAL,GOBP_NEGATIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_CONJUGATION_OR_REMOVAL
HP_LOW_ANTERIOR_HAIRLINE,HP_LOW_ANTERIOR_HAIRLINE
TBX1_TARGET_GENES,TBX1_TARGET_GENES
GCCNMSMYNTTG_UNKNOWN,GCCNMSMYNTTG_UNKNOWN
ARHGAP35_TARGET_GENES,ARHGAP35_TARGET_GENES
TIEN_INTESTINE_PROBIOTICS_6HR_UP,TIEN_INTESTINE_PROBIOTICS_6HR_UP
DIAZ_CHRONIC_MYELOGENOUS_LEUKEMIA_DN,DIAZ_CHRONIC_MYELOGENOUS_LEUKEMIA_DN
REACTOME_SIGNALING_BY_TGFB_FAMILY_MEMBERS,REACTOME_SIGNALING_BY_TGFB_FAMILY_MEMBERS
GSE2585,THYMIC_DC_VS_MTEC_UP,GSE2585,THYMIC_DC_VS_MTEC_UP
GOBP_MATURATION_OF_SSU_RRNA,GOBP_MATURATION_OF_SSU_RRNA
HP_WEBBED_NECK,HP_WEBBED_NECK
MULLIGHAN_NPM1_MUTATED_SIGNATURE_1_DN,MULLIGHAN_NPM1_MUTATED_SIGNATURE_1_DN
GOCC_SMALL_RIBOSOMAL_SUBUNIT,GOCC_SMALL_RIBOSOMAL_SUBUNIT
GRUTZMANN_PANCREATIC_CANCER_DN,GRUTZMANN_PANCREATIC_CANCER_DN
HP_THROMBOCYTOSIS,HP_THROMBOCYTOSIS
GSE22886,IGM_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_UP,GSE22886,IGM_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_UP
GOMB_TRANSLATION_INITIATION_FACTOR_ACTIVITY,GOMB_TRANSLATION_INITIATION_FACTOR_ACTIVITY
HP_CLEFT_LIP,HP_CLEFT_LIP
KEGG_ADHERENS_JUNCTION,KEGG_ADHERENS_JUNCTION
AIZARAN_LIVER_CL_NK1_NKT_CELLS_1,AIZARAN_LIVER_CL_NK1_NKT_CELLS_1
GSE10239,NAIVE_VS_MEMORY_CD8_TCELL_DN,GSE10239,NAIVE_VS_MEMORY_CD8_TCELL_DN
REACTOME_REGULATED_NECROSIS,REACTOME_REGULATED_NECROSIS
GSE2567,MPL_VS_MPL_AND_R848_STIM_BCELL_UP,GSE2567,MPL_VS_MPL_AND_R848_STIM_BCELL_UP
TRAVAGLINI_LUNG_CLUB_CELL,TRAVAGLINI_LUNG_CLUB_CELL
GSE13306,TREG_VS_TCONV_LAMINA_PROPRIA_DN,GSE13306,TREG_VS_TCONV_LAMINA_PROPRIA_DN
GSE589,WT_VS_IL10_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN,GSE589,WT_VS_IL10_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN
GSE20727,DNFB_ALLERGEN_TREATED_DC_UP,GSE20727,DNFB_ALLERGEN_TREATED_DC_UP
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS,GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS
GSE2405,HEAT_KILLED_VS_LIVE_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_24H_DN,GSE2405,HEAT_KILLED_VS_LIVE_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_24H_DN
GSE3920,JFN8_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP,GSE3920,JFN8_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP
GSE20727,CTRL_VS_DNFB_ALLERGEN_TREATED_DC_DN,GSE20727,CTRL_VS_DNFB_ALLERGEN_TREATED_DC_DN
