

GSE17974.0,HS_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_DN,GSE17974.0,HS_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_DN
GSE17974.0H_VS_48H_IN_VITRO_ACT_CD4_TCELL_DN,GSE17974.0H_VS_48H_IN_VITRO_ACT_CD4_TCELL_DN
GSE17974.1,SL_VS_72H_IL4_AND_ANTIL_IL12_ACT_CD4_TCELL_DN,GSE17974.1,SH_VS_72H_IL4_AND_ANTIL_IL12_ACT_CD4_TCELL_DN
GSE2959.1,HA_VS_72H_IL4_AND_ANTIL_IL12_ACT_CD4_TCELL_DN,GSE2959.1,HA_VS_72H_IL4_AND_ANTIL_IL12_ACT_CD4_TCELL_DN
GSE17974.2,SH_VS_72H_IL4_AND_ANTIL_IL12_ACT_CD4_TCELL_DN,GSE17974.2,SH_VS_72H_IL4_AND_ANTIL_IL12_ACT_CD4_TCELL_DN
GSE17974.0H_VS_2H_IN_VITRO_ACT_CD4_TCELL_UP,GSE17974.0H_VS_2H_IN_VITRO_ACT_CD4_TCELL_UP
GSE17721.PAMCSK4_VS_GADQUIMOD.12H.BMDC.UP,GSE17721.PAMCSK4_VS_GADQUIMOD.12H.BMDC.UP
GSE2967.CTRL_VS_DAY7_TIV_FLU_VACCINE_PBMC_2008.DN,GSE2967.CTRL_VS_DAY7_TIV_FLU_VACCINE_PBMC_2008.DN
GSE2409.PDL_IGANTON_VS_CTRL_IN_ACT_TCELL_LINE_UP,GSE2409.PDL_IGANTON_VS_CTRL_IN_ACT_TCELL_LINE_UP
GSE17721.CTRL_VS_PAMCSK4.6H.BMDC.UP,GSE17721.CTRL_VS_PAMCSK4.6H.BMDC.UP
GSE17974.CTRL_VS_ACT_IL4_AND_ANTIL_IL12.2H_CD4_TCELL_UP,GSE17974.CTRL_VS_ACT_IL4_AND_ANTIL_IL12.2H_CD4_TCELL_UP
GSE46606.IRF4_KO_VS_WT_CD40L_IL12_IL15_1DAY_STIMULATED_BCELL_UP,GSE46606.IRF4_KO_VS_WT_CD40L_IL12_IL15_1DAY_STIMULATED_BCELL_UP
GSE17974.VS_IL4_IN_VITRO_ACT_CD4_TCELL_UP,GSE17974.VS_IL4_IN_VITRO_ACT_CD4_TCELL_UP
GSE2585.THYMIC_MACROPHAGE_VS_MTEC.UP,GSE2585.THYMIC_MACROPHAGE_VS_MTEC.UP
GSE22601.DOUBLE_NEGATIVE_VS_DOUBLE_POSITIVE_THYMOCTYE.UP,GSE22601.DOUBLE_NEGATIVE_VS_DOUBLE_POSITIVE_THYMOCTYE.UP
HP_METABOLIC_ACIDOSIS,HP_METABOLIC_ACIDOSIS
GSE17974.CTRL_VS_ACT_IL4_AND_ANTIL_IL12.1H_CD4_TCELL_UP,GSE17974.CTRL_VS_ACT_IL4_AND_ANTIL_IL12.1H_CD4_TCELL_UP
MENSE_HYPOXIA,UP,MENSE_HYPOXIA,UP
GSE11961.GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7.UP,GSE11961.GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7.UP
GSE42021.TCONV_PLN_VS_CD24LO.TCONV_THYMUS.UP,GSE42021.TCONV_PLN_VS_CD24LO.TCONV_THYMUS.UP
GSE16490.CTRL_VS_IPNA.12H_STIM_IMMATURE_NEURON_CELL_LINE_DN,GSE16490.CTRL_VS_IPNA.12H_STIM_IMMATURE_NEURON_CELL_LINE_DN
REACTOME_METABOLISM_OF_NUCLEOTIDES,REACTOME_METABOLISM_OF_NUCLEOTIDES
GSE3982.EFF_MEMORY_VS_CENT_MEMORY_CD4_TCELL_UP,GSE3982.EFF_MEMORY_VS_CENT_MEMORY_CD4_TCELL_UP
E2F3_UP,V1_UP,E2F3_UP,V1_UP
SHAFFER_IRF4_TARGETS_IN_ACTIVATED_DENDRITIC_CELL,SHAFFER_IRF4_TARGETS_IN_ACTIVATED_DENDRITIC_CELL
HP_CARDIOMEGALY,HP_CARDIOMEGALY
GSE2967.DAY3_VS_DAY7_TIV_FLU_VACCINE_PBMC_2008.UP,GSE2967.DAY3_VS_DAY7_TIV_FLU_VACCINE_PBMC_2008.UP
HP_NEURODEGENERATION,HP_NEURODEGENERATION
