ARMSTRONG_EFFECTOR_CD8_TCELL_DN, GSE41867_DAY6_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN
GSE21360_SECONDARY_VS_TERTIARY_MEMORY_CD8_TCELL_UP, GSE21360_SECONDARY_VS_TERTIARY_MEMORY_CD8_TCELL_UP
GSE41867_LCMV_ARMSTRONG_VS_CLONE13_DAY6_EFFECTOR_CD8_TCELL_DN, GSE41867_LCMV_ARMSTRONG_VS_CLONE13_DAY6_EFFECTOR_CD8_TCELL_DN
GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_DN, GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_DN
GSE22432_PDC_VS_TGFB1_TREATEDCOMMON_DC_PROGENITOR_UP, GSE22432_PDC_VS_TGFB1_TREATEDCOMMON_DC_PROGENITOR_UP
GSE43863_NAIVE_VS_MEMORY_LY6C_INT_CXCR5POS_CD4_TCELL_D150_LCMV_DN, GSE43863_NAIVE_VS_MEMORY_LY6C_INT_CXCR5POS_CD4_TCELL_D150_LCMV_DN
GSE6674_ANTI_IGM_VS_PL2_3_STIM_BCELL_DN, GSE6674_ANTI_IGM_VS_PL2_3_STIM_BCELL_DN
GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_UP, GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_UP
GSE17721_CTRL_VS_POLYIC_4H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_4H_BMDC_UP
GSE24671_BAKIMULC_VS_SENDAI_VIRUS_INFECTED_MOUSE_SPLENOCYTES_DN, GSE24671_BAKIMULC_VS_SENDAI_VIRUS_INFECTED_MOUSE_SPLENOCYTES_DN
GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN, GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN
GSE17974_0H_VS_4H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_4H_IN_VITRO_ACT_CD4_TCELL_DN
REACTOME_NUCLEOTIDE_EXCISION_REPAIR, REACTOME_NUCLEOTIDE_EXCISION_REPAIR
GSE16385_MONOCYTE_VS_12H_IFNG_TNF_TREATED_MACROPHAGE_UP, GSE16385_MONOCYTE_VS_12H_IFNG_TNF_TREATED_MACROPHAGE_UP
GSE33424_CD161_INT_VS_NEG_CD8_TCELL_UP, GSE33424_CD161_INT_VS_NEG_CD8_TCELL_UP
GSE3982_MAST_CELL_VS_NKCELL_UP, GSE3982_MAST_CELL_VS_NKCELL_UP
GSE37301_LYMPHOID_PRIMED_MPP_VS_RAG2_KO_NK_CELL_UP, GSE37301_LYMPHOID_PRIMED_MPP_VS_RAG2_KO_NK_CELL_UP
GSE30083_SP1_VS_SP3_THYMOCYTE_UP, GSE30083_SP1_VS_SP3_THYMOCYTE_UP
GSE21774_CD56_BRIGHT_VS_DIM_CD62L_POSITIVE_NK_CELL_DN, GSE21774_CD56_BRIGHT_VS_DIM_CD62L_POSITIVE_NK_CELL_DN
GSE40277_EOS_AND_LEF1_TRANSDUCED_VS_GATA1_AND_SATB1_TRANSDUCED_CD4_TCELL_DN, GSE40277_EOS_AND_LEF1_TRANSDUCED_VS_GATA1_AND_SATB1_TRANSDUCED_CD4_TCELL_DN
REACTOME_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR_GG_NER, REACTOME_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR_GG_NER
GSE11924_TFH_VS_TH1_CD4_TCELL_DN, GSE11924_TFH_VS_TH1_CD4_TCELL_DN
GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_DN, GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_DN
GSE23321_CD8_STEM_CELL_MEMORY_VS_EFFECTOR_MEMORY_CD8_TCELL_UP, GSE23321_CD8_STEM_CELL_MEMORY_VS_EFFECTOR_MEMORY_CD8_TCELL_UP
GSE17721_0.5H_VS_4H_POLYIC_BMDC_UP, GSE17721_0.5H_VS_4H_POLYIC_BMDC_UP
GSE30971_2H_VS_4H_LPS_STIM_MACROPHAGE_WBP7_HET_UP, GSE30971_2H_VS_4H_LPS_STIM_MACROPHAGE_WBP7_HET_UP
REACTOME_CELLULAR_RESPONSE_TO_HEAT_STRESS, REACTOME_CELLULAR_RESPONSE_TO_HEAT_STRESS
HP_DECREASED_ACTIVITY_OF_MITOCHONDRIAL_COMPLEX_I, HP_DECREASED_ACTIVITY_OF_MITOCHONDRIAL_COMPLEX_I
GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT, GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT
REACTOME_REGULATION_OF_HSF1_MEDIATED_HEAT_SHOCK_RESPONSE, REACTOME_REGULATION_OF_HSF1_MEDIATED_HEAT_SHOCK_RESPONSE
REACTOME_TRNA_PROCESSING_IN_THE_NUCLEUS, REACTOME_TRNA_PROCESSING_IN_THE_NUCLEUS
REACTOME_FORMATION_OF_TC_NER_PRE_INCISION_COMPLEX, REACTOME_FORMATION_OF_TC_NER_PRE_INCISION_COMPLEX
GSE22443_NAIVE_VS_ACT_AND_IL12_TREATED_CD8_TCELL_DN, GSE22443_NAIVE_VS_ACT_AND_IL12_TREATED_CD8_TCELL_DN
GOBP_CELLULAR_RESPONSE_TO_HEAT, GOBP_CELLULAR_RESPONSE_TO_HEAT
GSE14308_NAIVE_CD4_TCELL_VS_NATURAL_TREG_UP, GSE14308_NAIVE_CD4_TCELL_VS_NATURAL_TREG_UP
GSE22282_HYPOXIA_VS_NORMOXIA_MYELOID_DC_UP, GSE22282_HYPOXIA_VS_NORMOXIA_MYELOID_DC_UP
GOBP_PROTEIN_TRANSMEMBRANE_TRANSPORT, GOBP_PROTEIN_TRANSMEMBRANE_TRANSPORT
GOBP_RESPONSE_TO_HEAT, GOBP_RESPONSE_TO_HEAT
GOBP_PROTEIN_INSERTION_INTO_MEMBRANE, GOBP_PROTEIN_INSERTION_INTO_MEMBRANE
GSE17721_CTRL_VS_CPG_12H_BMDC_DN, GSE17721_CTRL_VS_CPG_12H_BMDC_DN
GSE360_T_GONDII_VS_M_TUBERCULOSIS_DC_DN, GSE360_T_GONDII_VS_M_TUBERCULOSIS_DC_DN
REACTOME_SUMOYLATION_OF_RNA_BINDING_PROTEINS, REACTOME_SUMOYLATION_OF_RNA_BINDING_PROTEINS
REACTOME_VIRAL_MESSENGER_RNA_SYNTHESIS, REACTOME_VIRAL_MESSENGER_RNA_SYNTHESIS
REACTOME_SUMOYLATION_OF_DNA_REPLICATION_PROTEINS, REACTOME_SUMOYLATION_OF_DNA_REPLICATION_PROTEINS
REACTOME_SUMOYLATION_OF_UBIQUITINYLATION_PROTEINS, REACTOME_SUMOYLATION_OF_UBIQUITINYLATION_PROTEINS
REACTOME_DNA_DAMAGE_RECOGNITION_IN_GG_NER, REACTOME_DNA_DAMAGE_RECOGNITION_IN_GG_NER
MIR889_5P, MIR889_5P
GOCC_EXORIBONUCLEASE_COMPLEX, GOCC_EXORIBONUCLEASE_COMPLEX
RAX2_TARGET_GENES, RAX2_TARGET_GENES
REACTOME_SUMOYLATION_OF_SUMOYLATION_PROTEINS, REACTOME_SUMOYLATION_OF_SUMOYLATION_PROTEINS
REACTOME_HIV_ELONGATION_ARREST_AND_RECOVERY, REACTOME_HIV_ELONGATION_ARREST_AND_RECOVERY
GOCC_NUCLEAR_REPLICATION_FORK, GOCC_NUCLEAR_REPLICATION_FORK
REACTOME_INTERACTIONS_OF_VPR_WITH_HOST_CELLULAR_PROTEINS, REACTOME_INTERACTIONS_OF_VPR_WITH_HOST_CELLULAR_PROTEINS
GSE25846_IL10_POS_VS_NEG_CD8_TCELL_DAY7_POST_CORONAVIRUS_BRAIN_UP, GSE25846_IL10_POS_VS_NEG_CD8_TCELL_DAY7_POST_CORONAVIRUS_BRAIN_UP
REACTOME_EXPORT_OF_VIRAL_RIBONUCLEOPROTEINS_FROM_NUCLEUS, REACTOME_EXPORT_OF_VIRAL_RIBONUCLEOPROTEINS_FROM_NUCLEUS
REACTOME_POSTMITOTIC_NUCLEAR_PORE_COMPLEX_NPC_REFORMATION, REACTOME_POSTMITOTIC_NUCLEAR_PORE_COMPLEX_NPC_REFORMATION