GATGKMRGCG_UNKNOWN,GATGKMRGCG_UNKNOWN
REACTOME_E3_UBIQUITIN_LIGASES_UBIQUITINATE_TARGET_PROTEINS,REACTOME_E3_UBIQUITIN_LIGASES_UBIQUITINATE_TARGET_PROTEINS
LIL_NASOPHARYNGEAL_CARCINOMA_IL12_NASOPHARYNGEAL_CARCINOMA,IL12_NASOPHARYNGEAL_CARCINOMA
GOBP_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS,GOBP_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS
GSE24081,CONTROLLER_VS_PROGRESSOR_HIV_SPECIFIC_CD8_TCELL_UP,GSE24081,CONTROLLER_VS_PROGRESSOR_HIV_SPECIFIC_CD8_TCELL_UP
HP_ABNORMAL_RETICULOCYTE_MORPHOLOGY,HP_ABNORMAL_RETICULOCYTE_MORPHOLOGY
GOBP_CELL_DEATH_IN_RESPONSE_TO_OXIDATIVE_STRESS,GOBP_CELL_DEATH_IN_RESPONSE_TO_OXIDATIVE_STRESS
GOBP_NEGATIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CATABOLIC_PROCESS,GOBP_NEGATIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CATABOLIC_PROCESS
GSE1445,INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_UP,GSE1445,INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_UP
GSE10325,MYELOID_VS_LUPUS_MYELOID_UP,GSE10325,MYELOID_VS_LUPUS_MYELOID_UP
GOBP_RESPONSE_TO_HYDROGEN_PEROXIDE,GOBP_RESPONSE_TO_HYDROGEN_PEROXIDE
HP_HYDROPS_FETALIS,HP_HYDROPS_FETALIS
MODULE_192,MODULE_192
REACTOME_TGF_BETA_RECEPTOR_SIGNALING_ACTIVATES_SMADS,REACTOME_TGF_BETA_RECEPTOR_SIGNALING_ACTIVATES_SMADS
HP_ABNORMAL_SCAPULA_MORPHOLOGY,HP_ABNORMAL_SCAPULA_MORPHOLOGY
HP_MACROCYTIC_ANEMIA,HP_MACROCYTIC_ANEMIA
REACTOME_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEPTOR_NLR_SIGNALING_PATHWAYS,REACTOME_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEPTOR_NLR_SIGNALING_PATHWAYS
GSE3203,WT_VS_IFNARI_KO_INFLUENZA_INFECTED_IL_BCELL_DN,GSE3203,WT_VS_IFNARI_KO_INFLUENZA_INFECTED_IL_BCELL_DN
GOBP_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY,GOBP_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
HP_DUPLICATION_OF_THUMB_PHALANX,HP_DUPLICATION_OF_THUMB_PHALANX
GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_OXIDATIVE_STRESS,GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_OXIDATIVE_STRESS
HP_APLASIA_OF_THE_FINGERS,HP_APLASIA_OF_THE_FINGERS
REACTOME_AMYLOID_FIBER_FORMATION,REACTOME_AMYLOID_FIBER_FORMATION
WP_REGULATION_OF_ACTIN_CYTOSKELETON,WP_REGULATION_OF_ACTIN_CYTOSKELETON
REACTOME_RHO_GTPASES_ACTIVATE_PKNS,REACTOME_RHO_GTPASES_ACTIVATE_PKNS
CRGAARNNNNCGA_UNKNOWN,CRGAARNNNNCGA_UNKNOWN