HP_SKIN_ULCER,HP_SKIN_ULCER
GSE21927.SPLEEN_VS_4T1_TUMOR_MONOCTYE_BALBC.UP,GSE21927.SPLEEN_VS_4T1_TUMOR_MONOCTYE_BALBC.UP
GSE28574.KC16_LOW_THP_VS_NAIVE_CD4_TCELL_DN,GSE28574.KC16_LOW_THP_VS_NAIVE_CD4_TCELL_DN
HP_HYPERMELANOTIC_MACULE,HP_HYPERMELANOTIC_MACULE
MIR4794,MIR4794
WP_TYROBP_CAUSAL_NETWORK,WP_TYROBP_CAUSAL_NETWORK
HP_NAIL_DYSPLASIA,HP_NAIL_DYSPLASIA
HP_ARTERIOCLEROSIS,HP_ARTERIOCLEROSIS
GOCC_MICROBODY_MEMBRANE,GOCC_MICROBODY_MEMBRANE
GOBP_NUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS,GOBP_NUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS
HP_ABNORMALITY_OF_HAIR_PIGMENTATION,HP_ABNORMALITY_OF_HAIR_PIGMENTATION
GSE29618.LAIV_VS_TIV_FLU_VACCINE_DAY7_PDC_DN,GSE29618.LAIV_VS_TIV_FLU_VACCINE_DAY7_PDC_DN
GSE1566.WT_VS_EZH2_KO_LN_TCELL_UP,GSE1566.WT_VS_EZH2_KO_LN_TCELL_UP
SERVITTA_ISLET_HNF1A_TARGETS_UP,SERVITTA_ISLET_HNF1A_TARGETS_UP
GSE29949.MICROGLIA_BRAIN_VS_CD8_POS_DC_SPLEEN_DN,GSE29949.MICROGLIA_BRAIN_VS_CD8_POS_DC_SPLEEN_DN
HP_PREMATURE_GRAYING_OF_HAIR,HP_PREMATURE_GRAYING_OF_HAIR
GOCC_U12_TYPE_SPLICOSOMAL_COMPLEX,GOCC_U12_TYPE_SPLICOSOMAL_COMPLEX
HP_SQUAMOUS_CELL_CARCINOMA,HP_SQUAMOUS_CELL_CARCINOMA
GOMF_MODIFIED_AMINO_ACID_BINDING,GOMF_MODIFIED_AMINO_ACID_BINDING
MIR6847.3P,MIR6847.3P
HP_ABNORMAL_LARYNX_PHYSIOLOGY,HP_ABNORMAL_LARYNX_PHYSIOLOGY
GOBP_CELL_CELL_JUNCTION_ASSEMBLY,GOBP_CELL_CELL_JUNCTION_ASSEMBLY
HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_56D_TOP_100_DEG_AFTER_IN_VITRO_RE_STIMULATION.UP,HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_56D_TOP_100_DEG_AFTER_IN_VITRO_RE_STIMULATION.UP
HP_DICARBOXYLIC_ACIDURIA,HP_DICARBOXYLIC_ACIDURIA
GOBP_LIGASE_ACTIVITY_FORMING_CARBON_NITROGEN_BONDS,GOMF_LIGASE_ACTIVITY_FORMING_CARBON_NITROGEN_BONDS
WP_NUCLEOTIDEBINDING,OLIGOMERIZATION,DOMADIN,NOD_PATHWAY
GOBP_RIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS,GOBP_RIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS
HP_FATIGABLE_WEAKNESS_OF_SWALLOWING_MUSCLES,HP_FATIGABLE_WEAKNESS_OF_SWALLOWING_MUSCLES
MORI_IMMATURE_B_LYMPHOCYTE.UP,MORI_IMMATURE_B_LYMPHOCYTE.UP
LEONARD_HYPOXIA,LEONARD_HYPOXIA
PID_PRL_SIGNALING_EVENTS_PATHWAY,PID_PRL_SIGNALING_EVENTS_PATHWAY
HP_UNUSUAL_FUNGAL_INFECTION,HP_UNUSUAL_FUNGAL_INFECTION
MIR3136.5P,MIR3136.5P
DESCARTES_MAIN_FETAL_THYMOCTYES,DESCARTES_MAIN_FETAL_THYMOCTYES
GOBP_TELENCEPHALON_GLIAL_CELL_MIGRATION,GOBP_TELENCEPHALON_GLIAL_CELL_MIGRATION
MIR574C.3P,MIR574C.3P
VECCH1_GASTRIC_CANCER_ADVANCED_VS_EARLY_DN,VECCH1_GASTRIC_CANCER_ADVANCED_VS_EARLY_DN
GOBP_REGULATION_OF_PHAGOCYTOSIS,GOBP_REGULATION_OF_PHAGOCYTOSIS
