

GSE46606_IRF4HIGH_VS_WT_CD40L_IL12_IL5_DAY1_STIMULATED_BCELL_UP, GSE46606_IRF4HIGH_VS_WT_CD40L_IL12_IL5_DAY1_STIMULATED_BCELL_UP
GSE24726_WT_VS_E2_2_KO_PDC_UP, GSE24726_WT_VS_E2_2_KO_PDC_UP
GSE25085_FETAL_LIVER_VS_FETAL_BM.SP4_THYMIC_IMPLANT_UP, GSE25085_FETAL_LIVER_VS_FETAL_BM.SP4_THYMIC_IMPLANT_UP
GSE32986_CURDLAN_HIGHDOSE_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_UP, GSE32986_CURDLAN_HIGHDOSE_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_UP
GSE35685_CD34POS_CD10NEG_CD62LPOS_VS_CD34POS_CD10POS_BONE_MARROW_UP, GSE35685_CD34POS_CD10NEG_CD62LPOS_VS_CD34POS_CD10POS_BONE_MARROW_UP
GSE39864_WT_VS_GATA3_KO_TREG_DN, GSE39864_WT_VS_GATA3_KO_TREG_DN
GSE19923_WT_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP, GSE19923_WT_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP
GSE24292_WT_VS_PPARG_KO_MACROPHAGE_DN, GSE24292_WT_VS_PPARG_KO_MACROPHAGE_DN
GSE22935_UNSTIM_VS_12H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_DN, GSE22935_UNSTIM_VS_12H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_DN
E2F1_UP.V1_UP, E2F1_UP.V1_UP
GSE46606_IRF4_KO_VS_WT_UNSTIM_BCELL_UP, GSE46606_IRF4_KO_VS_WT_UNSTIM_BCELL_UP
GSE24726_WT_VS_E2_2_KO_PDC_DAY6_POST_DELETION_DN, GSE24726_WT_VS_E2_2_KO_PDC_DAY6_POST_DELETION_DN
GSE9650_NAIVE_VS_EFF_CD8_TCELL_DN, GSE9650_NAIVE_VS_EFF_CD8_TCELL_DN
GSE23568_ID3_KO_VS_WT_CD8_TCELL_DN, GSE23568_ID3_KO_VS_WT_CD8_TCELL_DN
GSE37301_PRO_BCELL_VS_RAG2_KO_NK_CELL_DN, GSE37301_PRO_BCELL_VS_RAG2_KO_NK_CELL_DN
GSE33424_CD161_HIGH_VS_NEG_CD8_TCELL_UP, GSE33424_CD161_HIGH_VS_NEG_CD8_TCELL_UP
MIR4423_3P, MIR4423_3P
GSE36891_POLYIC_TLR3_VS_PAM_TLR2_STIM_PERITONEAL_MACROPHAGE_DN, GSE36891_POLYIC_TLR3_VS_PAM_TLR2_STIM_PERITONEAL_MACROPHAGE_DN
GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_UP, GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_UP
GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP, GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP
RICKMAN_TUMOR_DIFFERENTIATED_WELL_VS_MODERATELY_UP, RICKMAN_TUMOR_DIFFERENTIATED_WELL_VS_MODERATELY_UP
RB_DN.V1_UP, RB_DN.V1_UP
MODULE_567, MODULE_567
PID_FANCONI_PATHWAY, PID_FANCONI_PATHWAY
GSE43955_10H_VS_30H_ACT_CD4_TCELL_DN, GSE43955_10H_VS_30H_ACT_CD4_TCELL_DN
GOBP_BASE_EXCISION_REPAIR, GOBP_BASE_EXCISION_REPAIR
GSE17721_LPS_VS_PAM3CSK4_0.5H_BMDC_DN, GSE17721_LPS_VS_PAM3CSK4_0.5H_BMDC_DN
MIR4784, MIR4784