REACTOME_TRANSPORT_OF_THE_SLBP_DEPENDANT_MATURE_MRNA, REACTOME_TRANSPORT_OF_THE_SLBP_DEPENDANT_MATURE_MRNA
GSE26488_WT_VS_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN, GSE26488_WT_VS_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN
GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_DAMAGE_RECOGNITION, GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_DAMAGE_RECOGNITION
REACTOME_NUCLEAR_PORE_COMPLEX_NPC_DISASSEMBLY, REACTOME_NUCLEAR_PORE_COMPLEX_NPC_DISASSEMBLY
GOBP_PIGMENT_METABOLIC_PROCESS, GOBP_PIGMENT_METABOLIC_PROCESS
GOBP_PIGMENT_BIOSYNTHETIC_PROCESS, GOBP_PIGMENT_BIOSYNTHETIC_PROCESS
GOCC_PRERIBOSOME_LARGE_SUBUNIT_PRECURSOR, GOCC_PRERIBOSOME_LARGE_SUBUNIT_PRECURSOR
GOCC_NUCLEAR_EXOSOME_RNASE_COMPLEX, GOCC_NUCLEAR_EXOSOME_RNASE_COMPLEX
GOBP_FATTY_ACYL_COA_METABOLIC_PROCESS, GOBP_FATTY_ACYL_COA_METABOLIC_PROCESS
GOBP_AXO_DENDRITIC_TRANSPORT, GOBP_AXO_DENDRITIC_TRANSPORT
MIR4687_3P, MIR4687_3P
GSE15930_STIM_VS_STIM_AND_IL12_72H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_IL12_72H_CD8_T_CELL_DN
HP_PROGRESSIVE_SPASTIC_PARAPLEGIA, HP_PROGRESSIVE_SPASTIC_PARAPLEGIA
MIR134_3P, MIR134_3P
GSE11884_WT_VS_FURIN_KO_NAIVE_CD4_TCELL_DN, GSE11884_WT_VS_FURIN_KO_NAIVE_CD4_TCELL_DN
REACTOME_CARGO_RECOGNITION_FOR_CLATHRIN_MEDIATED_ENDOCYTOSIS, REACTOME_CARGO_RECOGNITION_FOR_CLATHRIN_MEDIATED_ENDOCYTOSIS
GOBP_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS, GOBP_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS
REACTOME_TRISTETRAPROLIN_TTP_ZFP36_BINDS_AND_DESTABILIZES_MRNA, REACTOME_TRISTETRAPROLIN_TTP_ZFP36_BINDS_AND_DESTABILIZES_MRNA
TBK1.DN.48HRS_UP, TBK1.DN.48HRS_UP
GOCC_COP9_SIGNALOSOME, GOCC_COP9_SIGNALOSOME
GSE12707_AT16L1_HYPOMORPH_VS_WT_THYMUS_UP, GSE12707_AT16L1_HYPOMORPH_VS_WT_THYMUS_UP
MIR7155_3P, MIR7155_3P
MIR3136_3P, MIR3136_3P
GOBP_PURINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS, GOBP_PURINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS
GSE32423_CTRL_VS_IL7_MEMORY_CD8_TCELL_UP, GSE32423_CTRL_VS_IL7_MEMORY_CD8_TCELL_UP
MIR7974, MIR7974
REACTOME_REGULATION_OF_GLUCOKINASE_BY_GLUCOKINASE_REGULATORY_PROTEIN, REACTOME_REGULATION_OF_GLUCOKINASE_BY_GLUCOKINASE_REGULATORY_PROTEIN
GOBP_TETRAPYRROLE_METABOLIC_PROCESS, GOBP_TETRAPYRROLE_METABOLIC_PROCESS
HP_CEREBRAL_WHITE_MATTER_ATROPHY, HP_CEREBRAL_WHITE_MATTER_ATROPHY
HP_LOWER_LIMB_HYPERREFLEXIA, HP_LOWER_LIMB_HYPERREFLEXIA
GOBP_PORPHYRIN_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_PORPHYRIN_CONTAINING_COMPOUND_METABOLIC_PROCESS
GOBP_SULFUR_COMPOUND_CATABOLIC_PROCESS, GOBP_SULFUR_COMPOUND_CATABOLIC_PROCESS
REACTOME_BUTYRATE_RESPONSE_FACTOR_1_BRF1_BINDS_AND_DESTABILIZES_MRNA, REACTOME_BUTYRATE_RESPONSE_FACTOR_1_BRF1_BINDS_AND_DESTABILIZES_MRNA
HP_BRISK_REFLEXES, HP_BRISK_REFLEXES
HP_PREMATURE_LOSS_OF_TEETH, HP_PREMATURE_LOSS_OF_TEETH
HP_ABNORMALITY_OF_PAIN_SENSATION, HP_ABNORMALITY_OF_PAIN_SENSATION
GOBP_PROTEIN_REFOLDING, GOBP_PROTEIN_REFOLDING
MODULE_233, MODULE_233
GSE2585_CD80_HIGH_VS_LOW_MTEC_DN, GSE2585_CD80_HIGH_VS_LOW_MTEC_DN
GOBP_HEME_METABOLIC_PROCESS, GOBP_HEME_METABOLIC_PROCESS
KEGG_PROTEIN_EXPORT, KEGG_PROTEIN_EXPORT
HORTON_SREBF_TARGETS, HORTON_SREBF_TARGETS
REACTOME_PROTEIN_METHYLATION, REACTOME_PROTEIN_METHYLATION
RASHI_RESPONSE_TO_IONIZING_RADIATION_1, RASHI_RESPONSE_TO_IONIZING_RADIATION_1
HP_IMPAIRED_PAIN_SENSATION, HP_IMPAIRED_PAIN_SENSATION
GSE10463_CD40L_AND_VA347_VS_CD40L_IN_DC_DN, GSE10463_CD40L_AND_VA347_VS_CD40L_IN_DC_DN
RIZKI_TUMOR_INVASIVENESS_2D_UP, RIZKI_TUMOR_INVASIVENESS_2D_UP
GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_DN, GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_DN
HP_HYPERACTIVE_PATELLAR_REFLEX, HP_HYPERACTIVE_PATELLAR_REFLEX
GOMF_MISFOLDED_PROTEIN_BINDING, GOMF_MISFOLDED_PROTEIN_BINDING
GOBP_TETRAPYRROLE_BIOSYNTHETIC_PROCESS, GOBP_TETRAPYRROLE_BIOSYNTHETIC_PROCESS
GOMF_TUMOR_NECROSIS_FACTOR_RECEPTOR_BINDING, GOMF_TUMOR_NECROSIS_FACTOR_RECEPTOR_BINDING
BLANCO_MELO_INFLUENZA_A_INFECTION_A594_CELLS_UP, BLANCO_MELO_INFLUENZA_A_INFECTION_A594_CELLS_UP
GOMF_PROTEIN_TRANSPORTER_ACTIVITY, GOMF_PROTEIN_TRANSPORTER_ACTIVITY
GOBP_GLUTATHIONE_METABOLIC_PROCESS, GOBP_GLUTATHIONE_METABOLIC_PROCESS
GOBP_REGULATION_OF_RRNA_PROCESSING, GOBP_REGULATION_OF_RRNA_PROCESSING
WAKABAYASHI_ADIPOGENESIS_PPARG_BOUND_36HR, WAKABAYASHI_ADIPOGENESIS_PPARG_BOUND_36HR
HP_PREMATURE_LOSS_OF_PRIMARY_TEETH, HP_PREMATURE_LOSS_OF_PRIMARY_TEETH
GOBP_FATTY_ACID_DERIVATIVE_METABOLIC_PROCESS, GOBP_FATTY_ACID_DERIVATIVE_METABOLIC_PROCESS
GOMF_STRUCTURAL_CONSTITUENT_OF_NUCLEAR_PORE, GOMF_STRUCTURAL_CONSTITUENT_OF_NUCLEAR_PORE
GOMF_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS, GOMF_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS
GOBP_REGULATION_OF_TRANSLATIONAL_ELONGATION, GOBP_REGULATION_OF_TRANSLATIONAL_ELONGATION
GSE43863_NAIVE_VS_LY6C_INT_CXCR5POS_CD4_EFF_TCELL_D6_LCMV_DN, GSE43863_NAIVE_VS_LY6C_INT_CXCR5POS_CD4_EFF_TCELL_D6_LCMV_DN
GOMF_MACROMOLECULE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_MACROMOLECULE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_RRNA_PROCESSING, GOBP_POSITIVE_REGULATION_OF_RRNA_PROCESSING
HP_DECREASED_ABSENT_ANKLE_REFLEXES, HP_DECREASED_ABSENT_ANKLE_REFLEXES
GOBP_RETROGRADE_AXONAL_TRANSPORT, GOBP_RETROGRADE_AXONAL_TRANSPORT
HP_PROGRESSIVE_HEARING_IMPAIRMENT, HP_PROGRESSIVE_HEARING_IMPAIRMENT
WP_PHOTODYNAMIC_THERAPYINDUCED_UNFOLDED_PROTEIN_RESPONSE, WP_PHOTODYNAMIC_THERAPYINDUCED_UNFOLDED_PROTEIN_RESPONSE
