

GSE27786.LSK.VS.NKCELL.UP,GSE27786.LSK.VS.NKCELL.UP
GSE2405.S.AURUS.VS.UNTREATED_NEUTROPHIL.UP,GSE2405.S.AURUS.VS.UNTREATED_NEUTROPHIL.UP
GSE5455.IX.VIVO.VS.POST_24H_INCUBATION_MONOCYTES_FROM_TUMOR_BEARING_MOUSE.UP,GSE5455.IX.VIVO.VS.POST_24H_INCUBATION_MONOCYTES_FROM_TUMOR_BEARING_MOUSE.UP
GSE1622.ANTL.CDCDCD28.STIM.VS.UNSTIM_MEMORY_CD8_TCELL.UP,GSE1622.ANTL.CDCDCD28.STIM.VS.UNSTIM_MEMORY_CD8_TCELL.UP
GSE8621.UNSTIM.VS.LPS.PRIMED.UNSTIM_MACROPHAGE.UP,GSE8621.UNSTIM.VS.LPS.PRIMED.UNSTIM_MACROPHAGE.UP
GSE5589.WT.VS_IL10_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN.UP,GSE5589.WT.VS_IL10_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN.UP
GSE7460.FOXP3.MUT.VS.WT.ACT_TCONV.UP,GSE7460.FOXP3.MUT.VS.WT.ACT_TCONV.UP
HP.INFLAMMATORY_ABNORMALITY_OF_THE_EYE,HP.INFLAMMATORY_ABNORMALITY_OF_THE_EYE
GSE21063.3H.VS_16H.ANTI_ICM_STIM_NFATC1_KORCELL.UP,GSE21063.3H.VS_16H.ANTI_ICM_STIM_NFATC1_KORCELL.UP
GSE2567.MPL.VS.R848.STIM_BCELL.UP,GSE2567.MPL.VS.R848.STIM_BCELL.UP
HP.ABNORMAL_CONJUNCTIVA_MORPHOLOGY,HP.ABNORMAL_CONJUNCTIVA_MORPHOLOGY
GSE36078.UNTREATED.VS_ADS_T425A_HEXON_INF_IL1R_KO_MOUSE_LUNG_DC_DN,GSE36078.UNTREATED.VS_ADS_T425A_HEXON_INF_IL1R_KO_MOUSE_LUNG_DC_DN
GSE5099.MONOCYTE.VS.ALTERNATIVE_M2_MACROPHAGE.UP,GSE5099.MONOCYTE.VS.ALTERNATIVE_M2_MACROPHAGE.UP
GSE1029.MEMORY.VS.KLRG1NT_EHF_CD8_TCELL.UP,GSE1029.MEMORY.VS.KLRG1NT_EHF_CD8_TCELL.UP
GSE37533.UNTREATED.VS.PI0GLIZATONE_TREATED_CD4_TCELL_FOXP3_TRANSDUCED_CD4_TCELL_DN,GSE37533.UNTREATED.VS.PI0GLIZATONE_TREATED_CD4_TCELL_FOXP3_TRANSDUCED_CD4_TCELL_DN
HP.RED_EYE,HP.RED_EYE
BUPFA_HYPONIA_METAGENE,BUPFA_HYPONIA_METAGENE
SHAFFER_IRF4_TARGETS_IN_PLASMA_CELL_VS_MATURE_B_LYMPHOCYTE,SHAFFER_IRF4_TARGETS_IN_PLASMA_CELL_VS_MATURE_B_LYMPHOCYTE
GGGNRMNNYCAT_UNKNOWN,GGGNRMNNYCAT_UNKNOWN
HP.CONFUSION,HP.CONFUSION
MIR891B,MIR891B
GSE2405.0H.VS_6H_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_DN,GSE2405.0H.VS_6H_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_DN
GOBP_PROTEIN_K11_LINKED_UBIQUITINATION,GOBP_PROTEIN_K11_LINKED_UBIQUITINATION
GSE5589.WT.VS_IL6_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_DN,GSE5589.WT.VS_IL6_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_DN
GSE3203.WT.VS_JFNARI_KO_INFLUENZA_INFECTED_LN_BCELL_DN,GSE3203.WT.VS_JFNARI_KO_INFLUENZA_INFECTED_LN_BCELL_DN