GOBP_BLASTOCYST_DEVELOPMENT, GOBP_BLASTOCYST_DEVELOPMENT
GOMF_GDP_BINDING, GOMF_GDP_BINDING
MIR3150B_3P, MIR3150B_3P
REACTOME_HDR_THROUGH_SINGLE_STRAND_ANNHEALING_SSA, REACTOME_HDR_THROUGH_SINGLE_STRAND_ANNHEALING_SSA
WCTCNATGGY_UNKNOWN, WCTCNATGGY_UNKNOWN
HP_SUBMUCOUS_CLEFT_OF_SOFT_AND_HARD_PALATE, HP_SUBMUCOUS_CLEFT_OF_SOFT_AND_HARD_PALATE
GSE16385_ROSIGLITAZONE_IL4_VS_ROSIGLITAZONE_ALONE_STIM_MACROPHAGE_UP, GSE16385_ROSIGLITAZONE_IL4_VS_ROSIGLITAZONE_ALONE_STIM_MACROPHAGE_UP
HP_OP15THOTONUS, HP_OP15THOTONUS
HP_ADENOCARCINOMA_OF_THE_INTESTINES, HP_ADENOCARCINOMA_OF_THE_INTESTINES
GSE13522_WT_VS_JFNG_KO_SKING_T_CRUZI_Y_STRAIN_INF_DN, GSE13522_WT_VS_JFNG_KO_SKING_T_CRUZI_Y_STRAIN_INF_DN
GSE39152_SPLEEN_CD103_NEG_VS_BRAIN_CD103_POS_MEMORY_CD8_TCELL_DN, GSE39152_SPLEEN_CD103_NEG_VS_BRAIN_CD103_POS_MEMORY_CD8_TCELL_DN
MIR4445_5P, MIR4445_5P
GSE7460_FOXP3_MUT_VS_HET_ACT_TCONV_DN, GSE7460_FOXP3_MUT_VS_HET_ACT_TCONV_DN
HP_FOCAL_MOTOR_SEIZURE, HP_FOCAL_MOTOR_SEIZURE
HP_ADENOCARCINOMA_OF_THE_LARGE_INTESTINE, HP_ADENOCARCINOMA_OF_THE_LARGE_INTESTINE
GOBP_MITOTIC_G2_M_TRANSITION_CHECKPOINT, GOBP_MITOTIC_G2_M_TRANSITION_CHECKPOINT
GSE3994_WT_VS_PAC1_KO_ACTIVATED_MAST_CELL_UP, GSE3994_WT_VS_PAC1_KO_ACTIVATED_MAST_CELL_UP
ATAACCT_MIR154, ATAACCT_MIR154
LEE_CALORIE_RESTRICTION_MUSCLE_DN, LEE_CALORIE_RESTRICTION_MUSCLE_DN
HP_TRUNCAL_ATAXIA, HP_TRUNCAL_ATAXIA
HP_THICKENED_HELICES, HP_THICKENED_HELICES
BIOCARTA_CALCINEURIN_PATHWAY, BIOCARTA_CALCINEURIN_PATHWAY
WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_AUTOPHAGY, WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_AUTOPHAGY
HP_MALIGNANT_GENITOURINARY_TRACT_TUMOR, HP_MALIGNANT_GENITOURINARY_TRACT_TUMOR
BIOCARTA_MEF2D_PATHWAY, BIOCARTA_MEF2D_PATHWAY
GOBP_POSITIVE_REGULATION_OF_CHROMOSOME_SEPARATION, GOBP_POSITIVE_REGULATION_OF_CHROMOSOME_SEPARATION
AKL_HITLV1_INFECTION_DN, AKL_HITLV1_INFECTION_DN
GOMF_ACETYLGALUCOSAMINYTRANSFERASE_ACTIVITY, GOMF_ACETYLGALUCOSAMINYTRANSFERASE_ACTIVITY
CHEN_LVAD_SUPPORT_OF_FAILING_HEART_DN, CHEN_LVAD_SUPPORT_OF_FAILING_HEART_DN
HP_FOCAL_EMOTIONAL_SEIZURE, HP_FOCAL_EMOTIONAL_SEIZURE
HP_CENTRALLY_NUCLEATED_SKELETAL_MUSCLE_FIBERS, HP_CENTRALLY_NUCLEATED_SKELETAL_MUSCLE_FIBERS
MIR4777_3P, MIR4777_3P
HP_TYPICAL_ABSENCE_SEIZURE, HP_TYPICAL_ABSENCE_SEIZURE
HP_BIRTH_LENGTH_LESS_THAN_3RD_PERCENTILE, HP_BIRTH_LENGTH_LESS_THAN_3RD_PERCENTILE