MIR3085_5P, MIR3085_5P
GOBP_RNA_5_END_PROCESSING, GOBP_RNA_5_END_PROCESSING
GOBP_POSITIVE_REGULATION_OF_OSTEOCLAST_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_OSTEOCLAST_DIFFERENTIATION
GOBP_ATTACHMENT_OF_MITOTIC_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GOBP_ATTACHMENT_OF_MITOTIC_SPINDLE_MICROTUBULES_TO_KINETOCHORE
GOMF_GTPASE_INHIBITOR_ACTIVITY, GOMF_GTPASE_INHIBITOR_ACTIVITY
GOBP_CELLULAR_IRON_ION_HOMEOSTASIS, GOBP_CELLULAR_IRON_ION_HOMEOSTASIS
HP_CUBITUS_VALGUS, HP_CUBITUS_VALGUS
GOBP_SULFUR_AMINO_ACID_METABOLIC_PROCESS, GOBP_SULFUR_AMINO_ACID_METABOLIC_PROCESS
HP_PSEUDOBULBAR_SIGNS, HP_PSEUDOBULBAR_SIGNS
LEE_AGING_MUSCLE_DN, LEE_AGING_MUSCLE_DN
GOBP_POSITIVE_REGULATION_OF_RNA_SPLICING, GOBP_POSITIVE_REGULATION_OF_RNA_SPLICING
GOBP_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_INNER_MEMBRANE, GOBP_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_INNER_MEMBRANE
ZHONG_PFC_C1_ASTROCYTE, ZHONG_PFC_C1_ASTROCYTE
PID_HIF2PATHWAY, PID_HIF2PATHWAY
REACTOME_PHASE_II_CONJUGATION_OF_COMPOUNDS, REACTOME_PHASE_II_CONJUGATION_OF_COMPOUNDS
GOBP_REGULATION_OF_DNA_DAMAGE_CHECKPOINT, GOBP_REGULATION_OF_DNA_DAMAGE_CHECKPOINT
VILLANUEVA_LIVER_CANCER_KRT19_DN, VILLANUEVA_LIVER_CANCER_KRT19_DN
GOMF_PROTEIN_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_PROTEIN_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
MIR6870_3P, MIR6870_3P
GOBP_LONG_CHAIN_FATTY_ACYL_COA_METABOLIC_PROCESS, GOBP_LONG_CHAIN_FATTY_ACYL_COA_METABOLIC_PROCESS
GOBP_PROTEIN_DENEDDYLATION, GOBP_PROTEIN_DENEDDYLATION
REACTOME_TNFS_BIND_THEIR_PHYSIOLOGICAL_RECEPTORS, REACTOME_TNFS_BIND_THEIR_PHYSIOLOGICAL_RECEPTORS
GOBP_POSTTRANSLATIONAL_PROTEIN_TARGETING_TO_ENDOPLASMIC_RETICULUM_MEMBRANE, GOBP_POSTTRANSLATIONAL_PROTEIN_TARGETING_TO_ENDOPLASMIC_RETICULUM_MEMBRANE
GOMF_THIOLESTER_HYDROLASE_ACTIVITY, GOMF_THIOLESTER_HYDROLASE_ACTIVITY
GOBP_CYTOPLASMIC_TRANSLATIONAL_ELONGATION, GOBP_CYTOPLASMIC_TRANSLATIONAL_ELONGATION
GOBP_BLASTOCYST_FORMATION, GOBP_BLASTOCYST_FORMATION
KIM_TIAL1_TARGETS, KIM_TIAL1_TARGETS
REACTOME_SULFUR_AMINO_ACID_METABOLISM, REACTOME_SULFUR_AMINO_ACID_METABOLISM
GOBP_SULFUR_AMINO_ACID_BIOSYNTHETIC_PROCESS, GOBP_SULFUR_AMINO_ACID_BIOSYNTHETIC_PROCESS
PUJANA_BREAST_CANCER_WITH_BRCA1_MUTATED_DN, PUJANA_BREAST_CANCER_WITH_BRCA1_MUTATED_DN
KEGG_BIOSYNTHESIS_OF_UNSATURATED_FATTY_ACIDS, KEGG_BIOSYNTHESIS_OF_UNSATURATED_FATTY_ACIDS
GOBP_ACETYL_COA_METABOLIC_PROCESS, GOBP_ACETYL_COA_METABOLIC_PROCESS
GOBP_LONG_CHAIN_FATTY_ACID_BIOSYNTHETIC_PROCESS, GOBP_LONG_CHAIN_FATTY_ACID_BIOSYNTHETIC_PROCESS
PID_CIRCADIAN_PATHWAY, PID_CIRCADIAN_PATHWAY
REACTOME_DNA_DAMAGE_REVERSAL, REACTOME_DNA_DAMAGE_REVERSAL
HP_PROGRESSIVE_SENSORINEURAL_HEARING_IMPAIRMENT, HP_PROGRESSIVE_SENSORINEURAL_HEARING_IMPAIRMENT
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_ALDEHYDE_OR_OXO_GROUP_OF_DONORS, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_ALDEHYDE_OR_OXO_GROUP_OF_DONORS
GOBP_GRANULOCYTE_DIFFERENTIATION, GOBP_GRANULOCYTE_DIFFERENTIATION
LIAO_HAVE_SOX4_BINDING_SITES, LIAO_HAVE_SOX4_BINDING_SITES
WP_HIJACK_OF_UBIQUITINATION_BY_SARSCOV2, WP_HIJACK_OF_UBIQUITINATION_BY_SARSCOV2
HP_NARROW_NASAL_RIDGE, HP_NARROW_NASAL_RIDGE
GOBP_PEPTIDYL_HISTIDINE_MODIFICATION, GOBP_PEPTIDYL_HISTIDINE_MODIFICATION
REACTOME_REGULATION_OF_LOCALIZATION_OF_FOXO_TRANSCRIPTION_FACTORS, REACTOME_REGULATION_OF_LOCALIZATION_OF_FOXO_TRANSCRIPTION_FACTORS
WP_ONE_CARBON_METABOLISM, WP_ONE_CARBON_METABOLISM
GOCC_RIBONUCLEASE_P_COMPLEX, GOCC_RIBONUCLEASE_P_COMPLEX
GOBP_NEGATIVE_REGULATION_OF_ATPASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_ATPASE_ACTIVITY
HP_GENETIC_ANTICIPATION, HP_GENETIC_ANTICIPATION
GOBP_METHIONINE_METABOLIC_PROCESS, GOBP_METHIONINE_METABOLIC_PROCESS
GOMF_TRNA_SPECIFIC_RIBONUCLEASE_ACTIVITY, GOMF_TRNA_SPECIFIC_RIBONUCLEASE_ACTIVITY
GOBP_POSTTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE_TRANSLOCATION, GOBP_POSTTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE_TRANSLOCATION
GOBP_S_ADENOSYLMETHIONINE_CYCLE, GOBP_S_ADENOSYLMETHIONINE_CYCLE
GOBP_POSITIVE_REGULATION_BY_HOST_OF_VIRAL_GENOME_REPLICATION, GOBP_POSITIVE_REGULATION_BY_HOST_OF_VIRAL_GENOME_REPLICATION
GOBP_POLYADENYLATION_DEPENDENT_SNORNA_3_END_PROCESSING, GOBP_POLYADENYLATION_DEPENDENT_SNORNA_3_END_PROCESSING
GOBP_NEGATIVE_REGULATION_OF_CELL_CYCLE_CHECKPOINT, GOBP_NEGATIVE_REGULATION_OF_CELL_CYCLE_CHECKPOINT
HP_RESPIRATORY_FAILURE_REQUIRING_ASSISTED_VENTILATION, HP_RESPIRATORY_FAILURE_REQUIRING_ASSISTED_VENTILATION
HP_HYPOCHOLESTEROLEMIA, HP_HYPOCHOLESTEROLEMIA
HP_OSTEOMYELITIS, HP_OSTEOMYELITIS
GAUCHER_PBMC_YF_VAX_STAMARIL_UNKNOWN_AGE_60DY_DN, GAUCHER_PBMC_YF_VAX_STAMARIL_UNKNOWN_AGE_60DY_DN
HP_METAPHYSEAL_DYSPLASIA, HP_METAPHYSEAL_DYSPLASIA
GOBP_IMP_BIOSYNTHETIC_PROCESS, GOBP_IMP_BIOSYNTHETIC_PROCESS
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H_HEME_PROTEIN_AS_ACCEPTOR, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H_HEME_PROTEIN_AS_ACCEPTOR
GOBP_NEGATIVE_REGULATION_OF_B_CELL_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_B_CELL_APOPTOTIC_PROCESS
HP_HIATUS_HERNIA, HP_HIATUS_HERNIA
GOBP_NUCLEAR_BODY_ORGANIZATION, GOBP_NUCLEAR_BODY_ORGANIZATION
GOCC_ALPHA_DNA_POLYMERASE_PRIMASE_COMPLEX, GOCC_ALPHA_DNA_POLYMERASE_PRIMASE_COMPLEX
GOCC_MITOCHONDRIAL_INTERMEMBRANE_SPACE_PROTEIN_TRANSPORTER_COMPLEX, GOCC_MITOCHONDRIAL_INTERMEMBRANE_SPACE_PROTEIN_TRANSPORTER_COMPLEX