GNP2_FGR,GNP2_FGR
HP_CAPILLARY_HEMANGIOMA,HP_CAPILLARY_HEMANGIOMA
GOMF_PHOSPHATIDYLSELINE_BINDING,GOMF_PHOSPHATIDYLSELINE_BINDING
HP_ATROPHY_DEGENERATION_AFFECTING_THE_BRAINSTEM,HP_ATROPHY_DEGENERATION_AFFECTING_THE_BRAINSTEM
MIR439,MIR439
GOBP_NUCLEOBASE_METABOLIC_PROCESS,GOBP_NUCLEOBASE_METABOLIC_PROCESS
GOBP_GLIAL_CELL_MIGRATION,GOBP_GLIAL_CELL_MIGRATION
GOBP_NUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS,GOBP_NUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS
HP_PROLONGED_NEONATAL_JAUNDICE,HP_PROLONGED_NEONATAL_JAUNDICE
GOBP_PURINE_NUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS,GOBP_PURINE_NUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS
MIR6836.3P,MIR6836.3P
KEGG_NICOTINATE_AND_NICOTINAMIDE_METABOLISM,KEGG_NICOTINATE_AND_NICOTINAMIDE_METABOLISM
MORI_SMALL_PRE_BIL_LYMPHOCYTE.UP,MORI_SMALL_PRE_BIL_LYMPHOCYTE.UP
MIR3650,MIR3650
HP_OSTEOMYELITIS,HP_OSTEOMYELITIS
HP_PROKERIOD_FACIAL_APPEARANCE,HP_PROKERIOD_FACIAL_APPEARANCE
HP_HYPERGONADOTROPIC_HYPOGONADISM,HP_HYPERGONADOTROPIC_HYPOGONADISM
MIR6.3P,MIR6.3P
GOBP_RIBONUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS,GOBP_RIBONUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS
NAGASHIMA_NKCL_SIGNALING_DN,NAGASHIMA_NKCL_SIGNALING_DN
HP_MEDIASTINAL_LYMPHADENOPATHY,HP_MEDIASTINAL_LYMPHADENOPATHY
GOMF_PEPTIDASE_ACTIVATOR_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS,GOMF_PEPTIDASE_ACTIVATOR_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_INSULIN_SECRETION,GOBP_POSITIVE_REGULATION_OF_INSULIN_SECRETION
GOBP_PROTEIN_HYDROXYLATION,GOBP_PROTEIN_HYDROXYLATION
MIR4793.3P,MIR4793.3P
REACTOME_DISEASES_ASSOCIATED_WITH_GLYCOSYLATION_PRECURSOR_BIOSYNTHESIS,REACTOME_DISEASES_ASSOCIATED_WITH_GLYCOSYLATION_PRECURSOR_BIOSYNTHESIS
MIR3975,MIR3975
GOBP_POSITIVE_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_BY_HOST,GOBP_POSITIVE_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_BY_HOST
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_6,YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_6
HP_CORNEAL_EROSION,HP_CORNEAL_EROSION
HP_ERYTHEMA,HP_ERYTHEMA
HP_ACROCYANOSIS,HP_ACROCYANOSIS
GOBP_REGULATION_OF_B_CELL_PROLIFERATION,GOBP_REGULATION_OF_B_CELL_PROLIFERATION
WP_NAD_BIOSYNTHETIC_PATHWAYS,WP_NAD_BIOSYNTHETIC_PATHWAYS
HP_ABNORMAL_CORPUS_STRIATUM_MORPHOLOGY,HP_ABNORMAL_CORPUS_STRIATUM_MORPHOLOGY
MODULE_102,MODULE_102
WIEGEL_OXIDATIVE_STRESS_RESPONSE,WIEGEL_OXIDATIVE_STRESS_RESPONSE
HP_EXTRAPYRAMIDAL_MUSCULAR_RIGIDITY,HP_EXTRAPYRAMIDAL_MUSCULAR_RIGIDITY
GOBP_NAD_BIOSYNTHETIC_PROCESS,GOBP_NAD_BIOSYNTHETIC_PROCESS