GOBP_NEGATIVE_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS,GOBP_NEGATIVE_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
REACTOME_FCGAMMA_RECEPTOR_FCGR_DEPENDENT_PHAGOCYTOSIS,REACTOME_FCGAMMA_RECEPTOR_FCGR_DEPENDENT_PHAGOCYTOSIS
MODULE_115,MODULE_115
GOBP_MYD88_DEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY,GOBP_MYD88_DEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY
GOBP_FCR_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY,GOBP_FCR_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY
TONG_INTERACT_WITH_PTGI1,TONG_INTERACT_WITH_PTGI1
HP_RETICULOCYTOPENIA,HP_RETICULOCYTOPENIA
HP_MUTISM,HP_MUTISM
GOBP_NEGATIVE_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS,GOBP_NEGATIVE_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS
HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_HAND,HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_HAND
ZA_UNIPROT_QUM89_UNREVIEWED_TARGET_GENES,ZA_UNIPROT_QUM89_UNREVIEWED_TARGET_GENES
WP_G13_SIGNALING_PATHWAY,WP_G13_SIGNALING_PATHWAY
GOBP_VIRION_ASSEMBLY,GOBP_VIRION_ASSEMBLY
WANG_PROSTATE_CANCER_ANDROGEN_INDEPENDENT,WANG_PROSTATE_CANCER_ANDROGEN_INDEPENDENT
GSE10325,BCELL_VS_LUPUS_BCELL_UP,GSE10325,BCELL_VS_LUPUS_BCELL_UP
REACTOME_RIPK1_MEDIATED_REGULATED_NECROSIS,REACTOME_RIPK1_MEDIATED_REGULATED_NECROSIS
MIR3672,MIR3672
GOMB_TRANSLATION_INITIATION_FACTOR_BINDING,GOMB_TRANSLATION_INITIATION_FACTOR_BINDING
HP_ABSENT_THUMB,HP_ABSENT_THUMB
REACTOME_DISEASES_OF_CARBOHYDRATE_METABOLISM,REACTOME_DISEASES_OF_CARBOHYDRATE_METABOLISM
HY2_TARGET_GENES,HY2_TARGET_GENES
REACTOME_INTERLEUKIN_3_INTERLEUKIN_5_AND_GM-CSF_SIGNALING,REACTOME_INTERLEUKIN_3_INTERLEUKIN_5_AND_GM-CSF_SIGNALING
GNATENKO_PLATELET_SIGNATURE,GNATENKO_PLATELET_SIGNATURE
REACTOME_SIGNALING_BY_NOTCH1_FEST_DOMAIN_MUTANTS_IN_CANCER,REACTOME_SIGNALING_BY_NOTCH1_FEST_DOMAIN_MUTANTS_IN_CANCER
HP_AVASCULAR_NECROSIS,HP_AVASCULAR_NECROSIS
REACTOME_METALLOPROTEASE_DUBS,REACTOME_METALLOPROTEASE_DUBS
HP_THERMOKINESIS,HP_THERMOKINESIS
REACTOME_P75NTR_SIGNALS_VIA_NF_KB,REACTOME_P75NTR_SIGNALS_VIA_NF_KB
ANDERSEN_LIVER_CANCER_KRT19_UP,ANDERSEN_LIVER_CANCER_KRT19_UP
MAPK3_TARGET_GENES,MAPK3_TARGET_GENES
REACTOME_SEMAPHORIN_INTERACTIONS,REACTOME_SEMAPHORIN_INTERACTIONS
REACTOME_HSF1_DEPENDENT_TRANSACTIVATION,REACTOME_HSF1_DEPENDENT_TRANSACTIVATION
REACTOME_DHSF1_DEPENDENT_TRANSACTIVATION,REACTOME_DHSF1_DEPENDENT_TRANSACTIVATION
