ELL_VS_NEUTROPHIL_UP, GSE27786_BCELL_VS_NEUTROPHIL_UP

GSE27786_LSK_VS_MONO_MAC_UP, GSE27786_LSK_VS_MONO_MAC_UP
GSE27786_CD8_TCELL_VS_ERYTHROBLAST_UP, GSE27786_CD8_TCELL_VS_ERYTHROBLAST_UP
GSE27786_CD4_TCELL_VS_MONO_MAC_UP, GSE27786_CD4_TCELL_VS_MONO_MAC_UP
GSE27786_LSK_VS_NEUTROPHIL_UP, GSE27786_LSK_VS_NEUTROPHIL_UP
GSE27786_CD4_TCELL_VS_NEUTROPHIL_UP, GSE27786_CD4_TCELL_VS_NEUTROPHIL_UP
GSE27786_NKTCELL_VS_MONO_MAC_UP, GSE27786_NKTCELL_VS_MONO_MAC_UP
GSE27786_NKTCELL_VS_NEUTROPHIL_UP, GSE27786_NKTCELL_VS_NEUTROPHIL_UP
GSE27786_LIN_NEG_VS_NEUTROPHIL_UP, GSE27786_LIN_NEG_VS_NEUTROPHIL_UP
GSE5503_LIVER_DC_VS_PLN_DC_ACTIVATED_ALLOGENIC_TCELL_DN, GSE5503_LIVER_DC_VS_PLN_DC_ACTIVATED_ALLOGENIC_TCELL_DN
GSE27786_CD8_TCELL_VS_MONO_MAC_UP, GSE27786_CD8_TCELL_VS_MONO_MAC_UP
GSE27786_NKCELL_VS_NEUTROPHIL_UP, GSE27786_NKCELL_VS_NEUTROPHIL_UP
GSE27786_BCELL_VS_ERYTHROBLAST_UP, GSE27786_BCELL_VS_ERYTHROBLAST_UP
GSE20198_IL12_IL18_VS_IFNA_TREATED_ACT_CD4_TCELL_DN, GSE20198_IL12_IL18_VS_IFNA_TREATED_ACT_CD4_TCELL_DN
GSE27786_LIN_NEG_VS_CD8_TCELL_UP, GSE27786_LIN_NEG_VS_CD8_TCELL_UP
GSE2197_IMMUNOSUPPRESSIVE_DNA_VS_UNTREATEDIN_DC_DN, GSE2197_IMMUNOSUPPRESSIVE_DNA_VS_UNTREATEDIN_DC_DN
GSE27786_BCELL_VS_NKCELL_UP, GSE27786_BCELL_VS_NKCELL_UP
GSE27786_BCELL_VS_NKTCELL_UP, GSE27786_BCELL_VS_NKTCELL_UP
GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_NFATC1_KO_16H_DN, GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_NFATC1_KO_16H_DN
GSE20366_EX_VIVO_VS_HOMEOSTATIC_CONVERSION_NAIVE_CD4_TCELL_UP, GSE20366_EX_VIVO_VS_HOMEOSTATIC_CONVERSION_NAIVE_CD4_TCELL_UP
GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_DN, GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_DN
GSE15750_DAY6_VS_DAY10_EFF_CD8_TCELL_DN, GSE15750_DAY6_VS_DAY10_EFF_CD8_TCELL_DN
GSE6259_33D1_POS_VS_DEC205_POS_FLT3L_INDUCED_SPLENIC_DC_DN, GSE6259_33D1_POS_VS_DEC205_POS_FLT3L_INDUCED_SPLENIC_DC_DN
GSE17721_0.5H_VS_8H_PAM3CSK4_BMDC_UP, GSE17721_0.5H_VS_8H_PAM3CSK4_BMDC_UP
GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_UP
GSE17721_POLYIC_VS_CPG_24H_BMDC_UP, GSE17721_POLYIC_VS_CPG_24H_BMDC_UP
GSE32423_CTRL_VS_IL7_IL4_MEMORY_CD8_TCELL_DN, GSE32423_CTRL_VS_IL7_IL4_MEMORY_CD8_TCELL_DN
GSE5099_UNSTIM_VS_MCSF_TREATED_MONOCYTE_DAY7_DN, GSE5099_UNSTIM_VS_MCSF_TREATED_MONOCYTE_DAY7_DN
BROWNE_HCMV_INFECTION_24HR_UP, BROWNE_HCMV_INFECTION_24HR_UP
GSE32423_IL7_VS_IL7_IL4_MEMORY_CD8_TCELL_DN, GSE32423_IL7_VS_IL7_IL4_MEMORY_CD8_TCELL_DN
GCM_DENR, GCM_DENR
MORF_PPP5C, MORF_PPP5C
GOBP_MRNA_SPLICE_SITE_SELECTION, GOBP_MRNA_SPLICE_SITE_SELECTION
MIR4302, MIR4302
GOBP_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE, GOBP_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE
GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_UP, GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_UP
GSE14308_TH17_VS_NATURAL_TREG_DN, GSE14308_TH17_VS_NATURAL_TREG_DN
MIR4318, MIR4318
HP_ABNORMAL_CIRCULATING_PROTEINOGENIC_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_PROTEINOGENIC_AMINO_ACID_CONCENTRATION
GSE22025_UNTREATED_VS_TGFB1_TREATED_CD4_TCELL_DN, GSE22025_UNTREATED_VS_TGFB1_TREATED_CD4_TCELL_DN
MIR2278, MIR2278
MODULE_331, MODULE_331
GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_24H_UP, GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_24H_UP
GSE42021_CD24INT_TREG_VS_CD24INT_TCONV_THYMUS_DN, GSE42021_CD24INT_TREG_VS_CD24INT_TCONV_THYMUS_DN
HP_DELAYED_CNS_MYELINATION, HP_DELAYED_CNS_MYELINATION
LI_INDUCED_T_TO_NATURAL_KILLER_DN, LI_INDUCED_T_TO_NATURAL_KILLER_DN
HP_BASAL_CELL_CARCINOMA, HP_BASAL_CELL_CARCINOMA
MIR3664_5P, MIR3664_5P
GOBP_INTRACILIARY_TRANSPORT_INVOLVED_IN_CILIUM_ASSEMBLY, GOBP_INTRACILIARY_TRANSPORT_INVOLVED_IN_CILIUM_ASSEMBLY
GOMF_NUCLEAR_RECEPTOR_COACTIVATOR_ACTIVITY, GOMF_NUCLEAR_RECEPTOR_COACTIVATOR_ACTIVITY
GOBP_BRANCHING_MORPHOGENESIS_OF_AN_EPITHELIAL_TUBE, GOBP_BRANCHING_MORPHOGENESIS_OF_AN_EPITHELIAL_TUBE
REACTOME_INTRAFLAGELLAR_TRANSPORT, REACTOME_INTRAFLAGELLAR_TRANSPORT
GOBP_INTRACILIARY_TRANSPORT, GOBP_INTRACILIARY_TRANSPORT
AAAGGAT_MIR501, AAAGGAT_MIR501
HP_MALIGNANT_NEOPLASM_OF_THE_CENTRAL_NERVOUS_SYSTEM, HP_MALIGNANT_NEOPLASM_OF_THE_CENTRAL_NERVOUS_SYSTEM
REACTOME_REGULATION_OF_TNFR1_SIGNALING, REACTOME_REGULATION_OF_TNFR1_SIGNALING
GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG2_AND_FOXP3_TRASDUCED_UP, GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG2_AND_FOXP3_TRASDUCED_UP
GRE_C, GRE_C
MIR4633_5P, MIR4633_5P
HP_ABNORMAL_ASTROCYTE_MORPHOLOGY, HP_ABNORMAL_ASTROCYTE_MORPHOLOGY
GOBP_PLACENTA_DEVELOPMENT, GOBP_PLACENTA_DEVELOPMENT
SCHEIDEREIT_IKK_INTERACTING_PROTEINS, SCHEIDEREIT_IKK_INTERACTING_PROTEINS
MIR3132, MIR3132
GOBP_ENDODERM_DEVELOPMENT, GOBP_ENDODERM_DEVELOPMENT
HP_HAMARTOMA, HP_HAMARTOMA
HP_NEOPLASM_OF_THE_SMALL_INTESTINE, HP_NEOPLASM_OF_THE_SMALL_INTESTINE
GCM_CSNK1D, GCM_CSNK1D
GOBP_ENDODERM_FORMATION, GOBP_ENDODERM_FORMATION
GOBP_GOLGI_TO_PLASMA_MEMBRANE_TRANSPORT, GOBP_GOLGI_TO_PLASMA_MEMBRANE_TRANSPORT
GOCC_CILIARY_TIP, GOCC_CILIARY_TIP
NAGASHIMA_NRG1_SIGNALING_DN, NAGASHIMA_NRG1_SIGNALING_DN
GOBP_EMBRYONIC_PLACENTA_DEVELOPMENT, GOBP_EMBRYONIC_PLACENTA_DEVELOPMENT
REACTOME_TNFR1_INDUCED_NFKAPPAB_SIGNALING_PATHWAY, REACTOME_TNFR1_INDUCED_NFKAPPAB_SIGNALING_PATHWAY
ZSCAN18_TARGET_GENES, ZSCAN18_TARGET_GENES
MIR6529_5P, MIR6529_5P
MIR650, MIR650
HP_TOE_WALKING, HP_TOE_WALKING
