```
GSE22935 24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_DN, GSE22935 24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_DN
 GSE35685_CD34POS_CD38NEG_VS_CD34POS_CD10NEG_CD62LPOS_BONE_MARROW_UP, GSE35685_CD34POS_CD38NEG_VS_CD34POS_CD10NEG_CD62LPOS_BONE_MARROW_UP
GSE21063 WT VS NFATC1 KO 8H ANTI IGM STIM BCELL DN, GSE21063 WT VS NFATC1 KO 8H ANTI IGM STIM BCELL DN
GSE17721_POLYIC_VS_CPG_2H_BMDC_UP, GSE17721_POLYIC_VS_CPG_2H_BMDC_UP
GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP
GSE17721_4H_VS_24H_POLYIC_BMDC_DN, GSE17721_4H_VS_24H_POLYIC_BMDC_DN
GOLDRATH_EFF_VS_MEMORY_CD8_TCELL_DN, GOLDRATH_EFF_VS_MEMORY_CD8_TCELL_DN
GSE14908_RESTING_VS_HDM_STIM_CD4_TCELL_ATOPIC_PATIENT_DN, GSE14908_RESTING_VS_HDM_STIM_CD4_TCELL_ATOPIC_PATIENT_DN
GSE17721_PAM3CSK4_VS_CPG_4H_BMDC_UP, GSE17721_PAM3CSK4_VS_CPG_4H_BMDC_UP
GSE17721_4_VS_24H_CPG_BMDC_DN, GSE17721_4_VS_24H_CPG_BMDC_DN
GSE17721_CPG_VS_GARDIQUIMOD_6H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_6H_BMDC_DN
GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN
GSE43956_WT_VS_SGK1_KO_TH17_DIFFERENTIATED_CD4_TCELL_DN, GSE43956_WT_VS_SGK1_KO_TH17_DIFFERENTIATED_CD4_TCELL_DN
GSE46606_UNSTIM_VS_CD40L_IL2_IL5_3DAY_STIMULATED_IRF4HIGH_SORTED_BCELL_UP, GSE46606_UNSTIM_VS_CD40L_IL2_IL5_3DAY_STIMULATED_IRF4HIGH_SORTED_BCELL_UP
GSE43955_TGFB_IL6_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_DN, GSE43955_TGFB_IL6_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_DN
GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_VS_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_INH_PRETREAT_AND_ACT_BY_A3R_INH_PRETREAT_AND_ACT_BY_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_PRETREAT_AND_ACT_BY_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE1988_ADENOSINE_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE1988_ADENOSINE_A3R_INH_AND_TCELL_MEMBRANES_ACT_MAST_CELL_DN, GSE1988_ADENOSINE_A3R_INH_AND_TCELL_MEMBRANES_ACT_A
GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_2H_DN
GSE7568_CTRL_VS_24H_TGFB_TREATED_MACROPHAGES_WITH_IL4_AND_DEXAMETHASONE_UP, GSE7568_CTRL_VS_24H_TGFB_TREATED_MACROPHAGES_WITH_IL4_AND_DEXAMETHASONE_UP
GSE32128_INOS_DEPENDENT_VS_INOS_INDEPENDENT_ACTIVATED_TCELL_UP, GSE32128_INOS_DEPENDENT_VS_INOS_INDEPENDENT_ACTIVATED_TCELL_UP
GSE44649_WT_VS_MIR155_KO_NAIVE_CD8_TCELL_DN, GSE44649_WT_VS_MIR155_KO_NAIVE_CD8_TCELL_DN
GOBP_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI, GOBP_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI
GSE17721_POLYIC_VS_GARDIQUIMOD_1H_BMDC_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_1H_BMDC_UP
GSE5503_MLN_DC_VS_PLN_DC_ACTIVATED_ALLOGENIC_TCELL_DN, GSE5503_MLN_DC_VS_PLN_DC_ACTIVATED_ALLOGENIC_TCELL_DN