GSE24210_IL35_TREATED_VS_RESTING_TREG_DN, GSE24210_IL35_TREATED_VS_RESTING_TREG_DN
GOBP_MATURATION_OF_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA, GOBP_MATURATION_OF_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA
GOBP_POSITIVE_REGULATION_OF_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE, GOBP_POSITIVE_REGULATION_OF_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE
WP_MITOCHONDRIAL_GENE_EXPRESSION, WP_MITOCHONDRIAL_GENE_EXPRESSION
HP_EYELID_MYOCLOLUS, HP_EYELID_MYOCLOLUS
GOBP_B_CELL_APOPTOTIC_PROCESS, GOBP_B_CELL_APOPTOTIC_PROCESS
SHETH_LIVER_CANCER_VS_TXNIP_LOSS_PAM6, SHETH_LIVER_CANCER_VS_TXNIP_LOSS_PAM6
CHEOK_RESPONSE_TO_MERCAPTOPURINE_AND_LD_MTX_DN, CHEOK_RESPONSE_TO_MERCAPTOPURINE_AND_LD_MTX_DN
HP_RETICULOCYTOPENIA, HP_RETICULOCYTOPENIA
ONO_AML1_TARGETS_DN, ONO_AML1_TARGETS_DN
HP_FOCAL_NON_MOTOR_SEIZURE, HP_FOCAL_NON_MOTOR_SEIZURE
GOBP_NEGATIVE_REGULATION_OF_LYMPHCYTE_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_LYMPHCYTE_APOPTOTIC_PROCESS
HP_ABNORMAL_URINE_CARBOHYDRATE_LEVEL, HP_ABNORMAL_URINE_CARBOHYDRATE_LEVEL
GOBP_HYPEROSMOTIC_RESPONSE, GOBP_HYPEROSMOTIC_RESPONSE
GSE13762_CTRL_VS_125_VITAMIND_DAYS_DC_DN, GSE13762_CTRL_VS_125_VITAMIND_DAYS_DC_DN
GOBP_REGULATION_OF_LYMPHCYTE_APOPTOTIC_PROCESS, GOBP_REGULATION_OF_LYMPHCYTE_APOPTOTIC_PROCESS
WP_GLYCEROLIPIDS_AND_GLYCEROPHOSPHOLIPIDS, WP_GLYCEROLIPIDS_AND_GLYCEROPHOSPHOLIPIDS
GOMF_5_3_EXONUCLEASE_ACTIVITY, GOMF_5_3_EXONUCLEASE_ACTIVITY
HP_CLITORAL_HYPOPLASIA, HP_CLITORAL_HYPOPLASIA
GOMF_ANNHEALING_ACTIVITY, GOMF_ANNHEALING_ACTIVITY
KEGG_NON_HOMOLOGOUS_END_JOINING, KEGG_NON_HOMOLOGOUS_END_JOINING
HP_PARTIAL_DUPLICATION_OF_THE_PHALANX_OF_HAND, HP_PARTIAL_DUPLICATION_OF_THE_PHALANX_OF_HAND
HP_APLASTIC_CLAVICLE, HP_APLASTIC_CLAVICLE
HP_HYPOPLASTIC_LABIA_MINORA, HP_HYPOPLASTIC_LABIA_MINORA
HP_METAPHYSEAL_IRREGULARITY, HP_METAPHYSEAL_IRREGULARITY
GOBP_POSITIVE_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION
GSE9946_LISTERIA_INF_MATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_DN, GSE9946_LISTERIA_INF_MATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_DN
chr9p24, chr9p24
HP_PREMATURE_THELARCHIE, HP_PREMATURE_THELARCHIE
CARDOSO_RESPONSE_TO_GAMMA_RADIATION_AND_3AB, Cardoso_RESPONSE_TO_GAMMA_RADIATION_AND_3AB
GOBP_RESPONSE_TO_HYDROPEROXIDE, GOBP_RESPONSE_TO_HYDROPEROXIDE
REACTOME_HDR_THROUGH_MMEJ_ALT_NHEJ, REACTOME_HDR_THROUGH_MMEJ_ALT_NHEJ