CHEN_NEUROBLASTOMA_COPY_NUMBER_GAINS,CHEN_NEUROBLASTOMA_COPY_NUMBER_GAINS
MODULE_40,MODULE_40
BOYLAN_MULTIPLE_MYELOMA_C_CLUSTER_UP,BOYLAN_MULTIPLE_MYELOMA_C_CLUSTER_UP
HP_CONGENITAL_MICROCEPHALY,HP_CONGENITAL_MICROCEPHALY
HP_DECREASED_PLASMA_CARNITINE,HP_DECREASED_PLASMA_CARNITINE
KOHOUTEK_CCNT1_TARGETS,KOHOUTEK_CCNT1_TARGETS
VILLANUEVA_LIVER_CANCER_KRT19_DN,VILLANUEVA_LIVER_CANCER_KRT19_DN
ZHAN_V1_LATE_DIFFERENTIATION_GENES_UP,ZHAN_V1_LATE_DIFFERENTIATION_GENES_UP
GOCC_INTERCALATED_DISC,GOCC_INTERCALATED_DISC
ANDERSON_BLOOD_CN54GPH40_ADIUVANTED_WITH_GLA_AF_AGE_18_45YO_IDY_UP,ANDERSON_BLOOD_CN54GPH40_ADIUVANTED_WITH_GLA_AF_AGE_18_45YO_IDY_UP
XU_GH1_EXOGENOUS_TARGETS_DN,XU_GH1_EXOGENOUS_TARGETS_DN
BIOCARTA_CELLCELL_PATHWAY,BIOCARTA_CELLCELL_PATHWAY
GOBP_CEREBRAL_CORTEX_CELL_MIGRATION,GOBP_CEREBRAL_CORTEX_CELL_MIGRATION
GOBP_APICAL_JUNCTION_ASSEMBLY,GOBP_APICAL_JUNCTION_ASSEMBLY
KANG_FLUOROURACIL_RESISTANCE_UP,KANG_FLUOROURACIL_RESISTANCE_UP
WP_TRYPTOPHAN_CATABOLISM_LEADING_TO_NAD_PRODUCTION,WP_TRYPTOPHAN_CATABOLISM_LEADING_TO_NAD_PRODUCTION
HP_INTESTINAL_FISTULA,HP_INTESTINAL_FISTULA
KEEG_CYTOSOLIC_DNA_SENSING_PATHWAY,KEEG_CYTOSOLIC_DNA_SENSING_PATHWAY
GOBP_CEREBRAL_CORTEX_RADIALY_ORIENTED_CELL_MIGRATION,GOBP_CEREBRAL_CORTEX_RADIALY_ORIENTED_CELL_MIGRATION
GOBP_ETHER_METABOLIC_PROCESS,GOBP_ETHER_METABOLIC_PROCESS
GOBP_SULFUR_COMPOUND_CATABOLIC_PROCESS,GOBP_SULFUR_COMPOUND_CATABOLIC_PROCESS
REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_G1_CELL_CYCLE_ARREST,REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_G1_CELL_CYCLE_ARREST
HP_ABNORMAL_CIRCULATING_FATTY_ACID_ANION_CONCENTRATION,HP_ABNORMAL_CIRCULATING_FATTY_ACID_ANION_CONCENTRATION
HP_PERIODONTITIS,HP_PERIODONTITIS
GOBP_NUCLEOBASE_BIOSYNTHETIC_PROCESS,GOBP_NUCLEOBASE_BIOSYNTHETIC_PROCESS
REACTOME_NICOTINATE_METABOLISM,REACTOME_NICOTINATE_METABOLISM
GOMF_ENDOXYRIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOSTERS,GOMF_ENDOXYRIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOSTERS
GOBP_FOREBRAIN_CELL_MIGRATION,GOBP_FOREBRAIN_CELL_MIGRATION
MIR1225.5P,MIR1225.5P
MIR859,MIR859
HP_MACROVESICULAR_HEPATIC_STEATOSIS,HP_MACROVESICULAR_HEPATIC_STEATOSIS
HP_MUSCLE_STIFFNESS,HP_MUSCLE_STIFFNESS
HP_GINGIVITIS,HP_GINGIVITIS
GOBP_POSITIVE_REGULATION_OF_B_CELL_MEDIATED_IMMUNITY,GOBP_POSITIVE_REGULATION_OF_B_CELL_MEDIATED_IMMUNITY
GOBP_PALMITOYLTRANSFERASE_ACTIVITY,GOMF_PALMITOYLTRANSFERASE_ACTIVITY
HP_ABNORMAL_CIRCULATING_CREATININE_CONCENTRATION,HP_ABNORMAL_CIRCULATING_CREATININE_CONCENTRATION