GOBP_S_ADENOSYLMETHIONINE_METABOLIC_PROCESS, GOBP_S_ADENOSYLMETHIONINE_METABOLIC_PROCESS
GOBP_TRNA_5_LEADER_REMOVAL, GOBP_TRNA_5_LEADER_REMOVAL
WP_OMEGA9_FA_SYNTHESIS, WP_OMEGA9_FA_SYNTHESIS
HP_ABNORMAL_CIRCULATING_SULFUR_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_SULFUR_AMINO_ACID_CONCENTRATION
HP_CYTOCHROME_C_OXIDASE_NEGATIVE_MUSCLE_FIBERS, HP_CYTOCHROME_C_OXIDASE_NEGATIVE_MUSCLE_FIBERS
REACTOME_SIGNALING_BY_FGFR2_IN_DISEASE, REACTOME_SIGNALING_BY_FGFR2_IN_DISEASE
GOBP_PURINE_NUCLEOSIDE_CATABOLIC_PROCESS, GOBP_PURINE_NUCLEOSIDE_CATABOLIC_PROCESS
GOBP_DEADENYLATION_DEPENDENT_DECAPPING_OF_NUCLEAR_TRANSCRIBED_MRNA, GOBP_DEADENYLATION_DEPENDENT_DECAPPING_OF_NUCLEAR_TRANSCRIBED_MRNA
HP_DISTAL_LOWER_LIMB_MUSCLE_WEAKNESS, HP_DISTAL_LOWER_LIMB_MUSCLE_WEAKNESS
GOMF_PHOSPHATIDYLINOSITOL_3_5_BISPHOSPHATE_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_3_5_BISPHOSPHATE_PHOSPHATASE_ACTIVITY
BERTUCCI_INVASIVE_CARCINOMA_DUCTAL_VS_LOBULAR_UP, BERTUCCI_INVASIVE_CARCINOMA_DUCTAL_VS_LOBULAR_UP
HP_INCREASED_MUSCLE_LIPID_CONTENT, HP_INCREASED_MUSCLE_LIPID_CONTENT
JEON_SMAD6_TARGETS_DN, JEON_SMAD6_TARGETS_DN
GOBP_ALDEHYDE_CATABOLIC_PROCESS, GOBP_ALDEHYDE_CATABOLIC_PROCESS
HP_BACTERIAL_ENDOCARDITIS, HP_BACTERIAL_ENDOCARDITIS
GOCC_PARASPECKLES, GOCC_PARASPECKLES
GOMF_TUMOR_NECROSIS_FACTOR_RECEPTOR_SUPERFAMILY_BINDING, GOMF_TUMOR_NECROSIS_FACTOR_RECEPTOR_SUPERFAMILY_BINDING
GOCC_RIBONUCLEASE_MRP_COMPLEX, GOCC_RIBONUCLEASE_MRP_COMPLEX
HP_HOMOCYSTINURIA, HP_HOMOCYSTINURIA
HP_HYPERHOMOCYSTINEMIA, HP_HYPERHOMOCYSTINEMIA
NAKAYA_PBMC_FLUMIST_AGE_18_50YO_7DY_IFN_SUBSET_DN, NAKAYA_PBMC_FLUMIST_AGE_18_50YO_7DY_IFN_SUBSET_DN
GOBP_COENZYME_A_BIOSYNTHETIC_PROCESS, GOBP_COENZYME_A_BIOSYNTHETIC_PROCESS
GOCC_SIGNAL_RECOGNITION_PARTICLE, GOCC_SIGNAL_RECOGNITION_PARTICLE
GOBP_TRNA_5_END_PROCESSING, GOBP_TRNA_5_END_PROCESSING
MANN_RESPONSE_TO_AMIFOSTINE_UP, MANN_RESPONSE_TO_AMIFOSTINE_UP
GOBP_RRNA_BASE_METHYLATION, GOBP_RRNA_BASE_METHYLATION
HP_ABNORMAL_CIRCULATING_ASPARTATE_FAMILY_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_ASPARTATE_FAMILY_AMINO_ACID_CONCENTRATION
HP_PSEUDOBULBAR_PARALYSIS, HP_PSEUDOBULBAR_PARALYSIS
HP_ENLARGEMENT_OF_THE_COSTOCHONDRAL_JUNCTION, HP_ENLARGEMENT_OF_THE_COSTOCHONDRAL_JUNCTION
REACTOME_NR1H2_NR1H3_REGULATE_GENE_EXPRESSION_LINKED_TO_LIPOGENESIS, REACTOME_NR1H2_NR1H3_REGULATE_GENE_EXPRESSION_LINKED_TO_LIPOGENESIS
GOBP_PROTEIN_LOCALIZATION_TO_NUCLEAR_ENVELOPE, GOBP_PROTEIN_LOCALIZATION_TO_NUCLEAR_ENVELOPE
AMIT_EGF_RESPONSE_20_MCF10A, AMIT_EGF_RESPONSE_20_MCF10A
MIR210_3P, MIR210_3P
MIR181C_3P, MIR181C_3P
GOMF_RIBONUCLEASE_P_ACTIVITY, GOMF_RIBONUCLEASE_P_ACTIVITY
GOBP_OXIDATIVE_RNA_DEMETHYLATION, GOBP_OXIDATIVE_RNA_DEMETHYLATION
GOMF_OXIDATIVE_RNA_DEMETHYLASE_ACTIVITY, GOMF_OXIDATIVE_RNA_DEMETHYLASE_ACTIVITY
HP_DECREASED_METHYLCOBALAMIN, HP_DECREASED_METHYLCOBALAMIN
HP_ABNORMAL_CSF_CARBOXYLIC_ACID_CONCENTRATION, HP_ABNORMAL_CSF_CARBOXYLIC_ACID_CONCENTRATION
GOBP_PEPTIDYL_DIPHTHAMIDE_BIOSYNTHETIC_PROCESS_FROM_PEPTIDYL_HISTIDINE, GOBP_PEPTIDYL_DIPHTHAMIDE_BIOSYNTHETIC_PROCESS_FROM_PEPTIDYL_HISTIDINE
REACTOME_SYNTHESIS_OF_DIPHTHAMIDE_EEF2, REACTOME_SYNTHESIS_OF_DIPHTHAMIDE_EEF2
MIR1843, MIR1843
chr20p13, chr20p13
KEGG_LIMONENE_AND_PINENE_DEGRADATION, KEGG_LIMONENE_AND_PINENE_DEGRADATION
REACTOME_ATF6_ATF6_ALPHA_ACTIVATES_CHAPERONE_GENES, REACTOME_ATF6_ATF6_ALPHA_ACTIVATES_CHAPERONE_GENES
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_A_SULFUR_GROUP_OF_DONORS_DISULFIDE_AS_ACCEPTOR, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_A_SULFUR_GROUP_OF_DONORS_DISULFIDE_AS_ACCEPTOR
MIR371A_3P, MIR371A_3P
WP_AUTOSOMAL_RECESSIVE_OSTEOPETROSIS_PATHWAYS, WP_AUTOSOMAL_RECESSIVE_OSTEOPETROSIS_PATHWAYS
WHITESIDE_CISPLATIN_RESISTANCE_UP, WHITESIDE_CISPLATIN_RESISTANCE_UP
GOBP_TOOTH_ERUPTION, GOBP_TOOTH_ERUPTION
HP_OPSOCLONUS, HP_OPSOCLONUS
GOMF_UBIQUITIN_DEPENDENT_PROTEIN_BINDING, GOMF_UBIQUITIN_DEPENDENT_PROTEIN_BINDING
HP_IMPAIRED_DISTAL_TACTILE_SENSATION, HP_IMPAIRED_DISTAL_TACTILE_SENSATION
GOBP_CHAPERONE_MEDIATED_PROTEIN_TRANSPORT, GOBP_CHAPERONE_MEDIATED_PROTEIN_TRANSPORT
HP_HYPOMETHIONINEMIA, HP_HYPOMETHIONINEMIA
HP_SLENDER_TOE, HP_SLENDER_TOE
HP_MUSCULAR_VENTRICULAR_SEPTAL_DEFECT, HP_MUSCULAR_VENTRICULAR_SEPTAL_DEFECT
GOBP_DE_NOVO_IMP_BIOSYNTHETIC_PROCESS, GOBP_DE_NOVO_IMP_BIOSYNTHETIC_PROCESS
GOBP_NUCLEOSIDE_BISPHOSPHATE_CATABOLIC_PROCESS, GOBP_NUCLEOSIDE_BISPHOSPHATE_CATABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_TOLL_LIKE_RECEPTOR_2_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_TOLL_LIKE_RECEPTOR_2_SIGNALING_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY
BIOCARTA_RANKL_PATHWAY, BIOCARTA_RANKL_PATHWAY
REACTOME_GLUTAMATE_AND_GLUTAMINE_METABOLISM, REACTOME_GLUTAMATE_AND_GLUTAMINE_METABOLISM
GOBP_MEDIUM_CHAIN_FATTY_ACID_METABOLIC_PROCESS, GOBP_MEDIUM_CHAIN_FATTY_ACID_METABOLIC_PROCESS
RYAAAKNNNNNNTTGW_UNKNOWN, RYAAAKNNNNNNTTGW_UNKNOWN
MIR4802_5P, MIR4802_5P
DPPA3_TARGET_GENES, DPPA3_TARGET_GENES
GOBP_REGULATION_OF_CELL_CELL_ADHESION_MEDIATED_BY_INTEGRIN, GOBP_REGULATION_OF_CELL_CELL_ADHESION_MEDIATED_BY_INTEGRIN
HP_ABNORMAL_MUSCLE_GLYCOGEN_CONTENT, HP_ABNORMAL_MUSCLE_GLYCOGEN_CONTENT
HP_TIP_TOE_GAIT, HP_TIP_TOE_GAIT
GOMF_PROTEIN_KINASE_B_BINDING, GOMF_PROTEIN_KINASE_B_BINDING