GSE37301_MULTIPOTENT_PROGENITOR_VS_PRO_BCELL_UP, GSE37301_MULTIPOTENT_PROGENITOR_VS_PRO_BCELL_UP
GSE17721 LPS_VS_CPG_1H_BMDC_DN, GSE17721 LPS_VS_CPG_1H_BMDC_DN
PECE_MAMMARY_STEM_CELL_DN, PECE_MAMMARY_STEM_CELL_DN
GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_60H_UP, GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_60H_UP
HP_HIGH_NARROW_PALATE, HP_HIGH_NARROW_PALATE
GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN, GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN
GSE21546_WT_VS_ELK1_KO_ANTI_CD3_STIM_DP_THYMOCYTES_UP, GSE21546_WT_VS_ELK1_KO_ANTI_CD3_STIM_DP_THYMOCYTES_UP
MIR3692_5P, MIR3692_5P
HP_THICK_LOWER_LIP_VERMILION, HP_THICK_LOWER_LIP_VERMILION
ATAAGCT_MIR21, ATAAGCT_MIR21
MIR503_5P, MIR503_5P
GSE41978_WT_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN, GSE41978_WT_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN
GSE45365_CD8A_DC_VS_CD11B_DC_IFNAR_KO_MCMV_INFECTION_UP, GSE45365_CD8A_DC_VS_CD11B_DC_IFNAR_KO_MCMV_INFECTION_UP
CHAF1A_TARGET_GENES, CHAF1A_TARGET_GENES
MIR216B_3P, MIR216B_3P
MCBRYAN_PUBERTAL_BREAST_5_6WK_UP, MCBRYAN_PUBERTAL_BREAST_5_6WK_UP
HP_SEVERE_SHORT_STATURE, HP_SEVERE_SHORT_STATURE
AGGGCCA_MIR328, AGGGCCA_MIR328
REACTOME_RUNX1_INTERACTS_WITH_CO_FACTORS_WHOSE_PRECISE_EFFECT_ON_RUNX1_TARGETS_IS_NOT_KNOWN, REACTOME_RUNX1_INTERACTS_IS_NOT_KNOWN, REACTOME_RUNX1_INTERACTS_IS
GOBP_ENDOCYTIC_RECYCLING, GOBP_ENDOCYTIC_RECYCLING
GSE17721_CTRL_VS_CPG_1H_BMDC_DN, GSE17721_CTRL_VS_CPG_1H_BMDC_DN
GSE37416_CTRL_VS_0H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_CTRL_VS_0H_F_TULARENSIS_LVS_NEUTROPHIL_UP
WP_SMALL_CELL_LUNG_CANCER, WP_SMALL_CELL_LUNG_CANCER
HAY_BONE_MARROW_CD34_POS_PRE_B, HAY_BONE_MARROW_CD34_POS_PRE_B
REACTOME SYNTHESIS OF PIPS AT THE PLASMA MEMBRANE, REACTOME SYNTHESIS OF PIPS AT THE PLASMA MEMBRANE
GOBP_HISTONE_H3_K9_METHYLATION, GOBP_HISTONE_H3_K9_METHYLATION
CYCLIN_D1_UP.V1_DN, CYCLIN_D1_UP.V1_DN
REACTOME_NOD1_2_SIGNALING_PATHWAY, REACTOME_NOD1_2_SIGNALING_PATHWAY
GOBP_POSITIVE_REGULATION_OF_ENDOCYTOSIS, GOBP_POSITIVE_REGULATION_OF_ENDOCYTOSIS
KRAS.DF.V1_DN, KRAS.DF.V1_DN
GOBP_OSTEOCLAST_DIFFERENTIATION, GOBP_OSTEOCLAST_DIFFERENTIATION
MIR6893_3P, MIR6893_3P
MIR370 3P, MIR370 3P
GOBP_REGULATION_OF_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION, GOBP_REGULATION_OF_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION
MIR6818_5P, MIR6818_5P
HP_ABNORMAL_PULMONARY_THORACIC_IMAGING_FINDING, HP_ABNORMAL_PULMONARY_THORACIC_IMAGING_FINDING
GOBP_GOLGI_TO_ENDOSOME_TRANSPORT, GOBP_GOLGI_TO_ENDOSOME_TRANSPORT
 HP_SHORT_PALM, HP_SHORT_PALM
GOBP_HISTONE_H3_K9_MODIFICATION, GOBP_HISTONE_H3_K9_MODIFICATION
MIR219B_3P, MIR219B_3P
GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_UP, GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_UP