HP_NORMOCHROMIC_ANEMIA,HP_NORMOCHROMIC_ANEMIA
REACTOME_SIGNALING_BY_CSF3_G-CSF,REACTOME_SIGNALING_BY_CSF3_G-CSF
BILANGES_RAPAMYCIN_SENSITIVE_GENES,BILANGES_RAPAMYCIN_SENSITIVE_GENES
HP_PERSISTENCE_OF_HEMOGLOBIN_F,HP_PERSISTENCE_OF_HEMOGLOBIN_F
GSE18148,CBF8_KO_VS_WT_TREG_DN,GSE18148,CBF8_KO_VS_WT_TREG_DN
MIR6643_3P,MIR6643_3P
MODULE_29,MODULE_29
TIEN_INTESTINE_PROBIOTICS_2HR_UP,TIEN_INTESTINE_PROBIOTICS_2HR_UP
GSE13547,CTRL_VS_ANTI_ILGM_STIM_BCELL_2H_DN,GSE13547,CTRL_VS_ANTI_ILGM_STIM_BCELL_2H_DN
REACTOME_FLT3_SIGNALING_IN_DISEASE,REACTOME_FLT3_SIGNALING_IN_DISEASE
GSE13485,DAY7_VS_DAY21_YF17D_VACCINE_PBMIC_DN,GSE13485,DAY7_VS_DAY21_YF17D_VACCINE_PBMIC_DN
REACTOME_REGULATION_OF_TNFR1_SIGNALING,REACTOME_REGULATION_OF_TNFR1_SIGNALING
GOBP_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY,GOBP_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
REACTOME_UPTAKE_OF_BACTERIAL_TOXINS,REACTOME_UPTAKE_OF_BACTERIAL_TOXINS
GOBP_NEGATIVE_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS,GOBP_NEGATIVE_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
HP_PARTIAL_DUPLICATION_OF_THE_PHALANX_OF_HAND,HP_PARTIAL_DUPLICATION_OF_THE_PHALANX_OF_HAND
GSE46606,UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4MID_SORTED_BCELL_DN,GSE46606,UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4MID_SORTED_BCELL_DN
ZN777_TARGET_GENES,ZN777_TARGET_GENES
REACTOME_NOD1_2_SIGNALING_PATHWAY,REACTOME_NOD1_2_SIGNALING_PATHWAY
GOBP_RESPONSE_TO_NERVE_GROWTH_FACTOR,GOBP_RESPONSE_TO_NERVE_GROWTH_FACTOR
GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE,GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE
MIR6748_3P,MIR6748_3P
TRAVAGLINI_LUNG_CD8_MEMORY_EFFECTOR_T_CELL,TRAVAGLINI_LUNG_CD8_MEMORY_EFFECTOR_T_CELL
REACTOME_REGULATION_OF_SIGNALING_BY_CBL,REACTOME_REGULATION_OF_SIGNALING_BY_CBL
TERT_TARGET_GENES,TERT_TARGET_GENES
HP_IL12_ENDOGENOUS_PATHWAY,HP_IL12_ENDOGENOUS_PATHWAY
GSE1047,IL2_IL5_IL2_AND_HPV17_STIM_PDC_DN,GSE1047,IL2_IL5_IL2_AND_HPV17_STIM_PDC_DN
GOCC_CYTOSOLIC_SMALL_RIBOSOMAL_SUBUNIT,GOCC_CYTOSOLIC_SMALL_RIBOSOMAL_SUBUNIT
HP_INCREASED_MEAN_CORPUSCULAR_VOLUME,HP_INCREASED_MEAN_CORPUSCULAR_VOLUME
REACTOME_NEGATIVE_REGULATION_OF_MET_ACTIVITY,REACTOME_NEGATIVE_REGULATION_OF_MET_ACTIVITY
REACTOME_GLYCOGEN_METABOLISM,REACTOME_GLYCOGEN_METABOLISM