HP_DECREASED_NUMBER_OF_PERIPHERAL_MYELINATED_NERVE_FIBERS,HP_DECREASED_NUMBER_OF_PERIPHERAL_MYELINATED_NERVE_FIBERS
GAO_SMALL_INTESTINE_24W_C2_PROCRPOS_PROGENTOR,GAO_SMALL_INTESTINE_24W_C2_PROCRPOS_PROGENTOR
WP_MITOCHONDRIAL_LCFATTY_ACID_BETAOXIDATION,WP_MITOCHONDRIAL_LCFATTY_ACID_BETAOXIDATION
GSE29618.LAIV_VS_TIV_FLU_VACCINE_DAY7_BCELL_UP,GSE29618.LAIV_VS_TIV_FLU_VACCINE_DAY7_BCELL_UP
HP_YAGINAL_FISTULA,HP_YAGINAL_FISTULA
GOCC_SARCOPLASMIC_RETICULUM_MEMBRANE,GOCC_SARCOPLASMIC_RETICULUM_MEMBRANE
HP_PANNICULITIS,HP_PANNICULITIS
GOBP_EPITHELIAL_CELL_CELL_ADHESION,GOBP_EPITHELIAL_CELL_CELL_ADHESION
ROYALTY_LIVER_CANCER_SUBLASS_G23_DN,ROYALTY_LIVER_CANCER_SUBLASS_G23_DN
VIERRECCHIA_RESPONSE_TO_TGFB1_C2,VIERRECCHIA_RESPONSE_TO_TGFB1_C2
HP_RECTAL_FISTULA,HP_RECTAL_FISTULA
ZHENG_IL22_SIGNALING_UP,ZHENG_IL22_SIGNALING_UP
GOBP_CARBHYDRATE_PHOSPHORYLATION,GOBP_CARBHYDRATE_PHOSPHORYLATION
GOBP_PURINE_NUCLEOBASE_BIOSYNTHETIC_PROCESS,GOBP_PURINE_NUCLEOBASE_BIOSYNTHETIC_PROCESS
GOBP_REGULATION_OF_MEMBRANE_INVAGINATION,GOBP_REGULATION_OF_MEMBRANE_INVAGINATION
HP_ABNORMAL_SERUM_INTERFERON_LEVEL,HP_ABNORMAL_SERUM_INTERFERON_LEVEL
WANG_METASTASIS_OF_BREAST_CANCER_ESR1_DN,WANG_METASTASIS_OF_BREAST_CANCER_ESR1_DN
HP_HYPERURICEMIA,HP_HYPERURICEMIA
GOBP_GLYCEROL_ETHER_METABOLIC_PROCESS,GOBP_GLYCEROL_ETHER_METABOLIC_PROCESS
GOBP_REGULATION_OF_LEUKOCYTE_DEGRANULATION,GOBP_REGULATION_OF_LEUKOCYTE_DEGRANULATION
GOBP_PURINE_NUCLEOBASE_METABOLIC_PROCESS,GOBP_PURINE_NUCLEOBASE_METABOLIC_PROCESS
EHLEBS_ANEUPLOIDY_DN,EHLEBS_ANEUPLOIDY_DN
GOBP_NEGATIVE_REGULATION_OF_PLASMA_MEMBRANE_BOUNDED_CELL_PROJECTION_ASSEMBLY,GOBP_NEGATIVE_REGULATION_OF_PLASMA_MEMBRANE_BOUNDED_CELL_PROJECTION_ASSEMBLY
HP_PROTUBERANT_ABDOMEN,HP_PROTUBERANT_ABDOMEN
HP_PSYCHOTIC_MENTATION,HP_PSYCHOTIC_MENTATION
HP_ZONULAR_CATARACT,HP_ZONULAR_CATARACT
OHM_METHYLATED_IN_ADULT_CANCERS,OHM_METHYLATED_IN_ADULT_CANCERS
MODULE_343,MODULE_343
HP_CENTRALLY_NUCLEATED_SKELETAL_MUSCLE_FIBERS,HP_CENTRALLY_NUCLEATED_SKELETAL_MUSCLE_FIBERS
GOMF_CARBOHYDRATE_KINASE_ACTIVITY,GOMF_CARBOHYDRATE_KINASE_ACTIVITY
GOBP_IMP_METABOLIC_PROCESS,GOBP_IMP_METABOLIC_PROCESS
GOCC_NLRP1_NFAMASOMAL_COMPLEX,GOCC_NLRP1_NFAMASOMAL_COMPLEX
GOBP_ABNORMAL_CIRCULATING_NUCLEOBASE_CONCENTRATION,HP_ABNORMAL_CIRCULATING_NUCLEOBASE_CONCENTRATION
GOMF_14_3_3_PROTEIN_BINDING,GOMF_14_3_3_PROTEIN_BINDING
GOBP_PURINE_NUCLEOSIDE_BIOSYNTHETIC_PROCESS,GOBP_PURINE_NUCLEOSIDE_BIOSYNTHETIC_PROCESS
GOBP_PEPITIDYL_LYSINE_HYDROXYLATION,GOBP_PEPITIDYL_LYSINE_HYDROXYLATION
GOBP_TOLL_LIKE_RECEPTOR_4_SIGNALING_PATHWAY,GOBP_TOLL_LIKE_RECEPTOR_4_SIGNALING_PATHWAY