GOBP_POSITIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION
MIR3199, MIR3199
 GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_DERIVED_MACROPHAGE_UP, GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_DERIVED_MACROPHAGE_UP
GOBP_REGULATION_OF_RECEPTOR_RECYCLING, GOBP_REGULATION_OF_RECEPTOR_RECYCLING
 HP_BREAST_APLASIA, HP_BREAST_APLASIA
 MIR193B_5P, MIR193B_5P
 WANG_IMMORTALIZED_BY_HOXA9_AND_MEIS1_DN, WANG_IMMORTALIZED_BY_HOXA9_AND_MEIS1_DN
 KANG_CISPLATIN_RESISTANCE_UP, KANG_CISPLATIN_RESISTANCE_UP
 ZWANG_EGF_PERSISTENTLY_DN, ZWANG_EGF_PERSISTENTLY_DN
 MIR6808_3P, MIR6808_3P
 GOBP_MYELOID_CELL_DEVELOPMENT, GOBP_MYELOID_CELL_DEVELOPMENT
 MIR8052, MIR8052
 GOBP_REGULATION_OF_NUCLEOTIDE_BINDING_OLIGOMERIZATION_DOMAIN_CONTAINING_SIGNALING_PATHWAY, GOBP_REGULATION_OF_NUCLEOTIDE_BINDING_OLIGOMERIZATION_DOMAIN_CONTAINING_SIGNALING
 GOBP_POSITIVE_REGULATION_OF_RECEPTOR_RECYCLING, GOBP_POSITIVE_REGULATION_OF_RECEPTOR_RECYCLING
 WP_FATTY_ACID_BETA_OXIDATION, WP_FATTY_ACID_BETA_OXIDATION
 CYCLIN_D1_KE_.V1_DN, CYCLIN_D1_KE_.V1_DN
 GOBP_POSITIVE_REGULATION_OF_TISSUE_REMODELING, GOBP_POSITIVE_REGULATION_OF_TISSUE_REMODELING
 HP_ABNORMAL_LABIA_MINORA_MORPHOLOGY, HP_ABNORMAL_LABIA_MINORA_MORPHOLOGY
 GOBP_POSITIVE_REGULATION_OF_RECEPTOR_MEDIATED_ENDOCYTOSIS, GOBP_POSITIVE_REGULATION_OF_RECEPTOR_MEDIATED_ENDOCYTOSIS
 GROSS_ELK3_TARGETS_UP, GROSS_ELK3_TARGETS_UP
 HP_SUDDEN_EPISODIC_APNEA, HP_SUDDEN_EPISODIC_APNEA
 HP_MANDIBULAR_APLASIA, HP_MANDIBULAR_APLASIA
HP_APLASIA_HYPOPLASIA_OF_THE_TONGUE, HP_APLASIA_HYPOPLASIA_OF_THE_TONGUE
GOBP_MONOUBIQUITINATED_PROTEIN_DEUBIQUITINATION, GOBP_MONOUBIQUITINATED_PROTEIN_DEUBIQUITINATION
BIOCARTA_MITOCHONDRIA_PATHWAY, BIOCARTA_MITOCHONDRIA_PATHWAY
GOBP_PROTEIN_LOCALIZATION_TO_CILIARY_MEMBRANE, GOBP_PROTEIN_LOCALIZATION_TO_CILIARY_MEMBRANE
 REACTOME_TICAM1_RIP1_MEDIATED_IKK_COMPLEX_RECRUITMENT, REACTOME_TICAM1_RIP1_MEDIATED_IKK_COMPLEX_RECRUITMENT
 GOBP REGULATION_OF_MICROTUBULE_DEPOLYMERIZATION, GOBP_REGULATION_OF_MICROTUBULE_DEPOLYMERIZATION
 MIR503_3P, MIR503_3P
 GENTILE_UV_LOW_DOSE_UP, GENTILE_UV_LOW_DOSE_UP
 MIKKELSEN_PLURIPOTENT_STATE_DN, MIKKELSEN_PLURIPOTENT_STATE_DN
 GSE12839_CTRL_VS_IL12_TREATED_PBMC_DN, GSE12839_CTRL_VS_IL12_TREATED_PBMC_DN
 REACTOME SUMOYLATION OF DNA METHYLATION PROTEINS, REACTOME SUMOYLATION OF DNA METHYLATION PROTEINS
 HP_BRUXISM, HP_BRUXISM
 PAPASPYRIDONOS_UNSTABLE_ATEROSCLEROTIC_PLAQUE_DN, PAPASPYRIDONOS_UNSTABLE_ATEROSCLEROTIC_PLAQUE_DN
 GOBP_EXTRACELLULAR_MATRIX_ASSEMBLY, GOBP_EXTRACELLULAR_MATRIX_ASSEMBLY
 GOBP_NOTCH_RECEPTOR_PROCESSING, GOBP_NOTCH_RECEPTOR_PROCESSING
 GOBP_REGULATION_OF_TISSUE_REMODELING, GOBP_REGULATION_OF_TISSUE_REMODELING
 MODULE_537, MODULE_537
 HP_HYPOPLASTIC_LABIA_MINORA, HP_HYPOPLASTIC_LABIA_MINORA