REACTOME_ACTIVATED_NOTCH1_TRANSMITS_SIGNAL_TO_THE_NUCLEUS,REACTOME_ACTIVATED_NOTCH1_TRANSMITS_SIGNAL_TO_THE_NUCLEUS
HP_SPRENGEL_ANOMALY,HP_SPRENGEL_ANOMALY
GOBP_CENTRAL_NERVOUS_SYSTEM_NEURON_AXONOGENESIS,GOBP_CENTRAL_NERVOUS_SYSTEM_NEURON_AXONOGENESIS
GOBP_RIBOSOMAL_SMALL_SUBUNIT_ASSEMBLY,GOBP_RIBOSOMAL_SMALL_SUBUNIT_ASSEMBLY
HP_SUPERNUMERARY_NIPPLE,HP_SUPERNUMERARY_NIPPLE
KEGG_VIRAL_MYOCARDITIS,KEGG_VIRAL_MYOCARDITIS
REACTOME_MAP3K8_TPL2_DEPENDENT_MAPK1_3_ACTIVATION,REACTOME_MAP3K8_TPL2_DEPENDENT_MAPK1_3_ACTIVATION
REACTOME_ADHERENS_JUNCTION_INTERACTIONS,REACTOME_ADHERENS_JUNCTIONS_INTERACTIONS
CDX1_TARGET_GENES,CDX1_TARGET_GENES
HP_FATIGABLE_WEAKNESS_OF_SWALLOWING_MUSCLES,HP_FATIGABLE_WEAKNESS_OF_SWALLOWING_MUSCLES
REACTOME_LISTERIA_MONOCYTOGENES_ENTRY_INTO_HOST_CELLS,REACTOME_LISTERIA_MONOCYTOGENES_ENTRY_INTO_HOST_CELLS
GSE10422,WT_VS_BAFF_TRANSGENIC_ILN_BCELL_DN,GSE10422,WT_VS_BAFF_TRANSGENIC_ILN_BCELL_DN
REACTOME_DOWNRREGULATION_OF_SMA2_3_SMA24_TRANSCRIPTIONAL_ACTIVITY,REACTOME_DOWNRREGULATION_OF_SMA2_3_SMA24_TRANSCRIPTIONAL_ACTIVITY
REACTOME_INACTIVATION_OF_CSF3_G-CSF_SIGNALING,REACTOME_INACTIVATION_OF_CSF3_G-CSF_SIGNALING
REACTOME_CELL_EXTRACELLULAR_MATRIX_INTERACTIONS,REACTOME_CELL_EXTRACELLULAR_MATRIX_INTERACTIONS
QLPBMC_ZOSTAVAX_AGE_50_75YO_CORRELATED_WITH_T_CELL_RESPONSES_ID_INFORMATIVE_OF_LONG_TERM_RESPONSES,QLPBMC_ZOSTAVAX_AGE_50_75YO_CORRELATED_WITH_T_CELL_RESPONSES_ID_INFORMATIVE_OF_LONG_TERM_RESPONSES
REACTOME_NOTCH1_INTRACELLULAR_DOMAIN_REGULATES_TRANSCRIPTION,REACTOME_NOTCH1_INTRACELLULAR_DOMAIN_REGULATES_TRANSCRIPTION
REACTOME_P75NTR_RECRUITS_SIGNALLING_COMPLEXES,REACTOME_P75NTR_RECRUITS_SIGNALLING_COMPLEXES
REACTOME_IKK_COMPLEX_RECRUITMENT_MEDIATED_BY_RIP1,REACTOME_IKK_COMPLEX_RECRUITMENT_MEDIATED_BY_RIP1
ZHAN_MULTIPLE_MYELOMA_HP_UP,ZHAN_MULTIPLE_MYELOMA_HP_UP
ABE_INNER_EAR,ABE_INNER_EAR
LEE_LIVER_CANCER_MYC_TGFA_UP,LEE_LIVER_CANCER_MYC_TGFA_UP
GOBP_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEPTOR_SIGNALING_PATHWAY,GOBP_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEPTOR_SIGNALING_PATHWAY
REACTOME_OVARIAN_TUMOR_DOMAIN_PROTEASES,REACTOME_OVARIAN_TUMOR_DOMAIN_PROTEASES
REACTOME_SIGNALING_BY_ERBB4,REACTOME_SIGNALING_BY_ERBB4
REACTOME_NOTCH2_ACTIVATION_AND_TRANSMISSION_OF_SIGNAL_TO_THE_NUCLEUS,REACTOME_NOTCH2_ACTIVATION_AND_TRANSMISSION_OF_SIGNAL_TO_THE_NUCLEUS