HP.KERATITIS,HP.KERATITIS
GOBP_FATTY_ACID_BIOSYNTHETIC_PROCESS,GOBP_FATTY_ACID_BIOSYNTHETIC_PROCESS
GOBP_EXIT_FROM_MITOSIS,GOBP_EXIT_FROM_MITOSIS
ZHOUL_TNF_SIGNALING_30MIN,ZHOUL_TNF_SIGNALING_30MIN
GOMF_PROTEIN_TYROSINE_PHOSPHATASE_ACTIVITY,GOMF_PROTEIN_TYROSINE_PHOSPHATASE_ACTIVITY
GOBP_PEPITIDYL_TYROSINE_DEPHOSPHORYLATION,GOBP_PEPITIDYL_TYROSINE_DEPHOSPHORYLATION
MIR4722.3P,MIR4722.3P
P53_DN.V1.UP,P53_DN.V1.UP
GSE10325.BCELL.VS_LUPUS_BCELL_UP,GSE10325.BCELL.VS_LUPUS_BCELL_UP
GSE19930.STIM.VS.STIM_AND_IL12_24H_CD8_T_CELL_DN,GSE19930.STIM.VS.STIM_AND_IL12_24H_CD8_T_CELL_DN
YANG.BREAST_CANCER_ESR1_DN,YANG.BREAST_CANCER_ESR1_DN
GOMF_PROTEIN_TYROSINE_SERINE_THREONINE_PHOSPHATASE_ACTIVITY,GOMF_PROTEIN_TYROSINE_SERINE_THREONINE_PHOSPHATASE_ACTIVITY
GSE37301.COMMON_LYMPHOID_PROGENITOR_VS_GRAN_MONO_PROGENITOR_DN,GSE37301.COMMON_LYMPHOID_PROGENITOR_VS_GRAN_MONO_PROGENITOR_DN
MIR4774.3P,MIR4774.3P
HP.BLEPHARITIS,HP.BLEPHARITIS
RBL1_TARGET_GENES,RBL1_TARGET_GENES
WP_SEROTONIN_RECEPTOR_2_AND_ELSRFGATA4_SIGNALING,WP_SEROTONIN_RECEPTOR_2_AND_ELSRFGATA4_SIGNALING
SHETH_LIVER_CANCER_VS_TXNIP_LOSS_PAM3,SHETH_LIVER_CANCER_VS_TXNIP_LOSS_PAM3
GSE17974.II4_AND_ANTL_IL12_VS_UNTREATED_2H_ACT_CD4_TCELL_DN,GSE17974.II4_AND_ANTL_IL12_VS_UNTREATED_2H_ACT_CD4_TCELL_DN
GSE10325.MYELOID.VS_LUPUS_MYELOID.UP,GSE10325.MYELOID.VS_LUPUS_MYELOID.UP
HP.PULMONARY_EMBOLISM,HP.PULMONARY_EMBOLISM
HP_KERATOCONJUNCTIVITIS,HP_KERATOCONJUNCTIVITIS
GOMF_MONOCARBOXYLIC_ACID_BINDING,GOMF_MONOCARBOXYLIC_ACID_BINDING
GOBP_REGULATION_OF_POSTSYNAPSE_ORGANIZATION,GOBP_REGULATION_OF_POSTSYNAPSE_ORGANIZATION
WP_PHOTODYNAMIC_THERAPYINDUCED_UNFOLDED_PROTEIN_RESPONSE,WP_PHOTODYNAMIC_THERAPYINDUCED_UNFOLDED_PROTEIN_RESPONSE
MIR924,MIR924
MORF_TNFRSF6,MORF_TNFRSF6
GOBP_ENTRAINMENT_OF_CIRCADIAN_CLOCK,GOBP_ENTRAINMENT_OF_CIRCADIAN_CLOCK
GOBP_RESPONSE_TO_FATTY_ACID,GOBP_RESPONSE_TO_FATTY_ACID
DESCARTES_FETAL_PLACENTA_AFP_ALB_POSITIVE_CELLS,DESCARTES_FETAL_PLACENTA_AFP_ALB_POSITIVE_CELLS
GOBP_RNA_5_END_PROCESSING,GOBP_RNA_5_END_PROCESSING
GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_STRESS,GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_STRESS
LEE_LIVER_CANCER_MYC_TGFA_DN,LEE_LIVER_CANCER_MYC_TGFA_DN
AKL_HTLV1_INFECTION.UP,AKL_HTLV1_INFECTION.UP