 GOBP_CELLULAR_RESPONSE_TO_VITAMIN_D, GOBP_CELLULAR_RESPONSE_TO_VITAMIN_D
 GOBP_OSTEOCLAST_DEVELOPMENT, GOBP_OSTEOCLAST_DEVELOPMENT
ZNF418_TARGET_GENES, ZNF418_TARGET_GENES
 GCGCCTT_MIR525_MIR524, GCGCCTT_MIR525_MIR524
 GOBP_POSITIVE_REGULATION_OF_MYELOID_LEUKOCYTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MYELOID_LEUKOCYTE_DIFFERENTIATION
 HP_MACROTHROMBOCYTOPENIA, HP_MACROTHROMBOCYTOPENIA
 GOBP_REGULATION_OF_IRON_ION_TRANSPORT, GOBP_REGULATION_OF_IRON_ION_TRANSPORT
 GOBP REGULATION OF OSTEOCLAST DIFFERENTIATION, GOBP REGULATION OF OSTEOCLAST DIFFERENTIATION
REACTOME_TNF_RECEPTOR_SUPERFAMILY_TNFSF_MEMBERS_MEDIATING_NON_CANONICAL_NF_KB_PATHWAY, REACTOME_TNF_RECEPTOR_SUPERFAMILY_TNFSF_MEMBERS_MEDIATING_NON_CANONICAL_NF_KB_PATHWAY
MIR2277_3P, MIR2277_3P
GOBP_STRESS_RESPONSE_TO_METAL_ION, GOBP_STRESS_RESPONSE_TO_METAL_ION
GOBP CELLULAR RESPONSE TO NUTRIENT, GOBP CELLULAR RESPONSE TO NUTRIENT
GOBP_CELLULAR_RESPONSE_TO_VITAMIN, GOBP_CELLULAR_RESPONSE_TO_VITAMIN
 HP_URETERAL_OBSTRUCTION, HP_URETERAL_OBSTRUCTION
HP_FASTING_HYPOGLYCEMIA, HP_FASTING_HYPOGLYCEMIA
HP_ABNORMAL_URINE_MAGNESIUM_CONCENTRATION, HP_ABNORMAL_URINE_MAGNESIUM_CONCENTRATION
GOBP_CELLULAR_RESPONSE_TO_COPPER_ION, GOBP_CELLULAR_RESPONSE_TO_COPPER_ION
HP_HORIZONTAL_EYEBROW, HP_HORIZONTAL_EYEBROW
HP_SPONTANEOUS_HEMATOMAS, HP_SPONTANEOUS_HEMATOMAS
GOBP_BONE_CELL_DEVELOPMENT, GOBP_BONE_CELL_DEVELOPMENT
DESCARTES_MAIN_FETAL_UNIPOLAR_BRUSH_CELLS, DESCARTES_MAIN_FETAL_UNIPOLAR_BRUSH_CELLS
REACTOME_SIGNALING_BY_NOTCH1_T_7_9_NOTCH1_M1580_K2555_TRANSLOCATION_MUTANT, REACTOME_SIGNALING_BY_NOTCH1_T_7_9_NOTCH1_M1580_K2555_TRANSLOCATION_MUTANT
GOBP_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS, GOBP_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS
GOBP_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE, GOBP_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE
GOBP_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE, GOBP_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE
GOBP_REGULATION_OF_VIRAL_INDUCED_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_VIRAL_INDUCED_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY.
HP_BIFID_TONGUE, HP_BIFID_TONGUE
ERCC8_TARGET_GENES, ERCC8_TARGET_GENES
REACTOME_REGULATED_PROTEOLYSIS_OF_P75NTR, REACTOME_REGULATED_PROTEOLYSIS_OF_P75NTR
HP_OLIGURIA, HP_OLIGURIA
GOBP_ENDODERMAL_CELL_FATE_SPECIFICATION, GOBP_ENDODERMAL_CELL_FATE_SPECIFICATION
REACTOME_MECP2_REGULATES_TRANSCRIPTION_FACTORS, REACTOME_MECP2_REGULATES_TRANSCRIPTION_FACTORS
HP_CAPITATE_HAMATE_FUSION, HP_CAPITATE_HAMATE_FUSION
GOBP_MEMBRANE_RAFT_ORGANIZATION, GOBP_MEMBRANE_RAFT_ORGANIZATION
HOFMANN_MYELODYSPLASTIC_SYNDROM_LOW_RISK_UP, HOFMANN_MYELODYSPLASTIC_SYNDROM_LOW_RISK_UP