HP_WHITE_HAIR,HP_WHITE_HAIR
GOBP_IMP_BIOSYNTHETIC_PROCESS,GOBP_IMP_BIOSYNTHETIC_PROCESS
HP_LIVER_ABSCESS,HP_LIVER_ABSCESS
GOBP_PROTEIN_MATURATION_BY_IRON_SULFUR_CLUSTER_TRANSFER,GOBP_PROTEIN_MATURATION_BY_IRON_SULFUR_CLUSTER_TRANSFER
GOBP_CELL_MOTILITY_INVOLVED_IN_CEREBRAL_CORTEX_RADIAL_GLIA_GUIDED_MIGRATION,GOBP_CELL_MOTILITY_INVOLVED_IN_CEREBRAL_CORTEX_RADIAL_GLIA_GUIDED_MIGRATION
REACTOME_NUCLEOBASE_BIOSYNTHESIS,REACTOME_NUCLEOBASE_BIOSYNTHESIS
POS_RESPONSE_TO_HISTAMINE_DN,POS_RESPONSE_TO_HISTAMINE_DN
HP_JUVENILE_CATARACT,HP_JUVENILE_CATARACT
GOBP_GALACTOSE_CATABOLIC_PROCESS,GOBP_GALACTOSE_CATABOLIC_PROCESS
HP_NEONATAL_HYPOGLYCEMIA,HP_NEONATAL_HYPOGLYCEMIA
SILIGAN_TARGETS_OF_EWS_FLI1_FUSION_UP,SILIGAN_TARGETS_OF_EWS_FLI1_FUSION_UP
GOMF_ACID_AMINO_ACID_LIGASE_ACTIVITY,GOMF_ACID_AMINO_ACID_LIGASE_ACTIVITY
MIR514.5P,MIR514.5P
HP_CHERRY_RED_SPOT_OF_THE_MACULA,HP_CHERRY_RED_SPOT_OF_THE_MACULA
GOBP_REGULATION_OF_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY,GOBP_REGULATION_OF_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY
HP_GOUT,HP_GOUT
GOBP_MAINTENANCE_OF_APICAL_BASAL_CELL_POLARITY,GOBP_MAINTENANCE_OF_APICAL_BASAL_CELL_POLARITY
GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_OSMOTIC_STRESS,GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_OSMOTIC_STRESS
GOBP_GLYCOLIPID_TRANSPORT,GOBP_GLYCOLIPID_TRANSPORT
HP_MOYAMOYA_PHENOMENON,HP_MOYAMOYA_PHENOMENON
REACTOME_PURINE_RIBONUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHESIS,REACTOME_PURINE_RIBONUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHESIS
REACTOME_RNA_MODIFICATION_IN_THE_MITOCHONDRION,REACTOME_RNA_MODIFICATION_IN_THE_MITOCHONDRION
GOMF_PROCOLLAGEN_PROLINE_DIOXYGENASE_ACTIVITY,GOMF_PROCOLLAGEN_PROLINE_DIOXYGENASE_ACTIVITY
GOBP_NEGATIVE_REGULATION_OF_PHAGOCYTOSIS,GOBP_NEGATIVE_REGULATION_OF_PHAGOCYTOSIS
OVSYANNIKOVA_PBMC_FLUARIX_AGE_55_64YO_RESPONDERS_VS_NONRESPONDERS_IDY_DN,OVSYANNIKOVA_PBMC_FLUARIX_AGE_55_64YO_RESPONDERS_VS_NONRESPONDERS_IDY_DN
BIOCARTA_FREI_PATHWAY,BIOCARTA_FREI_PATHWAY
HP_HYPOKETOTIC_HYPOGLYCEMIA,HP_HYPOKETOTIC_HYPOGLYCEMIA
GOBP_APOPTOTIC_DNA_FRAGMENTATION,GOBP_APOPTOTIC_DNA_FRAGMENTATION
BIOCARTA_HELPER_PATHWAY,BIOCARTA_HELPER_PATHWAY
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_SINGLE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN,GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_SINGLE_DONORS_WITH_INCORPORATION_OF_MOLECULAR_OXYGEN
GOMF_VOLTAGE_GATED_CHLORIDE_CHANNEL_ACTIVITY,GOBP_VOLTAGE_GATED_CHLORIDE_CHANNEL_ACTIVITY