REACTOME_CARBOXYTERMINAL_POST_TRANSLATIONAL_MODIFICATIONS_OF_TUBULIN,REACTOME_CARBOXYTERMINAL_POST_TRANSLATIONAL_MODIFICATIONS_OF_TUBULIN
GOBP_GLUCOSE_6_PHOSPHATE_METABOLIC_PROCESS,GOBP_GLUCOSE_6_PHOSPHATE_METABOLIC_PROCESS
HP_CENTRALLY_NUCLEATED_SKELETAL_MUSCLE_FIBERS,HP_CENTRALLY_NUCLEATED_SKELETAL_MUSCLE_FIBERS
GOBP_FATTY_ACYL_COA_METABOLIC_PROCESS,GOBP_FATTY_ACYL_COA_METABOLIC_PROCESS
GOMF_THIOLESTER_HYDROLASE_ACTIVITY,GOMF_THIOLESTER_HYDROLASE_ACTIVITY
REACTOME_PENTOSE_PHOSPHATE_PATHWAY,REACTOME_PENTOSE_PHOSPHATE_PATHWAY
MIR6734.5P,MIR6734.5P
CHO_NR4A1_TARGETS,CHO_NR4A1_TARGETS
LEE_LIVER_CANCER_MYC_E2F1_DN,LEE_LIVER_CANCER_MYC_E2F1_DN
MIR4714.3P,MIR4714.3P
MODULE_541,MODULE_541
GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS,GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
GSE12963.UNINF.VS_ENV_AND_NEF_AND_VPR_DEFICIENT_HIV1_INF_CD4_TCELL_DN,GSE12963.UNINF.VS_ENV_AND_NEF_AND_VPR_DEFICIENT_HIV1_INF_CD4_TCELL_DN
MIR108.3P,MIR108.3P
MODULE_481,MODULE_481
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_RED.UP,GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_RED.UP
GOMF_O_ACYLTRANSFERASE_ACTIVITY,GOMF_O_ACYLTRANSFERASE_ACTIVITY
SILGAN_TARGETS_OF_EWS_FL1_FUSION_DN,SILGAN_TARGETS_OF_EWS_FL1_FUSION_DN
CUI_DEVELOPING_HEART_5TH_WEEK_VENTRICULAR_CARDIOMYOCYTE,CUI_DEVELOPING_HEART_5TH_WEEK_VENTRICULAR_CARDIOMYOCYTE
GOMF_CARBON_NITROGEN_LIGASE_ACTIVITY_WITH_GLUTAMINE_AS_AMIDO_N_DONOR,GOMF_CARBON_NITROGEN_LIGASE_ACTIVITY_WITH_GLUTAMINE_AS_AMIDO_N_DONOR
WANG_CLASSIC_ADIPOGENIC_TARGETS_OF_PPARG,WANG_CLASSIC_ADIPOGENIC_TARGETS_OF_PPARG
MURARO_PANCREAS_PANCREATIC_POLYPEPTIDE_CELL,MURARO_PANCREAS_PANCREATIC_POLYPEPTIDE_CELL
GOBP_REGULATION_OF_EXIT_FROM_MITOSIS,GOBP_REGULATION_OF_EXIT_FROM_MITOSIS
YAGUE_PREFUMOR_DRUG_RESISTANCE_DN,YAGUE_PREFUMOR_DRUG_RESISTANCE_DN
LEIN_OLIGODENDROCYTE_MARKERS,LEIN_OLIGODENDROCYTE_MARKERS
RUAN_RESPONSE_TO_TROGLITAZONE.UP,RUAN_RESPONSE_TO_TROGLITAZONE.UP
GOBP_SYNAPSE_ASSEMBLY,GOBP_SYNAPSE_ASSEMBLY
GOBP_PRIMARY_ALCOHOL_METABOLIC_PROCESS,GOBP_PRIMARY_ALCOHOL_METABOLIC_PROCESS
MORF_CD8A,MORF_CD8A
ESC_V6.5_UP_LATE.V1.UP,ESC_V6.5_UP_LATE.V1.UP
THEODOROU_MAMMARY_TUMORIGENESIS,THEODOROU_MAMMARY_TUMORIGENESIS
GOBP_REGULATION_OF_PHOSPHOLIPASE_C_ACTIVITY,GOBP_REGULATION_OF_PHOSPHOLIPASE_C_ACTIVITY
