

vs_NEUTROPHIL_UP, GSE3982_DC_vs_NEUTROPHIL_UP

GSE3982_NEUTROPHIL_VS_TH2_DN, GSE3982_NEUTROPHIL_VS_TH2_DN
GSE2286_DAY1_VS_DAY7_MONOCYTE_IN_CULTURE_DN, GSE2286_DAY1_VS_DAY7_MONOCYTE_IN_CULTURE_DN
GSE3982_NEUTROPHIL_VS_BCELL_DN, GSE3982_NEUTROPHIL_VS_BCELL_DN
GSE1112_OT1_VS_HY_CD8AB_THYMOCTE_RTQC_CULTURE_DN, GSE1112_OT1_VS_HY_CD8AB_THYMOCTE_RTQC_CULTURE_DN
GSE20152_SPHK1_VS_WT_HTNFA_OVEREXPRESS_ANGLE_DN, GSE20152_SPHK1_KO_VS_WT_HTNFA_OVEREXPRESS_ANGLE_DN
GSE3982_DC_VS_NEUTROPHIL_LPS_STIM_UP, GSE3982_DC_VS_NEUTROPHIL_LPS_STIM_UP
GSE2286_DAY0_VS_DAY7_MONOCYTE_IN_CULTURE_DN, GSE2286_DAY0_VS_DAY7_MONOCYTE_IN_CULTURE_DN
GSE3982_EOSINOPHIL_VS_DC_DN, GSE3982_EOSINOPHIL_VS_DC_DN
GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP
GSE360_CTRL_VS_M_TUBERCULOSIS_DC_UP, GSE360_CTRL_VS_M_TUBERCULOSIS_DC_UP
GSE2648_PRL_VS_DAY7_FLU_VACCINE_PDC_DN, GSE2648_PRL_VS_DAY7_FLU_VACCINE_PDC_DN
GSE3982_NEUTROPHIL_VS_CENT_MEMORY_CD4_TCELL_DN, GSE3982_NEUTROPHIL_VS_CENT_MEMORY_CD4_TCELL_DN
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP
GSE3982_MAC_VS_NKCELL_UP, GSE3982_MAC_VS_NKCELL_UP
GSE14769_20MIN_VS_360MIN_LPS_BMDM_UP, GSE14769_20MIN_VS_360MIN_LPS_BMDM_UP
HALLMARK_FATTY_ACID_METABOLISM, HALLMARK_FATTY_ACID_METABOLISM
GSE3982_DC_VS_NKCELL_UP, GSE3982_DC_VS_NKCELL_UP
GSE13493_DP_VS_CD8POS_THYMOCTE_DN, GSE13493_DP_VS_CD8POS_THYMOCTE_DN
GSE4606_IRF4_KO_VS_WT_CD40L_IL2_IL5_3DAY_STIMULATED_BCELL_UP, GSE4606_IRF4_KO_VS_WT_CD40L_IL2_IL5_3DAY_STIMULATED_BCELL_UP
GSE25123_CTRL_VS_IL4_STIM_MACROPHAGE_DN, GSE25123_CTRL_VS_IL4_STIM_MACROPHAGE_DN
GSE3982_EOSINOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN, GSE3982_EOSINOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN
GSE1685_MONOCYTE_VS_12H_ROSILITAZONE_TREATED_MACROPHAGE_UP, GSE1685_MONOCYTE_VS_12H_ROSILITAZONE_TREATED_MACROPHAGE_UP
GSE2648_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMOCTE_DN, GSE2648_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMOCTE_DN
GSE3982_DC_VS_MAC_IL6_STIM_UP, GSE3982_DC_VS_MAC_IL6_STIM_UP
GSE2648_WT_VS_HDAC7_DELTAP_TG_OT2_THYMOCTE_WITH_PEPTIDE_INJECTION_DN, GSE2648_WT_VS_HDAC7_DELTAP_TG_OT2_THYMOCTE_WITH_PEPTIDE_INJECTION_DN
GSE23505_IL6_IL12_IL2_VS_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP, GSE23505_IL6_IL12_IL2_VS_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP
GSE10325_LUPUS_CD4_TCELL_VS_LUPUS_MYELOID_UP, GSE10325_LUPUS_CD4_TCELL_VS_LUPUS_MYELOID_UP
GSE1925_CTRL_VS_3H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_UP, GSE1925_CTRL_VS_3H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_UP
GSE1112_OT1_VS_HY_CD8AB_THYMOCTE_RTQC_CULTURE_UP, GSE1112_OT1_VS_HY_CD8AB_THYMOCTE_RTQC_CULTURE_UP
GSE360_CTRL_VS_L_MAJOR_DC_UP, GSE360_CTRL_VS_L_MAJOR_DC_UP
LIU_CNV1_TARGETS_UP, LIU_CNV1_TARGETS_UP
GSE1440_DP_THYMOCTE_VS_THYMIC_STROMAL_CELL_DN, GSE1440_DP_THYMOCTE_VS_THYMIC_STROMAL_CELL_DN