GOBP_REGULATION_OF_B_CELL_APOPTOTIC_PROCESS, GOBP_REGULATION_OF_B_CELL_APOPTOTIC_PROCESS
HP_ABNORMAL_GREATER_SCIATIC_NOTCH_MORPHOLOGY, HP_ABNORMAL_GREATER_SCIATIC_NOTCH_MORPHOLOGY
HP_APLASIA_HYPOPLASIA_OF_THE_TIBIA, HP_APLASIA_HYPOPLASIA_OF_THE_TIBIA
GOBP_MITOTIC_G2_DNA_DAMAGE_CHECKPOINT, GOBP_MITOTIC_G2_DNA_DAMAGE_CHECKPOINT
GOCC_ALVEOLAR_LAMELLAR_BODY, GOCC_ALVEOLAR_LAMELLAR_BODY
MODULE_375, MODULE_375
GOBP_BASE_EXCISION_REPAIR_AP_SITE_FORMATION, GOBP_BASE_EXCISION_REPAIR_AP_SITE_FORMATION
GOMF_CYTIDYLYLTRANSFERASE_ACTIVITY, GOMF_CYTIDYLYLTRANSFERASE_ACTIVITY
HP_BREAST_APLASIA, HP_BREAST_APLASIA
NOUSHMEHR_GBM_GERMLINE_MUTATED, NOUSHMEHR_GBM_GERMLINE_MUTATED
GOBP_POSITIVE_REGULATION_OF_MITOTIC_NUCLEAR_DIVISION, GOBP_POSITIVE_REGULATION_OF_MITOTIC_NUCLEAR_DIVISION
HP_ACROCYANOSIS, HP_ACROCYANOSIS
GOBP_GLUCCOSAMINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS, GOBP_GLUCCOSAMINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS
HP_MOVEMENT_ABNORMALITY_OF_THE_TONGUE, HP_MOVEMENT_ABNORMALITY_OF_THE_TONGUE
GOCC_MULTIVESICULAR_BODY_LUMEN, GOCC_MULTIVESICULAR_BODY_LUMEN
HP_Y_SHAPED_METACARPALS, HP_Y_SHAPED_METACARPALS
HP_RECURRENT_BACTERIAL_SKIN_INFECTIONS, HP_RECURRENT_BACTERIAL_SKIN_INFECTIONS
WP_SULINDAC_METABOLIC_PATHWAY, WP_SULINDAC_METABOLIC_PATHWAY
HP_HYPERMETHIONINEMIA, HP_HYPERMETHIONINEMIA
HP_APLASIA_HYPOPLASIA_OF_THE_BREASTS, HP_APLASIA_HYPOPLASIA_OF_THE_BREASTS
GOBP_REGULATION_OF_VESICLE_FUSION, GOBP_REGULATION_OF_VESICLE_FUSION
HP_ABNORMALITY_OF_CARTILAGE_OF_EXTERNAL_EAR, HP_ABNORMALITY_OF_CARTILAGE_OF_EXTERNAL_EAR
GOCC_FACIT_COLLAGEN_TRIMER, GOCC_FACIT_COLLAGEN_TRIMER
MIR4321, MIR4321
GOBP_DNA_REPLICATION_SYNTHESIS_OF_RNA_PRIMER, GOBP_DNA_REPLICATION_SYNTHESIS_OF_RNA_PRIMER
KANG_CISPLATIN_RESISTANCE_DN, KANG_CISPLATIN_RESISTANCE_DN
HP_SMALL_CEREBRAL_CORTEX, HP_SMALL_CEREBRAL_CORTEX
GOBP_POSITIVE_REGULATION_OF_LYMPHCYTE_MIGRATION, GOBP_POSITIVE_REGULATION_OF_LYMPHCYTE_MIGRATION
WP_OXIDATIVE_STRESS, WP_OXIDATIVE_STRESS
GOMF_CYCLOSPORIN_A_BINDING, GOMF_CYCLOSPORIN_A_BINDING
HP_CENTRAL_Y_SHAPED_METACARPAL, HP_CENTRAL_Y_SHAPED_METACARPAL
HP_HYPOPLASIA_OF_OLFACTORY_TRACT, HP_HYPOPLASIA_OF_OLFACTORY_TRACT
HP_DELAYED_MENARCHE, HP_DELAYED_MENARCHE
GOBP_POSITIVE_REGULATION_OF_AXON_EXTENSION_INVOLVED_IN_AXON_GUIDANCE, GOBP_POSITIVE_REGULATION_OF_AXON_EXTENSION_INVOLVED_IN_AXON_GUIDANCE