GOBP_PENTOSE_METABOLIC_PROCESS,GOBP_PENTOSE_METABOLIC_PROCESS
HP_ENCEPHALITIS,HP_ENCEPHALITIS
REACTOME_DEFECTS_IN_VITAMIN_AND_COFACTOR_METABOLISM,REACTOME_DEFECTS_IN_VITAMIN_AND_COFACTOR_METABOLISM
MIR7855.5P,MIR7855.5P
GOBP_RESPONSE_TO_NITROSATIVE_STRESS,GOBP_RESPONSE_TO_NITROSATIVE_STRESS
GSE25123.ROSIGLITAZONE.VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_DN,GSE25123.ROSIGLITAZONE.VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_DN
GOBP_PROTEIN_GERANYLERANYLATION,GOBP_PROTEIN_GERANYLERANYLATION
REACTOME_INTERLEUKIN_36_PATHWAY,REACTOME_INTERLEUKIN_36_PATHWAY
HP_PROGRESSIVE_HEARING_IMPAIRMENT,HP_PROGRESSIVE_HEARING_IMPAIRMENT
GOMF_FATTY_ACID_BINDING,GOMF_FATTY_ACID_BINDING
VALK_AML_CLUSTER_15,VALK_AML_CLUSTER_15
KEGG_NITROGEN_METABOLISM,KEGG_NITROGEN_METABOLISM
HP_OROMANDIBULAR_DYSTONIA,HP_OROMANDIBULAR_DYSTONIA
KEGG_TYROSINE_METABOLISM,KEGG_TYROSINE_METABOLISM
HP_ESOPHAGEAL_STRICTURE,HP_ESOPHAGEAL_STRICTURE
GOMF_DIPEPTIDYL_PEPTIDASE_ACTIVITY,GOMF_DIPEPTIDYL_PEPTIDASE_ACTIVITY
MIR4513,MIR4513
GOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_5_ETTS_OF_TRICSTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_18S_RRNA,GOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_5_ETTS_OF_TRICSTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_18S_RRNA
MIR4537,MIR4537
GOCC_THO_COMPLEX,GOCC_THO_COMPLEX
GOMF_CAMP_RESPONSE_ELEMENT_BINDING,GOMF_CAMP_RESPONSE_ELEMENT_BINDING
GOUYER_TATL_TARGETS_DN,GOUYER_TATL_TARGETS_DN
HP_NECK_FLEXOR_WEAKNESS,HP_NECK_FLEXOR_WEAKNESS
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_4,YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_4
HP_VENOUS_MALFORMATION,HP_VENOUS_MALFORMATION
GOBP_DENSE_CORE_GRANULE_TRANSPORT,GOBP_DENSE_CORE_GRANULE_TRANSPORT
BIOCARTA_LONGEVITY_PATHWAY,BIOCARTA_LONGEVITY_PATHWAY
HP_HYPOPHOSPHATEMIA,HP_HYPOPHOSPHATEMIA
MORE_IL4,MORE_IL4
HP_AXIAL_DYSTONIA,HP_AXIAL_DYSTONIA
GTCTACC_MIR379,GTCTACC_MIR379
HP_ABNORMAL_URINE_PHOSPHATE_CONCENTRATION,HP_ABNORMAL_URINE_PHOSPHATE_CONCENTRATION
BIOCARTA_CXCR4_PATHWAY,BIOCARTA_CXCR4_PATHWAY
MIR678,MIR678
GAO_STOMACH_24W_C5_PUTATIVE_PIT_CELL_PROGENITOR,GAO_STOMACH_24W_C5_PUTATIVE_PIT_CELL_PROGENITOR
ZN774_TARGET_GENES,ZN774_TARGET_GENES
HP_ACUTE_ENCEPHALOPATHY,HP_ACUTE_ENCEPHALOPATHY