REACTOME_FORMATION_OF_APOPTOSIS_REACTOME_FORMATION_OF_APOPTOSIS
REACTOME_PLATELET_CALCIIUM_HOMEOSTASIS,REACTOME_PLATELET_CALCIIUM_HOMEOSTASIS
HP_ONYCHOMYCOSIS,HP_ONYCHOMYCOSIS
GOBP_GALACTOSE_METABOLIC_PROCESS,GOBP_GALACTOSE_METABOLIC_PROCESS
LAX1B_TARGET_GENES_LAX1B_TARGET_GENES
GOMF_PROCOLLAGEN_PROLINE_4_DIOXYGENASE_ACTIVITY,GOMF_PROCOLLAGEN_PROLINE_4_DIOXYGENASE_ACTIVITY
REACTOME_GALACTOSE_CATABOLISM,REACTOME_GALACTOSE_CATABOLISM
BIOCARTA_NEUTROPHIL_PATHWAY,BIOCARTA_NEUTROPHIL_PATHWAY
HP_ABNORMALITY_OF_BODY_MASS_INDEX,HP_ABNORMALITY_OF_BODY_MASS_INDEX
GOCC_CHROMAFFIN GRANULE_MEMBRANE,GOCC_CHROMAFFIN GRANULE_MEMBRANE
GOBP_B_CELL_PROLIFERATION,GOBP_B_CELL_PROLIFERATION
GOBP_POSITIVE_REGULATION_OF_PPTIDE_HORMONE_SECRETION,GOBP_POSITIVE_REGULATION_OF_PPTIDE_HORMONE_SECRETION
REACTOME_MYOGENESIS,REACTOME_MYOGENESIS
WP_CYSINE_AND METHIONINE_CATABOLISM,WP_CYSINE_AND METHIONINE_CATABOLISM
GARCIA_PNERIS_PBMC_HPV_16_L1_VLP_AGE_18_25YO_12MO_CORRELATED_WITH_ANTI BODY_RESPONSE_POSITIVE,GARCIA_PNERIS_PBMC_HPV_16_L1_VLP_AGE_18_25YO_12MO_CORRELATED_WITH_ANTI BODY_RESPONSE_POSITIVE
GOCC_CHROMAFFIN GRANULE,GOCC_CHROMAFFIN GRANULE
GOBP_REGULATION_OF_NEUTROPHIL_ACTIVATION,GOBP_REGULATION_OF_NEUTROPHIL_ACTIVATION
GOBP_NEGATIVE_REGULATION_OF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY,GOBP_NEGATIVE_REGULATION_OF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
REACTOME_C127.7A_NFAMASOMAL_PATHWAY,REACTOME_C127.7A_NFAMASOMAL_PATHWAY
GOBP_POSITIVE_REGULATION_OF_INSULIN_SECRETION_INVOLVED_IN_CELLULAR_RESPONSE_TO_GLUCOSE_STIMULUS,GOBP_POSITIVE_REGULATION_OF_INSULIN_SECRETION_INVOLVED_IN_CELLULAR_RESPONSE_TO_GLUCOSE_STIMULUS
HP_IMPAIRED_TACTILE_SENSATION,HP_IMPAIRED_TACTILE_SENSATION
GOBP_NUCLEOSIDE_TRIPHOSPHATE_CATABOLIC_PROCESS,GOBP_NUCLEOSIDE_TRIPHOSPHATE_CATABOLIC_PROCESS
SBOBLEV_PBMC_PANDEMRIX_AGE_18_64YO_RESPONDERS_VS_NONRESPONDERS_7DY_UP,SBOBLEV_PBMC_PANDEMRIX_AGE_18_64YO_RESPONDERS_VS_NONRESPONDERS_7DY_UP
NKX2.8_TARGET_GENES,NKX2.8_TARGET_GENES
HP_MACROSCOPIC_HEMATURIA,HP_MACROSCOPIC_HEMATURIA
GOMF_MYOISIN_IL_BINDING,GOMF_MYOISIN_IL_BINDING
GOBP_REGULATION_OF_SKELETAL_MUSCLE_FIBER_DEVELOPMENT,GOBP_REGULATION_OF_SKELETAL_MUSCLE_FIBER_DEVELOPMENT
HP_EXERCISE_INDUCED_MYALGIA,HP_EXERCISE_INDUCED_MYALGIA
GOBP_PEPITIDYL_PROLINE_HYDROXYLATION,GOBP_PEPITIDYL_PROLINE_HYDROXYLATION
BIOCARTA_MONOCYTE_PATHWAY,BIOCARTA_MONOCYTE_PATHWAY
GAO_SMALL_INTESTINE_24W_C1_TUFT_PROGENTOR,GAO_SMALL_INTESTINE_24W_C1_TUFT_PROGENTOR
GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_CALCIIUM_ION_CONCENTRATION,GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_CALCIIUM_ION_CONCENTRATION