GOBP_RIBONUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS, GOBP_RIBONUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS
REACTOME_APEXI_INDEPENDENT_RESOLUTION_OF_AP_SITES_VIA_THE_SINGLE_NUCLEOTIDE_REPLACEMENT_PATHWAY, REACTOME_APEXI_INDEPENDENT_RESOLUTION_OF_AP_SITES_VIA_THE_SINGLE_NUCLEOTIDE_REPLACEMENT_PATHWAY
MODULE_289, MODULE_289
HP_LOBULATED_TONGUE, HP_LOBULATED_TONGUE
GOBP_POSITIVE_REGULATION_OF_T_CELL_MIGRATION, GOBP_POSITIVE_REGULATION_OF_T_CELL_MIGRATION
GOCC_LATE_ENDOSOME_LUMEN, GOCC_LATE_ENDOSOME_LUMEN
MIR1273C, MIR1273C
FRASOR_TAMOXIFEN_RESPONSE_DN, FRASOR_TAMOXIFEN_RESPONSE_DN
HP_POLYCLONAL_ELEVATION_OF_IGM, HP_POLYCLONAL_ELEVATION_OF_IGM
HP_OVERHANGING_NASAL_TIP, HP_OVERHANGING_NASAL_TIP
HP_BIFID_NOSE, HP_BIFID_NOSE
GOBP_REGULATION_OF_T_CELL_MIGRATION, GOBP_REGULATION_OF_T_CELL_MIGRATION
GOBP_IMMUNOLOGICAL_SYNAPSE_FORMATION, GOBP_IMMUNOLOGICAL_SYNAPSE_FORMATION
GCM_PTTRU, GCM_PTTRU
PID_VEGF_VEGFR_PATHWAY, PID_VEGF_VEGFR_PATHWAY
GOMF_SUPEROXIDE_GENERATING_NADPH_OXIDASE_ACTIVATOR_ACTIVITY, GOMF_SUPEROXIDE_GENERATING_NADPH_OXIDASE_ACTIVATOR_ACTIVITY
GOBP_NUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS, GOBP_NUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS
HP_HYPOTHALAMIC_HAMARTOMA, HP_HYPOTHALAMIC_HAMARTOMA
GOBP_N_ACETYLNEURAMINATE_CATABOLIC_PROCESS, GOBP_N_ACETYLNEURAMINATE_CATABOLIC_PROCESS
HP_DECREASED_HIP_ABDUCTION, HP_DECREASED_HIP_ABDUCTION
HP_MACULAR_ATROPHY, HP_MACULAR_ATROPHY
GSE13485_CTRL_VS_DAY21_YF17D_VACCINE_PBMC_DN, GSE13485_CTRL_VS_DAY21_YF17D_VACCINE_PBMC_DN
GOBP_AXONOGENESIS_INVOLVED_IN_INNervation, GOBP_AXONOGENESIS_INVOLVED_IN_INNervation
HP_ASTERIXIS, HP_ASTERIXIS
GOBP_DNA_LIGATION_INVOLVED_IN_DNA_REPAIR, GOBP_DNA_LIGATION_INVOLVED_IN_DNA_REPAIR
GOCC_ALPHA_DNA_POLYMERASE_PRIMASE_COMPLEX, GOCC_ALPHA_DNA_POLYMERASE_PRIMASE_COMPLEX
GOCC_AP_4_ADAPTOR_COMPLEX, GOCC_AP_4_ADAPTOR_COMPLEX
HP_ABNORMALITY_OF_THE_CRUS_OF_THE_HELIX, HP_ABNORMALITY_OF_THE_CRUS_OF_THE_HELIX
GOBP_PENETRATION_OF_ZONA_PELLUCIDA, GOBP_PENETRATION_OF_ZONA_PELLUCIDA
GOBP_POSITIVE_REGULATION_OF_RECEPTOR_SIGNALING_PATHWAY_VIA_STAT, GOBP_POSITIVE_REGULATION_OF_RECEPTOR_SIGNALING_PATHWAY_VIA_STAT
GSE11367_CTRL_VS_IL17_TREATED_SMOOTH_MUSCLE_CELL_UP, GSE11367_CTRL_VS_IL17_TREATED_SMOOTH_MUSCLE_CELL_UP
GOBP_LONG_CHAIN_FATTY_ACID_IMPORT_ACROSS_PLASMA_MEMBRANE, GOBP_LONG_CHAIN_FATTY_ACID_IMPORT_ACROSS_PLASMA_MEMBRANE