GOBP_REGULATION_OF_PROTEIN_ADP_RIBOSYLATION,GOBP_REGULATION_OF_PROTEIN_ADP_RIBOSYLATION
GOBP_POSITIVE_REGULATION_OF_PEROXISOME_PROLIFERATOR_ACTIVATED_RECEPTOR_SIGNALING_PATHWAY,GOBP_POSITIVE_REGULATION_OF_PEROXISOME_PROLIFERATOR_ACTIVATED_RECEPTOR_SIGNALING_PATHWAY
GOBP_ACYLGLYCEROL_ACYL_CHAIN_REMODELING,GOBP_ACYLGLYCEROL_ACYL_CHAIN_REMODELING
REACTOME_DNA_DAMAGE_REVERSAL,REACTOME_DNA_DAMAGE_REVERSAL
GOCC_SYMMETRIC_SYNAPSE,GOCC_SYMMETRIC_SYNAPSE
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_CH_GROUP_OF_DONORS_OXYGEN_AS_ACCEPTOR,GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_CH_GROUP_OF_DONORS_OXYGEN_AS_ACCEPTOR
HP_ALLERGIC_RHINITIS,HP_ALLERGIC_RHINITIS
GOBP_SECRETORY_GROANULE_LOCALIZATION,GOBP_SECRETORY_GROANULE_LOCALIZATION
GOMF_AMINOACYLASE_ACTIVITY,GOMF_AMINOACYLASE_ACTIVITY
GOMF_UBIQUITIN_PROTEIN_TRANSFERASE_INHIBITOR_ACTIVITY,GOMF_UBIQUITIN_PROTEIN_TRANSFERASE_INHIBITOR_ACTIVITY
MORF_MYL3,MORF_MYL3
HP_ABNORMAL_CIRCULATING_MONOCARBOXYLIC_ACID_COCONCENTRATION,HP_ABNORMAL_CIRCULATING_MONOCARBOXYLIC_ACID_COCONCENTRATION
GOBP_REGULATION_OF_PEROXISOME_PROLIFERATOR_ACTIVATED_RECEPTOR_SIGNALING_PATHWAY,GOBP_REGULATION_OF_PEROXISOME_PROLIFERATOR_ACTIVATED_RECEPTOR_SIGNALING_PATHWAY
MORF_THF3,MORF_THF3
HP_ABNORMAL_BLOOD_SODIUM_CONCENTRATION,HP_ABNORMAL_BLOOD_SODIUM_CONCENTRATION
GOBP_PRENYLATION,GOBP_PRENYLATION
GOBP_SEMAPHORIN_PLEXIN_SIGNALING_PATHWAY_INVOLVED_IN_AXON_GUIDANCE,GOBP_SEMAPHORIN_PLEXIN_SIGNALING_PATHWAY_INVOLVED_IN_AXON_GUIDANCE
GOBP_RESPONSE_TO_STEROL,GOBP_RESPONSE_TO_STEROL
GOMF_COBALAMIN_BINDING,GOMF_COBALAMIN_BINDING
HP_INCREASED_URINARY_POTASSIUM,HP_INCREASED_URINARY_POTASSIUM
WP_BIOTIN_METABOLISM_INCLUDING_IEMS,WP_BIOTIN_METABOLISM_INCLUDING_IEMS
ZN586_TARGET_GENES,ZN586_TARGET_GENES
GOBP_PARAXIAL_MESODERM_FORMATION,GOBP_PARAXIAL_MESODERM_FORMATION
GOBP_RESPONSE_TO_ISOLATION_STRESS,GOBP_RESPONSE_TO_ISOLATION_STRESS
BIOCARTA_REELIN_PATHWAY,BIOCARTA_REELIN_PATHWAY
HP_IMPAIRED_DISTAL_PROPRIOCEPTION,HP_IMPAIRED_DISTAL_PROPRIOCEPTION
GOMF_PROTEIN_PRENYLTRANSFERASE_ACTIVITY,GOMF_PROTEIN_PRENYLTRANSFERASE_ACTIVITY
REACTOME_SOS_MEDIATED_SIGNALLING,REACTOME_SOS_MEDIATED_SIGNALLING
REACTOME_ACYL_CHAIN_REMODELING_OF_DAG_AND_TAG,REACTOME_ACYL_CHAIN_REMODELING_OF_DAG_AND_TAG
GOBP_TERNARY_METABOLIC_PROCESS,GOBP_TERNARY_METABOLIC_PROCESS