GSE24634_IL4_VS_CTRL_TREATED_NAIVE_CD4_TCELL_DAY3_UP, GSE24634_IL4_VS_CTRL_TREATED_NAIVE_CD4_TCELL_DAY3_UP
GSE3688_UNTREATED_VS_IL2_TREATED_TCELL_2H_DN, GSE3688_UNTREATED_VS_IL2_TREATED_TCELL_2H_DN
GSE6674_UNSTIM_VS_CPG_STIM_BCELL_DN, GSE6674_UNSTIM_VS_CPG_STIM_BCELL_DN
GSE43955_TH0_VS_TGFB_IL6_THI7_ACT_CD4_TCELL_60H_DN, GSE43955_TH0_VS_TGFB_IL6_THI7_ACT_CD4_TCELL_60H_DN
GSE1306_RA_VS_UNTREATED_MEM_CD4_TCELL_DN, GSE1306_RA_VS_UNTREATED_MEM_CD4_TCELL_DN
MIR4727_5P, MIR4727_5P
GSE22045_TREG_VS_TCONV_DN, GSE22045_TREG_VS_TCONV_DN
AIZARANI_LIVER_C4_EPCAM_POS_BILE_DUCT_CELLS_1, AIZARANI_LIVER_C4_EPCAM_POS_BILE_DUCT_CELLS_1
GSE360_DC_VS_MAC_T_GONDIL_UP, GSE360_DC_VS_MAC_T_GONDIL_UP
KAECH_DAY8_EFF_VS_DAY15_EFF_CD8_TCELL_DN, KAECH_DAY8_EFF_VS_DAY15_EFF_CD8_TCELL_DN
HP_GLIC8S, HP_GLIC8S
MODULE_43, MODULE_43
GOBP_RNA_POLYADENYLATION, GOBP_RNA_POLYADENYLATION
GSE2826_WT_VS_XID_BCELL_DN, GSE2826_WT_VS_XID_BCELL_DN
REACTOME_FATTY_ACID_METABOLISM, REACTOME_FATTY_ACID_METABOLISM
GSE17721_PAM3CSK4_VS_GAD1QUIMOD_1H_BMDC_UP, GSE17721_PAM3CSK4_VS_GAD1QUIMOD_1H_BMDC_UP
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_3H_DN, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_3H_DN
DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_OLFACTORY_ENSHEATHING_GLIA, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_OLFACTORY_ENSHEATHING_GLIA
SWEET_KRAS_TARGETS_UP, SWEET_KRAS_TARGETS_UP
GSE3982_DC_VS_MAC_UP, GSE3982_DC_VS_MAC_UP
GOBP_CARDIAC_CHAMBER_DEVELOPMENT, GOBP_CARDIAC_CHAMBER_DEVELOPMENT
GOBP_CELL_SUBSTRATE_JUNCTION_ORGANIZATION, GOBP_CELL_SUBSTRATE_JUNCTION_ORGANIZATION
LENAOUR_DENDRITIC_CELL_MATURATION_UP, LENAOUR_DENDRITIC_CELL_MATURATION_UP
SISTO_RESPONSE_TO_DN_OR_SISTO_RESPONSE_TO_UV_CS
HP_ABNORMAL_RENAL_TUBULE_MORPHOLOGY, HP_ABNORMAL_RENAL_TUBULE_MORPHOLOGY
TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_LOBULAR_NORMAL_UP, TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_LOBULAR_NORMAL_UP
GOME_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRIED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN, GOME_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRIED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN
HP_ADENOCARCINOMA_OF_THE_INTESTINES, HP_ADENOCARCINOMA_OF_THE_INTESTINES
MODULE_40, MODULE_40
GSE34217_MIR17_92_OVEREXPRESS_VS_WT_ACT_CD8_TCELL_DN, GSE34217_MIR17_92_OVEREXPRESS_VS_WT_ACT_CD8_TCELL_DN
TSENG_IRS1_TARGETS_DN, TSENG_IRS1_TARGETS_DN
HP_ADENOCARCINOMA_OF_THE_LARGE_INTESTINE, HP_ADENOCARCINOMA_OF_THE_LARGE_INTESTINE
GOBP_NEURAL_TUBE_DEVELOPMENT, GOBP_NEURAL_TUBE_DEVELOPMENT
GSE40685_NAIVE_CD4_TCELL_VS_FOXP3_KO_TREG_PRECURSOR_DN, GSE40685_NAIVE_CD4_TCELL_VS_FOXP3_KO_TREG_PRECURSOR_DN
GOBP_CARDIAC_SEPTUM_DEVELOPMENT, GOBP_CARDIAC_SEPTUM_DEVELOPMENT