Jhe1521.sh1521
HP_XANTHELASMA,HP_XANTHELASMA
PASTURAL_RIZ1_TARGETS_UP,PASTURAL_RIZ1_TARGETS_UP
REACTOME_ACETYLCHOLINE_REGULATES_INSULIN_SECRETION,REACTOME_ACETYLCHOLINE_REGULATES_INSULIN_SECRETION
HP_ABNORMALITY_OF_THE_FIRST_METATARSAL_BONE,HP_ABNORMALITY_OF_THE_FIRST_METATARSAL_BONE
GOBP_REGULATION_OF_APOPTOTIC_DNA_FRAGMENTATION,GOBP_REGULATION_OF_APOPTOTIC_DNA_FRAGMENTATION
BIOCARTA_LYMPHOCYTE_PATHWAY,BIOCARTA_LYMPHOCYTE_PATHWAY
REACTOME_CREB3_FACTORS_ACTIVATE_GENES,REACTOME_CREB3_FACTORS_ACTIVATE_GENES
HP_IMPAIRED_NEUTROPHIL_BACTERICIDAL_ACTIVITY,HP_IMPAIRED_NEUTROPHIL_BACTERICIDAL_ACTIVITY
GOCC_ORIGIN_RECOGNITION_COMPLEX,GOCC_ORIGIN_RECOGNITION_COMPLEX
GOMF_PHOSPHATIDYLINOSITOL_3_5_BISPHOSPHATE_PHOSPHATASE_ACTIVITY,GOMF_PHOSPHATIDYLINOSITOL_3_5_BISPHOSPHATE_PHOSPHATASE_ACTIVITY
HP_TUBULOINTERSTITIAL_NEPHRITIS,HP_TUBULOINTERSTITIAL_NEPHRITIS
WP_NAD_METABOLISM,WP_NAD_METABOLISM
GOBP_PURINE_NUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS,GOBP_PURINE_NUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS
GOBP_ACTIVATION_OF_CYSINEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS_BY_CYTOCHROME_C,GOBP_ACTIVATION_OF_CYSINEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS_BY_CYTOCHROME_C
HP_FLARING_OF_RIB_CAGE,HP_FLARING_OF_RIB_CAGE
GOBP_TELOMERE_ASSEMBLY,GOBP_TELOMERE_ASSEMBLY
GOBP_REGULATION_OF_STORE_OPERATED_CALCIIUM_CHANNEL_ACTIVITY,GOBP_REGULATION_OF_STORE_OPERATED_CALCIIUM_CHANNEL_ACTIVITY
GOMF_LONG_CHAIN_FATTY_ACID_COA_BINDING,GOBP_LONG_CHAIN_FATTY_ACID_COA_BINDING
GOBP_PHOSPHATE_ION_TRANSMEMBRANE_TRANSPORT,GOBP_PHOSPHATE_ION_TRANSMEMBRANE_TRANSPORT
GOBP_REGULATION_OF_B_CELL_ACTIVATION,GOBP_REGULATION_OF_B_CELL_ACTIVATION
GSE3204.UNTREATED_VS_ROSGLIZATONE_TREATED_LY6C_LOW_MONOCTYE_UP,GSE3204.UNTREATED_VS_ROSGLIZATONE_TREATED_LY6C_LOW_MONOCTYE_UP
KEEG_RIBOFLAVIN_METABOLISM,KEEG_RIBOFLAVIN_METABOLISM
GOBP_CELL_VOLUME_HOMEOSTASIS,GOBP_CELL_VOLUME_HOMEOSTASIS
HP_NEONATAL_ALLOIMMUNE_THROMBOCYTOPENIA,HP_NEONATAL_ALLOIMMUNE_THROMBOCYTOPENIA
ONGUSAHA_BRCA1_TARGETS_UP,ONGUSAHA_BRCA1_TARGETS_UP
HP_AUTOAMPUTATION,HP_AUTOAMPUTATION
ACEVEDO_LIVER_CANCER_WITH_H3K9ME3_UP,ACEVEDO_LIVER_CANCER_WITH_H3K9ME3_UP
HP_EXAGGERATED_CUPID_S_BOW,HP_EXAGGERATED_CUPID_S_BOW
BIOCARTA_CBIR_PATHWAY,BIOCARTA_CBIR_PATHWAY
GOCC_DENDRITIC_SPINE_NECK,GOCC_DENDRITIC_SPINE_NECK
TERAMOTO_OPN_TARGETS_CLUSTER_5,MIR4801.3P,MIR4801.3P
FARDIN_HYPOXIA_9,FARDIN_HYPOXIA_9
HP_ULNAR_BOWING,HP_ULNAR_BOWING
SA_PROGRAMMED_CELL_DEATH,SA_PROGRAMMED_CELL_DEATH