REACTOME_SIGNALING_BY_PDGFRA_TRANSMEMBRANE_JUXTAMEMBRANE_AND_KINASE_DOMAIN_MUTANTS,REACTOME_SIGNALING_BY_PDGFRA_TRANSMEMBRANE_JUXTAMEMBRANE_AND_KINASE_DOMAIN_MUTANTS
GOBP_REGULATION_OF_HISTONE_DEACETYLASE_ACTIVITY,GOBP_REGULATION_OF_HISTONE_DEACETYLASE_ACTIVITY
HP_PAIN_INSENSITIVITY,HP_PAIN_INSENSITIVITY
GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_13_PRODUCTION,GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_13_PRODUCTION
GOCC_SEMAPHORIN_RECEPTOR_COMPLEX,GOCC_SEMAPHORIN_RECEPTOR_COMPLEX
REACTOME_SHC_RELATED_EVENTS_TRIGGERED_BY_FGFR,REACTOME_SHC_RELATED_EVENTS_TRIGGERED_BY_FGFR
HP_RESPONSE_TO_DRUGS_ACTING_ON_NEUROMUSCULAR_TRANSMISSION,HP_RESPONSE_TO_DRUGS_ACTING_ON_NEUROMUSCULAR_TRANSMISSION
GOBP_CELLULAR_TRIGLYCERIDE_HOMEOSTASIS,GOBP_CELLULAR_TRIGLYCERIDE_HOMEOSTASIS
GOBP_OPIOID_RECEPTOR_SIGNALING_PATHWAY,GOBP_OPIOID_RECEPTOR_SIGNALING_PATHWAY
GOMF_DIACYLGLYCEROL_O_ACYLTRANSFERASE_ACTIVITY,GOMF_DIACYLGLYCEROL_O_ACYLTRANSFERASE_ACTIVITY
HP_DEEP_SET_NAILS,HP_DEEP_SET_NAILS
GOBP_RRNA_5_END_PROCESSING,GOBP_RRNA_5_END_PROCESSING
HP_ABNORMALITY_OF_THE_PREMAXILLA,HP_ABNORMALITY_OF_THE_PREMAXILLA
GOMF_O_ACETYLTRANSFERASE_ACTIVITY,GOMF_O_ACETYLTRANSFERASE_ACTIVITY
HP_ABNORMAL_CIRCULATING_GLUTAMINE_CONCENTRATION,HP_ABNORMAL_CIRCULATING_GLUTAMINE_CONCENTRATION
GOBP_PEPITIDYL_DIPHTHAMIDE_BIOSYNTHETIC_PROCESS_FROM_PEPITIDYL_HISTIDINE,GOBP_PEPITIDYL_DIPHTHAMIDE_BIOSYNTHETIC_PROCESS_FROM_PEPITIDYL_HISTIDINE
GOBP_CD8_POSITIVE_ALPHA_BETA_T_CELL_PROLIFERATION,GOBP_CD8_POSITIVE_ALPHA_BETA_T_CELL_PROLIFERATION
REACTOME_SYNTHESIS_OF_DIPHTHAMIDE_EEF2,REACTOME_SYNTHESIS_OF_DIPHTHAMIDE_EEF2
GOBP_SERINE_FAMILY_AMINO_ACID_CATABOLIC_PROCESS,GOBP_SERINE_FAMILY_AMINO_ACID_CATABOLIC_PROCESS
SMID_BREAST_CANCER_RELAPSE_IN_BONE.UP,SMID_BREAST_CANCER_RELAPSE_IN_BONE.UP
GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_5_PRODUCTION,GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_5_PRODUCTION
REACTOME_UREA_CYCLE,REACTOME_UREA_CYCLE
MIR598.3P,MIR598.3P
GOCC_ENDOPLASMIC_RETICULUM_PLASMA_MEMBRANE_CONTACT_SITE,GOCC_ENDOPLASMIC_RETICULUM_PLASMA_MEMBRANE_CONTACT_SITE
GOBP_MONOVALENT_INORGANIC_ANION_HOMEOSTASIS,GOBP_MONOVALENT_INORGANIC_ANION_HOMEOSTASIS
GOBP_CELLULAR_RESPONSE_TO_STEROL,GOBP_CELLULAR_RESPONSE_TO_STEROL