GOBP_FOCAL_ADHESION_ASSEMBLY, GOBP_FOCAL_ADHESION_ASSEMBLY
VGCANTCCR_UNKNOWN, VGCANTCCR_UNKNOWN
RUSSLINGER_DUODENAL_TUFT_CELLS, RUSSLINGER_DUODENAL_TUFT_CELLS
HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_HAND, HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_HAND
TAAYNRNNTCC_UNKNOWN, TAAYNRNNTCC_UNKNOWN
GSE360_DC_VS_MAC_M_TUBERCULOSIS_UP, GSE360_DC_VS_MAC_M_TUBERCULOSIS_UP
GOBP_NEPHRON_DEVELOPMENT, GOBP_NEPHRON_DEVELOPMENT
GOME_FLAVIN_ADENINE_DINUCLEOTIDE_BINDING, GOME_FLAVIN_ADENINE_DINUCLEOTIDE_BINDING
DACCOSTA_UV_RESPONSE_VIA_ERCC_XPCS_DN, DACCOSTA_UV_RESPONSE_VIA_ERCC_XPCS_DN
GSE26488_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMOCTE_UP, GSE26488_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMOCTE_UP
LUCAS_HNF4A_TARGETS_UP, LUCAS_HNF4A_TARGETS_UP
MODULE_343, MODULE_343
HADDAD_T_LYMPHOCYTE_AND_NK_PROGENITOR_UP, HADDAD_T_LYMPHOCYTE_AND_NK_PROGENITOR_UP
GALLUZZI_PERMEABILIZE_MITOCHONDRIA, GALLUZZI_PERMEABILIZE_MITOCHONDRIA
GOBP_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING, GOBP_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING
HARALAMBIEVA_PBMCM_M_R_IL6_IL13_IJOY_STIMULATED_VS_UNSTIMULATED_3_TO_7YR_POST_TOP_DEG_UP, HARALAMBIEVA_PBMCM_M_R_IL6_IL13_IJOY_STIMULATED_VS_UNSTIMULATED_3_TO_7YR_POST_TOP_DEG_UP
ROSS_AML_OF_FAB_M7_TYPE, ROSS_AML_OF_FAB_M7_TYPE
GOBP_REGULATION_OF_MEMBRANE_INVAGINATION, GOBP_REGULATION_OF_MEMBRANE_INVAGINATION
GSE43565_NK_CELL_VS_BCELL_DN, GSE43565_NK_CELL_VS_BCELL_DN
VALK_AML_CLUSTER_9, VALK_AML_CLUSTER_9
GOBP_CARDIAC_SEPTUM_MORPHOGENESIS, GOBP_CARDIAC_SEPTUM_MORPHOGENESIS
WP_PROSTAGLANDIN_SYNTHESIS_AND_REGULATION, WP_PROSTAGLANDIN_SYNTHESIS_AND_REGULATION
HU_FETAL_RETINA_AMACRINE, HU_FETAL_RETINA_AMACRINE
MIR378F, MIR378F
MIR378E, MIR378E
MIR378A_3P, MIR378C, MIR378D, MIR378I, MIR378A_3P, MIR378C, MIR378D, MIR378I
BOYALT_LIVER_CANCER_SUBCLASS_G6_UP, BOYALT_LIVER_CANCER_SUBCLASS_G6_UP
HP_HEMANGIOBLASTOMA, HP_HEMANGIOBLASTOMA
MIR422A, MIR422A
REACTOME_COLLAGEN_FORMATION, REACTOME_COLLAGEN_FORMATION
MODULE_221, MODULE_221
HP_CEREBRAL_HEMORRHAGE, HP_CEREBRAL_HEMORRHAGE
GOCC_ACTOMYOSIN, GOCC_ACTOMYOSIN
HP_ABNORMALITY_OF_SKIN_ADNEXA_PHYSIOLOGY, HP_ABNORMALITY_OF_SKIN_ADNEXA_PHYSIOLOGY
HP_HYPERHIDROSIS, HP_HYPERHIDROSIS
MODULE_295, MODULE_295
LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_LARGE_VS_TINY_DN, LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_LARGE_VS_TINY_DN
GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_BCELL_UP, GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_BCELL_UP
ANDERSEN_LIVER_CANCER_KRT19_DN, ANDERSEN_LIVER_CANCER_KRT19_DN
HP_CHEST_PAIN, HP_CHEST_PAIN
MIR379B, MIR379B
ZHAN_MULTIPLE_MYELOMA_HP_DN, ZHAN_MULTIPLE_MYELOMA_HP_DN
GOBP_UNSATURATED_FATTY_ACID_METABOLIC_PROCESS, GOBP_UNSATURATED_FATTY_ACID_METABOLIC_PROCESS
GOBP_RIBONUCLEOTIDE_CATABOLIC_PROCESS, GOBP_RIBONUCLEOTIDE_CATABOLIC_PROCESS