HP_TRACHEAL_STENOSIS, HP_TRACHEAL_STENOSIS
GOBP_OSTEOCLAST_FUSION, GOBP_OSTEOCLAST_FUSION
HP_BIFID_UTERUS, HP_BIFID_UTERUS
GOMF_HMG_BOX_DOMAIN_BINDING, GOMF_HMG_BOX_DOMAIN_BINDING
ZHANG_GATA6_TARGETS_UP, ZHANG_GATA6_TARGETS_UP
CAR_TNFRSF25, CAR_TNFRSF25
LIU_TARGETS_OF_VMYB_VS_CMYB_UP, LIU_TARGETS_OF_VMYB_VS_CMYB_UP
WP_TCELL_RECEPTOR_AND_COSTIMULATORY_SIGNALING, WP_TCELL_RECEPTOR_AND_COSTIMULATORY_SIGNALING
GOCC_FILTRATION_DIAPHRAGM, GOCC_FILTRATION_DIAPHRAGM
GOBP_POSITIVE_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATION, GOBP_POSITIVE_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATION
GOMF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY, GOMF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY
GOBP_HISTAMINE_PRODUCTION_INVOLVED_IN_INFLAMMATORY_RESPONSE, GOBP_HISTAMINE_PRODUCTION_INVOLVED_IN_INFLAMMATORY_RESPONSE
HP_ABNORMALITY_OF_THE_DENTATE_NUCLEUS, HP_ABNORMALITY_OF_THE_DENTATE_NUCLEUS
HP_MONOCHROMACY, HP_MONOCHROMACY
HP_RECURRENT_ENTEROVIRAL_INFECTIONS, HP_RECURRENT_ENTEROVIRAL_INFECTIONS
GOBP_REGULATION_OF_RETINAL_GANGLION_CELL_AXON_GUIDANCE, GOBP_REGULATION_OF_RETINAL_GANGLION_CELL_AXON_GUIDANCE
GOMF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_BINDING, GOMF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_BINDING
GOBP_ASPARAGINE_METABOLIC_PROCESS, GOBP_ASPARAGINE_METABOLIC_PROCESS
GOBP_POLYPHOSPHATE_CATABOLIC_PROCESS, GOBP_POLYPHOSPHATE_CATABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_NLRP3_INFLAMMASOME_COMPLEX_ASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_NLRP3_INFLAMMASOME_COMPLEX_ASSEMBLY
GOMF_EXOPOLYPHOSPHATASE_ACTIVITY, GOMF_EXOPOLYPHOSPHATASE_ACTIVITY
HP_NEOPLASM_OF_THE_GALLBLADDER, HP_NEOPLASM_OF_THE_GALLBLADDER
GOBP_CORTICOSTERONE_SECRETION, GOBP_CORTICOSTERONE_SECRETION
REACTOME_SIGNALLING_TO_P38_VIA_RIT_AND_RIN, REACTOME_SIGNALLING_TO_P38_VIA_RIT_AND_RIN
GOBP_MULTINUCLEAR_OSTEOCLAST_DIFFERENTIATION, GOBP_MULTINUCLEAR_OSTEOCLAST_DIFFERENTIATION
HP_OSTEOCHONDROSIS, HP_OSTEOCHONDROSIS
REACTOME_METABOLISM_OF_INGESTED_SEMET_SEC_MESEC_INTO_H2SE, REACTOME_METABOLISM_OF_INGESTED_SEMET_SEC_MESEC_INTO_H2SE
REACTOME_TRANSPORT_OF_FATTY_ACIDS, REACTOME_TRANSPORT_OF_FATTY_ACIDS
HP_ABNORMAL_CIRCULATING_AROMATIC_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_AROMATIC_AMINO_ACID_CONCENTRATION
SHIRAISHI_PLZF_TARGETS_UP, SHIRAISHI_PLZF_TARGETS_UP