GOMF_STRUCTURAL_CONSTITUENT_OF_CYTOSKELETON,GOMF_STRUCTURAL_CONSTITUENT_OF_CYTOSKELETON
HP_LOCALIZED_HIRSUTISM,HP_LOCALIZED_HIRSUTISM
HP_ABNORMAL_CIRCULATING_PORPHYRIN_CONCENTRATION,HP_ABNORMAL_CIRCULATING_PORPHYRIN_CONCENTRATION
GOBP_PURINE_NUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS,GOBP_PURINE_NUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS
HP_PROXIMAL_RENAL_TUBULAR_ACIDOSIS,HP_PROXIMAL_RENAL_TUBULAR_ACIDOSIS
GOBP_CYTOPLASMIC_TRANSLATIONAL_ELONGATION,GOBP_CYTOPLASMIC_TRANSLATIONAL_ELONGATION
HP_GLOSTIS,HP_GLOSTIS
HP_UNUSUAL_SKIN_INFECTION,HP_UNUSUAL_SKIN_INFECTION
CAO_BLOOD_FLUZONE_AGE_05_14YO_1DY_DN,CAO_BLOOD_FLUZONE_AGE_05_14YO_1DY_DN
GOBP_FRUCTOSE_6_PHOSPHATE_METABOLIC_PROCESS,GOBP_FRUCTOSE_6_PHOSPHATE_METABOLIC_PROCESS
HP_PROGRESSIVE_SENSORINEURAL_HEARING_IMPAIRMENT,HP_PROGRESSIVE_SENSORINEURAL_HEARING_IMPAIRMENT
HARALAMBIEVA_P8MC_M_M_R_IL_AGE_11_22YO_VACCINATED_VS_UNVACCINATED_LOW_ANTI BODY_RESPONSES_TO_TREATMENT_7YR.UP,HARALAMBIEVA_P8MC_M_M_R_IL_AGE_11_22YO_VACCINATED_VS_UNVACCINATED_LOW_ANTI BODY_RESPONSES_TO_TREATMENT_7YR.UP
BIOCARTA_TRKA_PATHWAY,BIOCARTA_TRKA_PATHWAY
HOEMANN_MYELODYSPLASTIC_SYNDROM_RISK.UP,HOEMANN_MYELODYSPLASTIC_SYNDROM_RISK.UP
BIOCARTA_CCR3_PATHWAY,BIOCARTA_CCR3_PATHWAY
LOPEZ_EPITHELIOID_MESOTHELIOMA,LOPEZ_EPITHELIOID_MESOTHELIOMA
HP_PROMINENCE_OF_THE_PREMAXILLA,HP_PROMINENCE_OF_THE_PREMAXILLA
GOBP_PARANODAL_JUNCTION_ASSEMBLY,GOBP_PARANODAL_JUNCTION_ASSEMBLY
GOMF_BIOTIN_BINDING,GOMF_BIOTIN_BINDING
HP_SINGLE_FIBER_EMG_ABNORMALITY,HP_SINGLE_FIBER_EMG_ABNORMALITY
GOBP_ACTIN_MODIFICATION,GOBP_ACTIN_MODIFICATION
GOBP_HISTIDINE_CATABOLIC_PROCESS,GOBP_HISTIDINE_CATABOLIC_PROCESS
REACTOME_HISTIDINE_CATABOLISM,REACTOME_HISTIDINE_CATABOLISM
GOBP_CELLULAR DETOXIFICATION_OF_ALDEHYDE,GOBP_CELLULAR DETOXIFICATION_OF_ALDEHYDE
HP_MUSCLE_FIBER_TUBULAR_INCLUSIONS,HP_MUSCLE_FIBER_TUBULAR_INCLUSIONS
REACTOME_ACTIVATED_NTRK3_SIGNALS_THROUGH_RAS,REACTOME_ACTIVATED_NTRK3_SIGNALS_THROUGH_RAS
GOBP_RESPONSE_TO_FOLLICLE_STIMULATING_HORMONE,GOBP_RESPONSE_TO_FOLLICLE_STIMULATING_HORMONE
GOBP_PROTEIN_NITROSYLATION,GOBP_PROTEIN_NITROSYLATION
HP_APLASIA_OF_THE_MUSCULATURE,HP_APLASIA_OF_THE_MUSCULATURE
GOBP_DIACYLGLYCEROL_BIOSYNTHETIC_PROCESS,GOBP_DIACYLGLYCEROL_BIOSYNTHETIC_PROCESS
GOBP_RETROGRADE_TRANS_SYNAPTIC_SIGNALING,GOBP_RETROGRADE_TRANS_SYNAPTIC_SIGNALING