GOBP_REGULATION_OF_ACTIN_FILAMENT_BUNDLE_ASSEMBLY, GOBP_REGULATION_OF_ACTIN_FILAMENT_BUNDLE_ASSEMBLY
HP_ABNORMALITY_OF_PRIMARY_TEETH, HP_ABNORMALITY_OF_PRIMARY_TEETH
MIR68H_3P, MIR68H_3P
FOU372_01, FOU372_01
GOBP_VENTRICULAR_SEPTUM_DEVELOPMENT, GOBP_VENTRICULAR_SEPTUM_DEVELOPMENT
GOBP_ARTERY_DEVELOPMENT, GOBP_ARTERY_DEVELOPMENT
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_16, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_16
HP_ABNORMAL_CAUDATE_NUCLEUS_MORPHOLOGY, HP_ABNORMAL_CAUDATE_NUCLEUS_MORPHOLOGY
GOBP_MYELOID_CELL_APOPTOTIC_PROCESS, GOBP_MYELOID_CELL_APOPTOTIC_PROCESS
GOBP_ATRIOVENTRICULAR_VALVE_DEVELOPMENT, GOBP_ATRIOVENTRICULAR_VALVE_DEVELOPMENT
GOBP_REGULATION_OF_MRNA_POLYADENYLATION, GOBP_REGULATION_OF_MRNA_POLYADENYLATION
GOBP_NEPHRON_EPITHELIUM_DEVELOPMENT, GOBP_NEPHRON_EPITHELIUM_DEVELOPMENT
GOBP_AUTONOMIC_NERVOUS_SYSTEM_DEVELOPMENT, GOBP_AUTONOMIC_NERVOUS_SYSTEM_DEVELOPMENT
GAZDA_DIAMOND_BLACKFAN_ANEMIA_MYELOID_UP, GAZDA_DIAMOND_BLACKFAN_ANEMIA_MYELOID_UP
GOBP_ARTERY_MORPHOGENESIS, GOBP_ARTERY_MORPHOGENESIS
GOME_PROFILIN_BINDING, GOME_PROFILIN_BINDING
VISALA_RESPONSE_TO_HEAT_SHOCK_AND_AGING_UP, VISALA_RESPONSE_TO_HEAT_SHOCK_AND_AGING_UP
GOBP_AORTA_DEVELOPMENT, GOBP_AORTA_DEVELOPMENT
VANTVEER_BREAST_CANCER_METASTASIS_UP, VANTVEER_BREAST_CANCER_METASTASIS_UP
SUZUKI_RESPONSE_TO_TSA_AND_DECITABINE_1A, SUZUKI_RESPONSE_TO_TSA_AND_DECITABINE_1A
GOBP_NEGATIVE_REGULATION_OF_ANOIKS, GOBP_NEGATIVE_REGULATION_OF_ANOIKS
GOBP_REGULATION_OF_ACTOMYOSIN_STRUCTURE_ORGANIZATION, GOBP_REGULATION_OF_ACTOMYOSIN_STRUCTURE_ORGANIZATION
GOBP_SUBSTRATE_DEPENDENT_CELL_MIGRATION, GOBP_SUBSTRATE_DEPENDENT_CELL_MIGRATION
HP_MYOKYMIA, HP_MYOKYMIA
DOANE_BREAST_CANCER_CLASSES_UP, DOANE_BREAST_CANCER_CLASSES_UP
HP_SINUS_TACHYCARDIA, HP_SINUS_TACHYCARDIA
MELLMAN_TUT1_TARGETS_DN, MELLMAN_TUT1_TARGETS_DN
TRAVAGLINI_LUNG_MYELOID_DENDRITIC_TYPE_1_CELL, TRAVAGLINI_LUNG_MYELOID_DENDRITIC_TYPE_1_CELL
CLAUS_PCR_POSITIVE_MENINGIOMA_DN, CLAUS_PCR_POSITIVE_MENINGIOMA_DN
HALLMARK_HEDGEHOG_SIGNALING, HALLMARK_HEDGEHOG_SIGNALING
GOBP_KIDNEY_MORPHOGENESIS, GOBP_KIDNEY_MORPHOGENESIS
KEGG_TRYPTOPHAN_METABOLISM, KEGG_TRYPTOPHAN_METABOLISM
BOYALT_LIVER_CANCER_SUBCLASS_G123_DN, BOYALT_LIVER_CANCER_SUBCLASS_G123_DN
HAN_JNK_SINGALING_UP, HAN_JNK_SINGALING_UP
KEGG_PROXIMAL_TUBULE_BICARBONATE_RECLAMATION, KEGG_PROXIMAL_TUBULE_BICARBONATE_RECLAMATION
GOBP_REGULATION_OF_AXON_GUIDANCE, GOBP_REGULATION_OF_AXON_GUIDANCE
GOBP_ECTODERM_DEVELOPMENT, GOBP_ECTODERM_DEVELOPMENT
ABBUD_LIF_SIGNALING_2_UP, ABBUD_LIF_SIGNALING_2_UP
GOBP_SEMAPHORIN_PLEXIN_SIGNALING_PATHWAY_INVOLVED_IN_NEURON_PROJECTION_GUIDANCE, GOBP_SEMAPHORIN_PLEXIN_SIGNALING_PATHWAY_INVOLVED_IN_NEURON_PROJECTION_GUIDANCE
GOBP_ANATOMICAL_STRUCTURE_ARRANGEMENT, GOBP_ANATOMICAL_STRUCTURE_ARRANGEMENT