RAMPON_ENRICHED_LEARNING_ENVIRONMENT_EARLY_UP, RAMPON_ENRICHED_LEARNING_ENVIRONMENT_EARLY_UP
HP_DECREASED_CORNEAL_REFLEX, HP_DECREASED_CORNEAL_REFLEX
BIOCARTA_FREE_PATHWAY, BIOCARTA_FREE_PATHWAY
BRUNEAU_SEPTATION_VENTRICULAR, BRUNEAU_SEPTATION_VENTRICULAR
WP_TRANSSULFURATION_PATHWAY, WP_TRANSSULFURATION_PATHWAY
WP_LIVER_X_RECEPTOR_PATHWAY, WP_LIVER_X_RECEPTOR_PATHWAY
SIMBULAN_UV_RESPONSE_NORMAL_UP, SIMBULAN_UV_RESPONSE_NORMAL_UP
KRISHNAN_FURIN_TARGETS_UP, KRISHNAN_FURIN_TARGETS_UP
GOBP_RELAXATION_OF_SMOOTH_MUSCLE, GOBP_RELAXATION_OF_SMOOTH_MUSCLE
GOBP_REGULATION_OF_MULTICELLULAR_ORGANISM_GROWTH, GOBP_REGULATION_OF_MULTICELLULAR_ORGANISM_GROWTH
HP_OPTIC_NERVE_COMPRESSION, HP_OPTIC_NERVE_COMPRESSION
HP_PYODERMA_GANGRENOSUM, HP_PYODERMA_GANGRENOSUM
HP_RIGHT_AORTIC_ARCH_WITH_MIRROR_IMAGE_BRANCHING, HP_RIGHT_AORTIC_ARCH_WITH_MIRROR_IMAGE_BRANCHING
HP_PROXIMAL_FEMORAL_EPIPHYSIOLYSIS, HP_PROXIMAL_FEMORAL_EPIPHYSIOLYSIS
GOCC_NUCLEAR_PORE_CENTRAL_TRANSPORT_CHANNEL, GOCC_NUCLEAR_PORE_CENTRAL_TRANSPORT_CHANNEL
RBM17_TARGET_GENES, RBM17_TARGET_GENES
HP_SEVERE_LACTIC_ACIDOSIS, HP_SEVERE_LACTIC_ACIDOSIS
HP_REDUCTION_OF_NEUTROPHIL_MOTILITY, HP_REDUCTION_OF_NEUTROPHIL_MOTILITY
GOMF_PROTEIN_PHOSPHATASE_2B_BINDING, GOMF_PROTEIN_PHOSPHATASE_2B_BINDING
MIR147B_3P, MIR147B_3P
REACTOME_PROLONGED_ERK_ACTIVATION_EVENTS, REACTOME_PROLONGED_ERK_ACTIVATION_EVENTS
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_ALDEHYDE_OR_OXO_GROUP_OF_DONORS_NAD_OR_NADP_AS_ACCEPTOR, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_ALDEHYDE_OR_OXO_GROUP_OF_DONORS_NAD_OR_NADP_AS_ACCEPTOR
LEE_CALORIE_RESTRICTION_MUSCLE_UP, LEE_CALORIE_RESTRICTION_MUSCLE_UP
GOBP_KETONE_CATABOLIC_PROCESS, GOBP_KETONE_CATABOLIC_PROCESS
HP_APLASIA_HYPOPLASIA_INVOLVING_THE_CARPAL_BONES, HP_APLASIA_HYPOPLASIA_INVOLVING_THE_CARPAL_BONES
HP_AQUEDUCTAL_STENOSIS, HP_AQUEDUCTAL_STENOSIS
HP_ABNORMAL_MONOCYTE_MORPHOLOGY, HP_ABNORMAL_MONOCYTE_MORPHOLOGY
HP_DIAPHRAGMATIC_EVENTRATION, HP_DIAPHRAGMATIC_EVENTRATION
GOBP_REGULATION_OF_PROTEIN_COMPLEX_STABILITY, GOBP_REGULATION_OF_PROTEIN_COMPLEX_STABILITY
REACTOME_LIPOPHAGY, REACTOME_LIPOPHAGY
GOCC_SARCOPLASMIC_RETICULUM_LUMEN, GOCC_SARCOPLASMIC_RETICULUM_LUMEN
HP_CERVICAL_SPINE_INSTABILITY, HP_CERVICAL_SPINE_INSTABILITY
GOBP_FATTY_ACID_ELONGATION_SATURATED_FATTY_ACID, GOBP_FATTY_ACID_ELONGATION_SATURATED_FATTY_ACID
GOMF_3_OXO_ARACHIDOYL_COA_SYNTHASE_ACTIVITY, GOMF_3_OXO_ARACHIDOYL_COA_SYNTHASE_ACTIVITY
GOBP_PROTEIN_REPAIR, GOBP_PROTEIN_REPAIR
GOBP_REGULATION_OF_MITOCHONDRIAL_MRNA_STABILITY, GOBP_REGULATION_OF_MITOCHONDRIAL_MRNA_STABILITY
GOMF_PHOSPHATIDYLINOSITOL_3_5_BISPHOSPHATE_3_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_3_5_BISPHOSPHATE_3_PHOSPHATASE_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_OSTEOCLAST_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_OSTEOCLAST_DEVELOPMENT
GOBP_NEGATIVE_REGULATION_OF_IRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE, GOBP_NEGATIVE_REGULATION_OF_IRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE
GOMF_MRF_BINDING, GOMF_MRF_BINDING
HP_HYPOCHROMIA, HP_HYPOCHROMIA
GOBP_COP9_SIGNALOSOME_ASSEMBLY, GOBP_COP9_SIGNALOSOME_ASSEMBLY
GOBP_ENERGY_COUPLED_PROTON_TRANSMEMBRANE_TRANSPORT_AGAINST_ELECTROCHEMICAL_GRADIENT, GOBP_ENERGY_COUPLED_PROTON_TRANSMEMBRANE_TRANSPORT_AGAINST_ELECTROCHEMICAL_GRADIENT
GOBP_TRNA_THIO_MODIFICATION, GOBP_TRNA_THIO_MODIFICATION
GOCC_VCB_COMPLEX, GOCC_VCB_COMPLEX
HP_FACIAL_PARALYSIS, HP_FACIAL_PARALYSIS
MIR518A_3P_MIR518B_MIR518C_3P_MIR518D_3P_MIR518F_3P_MIR526A_3P, MIR518A_3P_MIR518B_MIR518C_3P_MIR518D_3P_MIR518F_3P_MIR526A_3P
GOBP_REGULATION_OF_ACYL_COA_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_ACYL_COA_BIOSYNTHETIC_PROCESS
GOCC_CENTRAL_ELEMENT, GOCC_CENTRAL_ELEMENT
ZEILSTRA_CD44_TARGETS_UP, ZEILSTRA_CD44_TARGETS_UP
WP_FTO_OBESITY_VARIANT_MECHANISM, WP_FTO_OBESITY_VARIANT_MECHANISM
WP_DUAL_HIJACK_MODEL_OF_VIF_IN_HIV_INFECTION, WP_DUAL_HIJACK_MODEL_OF_VIF_IN_HIV_INFECTION
HP_HYPERMETHIONINEMIA, HP_HYPERMETHIONINEMIA
GOBP_NEUTROPHIL_DIFFERENTIATION, GOBP_NEUTROPHIL_DIFFERENTIATION
GOBP_NEGATIVE_REGULATION_OF_CHOLESTEROL_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_CHOLESTEROL_METABOLIC_PROCESS
GOCC_ENDOPLASMIC_RETICULUM_PALMITOYLTRANSFERASE_COMPLEX, GOCC_ENDOPLASMIC_RETICULUM_PALMITOYLTRANSFERASE_COMPLEX
BOYLAN_MULTIPLE_MYELOMA_D_CLUSTER_UP, BOYLAN_MULTIPLE_MYELOMA_D_CLUSTER_UP
GOBP_GLYCINE_TRANSPORT, GOBP_GLYCINE_TRANSPORT
GOBP_CELLULAR_RESPONSE_TO_MANGANESE_ION, GOBP_CELLULAR_RESPONSE_TO_MANGANESE_ION
HP_LIMITED_EXTRAOCULAR_MOVEMENTS, HP_LIMITED_EXTRAOCULAR_MOVEMENTS
GOBP_REGULATION_OF_SULFUR_METABOLIC_PROCESS, GOBP_REGULATION_OF_SULFUR_METABOLIC_PROCESS
GOBP_HOMOCYSTEINE_METABOLIC_PROCESS, GOBP_HOMOCYSTEINE_METABOLIC_PROCESS
HP_ACHALASIA, HP_ACHALASIA
GOBP_RESPONSE_TO_HYDROXYUREA, GOBP_RESPONSE_TO_HYDROXYUREA
GOBP_INTERMEDIATE_FILAMENT_BASED_PROCESS, GOBP_INTERMEDIATE_FILAMENT_BASED_PROCESS
HP_HYPOKINESIA, HP_HYPOKINESIA
WP_VITAMIN_B12_METABOLISM, WP_VITAMIN_B12_METABOLISM
HP_ABNORMALITY_OF_THE_TRAGUS, HP_ABNORMALITY_OF_THE_TRAGUS
HP_ABNORMALITY_OF_THE_FORAMEN_MAGNUM, HP_ABNORMALITY_OF_THE_FORAMEN_MAGNUM
GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CENTROSOME, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CENTROSOME
GOCC_MULTIMERIC_RIBONUCLEASE_P_COMPLEX, GOCC_MULTIMERIC_RIBONUCLEASE_P_COMPLEX
GOMF_RIBONUCLEASE_P_RNA_BINDING, GOMF_RIBONUCLEASE_P_RNA_BINDING