WP_TNF_RELATED_WEAK_INDUCER_OF_APOPTOSIS_TWEAK_SIGNALING_PATHWAY, WP_TNF_RELATED_WEAK_INDUCER_OF_APOPTOSIS_TWEAK_SIGNALING_PATHWAY
GOBP_SERINE_FAMILY_AMINO_ACID_METABOLIC_PROCESS, GOBP_SERINE_FAMILY_AMINO_ACID_METABOLIC_PROCESS
ACATATC_MIR190, ACATATC_MIR190
GOBP_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION
GOBP_NEGATIVE_REGULATION_OF_VIRAL_PROCESS, GOBP_NEGATIVE_REGULATION_OF_VIRAL_PROCESS
HP_ABNORMALITY_OF_THE_CARPAL_BONES, HP_ABNORMALITY_OF_THE_CARPAL_BONES
GOBP_SPINAL_CORD_DEVELOPMENT, GOBP_SPINAL_CORD_DEVELOPMENT
MIR197_5P, MIR197_5P
GOBP_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY, GOBP_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY
NIKOLSKY_MUTATED_AND_AMPLIFIED_IN_BREAST_CANCER, NIKOLSKY_MUTATED_AND_AMPLIFIED_IN_BREAST_CANCER
ROY_WOUND_BLOOD_VESSEL_UP, ROY_WOUND_BLOOD_VESSEL_UP
GOMF_MISMATCHED_DNA_BINDING, GOMF_MISMATCHED_DNA_BINDING
HP_NEOPLASM_OF_THE_RESPIRATORY_SYSTEM, HP_NEOPLASM_OF_THE_RESPIRATORY_SYSTEM
GOBP_MISMATCH_REPAIR, GOBP_MISMATCH_REPAIR
HP_INTELLECTUAL_DISABILITY_PROGRESSIVE, HP_INTELLECTUAL_DISABILITY_PROGRESSIVE
GOBP_DOSAGE_COMPENSATION, GOBP_DOSAGE_COMPENSATION
CCATCCA_MIR432, CCATCCA_MIR432
MILI_PSEUDOPODIA, MILI_PSEUDOPODIA
GOBP_SALIVARY_GLAND_DEVELOPMENT, GOBP_SALIVARY_GLAND_DEVELOPMENT
GOBP_NEGATIVE_REGULATION_OF_VIRAL_GENOME_REPLICATION, GOBP_NEGATIVE_REGULATION_OF_VIRAL_GENOME_REPLICATION
TERT_TARGET_GENES, TERT_TARGET_GENES
MIR296_5P, MIR296_5P
GOBP_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY, GOBP_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY
HP_CONTRACTURES_INVOLVING_THE_JOINTS_OF_THE_FEET, HP_CONTRACTURES_INVOLVING_THE_JOINTS_OF_THE_FEET
GOMF_STEROL_BINDING, GOMF_STEROL_BINDING
FOXC1_TARGET_GENES, FOXC1_TARGET_GENES
WP_THE_OVERLAP_BETWEEN_SIGNAL_TRANSDUCTION_PATHWAYS_THAT_CONTRIBUTE_TO_A_RANGE_OF_LMNA_LAMINOPATHIES, WP_THE_OVERLAP_BETWEEN_SIGNAL_TRANSDUCTION_PATHWAYS_THAT_CONTRIBUTE_TO_A_RA
GOBP_REGULATION_OF_BICELLULAR_TIGHT_JUNCTION_ASSEMBLY, GOBP_REGULATION_OF_BICELLULAR_TIGHT_JUNCTION_ASSEMBLY
GOBP_REGULATION_OF_RAC_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_REGULATION_OF_RAC_PROTEIN_SIGNAL_TRANSDUCTION
GOBP_HISTONE_H4_K5_ACETYLATION, GOBP_HISTONE_H4_K5_ACETYLATION
REACTOME_NF_KB_ACTIVATION_THROUGH_FADD_RIP_1_PATHWAY_MEDIATED_BY_CASPASE_8_AND_10, REACTOME_NF_KB_ACTIVATION_THROUGH_FADD_RIP_1_PATHWAY_MEDIATED_BY_CASPASE_8_AND_10
GOBP_HISTONE_MONOUBIQUITINATION, GOBP_HISTONE_MONOUBIQUITINATION
HP_HAPPY_DEMEANOR, HP_HAPPY_DEMEANOR
KEGG_BASAL_CELL_CARCINOMA, KEGG_BASAL_CELL_CARCINOMA
YRCCAKNNGNCGC_UNKNOWN, YRCCAKNNGNCGC_UNKNOWN
MEIS1BHOXA9_01, MEIS1BHOXA9_01
YANAGIHARA_ESX1_TARGETS, YANAGIHARA_ESX1_TARGETS