HP_FLUSHING, HP_FLUSHING
GOBP_POSITIVE_REGULATION_OF_STRESS_FIBER_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_STRESS_FIBER_ASSEMBLY
GOBP_MITRAL_VALVE_DEVELOPMENT, GOBP_MITRAL_VALVE_DEVELOPMENT
GOCC_HEMIDESMOSOME, GOCC_HEMIDESMOSOME
MTHFD1_TARGET_GENES, MTHFD1_TARGET_GENES
HP_BARREL_SHAPED_CHEST, HP_BARREL_SHAPED_CHEST
GOBP_DIGESTIVE_TRACT_MORPHOGENESIS, GOBP_DIGESTIVE_TRACT_MORPHOGENESIS
WP_MITOCHONDRIAL_LCFATTY_ACID_BETAOXIDATION, WP_MITOCHONDRIAL_LCFATTY_ACID_BETAOXIDATION
GOBP_RENAL_TUBULE_DEVELOPMENT, GOBP_RENAL_TUBULE_DEVELOPMENT
GOME_MUSCLE_ALPHA_ACTININ_BINDING, GOME_MUSCLE_ALPHA_ACTININ_BINDING
GOBP_POSITIVE_REGULATION_OF_FOCAL_ADHESION_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_FOCAL_ADHESION_ASSEMBLY
GOBP_PEPTIDE_CATABOLIC_PROCESS, GOBP_PEPTIDE_CATABOLIC_PROCESS
DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_RESPIRATORY_COLUMNAR_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_RESPIRATORY_COLUMNAR_CELLS
GNP2_MATK, GNP2_MATK
WP_TRYPTOPHAN_CATABOLISM_LEADING_TO_NAD_PRODUCTION, WP_TRYPTOPHAN_CATABOLISM_LEADING_TO_NAD_PRODUCTION
HP_SPONDYLOLYSIS, HP_SPONDYLOLYSIS
GOBP_AORTA_MORPHOGENESIS, GOBP_AORTA_MORPHOGENESIS
BLANCO_MELO_SARS_COV_1_INFECTION_MCR5_CELLS_DN, BLANCO_MELO_SARS_COV_1_INFECTION_MCR5_CELLS_DN
GCM_LTK, GCM_LTK
MCDOWELL_ACUTE_LUNG_INJURY_DN, MCDOWELL_ACUTE_LUNG_INJURY_DN
GOME_SEMAPHORIN_RECEPTOR_ACTIVITY, GOME_SEMAPHORIN_RECEPTOR_ACTIVITY
HP_LENS_LUXATION, HP_LENS_LUXATION
GOBP_OUTFLOW_TRACT_SEPTUM_MORPHOGENESIS, GOBP_OUTFLOW_TRACT_SEPTUM_MORPHOGENESIS
LEE_LIVER_CANCER_DENA_DN, LEE_LIVER_CANCER_DENA_DN
MAINA_VHL_TARGETS_UP, MAINA_VHL_TARGETS_UP
HP_CORNEAL_SCARRING, HP_CORNEAL_SCARRING
HP_LOW_FRUSTRATION_TOLERANCE, HP_LOW_FRUSTRATION_TOLERANCE
GOBP_NEGATIVE_REGULATION_OF_EPITHELIAL_CELL_MIGRATION, GOBP_NEGATIVE_REGULATION_OF_EPITHELIAL_CELL_MIGRATION
GOCC_SEMAPHORIN_RECEPTOR_COMPLEX, GOCC_SEMAPHORIN_RECEPTOR_COMPLEX
MODULE_354, MODULE_354
NOUSHMEHR_GBM_SOMATIC_MUTATED, NOUSHMEHR_GBM_SOMATIC_MUTATED
GOBP_REGULATION_OF_EXTRACELLULAR_MATRIX_DISASSEMBLY, GOBP_REGULATION_OF_EXTRACELLULAR_MATRIX_DISASSEMBLY
GOCC_EQUATORIAL_MICROTUBULE_ORGANIZING_CENTER, GOCC_EQUATORIAL_MICROTUBULE_ORGANIZING_CENTER
HP_APLASIA_HYPOPLASIA_INVOLVING_THE_SINUSES, HP_APLASIA_HYPOPLASIA_INVOLVING_THE_SINUSES
GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS
HP_DISTAL_LOWER_LIMB_AMYOTROPHY, HP_DISTAL_LOWER_LIMB_AMYOTROPHY
GOBP_NAD_BIOSYNTHETIC_PROCESS, GOBP_NAD_BIOSYNTHETIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION
SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX5_UP, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX5_UP
GOBP_LINOLEIC_ACID_METABOLIC_PROCESS, GOBP_LINOLEIC_ACID_METABOLIC_PROCESS
HP_OVERLAPPING_FINGERS, HP_OVERLAPPING_FINGERS
HP_HYPOGLYCEMIC_COMA, HP_HYPOGLYCEMIC_COMA