GOBP_REGULATION_OF_FEEDING_BEHAVIOR, GOBP_REGULATION_OF_FEEDING_BEHAVIOR
GOBP_DETOXIFICATION_OF_INORGANIC_COMPOUND, GOBP_DETOXIFICATION_OF_INORGANIC_COMPOUND
HP_POSTERIOR_UVEITIS, HP_POSTERIOR_UVEITIS
HP_PILL_ROLLING_TREMOR, HP_PILL_ROLLING_TREMOR
GOMF_THROMBIN_ACTIVATED_RECEPTOR_ACTIVITY, GOMF_THROMBIN_ACTIVATED_RECEPTOR_ACTIVITY
ALONSO_METASTASIS_DN, ALONSO_METASTASIS_DN
HP_PANCREATIC_CALCIFICATION, HP_PANCREATIC_CALCIFICATION
MIR4730, MIR4730
HP_RETINAL_EXUDATE, HP_RETINAL_EXUDATE
GOBP_NEGATIVE_REGULATION_OF_PROGRAMMED_NECROTIC_CELL_DEATH, GOBP_NEGATIVE_REGULATION_OF_PROGRAMMED_NECROTIC_CELL_DEATH
HP_HAND_APRAXIA, HP_HAND_APRAXIA
HP_HYPERMAGNESIURIA, HP_HYPERMAGNESIURIA
STARK_BRAIN_22Q11_DELETION, STARK_BRAIN_22Q11_DELETION
GALI_TP53_TARGETS_APOPTOTIC_UP, GALI_TP53_TARGETS_APOPTOTIC_UP
GOBP_NEGATIVE_REGULATION_OF_NECROTIC_CELL_DEATH, GOBP_NEGATIVE_REGULATION_OF_NECROTIC_CELL_DEATH
HP_BROAD_CLAVICLES, HP_BROAD_CLAVICLES
HP_LONG_NECK, HP_LONG_NECK
GOBP_RECEPTOR_TRANSACTIVATION, GOBP_RECEPTOR_TRANSACTIVATION
GOBP_AMINO_ACID_NEUROTRANSMITTER_REUPTAKE, GOBP_AMINO_ACID_NEUROTRANSMITTER_REUPTAKE
BONOME_OVARIAN_CANCER_POOR_SURVIVAL_UP, BONOME_OVARIAN_CANCER_POOR_SURVIVAL_UP
MIR6125, MIR6125
GOBP_RESPONSE_TO_VITAMIN_D, GOBP_RESPONSE_TO_VITAMIN_D
GOMF_BILE_ACID_SODIUM_SYMPORTER_ACTIVITY, GOMF_BILE_ACID_SODIUM_SYMPORTER_ACTIVITY
HP_OSTEOMALACIA, HP_OSTEOMALACIA
HP_ABNORMAL_CIRCULATING_RENIN, HP_ABNORMAL_CIRCULATING_RENIN
GOBP_MESENCHYMAL_TO_EPITHELIAL_TRANSITION, GOBP_MESENCHYMAL_TO_EPITHELIAL_TRANSITION
HP_CHONDROCALCINOSIS, HP_CHONDROCALCINOSIS
HP NAIL PITS, HP NAIL PITS
REACTOME_DEFECTIVE_FACTOR_IX_CAUSES_HEMOPHILIA_B, REACTOME_DEFECTIVE_FACTOR_IX_CAUSES_HEMOPHILIA_B
GOBP_DETECTION_OF_TEMPERATURE_STIMULUS_INVOLVED_IN_SENSORY_PERCEPTION, GOBP_DETECTION_OF_TEMPERATURE_STIMULUS_INVOLVED_IN_SENSORY_PERCEPTION
GOBP_PROPRIOCEPTION, GOBP_PROPRIOCEPTION
GOBP_POSITIVE_REGULATION_OF_MESENCHYMAL_STEM_CELL_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_MESENCHYMAL_STEM_CELL_PROLIFERATION
GOBP_MESENCHYMAL_STEM_CELL_PROLIFERATION, GOBP_MESENCHYMAL_STEM_CELL_PROLIFERATION
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GOBP_REGULATION_OF_RECEPTOR_MEDIATED_ENDOCYTOSIS_INVOLVED_IN_CHOLESTEROL_TRANSPORT, GOBP_REGULATION_OF_RECEPTOR_MEDIATED_ENDOCYTOSIS_INVOLVED_IN_CHOLESTEROL_TRANSPORT

GOBP_DETECTION_OF_TEMPERATURE_STIMULUS, GOBP_DETECTION_OF_TEMPERATURE_STIMULUS

GOMF_BILE_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_BILE_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY

REACTOME_DEFECTIVE_F9_ACTIVATION, REACTOME_DEFECTIVE_F9_ACTIVATION

GOBP_CHEMOSENSORY_BEHAVIOR, GOBP_CHEMOSENSORY_BEHAVIOR

KO_VS_WT_NEUTROPHIL_DN, GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_DN