WORSCHECH_TUMOR_REJECTION_DN, WORSCHECH_TUMOR_REJECTION_DN
HOFMANN_MYELODYSPLASTIC_SYNDROM_HIGH_RISK_UP, HOFMANN_MYELODYSPLASTIC_SYNDROM_HIGH_RISK_UP
HP_PROXIMAL_FEMORAL_METAPHYSEAL_ABNORMALITY, HP_PROXIMAL_FEMORAL_METAPHYSEAL_ABNORMALITY
GOCC_NUCLEAR_PORE_OUTER_RING, GOCC_NUCLEAR_PORE_OUTER_RING
MIR3115, MIR3115
WP_CYSTEINE_AND_METHIONINE_CATABOLISM, WP_CYSTEINE_AND_METHIONINE_CATABOLISM
WP_MAPK_AND_NFKB_SIGNALLING_PATHWAYS_INHIBITED_BY_YERSINIA_YOPJ, WP_MAPK_AND_NFKB_SIGNALLING_PATHWAYS_INHIBITED_BY_YERSINIA_YOPJ
HP_CARPAL_BONE_HYPOPLASIA, HP_CARPAL_BONE_HYPOPLASIA
HP_ANISOCORIA, HP_ANISOCORIA
WP_METHIONINE_METABOLISM_LEADING_TO_SULPHUR_AMINO_ACIDS_AND_RELATED_DISORDERS, WP_METHIONINE_METABOLISM_LEADING_TO_SULPHUR_AMINO_ACIDS_AND_RELATED_DISORDERS
REACTOME_VITAMIN_D_CALCIFEROL_METABOLISM, REACTOME_VITAMIN_D_CALCIFEROL_METABOLISM
HP_CHRONIC_NEUTROPENIA, HP_CHRONIC_NEUTROPENIA
GOBP_INNER_CELL_MASS_CELL_DIFFERENTIATION, GOBP_INNER_CELL_MASS_CELL_DIFFERENTIATION
GOBP_POSITIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
HP_MACROVESICULAR_HEPATIC_STEATOSIS, HP_MACROVESICULAR_HEPATIC_STEATOSIS
MODULE_132, MODULE_132
HP_CYSTATHIONINEMIA, HP_CYSTATHIONINEMIA
HP_BIRTH_LENGTH_GREATER_THAN_97TH_PERCENTILE, HP_BIRTH_LENGTH_GREATER_THAN_97TH_PERCENTILE
GOBP_OUTER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GOBP_OUTER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
GOBP_DNA_ENDOREDUPLICATION, GOBP_DNA_ENDOREDUPLICATION
GOBP_NEGATIVE_REGULATION_OF_DNA_DAMAGE_CHECKPOINT, GOBP_NEGATIVE_REGULATION_OF_DNA_DAMAGE_CHECKPOINT
GOMF_RNA_DEPENDENT_ATPASE_ACTIVITY, GOMF_RNA_DEPENDENT_ATPASE_ACTIVITY
GOBP_TETRAHYDROBIOPTERIN_METABOLIC_PROCESS, GOBP_TETRAHYDROBIOPTERIN_METABOLIC_PROCESS
ZNF419_TARGET_GENES, ZNF419_TARGET_GENES
REACTOME_SIGNALING_BY_MRAS_COMPLEX_MUTANTS, REACTOME_SIGNALING_BY_MRAS_COMPLEX_MUTANTS
WP_INTERLEUKIN1_INDUCED_ACTIVATION_OF_NFKAPPAB, WP_INTERLEUKIN1_INDUCED_ACTIVATION_OF_NFKAPPAB
WP_FOLATE_METABOLISM, WP_FOLATE_METABOLISM
MODULE_514, MODULE_514
HP_HYPOPLASIA_OF_THE_EAR_CARTILAGE, HP_HYPOPLASIA_OF_THE_EAR_CARTILAGE
WP_APOE_AND_MIR146_IN_INFLAMMATION_AND_ATHEROSCLEROSIS, WP_APOE_AND_MIR146_IN_INFLAMMATION_AND_ATHEROSCLEROSIS
DER_IFN_BETA_RESPONSE_DN, DER_IFN_BETA_RESPONSE_DN
HP_IMPAIRED_VIBRATION_SENSATION_AT_ANKLES, HP_IMPAIRED_VIBRATION_SENSATION_AT_ANKLES
GOMF_GLYCINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_GLYCINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
HP_INCREASED_MUSCLE_GLYCOGEN_CONTENT, HP_INCREASED_MUSCLE_GLYCOGEN_CONTENT
GOBP_SRP_DEPENDENT_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE_SIGNAL_SEQUENCE_RECOGNITION, GOBP_SRP_DEPENDENT_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE_SIGNAL_SEQUENCE_RECOGNITION
GOBP_AMINO_ACID_BETAINE_BIOSYNTHETIC_PROCESS, GOBP_AMINO_ACID_BETAINE_BIOSYNTHETIC_PROCESS
GOCC_ENDOPLASMIC_RETICULUM_CHAPERONE_COMPLEX, GOCC_ENDOPLASMIC_RETICULUM_CHAPERONE_COMPLEX
REACTOME_ARMS_MEDIATED_ACTIVATION, REACTOME_ARMS_MEDIATED_ACTIVATION
GOBP_POSITIVE_REGULATION_OF_TOLL_LIKE_RECEPTOR_2_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_TOLL_LIKE_RECEPTOR_2_SIGNALING_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS
SUZUKI_CTCFL_TARGETS_UP, SUZUKI_CTCFL_TARGETS_UP
GOBP_POSITIVE_REGULATION_OF_MAST_CELL_CHEMOTAXIS, GOBP_POSITIVE_REGULATION_OF_MAST_CELL_CHEMOTAXIS
PETRETTO_LEFT_VENTRICLE_MASS_QTL_CIS_UP, PETRETTO_LEFT_VENTRICLE_MASS_QTL_CIS_UP
REACTOME_PROTEIN_REPAIR, REACTOME_PROTEIN_REPAIR
STONER_ESOPHAGEAL_CARCINOGENESIS_DN, STONER_ESOPHAGEAL_CARCINOGENESIS_DN
WP_FLUOROACETIC_ACID_TOXICITY, WP_FLUOROACETIC_ACID_TOXICITY
HP_ABNORMAL_CSF_PYRUVATE_FAMILY_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CSF_PYRUVATE_FAMILY_AMINO_ACID_CONCENTRATION
GOBP_OXIDATIVE_DNA_DEMETHYLATION, GOBP_OXIDATIVE_DNA_DEMETHYLATION
GOBP_RNA_GUANINE_N7_METHYLATION, GOBP_RNA_GUANINE_N7_METHYLATION
GOBP_MEDIUM_CHAIN_FATTY_ACID_BIOSYNTHETIC_PROCESS, GOBP_MEDIUM_CHAIN_FATTY_ACID_BIOSYNTHETIC_PROCESS
GOBP_VESICLE_UNCOATING, GOBP_VESICLE_UNCOATING
GOCC_EKC_KEOPS_COMPLEX, GOCC_EKC_KEOPS_COMPLEX
GOMF_DNA_METHYLTRANSFERASE_ACTIVITY, GOMF_DNA_METHYLTRANSFERASE_ACTIVITY
HP_FAT_MALABSORPTION, HP_FAT_MALABSORPTION
REACTOME_METHYLATION, REACTOME_METHYLATION
GOBP_MUSCLE_CELL_CELLULAR_HOMEOSTASIS, GOBP_MUSCLE_CELL_CELLULAR_HOMEOSTASIS
HP_ANARTHRIA, HP_ANARTHRIA
HP_DECREASED_CIRCULATING_ALDOSTERONE_LEVEL, HP_DECREASED_CIRCULATING_ALDOSTERONE_LEVEL
MODULE_354, MODULE_354
HP_ANTERIOR_RIB_CUPPING, HP_ANTERIOR_RIB_CUPPING
HP_URINARY_RETENTION, HP_URINARY_RETENTION
PHESSE_TARGETS_OF_APC_AND_MBD2_DN, PHESSE_TARGETS_OF_APC_AND_MBD2_DN
GOCC_PRESYNAPTIC_CYTOSOL, GOCC_PRESYNAPTIC_CYTOSOL
REACTOME_SUMOYLATION_OF_IMMUNE_RESPONSE_PROTEINS, REACTOME_SUMOYLATION_OF_IMMUNE_RESPONSE_PROTEINS
WP_PHOTODYNAMIC_THERAPYINDUCED_NFKB_SURVIVAL_SIGNALING, WP_PHOTODYNAMIC_THERAPYINDUCED_NFKB_SURVIVAL_SIGNALING
DESCARTES_MAIN_FETAL_AFP_ALB_POSITIVE_CELLS, DESCARTES_MAIN_FETAL_AFP_ALB_POSITIVE_CELLS
GOBP_REGULATION_OF_RESPIRATORY_SYSTEM_PROCESS, GOBP_REGULATION_OF_RESPIRATORY_SYSTEM_PROCESS
HP_PROGRESSIVE_PES_CAVUS, HP_PROGRESSIVE_PES_CAVUS
GOBP_SKELETAL_MUSCLE_TISSUE_GROWTH, GOBP_SKELETAL_MUSCLE_TISSUE_GROWTH
REACTOME_CHL1_INTERACTIONS, REACTOME_CHL1_INTERACTIONS
MIR10401_3P, MIR10401_3P
GOBP_LUTEOLYSIS, GOBP_LUTEOLYSIS
GOBP_PARACRINE_SIGNALING, GOBP_PARACRINE_SIGNALING