GOME_TRANSLATION_REPRESSOR_ACTIVITY, GOME_TRANSLATION_REPRESSOR_ACTIVITY
GOBP_INDOLE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_INDOLE_CONTAINING_COMPOUND_METABOLIC_PROCESS
GOBP_PHENOL_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_PHENOL_CONTAINING_COMPOUND_METABOLIC_PROCESS
GOBP_NEPHRON_MORPHOGENESIS, GOBP_NEPHRON_MORPHOGENESIS
GOBP_LIPOPHAGY, GOBP_LIPOPHAGY
GOME_LEUCINE_BINDING, GOME_LEUCINE_BINDING
GOBP_POSITIVE_REGULATION_OF_RECEPTOR_SIGNALING_PATHWAY_VIA_STAT, GOBP_POSITIVE_REGULATION_OF_RECEPTOR_SIGNALING_PATHWAY_VIA_STAT
GOBP_BENZENE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_BENZENE_CONTAINING_COMPOUND_METABOLIC_PROCESS
GOBP_HOMEOSTASIS_OF_NUMBER_OF_CELLS_WITHIN_A_TISSUE, GOBP_HOMEOSTASIS_OF_NUMBER_OF_CELLS_WITHIN_A_TISSUE
TONKS_TARGETS_OF_RUNX1_RUNX1_FUSION_SUSTAINED_IN GRANULOCYTE_UP, TONKS_TARGETS_OF_RUNX1_RUNX1_FUSION_SUSTAINED_IN GRANULOCYTE_UP
WP_SEROTONIN_TRANSPORTER_ACTIVITY, WP_SEROTONIN_TRANSPORTER_ACTIVITY
KEGG_DRUG_METABOLISM_CYTOCHROME_P450, KEGG_DRUG_METABOLISM_CYTOCHROME_P450
GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS
GOME_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY, GOME_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_CILIUM_MOVEMENT, GOBP_POSITIVE_REGULATION_OF_CILIUM_MOVEMENT
GOBP_AXONOGENESIS_INVOLVED_IN_INNervation, GOBP_AXONOGENESIS_INVOLVED_IN_INNervation
HP_OPEN_OPERCULUM, HP_OPEN_OPERCULUM
GOBP_REGULATION_OF_LIPOPHAGY, GOBP_REGULATION_OF_LIPOPHAGY
HP_HYPOVENTILATION, HP_HYPOVENTILATION
TAVAZOIE_METASTASIS, TAVAZOIE_METASTASIS
PID_VEGF_VEGFR_PATHWAY, PID_VEGF_VEGFR_PATHWAY
REACTOME_CRMP5_IN_SEMA3A_SIGNALING, REACTOME_CRMP5_IN_SEMA3A_SIGNALING
GOBP_CARDIAC_RIGHT_VENTRICLE_MORPHOGENESIS, GOBP_CARDIAC_RIGHT_VENTRICLE_MORPHOGENESIS
GOBP_MESONEPHRIC_TUBULE_MORPHOGENESIS, GOBP_MESONEPHRIC_TUBULE_MORPHOGENESIS
HP_SHOULDER_FLEXION_CONTRACTURE, HP_SHOULDER_FLEXION_CONTRACTURE
GOBP_MOTOR_NEURON_MIGRATION, GOBP_MOTOR_NEURON_MIGRATION
MIKKELSEN_IPS_LCP_WITH_H3K4ME3_AND_H3K27ME3, MIKKELSEN_IPS_LCP_WITH_H3K4ME3_AND_H3K27ME3
HP_FOCAL_CORTICAL_DYSPLASIA_TYPE_II, HP_FOCAL_CORTICAL_DYSPLASIA_TYPE_II
GOCC_GARP_COMPLEX, GOCC_GARP_COMPLEX
GOME_FIBROBLAST_GROWTH_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY, GOME_FIBROBLAST_GROWTH_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY
GOME_MITOGEN_ACTIVATED_PROTEIN_KINASE_P38_BINDING, GOME_MITOGEN_ACTIVATED_PROTEIN_KINASE_P38_BINDING
GOBP_MEDIAN_CHAIN_FATTY_ACID_COA_LIGASE_ACTIVITY, GOBP_MEDIAN_CHAIN_FATTY_ACID_COA_LIGASE_ACTIVITY
CAO_BLOOD_FLUMIST_AGE_05_14YO_1DY_DN, CAO_BLOOD_FLUMIST_AGE_05_14YO_1DY_DN
ZHONG_PFC_C3_HOPX_POS_OFC, ZHONG_PFC_C3_HOPX_POS_OFC
GOBP_SYMPATHETIC_NERVOUS_SYSTEM_DEVELOPMENT, GOBP_SYMPATHETIC_NERVOUS_SYSTEM_DEVELOPMENT
GOBP_GLYCOSIDE_METABOLIC_PROCESS, GOBP_GLYCOSIDE_METABOLIC_PROCESS