HP_IMPAIRED_VIBRYON_SENSATION_IN_THE_LOWER_LIMBS,HP_IMPAIRED_VIBRYON_SENSATION_IN_THE_LOWER_LIMBS
GOBP_MYD88_INDEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY,GOBP_MYD88_INDEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY
HAY_BONE_MARROW_CD34_POS_HSC,HAY_BONE_MARROW_CD34_POS_HSC
GSE43955,1H_VS_42H_ACT_CD4_TCELL_DN,GSE43955,1H_VS_42H_ACT_CD4_TCELL_DN
CCANNAGRGKGC_UNKNOWN,CCANNAGRGKGC_UNKNOWN
REACTOME_NRF1_SIGNALS_CELL_DEATH_FROM_THE_NUCLEUS,REACTOME_NRF1_SIGNALS_CELL_DEATH_FROM_THE_NUCLEUS
GOBP_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS,GOBP_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS
REACTOME_IL12B_MEDIATED_ENTRY_OF_LISTERIA_MONOCYTOGENES_INTO_HOST_CELL,REACTOME_IL12B_MEDIATED_ENTRY_OF_LISTERIA_MONOCYTOGENES_INTO_HOST_CELL
REACTOME_REGULATION_OF_INNATE_IMMUNE_RESPONSES_TO_CYTOSOLIC_DNA,REACTOME_REGULATION_OF_INNATE_IMMUNE_RESPONSES_TO_CYTOSOLIC_DNA
GOBP_NEGATIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_CELL_DEATH,GOBP_NEGATIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_CELL_DEATH
MTHFD1_TARGET_GENES,MTHFD1_TARGET_GENES
HP_PROMINENT_METOPIC RIDGE,HP_PROMINENT_METOPIC RIDGE
GOBP_POSITIVE_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS,GOBP_POSITIVE_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS
REACTOME_NEGATIVE_REGULATION_OF_FLT3,REACTOME_NEGATIVE_REGULATION_OF_FLT3
REACTOME_GLYCOGEN_SYNTHESIS,REACTOME_GLYCOGEN_SYNTHESIS
HP_ABNORMALITY_OF_THE_METOPIC_SUTURE,HP_ABNORMALITY_OF_THE_METOPIC_SUTURE
REACTOME_ALPHA_PROTEIN_KINASE_1_SIGNALING_PATHWAY,REACTOME_ALPHA_PROTEIN_KINASE_1_SIGNALING_PATHWAY
REACTOME_IRAK1_RECRUITS_IKK_COMPLEX,REACTOME_IRAK1_RECRUITS_IKK_COMPLEX
STN1_TARGET_GENES,STN1_TARGET_GENES
REACTOME_REGULATION_OF_BACH1_ACTIVITY,REACTOME_REGULATION_OF_BACH1_ACTIVITY
REACTOME_SIGNALING_BY_NOTCH2,REACTOME_SIGNALING_BY_NOTCH2
REACTOME_GLYCOGEN_STORAGE_DISEASES,REACTOME_GLYCOGEN_STORAGE_DISEASES
REACTOME_NOTCH4_ACTIVATION_AND_TRANSMISSION_OF_SIGNAL_TO_THE_NUCLEUS,REACTOME_NOTCH4_ACTIVATION_AND_TRANSMISSION_OF_SIGNAL_TO_THE_NUCLEUS
REACTOME_BUDDING_AND_MATURATION_OF_HIV_VIRION,REACTOME_BUDDING_AND_MATURATION_OF_HIV_VIRION
REACTOME_SIGNALING_BY_EGR1_IN_CANCER,REACTOME_SIGNALING_BY_EGR1_IN_CANCER
REACTOME_CONSTITUTIVE_SIGNALING_BY_LIGAND_RESPONSEIVE_EGFR_CANCER_VARIANTS,REACTOME_CONSTITUTIVE_SIGNALING_BY_LIGAND_RESPONSEIVE_EGFR_CANCER_VARIANTS