HP_ABNORMAL_LIGHT_ADAPTED_ELECTRORETINOGRAM,HP_ABNORMAL_LIGHT_ADAPTED_ELECTRORETINOGRAM
GOBP_OPIOID_RECEPTOR_ACTIVITY,GOMF_OPIOID_RECEPTOR_ACTIVITY
WP_TRIACYLGLYCERIDE_SYNTHESIS,WP_TRIACYLGLYCERIDE_SYNTHESIS
BIOCARTA_HBX_PATHWAY,BIOCARTA_HBX_PATHWAY
GOBP_TRANS_SYNAPTIC_SIGNALING_BY_LIPID,GOBP_TRANS_SYNAPTIC_SIGNALING_BY_LIPID
GOBP_RETINAL_METABOLIC_PROCESS,GOBP_RETINAL_METABOLIC_PROCESS
MODULE_227,MODULE_227
REACTOME_POST_CHAPERONIN_TUBULIN_FOLDING_PATHWAY,REACTOME_POST_CHAPERONIN_TUBULIN_FOLDING_PATHWAY
CASTELLANO_HRAS_TARGETS_DN,CASTELLANO_HRAS_TARGETS_DN
HP_NASAL_REGURITATION,HP_NASAL_REGURITATION
GOBP_UREA_CYCLE,GOBP_UREA_CYCLE
REACTOME_SHC1_EVENTS_IN_ERBB4_SIGNALING,REACTOME_SHC1_EVENTS_IN_ERBB4_SIGNALING
HP_ABSENT_FOVEAL_REFLEX,HP_ABSENT_FOVEAL_REFLEX
GAUSSMANN_MLL_AFL_FUSION_TARGETS_D_DN,GAUSSMANN_MLL_AFL_FUSION_TARGETS_D_DN
WP_POLATEALCORON_AND_CANCER_PATHWAY_HYPOTHESES,WP_POLATEALCORON_AND_CANCER_PATHWAY_HYPOTHESES
REACTOME_ACTIVATED_NTRK2_SIGNALS_THROUGH_RAS,REACTOME_ACTIVATED_NTRK2_SIGNALS_THROUGH_RAS
MIR4787.5P,MIR4787.5P
GOBP_RESPIRATORY_BURST_INVOLVED_IN_INFLAMMATORY_RESPONSE,GOBP_RESPIRATORY_BURST_INVOLVED_IN_INFLAMMATORY_RESPONSE
GOBP_MONOACYLGLYCEROL_BIOSYNTHETIC_PROCESS,GOBP_MONOACYLGLYCEROL_BIOSYNTHETIC_PROCESS
TUCAMISTO_TUMOR_SUPPRESSION_BY_COL13A1_DN,TUCAMISTO_TUMOR_SUPPRESSION_BY_COL13A1_DN
GOBP_POSITIVE_REGULATION_OF_FAT_CELL_DIFFERENTIATION,GOBP_POSITIVE_REGULATION_OF_FAT_CELL_DIFFERENTIATION
GOBP_CEREBRAL_CORTEX_RADIALY_ORIENTED_CELL_MIGRATION,GOBP_CEREBRAL_CORTEX_RADIALY_ORIENTED_CELL_MIGRATION
GOBP_INTERLEUKIN_21_MEDIATED_SIGNALING_PATHWAY,GOBP_INTERLEUKIN_21_MEDIATED_SIGNALING_PATHWAY
HP_PECTORAL_MUSCLE_HYPOPLASIA_APLASIA,HP_PECTORAL_MUSCLE_HYPOPLASIA_APLASIA
GOBP_UREA_METABOLIC_PROCESS,GOBP_UREA_METABOLIC_PROCESS
GOBP_CEREBRAL_CORTEX_CELL_MIGRATION,GOBP_CEREBRAL_CORTEX_CELL_MIGRATION
GOMF_COA_HYDROLASE_ACTIVITY,GOMF_COA_HYDROLASE_ACTIVITY
GOBP_RESPONSE_TO_GRAVITY,GOBP_RESPONSE_TO_GRAVITY
GOMF_NEGATIVE_REGULATION_OF_MACROPHAGE_MIGRATION,GOBP_NEGATIVE_REGULATION_OF_MACROPHAGE_MIGRATION
GOMF_ALCOHOL_DEHYDROGENASE_NAD_P_PLUS_ACTIVITY,GOMF_ALCOHOL_DEHYDROGENASE_NAD_P_PLUS_ACTIVITY
HP_HYPEREXTENSIBILITY_OF_THE_FINGER_JOINTS,HP_HYPEREXTENSIBILITY_OF_THE_FINGER_JOINTS