KORKOLA_EMBRYONIC_CARCINOMA_VS_SEMINOMA_DN, KORKOLA_EMBRYONIC_CARCINOMA_VS_SEMINOMA_DN
HP_ABNORMAL_SPEECH_INCENTERS, HP_ABNORMAL_SPEECH_INCENTERS
GOBP_LEARNED_VOICIALIZATION_BEHAVIOR_OR_VOICAL_LEARNING, GOBP_LEARNED_VOICIALIZATION_BEHAVIOR_OR_VOICAL_LEARNING
GOBP_POSITIVE_REGULATION_OF_AXON_EXTENSION_INVOLVED_IN_AXON_GUIDANCE, GOBP_POSITIVE_REGULATION_OF_AXON_EXTENSION_INVOLVED_IN_AXON_GUIDANCE
HP_ABSENT_PHALANGEAL_CREASE, HP_ABSENT_PHALANGEAL_CREASE
MIR933, MIR933
HP_POPLITEAL_PTERYGILUM, HP_POPLITEAL_PTERYGILUM
GOBP_NEURAL_CRIST_CELL_MIGRATION_INVOLVED_IN_AUTONOMIC_NERVOUS_SYSTEM_DEVELOPMENT, GOBP_NEURAL_CRIST_CELL_MIGRATION_INVOLVED_IN_AUTONOMIC_NERVOUS_SYSTEM_DEVELOPMENT
GOBP_CELL_FATE_SPECIFICATION, GOBP_CELL_FATE_SPECIFICATION
HP_SPONDYLOLISTHESIS, HP_SPONDYLOLISTHESIS
GOBP_CAMP_CATABOLIC_PROCESS, GOBP_CAMP_CATABOLIC_PROCESS
GOBP_GLIAL_CELL_FATE_COMMITMENT, GOBP_GLIAL_CELL_FATE_COMMITMENT
GOBP_DIPEPTIDE_TRANSMEMBRANE_TRANSPORT, GOBP_DIPEPTIDE_TRANSMEMBRANE_TRANSPORT
GOBP_DETECTION_OF_MUSCLE_STRETCH, GOBP_DETECTION_OF_MUSCLE_STRETCH
GOBP_REGULATION_OF_CDC42_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_REGULATION_OF_CDC42_PROTEIN_SIGNAL_TRANSDUCTION
chr4q32, chr4q32
TSENG_ADIPOGENIC_POTENTIAL_DN, TSENG_ADIPOGENIC_POTENTIAL_DN
GOBP_REGULATION_OF_GLOMERULAR_FILTRATION, GOBP_REGULATION_OF_GLOMERULAR_FILTRATION
GOBP_INDOLALKYLAMINE_METABOLIC_PROCESS, GOBP_INDOLALKYLAMINE_METABOLIC_PROCESS
HP_PERIODIC_PARALYSIS, HP_PERIODIC_PARALYSIS
GOBP_LYMPH_VESSEL_MORPHOGENESIS, GOBP_LYMPH_VESSEL_MORPHOGENESIS
GOBP_ASCENDING_AORTA_DEVELOPMENT, GOBP_ASCENDING_AORTA_DEVELOPMENT
GOME_LEUKOTRIENE_C4_SYNTHASE_ACTIVITY, GOME_LEUKOTRIENE_C4_SYNTHASE_ACTIVITY
GOME_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_BINDING, GOME_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_BINDING
GOBP_REGULATION_OF_RETINAL_GANGLION_CELL_AXON_GUIDANCE, GOBP_REGULATION_OF_RETINAL_GANGLION_CELL_AXON_GUIDANCE
GOBP_REGULATION_OF_LYMPHANGIOGENESIS, GOBP_REGULATION_OF_LYMPHANGIOGENESIS
FON2_TARGET_GENES, FON2_TARGET_GENES
GOBP_GLUTAMATE_BIOSYNTHETIC_PROCESS, GOBP_GLUTAMATE_BIOSYNTHETIC_PROCESS
GOBP_CARBON_DIOXIDE_TRANSPORT, GOBP_CARBON_DIOXIDE_TRANSPORT
GOBP_REGULATION_OF_INOSITOL_1_4_5_TRISPHOSPHATE_SENSITIVE_CALCIUM_RELEASE_CHANNEL_ACTIVITY, GOBP_REGULATION_OF_INOSITOL_1_4_5_TRISPHOSPHATE_SENSITIVE_CALCIUM_RELEASE_CHANNEL_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_POSITIVE_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
WP_FTO_OBESITY_VARIANT_MECHANISM_WP_FTO_OBESITY_VARIANT_MECHANISM
HP_AXILLARY_PTERYGILUM, HP_AXILLARY_PTERYGILUM
GOBP_DE_NOVO_NAD_BIOSYNTHETIC_PROCESS, GOBP_DE_NOVO_NAD_BIOSYNTHETIC_PROCESS
GOBP_DOPAMINE_CATABOLIC_PROCESS, GOBP_DOPAMINE_CATABOLIC_PROCESS
CCM_ATM, CCM_ATM
GOBP_TRIVALENT_INORGANIC_ANION_HOMEOSTASIS, GOBP_TRIVALENT_INORGANIC_ANION_HOMEOSTASIS