GOBP_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_GROWTH, GOBP_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_GROWTH
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN_REDUCED_PTERIDINE_AS_ONE_DONOR_AND_INCORPORATION_OF_ONE_ATOM_OF_OXYGEN, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN_REDUCED_PTERIDINE_AS_ONE_
HP_POLLAKISURIA, HP_POLLAKISURIA
GOBP_RELAXATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE, GOBP_RELAXATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE
HP_THORACOLUMBAR_KYPHOSCOLIOSIS, HP_THORACOLUMBAR_KYPHOSCOLIOSIS
HP_DECREASED_ACHILLES_REFLEX, HP_DECREASED_ACHILLES_REFLEX
HP_TRANSIENT_NEUTROPENIA, HP_TRANSIENT_NEUTROPENIA
GOBP_PROTON_TRANSPORTING_V_TYPE_ATPASE_COMPLEX_ASSEMBLY, GOBP_PROTON_TRANSPORTING_V_TYPE_ATPASE_COMPLEX_ASSEMBLY
GOCC_GLYCOGEN_GRANULE, GOCC_GLYCOGEN_GRANULE
GOMF_7S_RNA_BINDING, GOMF_7S_RNA_BINDING
GOBP_TYROSINE_METABOLIC_PROCESS, GOBP_TYROSINE_METABOLIC_PROCESS
GOMF_FATTY_ACID_SYNTHASE_ACTIVITY, GOMF_FATTY_ACID_SYNTHASE_ACTIVITY
HP_OVOID_VERTEBRAL_BODIES, HP_OVOID_VERTEBRAL_BODIES
HP_DIAPHYSEAL_SCLEROSIS, HP_DIAPHYSEAL_SCLEROSIS
HP_DEFORMED_RIB_CAGE, HP_DEFORMED_RIB_CAGE
HP_PROXIMAL_FEMORAL_METAPHYSEAL_IRREGULARITY, HP_PROXIMAL_FEMORAL_METAPHYSEAL_IRREGULARITY
HP_PERIPHERAL_OPACIFICATION_OF_THE_CORNEA, HP_PERIPHERAL_OPACIFICATION_OF_THE_CORNEA
HP_COMMON_ATRIUM, HP_COMMON_ATRIUM
GOBP_PROLINE_METABOLIC_PROCESS, GOBP_PROLINE_METABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_CELL_DEATH, GOBP_POSITIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_CELL_DEATH
chr6p24, chr6p24
LI_ADIPOGENESIS_BY_ACTIVATED_PPARG, LI_ADIPOGENESIS_BY_ACTIVATED_PPARG
REACTOME_PURINE_CATABOLISM, REACTOME_PURINE_CATABOLISM
GOBP_LACTATE_METABOLIC_PROCESS, GOBP_LACTATE_METABOLIC_PROCESS
GOCC_BLOC_1_COMPLEX, GOCC_BLOC_1_COMPLEX
GOBP_PRENYLATION, GOBP_PRENYLATION
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
GOCC_CD40_RECEPTOR_COMPLEX, GOCC_CD40_RECEPTOR_COMPLEX
GOBP_TYROSINE_CATABOLIC_PROCESS, GOBP_TYROSINE_CATABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_GROWTH, GOBP_POSITIVE_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_GROWTH
GOBP_DE_NOVO_CENTRIOLE_ASSEMBLY, GOBP_DE_NOVO_CENTRIOLE_ASSEMBLY
REACTOME_TYROSINE_CATABOLISM, REACTOME_TYROSINE_CATABOLISM
WP_SULINDAC_METABOLIC_PATHWAY, WP_SULINDAC_METABOLIC_PATHWAY
HP_HYPOPLASIA_OF_THE_FEMORAL_HEAD, HP_HYPOPLASIA_OF_THE_FEMORAL_HEAD
GOBP_ANTERIOR_HEAD_DEVELOPMENT, GOBP_ANTERIOR_HEAD_DEVELOPMENT
GOBP_NEGATIVE_REGULATION_OF_CERAMIDE_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_CERAMIDE_BIOSYNTHETIC_PROCESS
HP_DECREASED_ACTIVITY_OF_MITOCHONDRIAL_COMPLEX_II, HP_DECREASED_ACTIVITY_OF_MITOCHONDRIAL_COMPLEX_II
HP_ABNORMAL_CIRCULATING_AMYLASE_LEVEL, HP_ABNORMAL_CIRCULATING_AMYLASE_LEVEL
GOBP_HOMOSERINE_METABOLIC_PROCESS, GOBP_HOMOSERINE_METABOLIC_PROCESS
GOBP_EXOCRINE_PANCREAS_DEVELOPMENT, GOBP_EXOCRINE_PANCREAS_DEVELOPMENT
GOBP_VACUOLAR_PROTON_TRANSPORTING_V_TYPE_ATPASE_COMPLEX_ASSEMBLY, GOBP_VACUOLAR_PROTON_TRANSPORTING_V_TYPE_ATPASE_COMPLEX_ASSEMBLY
GOBP_CLATHRIN_COAT_DISASSEMBLY, GOBP_CLATHRIN_COAT_DISASSEMBLY
GOBP_REGULATION_OF_MOTOR_NEURON_APOPTOTIC_PROCESS, GOBP_REGULATION_OF_MOTOR_NEURON_APOPTOTIC_PROCESS
GOCC_PCAF_COMPLEX, GOCC_PCAF_COMPLEX
GOCC_PAM_COMPLEX_TIM23_ASSOCIATED_IMPORT_MOTOR, GOCC_PAM_COMPLEX_TIM23_ASSOCIATED_IMPORT_MOTOR
GOCC_NUCLEOLAR_RIBONUCLEASE_P_COMPLEX, GOCC_NUCLEOLAR_RIBONUCLEASE_P_COMPLEX
GOMF_GLYCOLIPID_MANNOSYLTRANSFERASE_ACTIVITY, GOMF_GLYCOLIPID_MANNOSYLTRANSFERASE_ACTIVITY
GOMF_RRNA_GUANINE_METHYLTRANSFERASE_ACTIVITY, GOMF_RRNA_GUANINE_METHYLTRANSFERASE_ACTIVITY
GOMF_SULFATIDE_BINDING, GOMF_SULFATIDE_BINDING
HP_MONOCYTOSIS, HP_MONOCYTOSIS
GOBP_PERIPHERAL_NERVOUS_SYSTEM_MYELIN_MAINTENANCE, GOBP_PERIPHERAL_NERVOUS_SYSTEM_MYELIN_MAINTENANCE
GOBP_REGULATION_OF_MAST_CELL_CHEMOTAXIS, GOBP_REGULATION_OF_MAST_CELL_CHEMOTAXIS
REACTOME_METABOLISM_OF_INGESTED_SEMET_SEC_MESEC_INTO_H2SE, REACTOME_METABOLISM_OF_INGESTED_SEMET_SEC_MESEC_INTO_H2SE
HP_CONGENITAL_HEMOLYTIC_ANEMIA, HP_CONGENITAL_HEMOLYTIC_ANEMIA
HP_ABNORMAL_RIB_OSSIFICATION, HP_ABNORMAL_RIB_OSSIFICATION
GOBP_NEUROFILAMENT_CYTOSKELETON_ORGANIZATION, GOBP_NEUROFILAMENT_CYTOSKELETON_ORGANIZATION
GOCC_GLYCOSYLPHOSPHATIDYLINOSITOL_N_ACETYLGLUCOSAMINYLTRANSFERASE_GPI_GNT_COMPLEX, GOCC_GLYCOSYLPHOSPHATIDYLINOSITOL_N_ACETYLGLUCOSAMINYLTRANSFERASE_GPI_GNT_COMPLEX
GOMF_DEAD_H_BOX_RNA_HELICASE_BINDING, GOMF_DEAD_H_BOX_RNA_HELICASE_BINDING
MODULE_326, MODULE_326
GOBP_CARNITINE_METABOLIC_PROCESS, GOBP_CARNITINE_METABOLIC_PROCESS
GOBP_REGULATION_OF_RESPIRATORY_GASEOUS_EXCHANGE, GOBP_REGULATION_OF_RESPIRATORY_GASEOUS_EXCHANGE
MOREIRA_RESPONSE_TO_TSA_DN, MOREIRA_RESPONSE_TO_TSA_DN
KEGG_PRION_DISEASES, KEGG_PRION_DISEASES
MODULE_101, MODULE_101
GOBP_RESPONSE_TO_CISPLATIN, GOBP_RESPONSE_TO_CISPLATIN
GOBP_FATTY_ACID_DERIVATIVE_CATABOLIC_PROCESS, GOBP_FATTY_ACID_DERIVATIVE_CATABOLIC_PROCESS
HP_STEATORRHEA, HP_STEATORRHEA
HP_PENOSCROTAL_HYPOSPADIAS, HP_PENOSCROTAL_HYPOSPADIAS
HP_LOW_BACK_PAIN, HP_LOW_BACK_PAIN
ZHONG_PFC_HES1_POS_C1_NPC, ZHONG_PFC_HES1_POS_C1_NPC
GOBP_NEGATIVE_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS
GOCC_GLYCINERGIC_SYNAPSE, GOCC_GLYCINERGIC_SYNAPSE
GOMF_C3HC4_TYPE_RING_FINGER_DOMAIN_BINDING, GOMF_C3HC4_TYPE_RING_FINGER_DOMAIN_BINDING
GOBP_OXYGEN_METABOLIC_PROCESS, GOBP_OXYGEN_METABOLIC_PROCESS