REACTOME_GAP_JUNCTION_TRAFFICKING_AND_REGULATION,REACTOME_GAP_JUNCTION_TRAFFICKING_AND_REGULATION
GOBP_NEGATIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY,GOBP_NEGATIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
VERRECCHIA_RESPONSE_TO_TGFB1_C3,VERRECCHIA_RESPONSE_TO_TGFB1_C3
KEGG_GAP_JUNCTION,KEGG_GAP_JUNCTION
HP_LOWER_LIMB_METAPHYSICAL_IRREGULARITY,HP_LOWER_LIMB_METAPHYSICAL_IRREGULARITY
GOBP_REGULATION_OF_TRANSLATIONAL_INITIATION_BY_EIF2_ALPHA_PHOSPHORYLATION,GOBP_REGULATION_OF_TRANSLATIONAL_INITIATION_BY_EIF2_ALPHA_PHOSPHORYLATION
REACTOME_SEMA3A_PAK_DEPENDENT_AXON_REPULSION,REACTOME_SEMA3A_PAK_DEPENDENT_AXON_REPULSION
REACTOME_TGFB_BETA_RECEPTOR_SIGNALING_IN_EMT_EPITHELIAL_TO_MESENCHYMAL_TRANSITION,REACTOME_TGFB_BETA_RECEPTOR_SIGNALING_IN_EMT_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
GOMB_STRUCTURAL_CONSTITUENT_OF_SYNAPE,GOMB_STRUCTURAL_CONSTITUENT_OF_SYNAPE
REACTOME_SIGNALING_BY_FGFR1,REACTOME_SIGNALING_BY_FGFR1
GOBP_NEGATIVE_REGULATION_OF_CENTROSOME_DUPLICATION,GOBP_NEGATIVE_REGULATION_OF_CENTROSOME_DUPLICATION
REACTOME_RHO_GTPASES_ACTIVATE_RCKOS,REACTOME_RHO_GTPASES_ACTIVATE_RCKOS
REACTOME_SIGNALING_BY_FGFR3,REACTOME_SIGNALING_BY_FGFR3
GOBP_TELOMERE_ASSEMBLY,GOBP_TELOMERE_ASSEMBLY
REACTOME_TICAM1_RIP1_MEDIATED_IKK_COMPLEX_RECRUITMENT,REACTOME_TICAM1_RIP1_MEDIATED_IKK_COMPLEX_RECRUITMENT
GOBP_POSTSYNAPTIC_CYTOSKELETON_ORGANIZATION,GOBP_POSTSYNAPTIC_CYTOSKELETON_ORGANIZATION
GOBP_AUTOPHAGIC_CELL_DEATH,GOBP_AUTOPHAGIC_CELL_DEATH
REACTOME_IL1A_MEDIATED_ENTRY_OF_LISTERIA_MONOCYTOGENES_INTO_HOST_CELLS,REACTOME_IL1A_MEDIATED_ENTRY_OF_LISTERIA_MONOCYTOGENES_INTO_HOST_CELLS
PIONTEK_PKD1_TARGETS_DN,PIONTEK_PKD1_TARGETS_DN
GOBP_REGULATION_OF_NITRIC_OXIDE_METABOLIC_PROCESS,GOBP_REGULATION_OF_NITRIC_OXIDE_METABOLIC_PROCESS
REACTOME_JNK_CJUN_KINASES_PHOSPHORYLATION_AND_ACTIVATION_MEDIATED_BY_ACTIVATED_HUMAN_TAK1,REACTOME_JNK_CJUN_KINASES_PHOSPHORYLATION_AND_ACTIVATION_MEDIATED_BY_ACTIVATED_HUMAN_TAK1
GOBP_POSITIVE_REGULATION_OF_EMBRYONIC_DEVELOPMENT,GOBP_POSITIVE_REGULATION_OF_EMBRYONIC_DEVELOPMENT
REACTOME_SIGNALING_BY_NOTCH1_HD_DOMAIN_MUTANTS_IN_CANCER,REACTOME_SIGNALING_BY_NOTCH1_HD_DOMAIN_MUTANTS_IN_CANCER
REACTOME_TRAF6_MEDIATED_INDUCTION_OF_TAK1_COMPLEX_WITHIN_TLR4_COMPLEX,REACTOME_TRAF6_MEDIATED_INDUCTION_OF_TAK1_COMPLEX_WITHIN_TLR4_COMPLEX