HP_RECURRENT_CANDIDA_INFECTIONS, HP_RECURRENT_CANDIDA_INFECTIONS
GOBP_BEHAVIORAL_RESPONSE_TO_PAIN, GOBP_BEHAVIORAL_RESPONSE_TO_PAIN
GOBP_NEURAL_CREST_CELL_MIGRATION_INVOLVED_IN_AUTONOMIC_NERVOUS_SYSTEM_DEVELOPMENT, GOBP_NEURAL_CREST_CELL_MIGRATION_INVOLVED_IN_AUTONOMIC_NERVOUS_SYSTEM_DEVELOPMENT
MIR95_3P, MIR95_3P
GOBP_POST_CHAPERONIN_TUBULIN_FOLDING_PATHWAY, GOBP_POST_CHAPERONIN_TUBULIN_FOLDING_PATHWAY
GOBP_PENTOSE_CATABOLIC_PROCESS, GOBP_PENTOSE_CATABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_INTERFERON_ALPHA_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_INTERFERON_ALPHA_PRODUCTION
GOBP_MEIOTIC_DNA_DOUBLE_STRAND_BREAK_FORMATION, GOBP_MEIOTIC_DNA_DOUBLE_STRAND_BREAK_FORMATION
REACTOME_PROTEIN_REPAIR, REACTOME_PROTEIN_REPAIR
MIR887_3P, MIR887_3P
GOBP_HISTAMINE_METABOLIC_PROCESS, GOBP_HISTAMINE_METABOLIC_PROCESS
GOBP_HYDROGEN_SULFIDE_BIOSYNTHETIC_PROCESS, GOBP_HYDROGEN_SULFIDE_BIOSYNTHETIC_PROCESS
GOBP_REGULATION_OF_TISSUE_REMODELING, GOBP_REGULATION_OF_TISSUE_REMODELING
WP_SEROTONIN_AND_ANGIETYRELATED_EVENTS, WP_SEROTONIN_AND_ANGIETYRELATED_EVENTS
GOBP_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATION, GOBP_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATION
HP_ABNORMAL_LIGHT_ADAPTED_ELECTRORETINOGRAM, HP_ABNORMAL_LIGHT_ADAPTED_ELECTRORETINOGRAM
HP_PROSTATITIS, HP_PROSTATITIS
HP_ECCENTRIC_VISUAL_FIXATION, HP_ECCENTRIC_VISUAL_FIXATION
HP_ABNORMAL_PUPILLARY_LIGHT_REFLEX, HP_ABNORMAL_PUPILLARY_LIGHT_REFLEX
GOBP_HISTAMINE_SECRETION, GOBP_HISTAMINE_SECRETION
HP_HYPOPLASIA_OF_THE_FOVEA, HP_HYPOPLASIA_OF_THE_FOVEA
GOBP_HISTAMINE_TRANSPORT, GOBP_HISTAMINE_TRANSPORT
GOBP_REGULATION_OF_CONNECTIVE_TISSUE_REPLACEMENT, GOBP_REGULATION_OF_CONNECTIVE_TISSUE_REPLACEMENT
GOBP_MOTOR_NEURON_MIGRATION, GOBP_MOTOR_NEURON_MIGRATION
HP_PYODERMA, HP_PYODERMA
HP_ACHROMATOPSIA, HP_ACHROMATOPSIA
GOBP_DENDRITE_ARBORIZATION, GOBP_DENDRITE_ARBORIZATION
ZELISTR_A_CD44_TARGETS_UP, ZELISTR_A_CD44_TARGETS_UP
HP_ABNORMALITY_OF_BODY_MASS_INDEX, HP_ABNORMALITY_OF_BODY_MASS_INDEX
GOBP_HYDROGEN_SULFIDE_METABOLIC_PROCESS, GOBP_HYDROGEN_SULFIDE_METABOLIC_PROCESS
WP_GANGLIOSPHINGOLIPID_METABOLISM, WP_GANGLIOSPHINGOLIPID_METABOLISM
GRAHAM_CML_DIVIDING_VS_NORMAL_DIVIDING_UP, GRAHAM_CML_DIVIDING_VS_NORMAL_DIVIDING_UP
GOMF_PHOSPHOTRANSFERASE_ACTIVITY_NITROGENOUS_GROUP_AS_ACCEPTOR, GOMF_PHOSPHOTRANSFERASE_ACTIVITY_NITROGENOUS_GROUP_AS_ACCEPTOR