GOMF_PHOSPHATIDYLINOSITOL_N_ACETYLGLUCOSAMINYLTRANSFERASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_N_ACETYLGLUCOSAMINYLTRANSFERASE_ACTIVITY
MODULE_280, MODULE_280
WP_DOPAMINE_METABOLISM, WP_DOPAMINE_METABOLISM
HP_EEG_WITH_MULTIFOCAL_SLOW_ACTIVITY, HP_EEG_WITH_MULTIFOCAL_SLOW_ACTIVITY
GOMF_COA_HYDROLASE_ACTIVITY, GOMF_COA_HYDROLASE_ACTIVITY
GOBP_REGULATION_OF_PERK_MEDIATED_UNFOLDED_PROTEIN_RESPONSE, GOBP_REGULATION_OF_PERK_MEDIATED_UNFOLDED_PROTEIN_RESPONSE
GOBP_DEOXYRIBONUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS, GOBP_DEOXYRIBONUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS
BIOCARTA_FLUMAZENIL_PATHWAY, BIOCARTA_FLUMAZENIL_PATHWAY
GOBP_MOTOR_NEURON_APOPTOTIC_PROCESS, GOBP_MOTOR_NEURON_APOPTOTIC_PROCESS
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_METAL_IONS_NAD_OR_NADP_AS_ACCEPTOR, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_METAL_IONS_NAD_OR_NADP_AS_ACCEPTOR
MIR3193, MIR3193
ZNF112_TARGET_GENES, ZNF112_TARGET_GENES
GOCC_RIBBON_SYNAPSE, GOCC_RIBBON_SYNAPSE
GOMF_LONG_CHAIN_FATTY_ACYL_COA_BINDING, GOMF_LONG_CHAIN_FATTY_ACYL_COA_BINDING
GOBP_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_INDEPENDENT_DECAY, GOBP_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_INDEPENDENT_DECAY
GOBP_POSITIVE_REGULATION_OF_CELL_GROWTH_INVOLVED_IN_CARDIAC_MUSCLE_CELL_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_CELL_GROWTH_INVOLVED_IN_CARDIAC_MUSCLE_CELL_DEVELOPMENT
HP_HYPOREFLEXIA_OF_LOWER_LIMBS, HP_HYPOREFLEXIA_OF_LOWER_LIMBS
REACTOME_NEGATIVE_REGULATION_OF_ACTIVITY_OF_TFAP2_AP_2_FAMILY_TRANSCRIPTION_FACTORS, REACTOME_NEGATIVE_REGULATION_OF_ACTIVITY_OF_TFAP2_AP_2_FAMILY_TRANSCRIPTION_FACTORS
GOBP_RHODOPSIN_MEDIATED_SIGNALING_PATHWAY, GOBP_RHODOPSIN_MEDIATED_SIGNALING_PATHWAY
GOBP_MAST_CELL_MIGRATION, GOBP_MAST_CELL_MIGRATION
GOBP_SULFUR_AMINO_ACID_CATABOLIC_PROCESS, GOBP_SULFUR_AMINO_ACID_CATABOLIC_PROCESS
GOMF_PALMITOYL_COA_HYDROLASE_ACTIVITY, GOMF_PALMITOYL_COA_HYDROLASE_ACTIVITY
HP_UNDERDEVELOPED_TRAGUS, HP_UNDERDEVELOPED_TRAGUS
GOBP_CARNITINE_BIOSYNTHETIC_PROCESS, GOBP_CARNITINE_BIOSYNTHETIC_PROCESS
GOCC_PHOTORECEPTOR_RIBBON_SYNAPSE, GOCC_PHOTORECEPTOR_RIBBON_SYNAPSE
HP_GLANDULAR_HYPOSPADIAS, HP_GLANDULAR_HYPOSPADIAS
HP_SKULL_ASYMMETRY, HP_SKULL_ASYMMETRY
GOBP_CELLULAR_SPHINGOLIPID_HOMEOSTASIS, GOBP_CELLULAR_SPHINGOLIPID_HOMEOSTASIS
GOCC_ELONGIN_COMPLEX, GOCC_ELONGIN_COMPLEX
GOBP_MULTI_CILIATED_EPITHELIAL_CELL_DIFFERENTIATION, GOBP_MULTI_CILIATED_EPITHELIAL_CELL_DIFFERENTIATION
GOBP_DNA_CATABOLIC_PROCESS_EXONUCLEOLYTIC, GOBP_DNA_CATABOLIC_PROCESS_EXONUCLEOLYTIC
GOBP_POSITIVE_REGULATION_OF_LIPOPOLYSACCHARIDE_MEDIATED_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_LIPOPOLYSACCHARIDE_MEDIATED_SIGNALING_PATHWAY
GOBP_HYDROGEN_SULFIDE_METABOLIC_PROCESS, GOBP_HYDROGEN_SULFIDE_METABOLIC_PROCESS
GOBP_ICOSANOID_CATABOLIC_PROCESS, GOBP_ICOSANOID_CATABOLIC_PROCESS
REACTOME_CHREBP_ACTIVATES_METABOLIC_GENE_EXPRESSION, REACTOME_CHREBP_ACTIVATES_METABOLIC_GENE_EXPRESSION
GOCC_CLATHRIN_SCULPTED_GAMMA_AMINOBUTYRIC_ACID_TRANSPORT_VESICLE, GOCC_CLATHRIN_SCULPTED_GAMMA_AMINOBUTYRIC_ACID_TRANSPORT_VESICLE
chr2q31, chr2q31
HP_ABNORMALITY_OF_VITAMIN_A_METABOLISM, HP_ABNORMALITY_OF_VITAMIN_A_METABOLISM
HP_INTRINSIC_HAND_MUSCLE_ATROPHY, HP_INTRINSIC_HAND_MUSCLE_ATROPHY
HP_ABNORMALITY_OF_VITAMIN_E_METABOLISM, HP_ABNORMALITY_OF_VITAMIN_E_METABOLISM
GOBP_NEGATIVE_REGULATION_OF_TOLL_LIKE_RECEPTOR_4_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_TOLL_LIKE_RECEPTOR_4_SIGNALING_PATHWAY
HP_PANCREATIC_HYPOPLASIA, HP_PANCREATIC_HYPOPLASIA
MATZUK_PREOVULATORY_FOLLICLE, MATZUK_PREOVULATORY_FOLLICLE
GOBP_TOXIN_METABOLIC_PROCESS, GOBP_TOXIN_METABOLIC_PROCESS
HP_ABNORMAL_ERYTHROCYTE_PHYSIOLOGY, HP_ABNORMAL_ERYTHROCYTE_PHYSIOLOGY
HP_APLASIA_HYPOPLASIA_OF_THE_TRAGUS, HP_APLASIA_HYPOPLASIA_OF_THE_TRAGUS
GOBP_CELLULAR_RESPONSE_TO_CISPLATIN, GOBP_CELLULAR_RESPONSE_TO_CISPLATIN
GOMF_GLYCERALDEHYDE_3_PHOSPHATE_DEHYDROGENASE_NADPLUS_NON_PHOSPHORYLATING_ACTIVITY, GOMF_GLYCERALDEHYDE_3_PHOSPHATE_DEHYDROGENASE_NADPLUS_NON_PHOSPHORYLATING_ACTIVITY
GOBP_PROLINE_BIOSYNTHETIC_PROCESS, GOBP_PROLINE_BIOSYNTHETIC_PROCESS
GOBP_HYDROGEN_SULFIDE_BIOSYNTHETIC_PROCESS, GOBP_HYDROGEN_SULFIDE_BIOSYNTHETIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_MUSCLE_TISSUE_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_MUSCLE_TISSUE_DEVELOPMENT
GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_4_SIGNALING_PATHWAY, GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_4_SIGNALING_PATHWAY
chr11p12, chr11p12
REACTOME_GABA_SYNTHESIS_RELEASE_REUPTAKE_AND_DEGRADATION, REACTOME_GABA_SYNTHESIS_RELEASE_REUPTAKE_AND_DEGRADATION
GOCC_CLATHRIN_SCULPTED_VESICLE, GOCC_CLATHRIN_SCULPTED_VESICLE
chr7p21, chr7p21
WP_EXTRACELLULAR_VESICLES_IN_THE_CROSSTALK_OF_CARDIAC_CELLS, WP_EXTRACELLULAR_VESICLES_IN_THE_CROSSTALK_OF_CARDIAC_CELLS
GOBP_POSITIVE_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_METAL_IONS, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_METAL_IONS
GOBP_POSITIVE_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_DEVELOPMENT
GOBP_STRESS_RESPONSE_TO_METAL_ION, GOBP_STRESS_RESPONSE_TO_METAL_ION
GOBP_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_DEVELOPMENT, GOBP_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_DEVELOPMENT
TAKADA_GASTRIC_CANCER_COPY_NUMBER_DN, TAKADA_GASTRIC_CANCER_COPY_NUMBER_DN