GOBP_CATECHOL_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_CATECHOL_CONTAINING_COMPOUND_METABOLIC_PROCESS
HP_PERIODIC_HYPOKALEMIC_PARESIS, HP_PERIODIC_HYPOKALEMIC_PARESIS
GOBP_TRYPTOPHAN_METABOLIC_PROCESS, GOBP_TRYPTOPHAN_METABOLIC_PROCESS
HP_HIGHLY_ELEVATED_CREATINE_KINASE, HP_HIGHLY_ELEVATED_CREATINE_KINASE
HP_ROUND_EAR, HP_ROUND_EAR
GOBP_REGULATION_OF_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI, GOBP_REGULATION_OF_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI
HP_ABNORMAL_PROTEIN_O_LINKED_GLYCOSYLATION, HP_ABNORMAL_PROTEIN_O_LINKED_GLYCOSYLATION
GOBP_VOICAL_LEARNING, GOBP_VOICAL_LEARNING
MAHADEVAN_IMATINIB_RESISTANCE_UP, MAHADEVAN_IMATINIB_RESISTANCE_UP
chr6q26, chr6q26
GOCC_MHC_CLASS_II_PROTEIN_COMPLEX, GOCC_MHC_CLASS_II_PROTEIN_COMPLEX
GOBP_SYMPATHETIC_GANGLION_DEVELOPMENT, GOBP_SYMPATHETIC_GANGLION_DEVELOPMENT
GOME_MHC_CLASS_II_RECEPTOR_ACTIVITY, GOME_MHC_CLASS_II_RECEPTOR_ACTIVITY
REACTOME_ERYTHROCYTES_TAKE_UP_OXYGEN_AND_RELEASE_CARBON_DIOXIDE, REACTOME_ERYTHROCYTES_TAKE_UP_OXYGEN_AND_RELEASE_CARBON_DIOXIDE
HP_FASTING_HYPERINSULINEMIA, HP_FASTING_HYPERINSULINEMIA
GOBP_NORADRENERGIC_NEURON_DIFFERENTIATION, GOBP_NORADRENERGIC_NEURON_DIFFERENTIATION
GOBP_RENAL_SYSTEM_PROCESS_INVOLVED_IN_REGULATION_OF_BLOOD_VOLUME, GOBP_RENAL_SYSTEM_PROCESS_INVOLVED_IN_REGULATION_OF_BLOOD_VOLUME
GOBP_MESODERMAL_CELL_DIFFERENTIATION, GOBP_MESODERMAL_CELL_DIFFERENTIATION
HP_SKULL_ASYMMETRY, HP_SKULL_ASYMMETRY
GOBP_SEPTUM_PRIMUM_DEVELOPMENT, GOBP_SEPTUM_PRIMUM_DEVELOPMENT
GOBP_TRYPTOPHAN_CATABOLIC_PROCESS, GOBP_TRYPTOPHAN_CATABOLIC_PROCESS, GOBP_TRYPTOPHAN_CATABOLIC_PROCESS, GOBP_TRYPTOPHAN_CATABOLIC_PROCESS
GOBP_TRYPTOPHAN_CATABOLIC_PROCESS, GOBP_TRYPTOPHAN_CATABOLIC_PROCESS
GOBP_PROTEIN_C_LINKED_GLYCOSYLATION, GOBP_PROTEIN_C_LINKED_GLYCOSYLATION
GOBP_NEGATIVE_REGULATION_OF_PROTAGLANDIN_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTAGLANDIN_BIOSYNTHETIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_CYTOSOLASMIC_TRANSLATION, GOBP_NEGATIVE_REGULATION_OF_CYTOSOLASMIC_TRANSLATION
GOBP_POSITIVE_REGULATION_OF_GLUTAMATE_SECRETION, GOBP_POSITIVE_REGULATION_OF_GLUTAMATE_SECRETION
GOBP_TRYPTOPHAN_CATABOLIC_PROCESS, GOBP_TRYPTOPHAN_CATABOLIC_PROCESS
GOBP_LYMPHANGIOGENESIS, GOBP_LYMPHANGIOGENESIS
GOBP_ACTIVATION_OF_JANUS_KINASE_ACTIVITY, GOBP_ACTIVATION_OF_JANUS_KINASE_ACTIVITY
GOBP_CHITIN_METABOLIC_PROCESS, GOBP_CHITIN_METABOLIC_PROCESS
GOME_ARYLESTERASE_ACTIVITY, GOME_ARYLESTERASE_ACTIVITY
GOBP_REGULATION_OF_GLUTAMATE_SECRETION, GOBP_REGULATION_OF_GLUTAMATE_SECRETION
GOBP_REGULATION_OF_BILE_ACID_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_BILE_ACID_BIOSYNTHETIC_PROCESS
REACTOME_PHOSPHOLIPASE_C_MEDIATED_CASCADE_FGR4, REACTOME_PHOSPHOLIPASE_C_MEDIATED_CASCADE_FGR4
WINNENPENNINGX_MELANOMA_METASTASIS_DN, WINNENPENNINGX_MELANOMA_METASTASIS_DN