GOBP_HISTONE_H4_K16_ACETYLATION, GOBP_HISTONE_H4_K16_ACETYLATION
HP_SKIN_APPENDAGE_NEOPLASM, HP_SKIN_APPENDAGE_NEOPLASM
GOBP_COMMISSURAL_NEURON_AXON_GUIDANCE, GOBP_COMMISSURAL_NEURON_AXON_GUIDANCE
GOBP_GOLGI_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT, GOBP_GOLGI_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT
GOBP_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_INNER_MEMBRANE, GOBP_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_INNER_MEMBRANE
GOBP_EMBRYONIC_CAMERA_TYPE_EYE_DEVELOPMENT, GOBP_EMBRYONIC_CAMERA_TYPE_EYE_DEVELOPMENT
HP_SYNOSTOSIS_OF_CARPAL_BONES, HP_SYNOSTOSIS_OF_CARPAL_BONES
WP_TOLLLIKE_RECEPTOR_SIGNALING_RELATED_TO_MYD88, WP_TOLLLIKE_RECEPTOR_SIGNALING_RELATED_TO_MYD88
GOBP_PLACENTA_BLOOD_VESSEL_DEVELOPMENT, GOBP_PLACENTA_BLOOD_VESSEL_DEVELOPMENT
GOBP_L_SERINE_METABOLIC_PROCESS, GOBP_L_SERINE_METABOLIC_PROCESS
HP_MASK_LIKE_FACIES, HP_MASK_LIKE_FACIES
GOBP_APICAL_JUNCTION_ASSEMBLY, GOBP_APICAL_JUNCTION_ASSEMBLY
HP_SPINAL_CORD_LESION, HP_SPINAL_CORD_LESION
GOBP_EMBRYONIC_PLACENTA_MORPHOGENESIS, GOBP_EMBRYONIC_PLACENTA_MORPHOGENESIS
GOBP_POSITIVE_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_BY_HOST, GOBP_POSITIVE_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_BY_HOST
WP_HEDGEHOG_SIGNALING_PATHWAY_NETPATH, WP_HEDGEHOG_SIGNALING_PATHWAY_NETPATH
GOCC_MISMATCH_REPAIR_COMPLEX, GOCC_MISMATCH_REPAIR_COMPLEX
GOBP_POSITIVE_REGULATION_OF_RAC_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_POSITIVE_REGULATION_OF_RAC_PROTEIN_SIGNAL_TRANSDUCTION
HP_CARCINOMA, HP_CARCINOMA
HP_ADENOMATOUS_COLONIC_POLYPOSIS, HP_ADENOMATOUS_COLONIC_POLYPOSIS
HP_INCREASED_SERINE_FAMILY_AMINO_ACID_IN_URINE, HP_INCREASED_SERINE_FAMILY_AMINO_ACID_IN_URINE
BIOCARTA_PITX2_PATHWAY, BIOCARTA_PITX2_PATHWAY
JAZAERI_BREAST_CANCER_BRCA1_VS_BRCA2_DN, JAZAERI_BREAST_CANCER_BRCA1_VS_BRCA2_DN
GOBP_HISTONE_H2A_MONOUBIQUITINATION, GOBP_HISTONE_H2A_MONOUBIQUITINATION
HP_CROSSED_FUSED_RENAL_ECTOPIA, HP_CROSSED_FUSED_RENAL_ECTOPIA
GOBP_POSITIVE_REGULATION_OF_PHOSPHATASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_PHOSPHATASE_ACTIVITY
ZNF776_TARGET_GENES, ZNF776_TARGET_GENES
GOCC_MRNA_EDITING_COMPLEX, GOCC_MRNA_EDITING_COMPLEX
GOBP_EMBRYONIC_EYE_MORPHOGENESIS, GOBP_EMBRYONIC_EYE_MORPHOGENESIS
GOBP_INTERFERON_ALPHA_PRODUCTION, GOBP_INTERFERON_ALPHA_PRODUCTION
GOBP_EMBRYONIC_CAMERA_TYPE_EYE_MORPHOGENESIS, GOBP_EMBRYONIC_CAMERA_TYPE_EYE_MORPHOGENESIS
GOBP_BRANCHING_INVOLVED_IN_BLOOD_VESSEL_MORPHOGENESIS, GOBP_BRANCHING_INVOLVED_IN_BLOOD_VESSEL_MORPHOGENESIS
GOCC_GLYCOPROTEIN_COMPLEX, GOCC_GLYCOPROTEIN_COMPLEX
WP_EDA_SIGNALLING_IN_HAIR_FOLLICLE_DEVELOPMENT, WP_EDA_SIGNALLING_IN_HAIR_FOLLICLE_DEVELOPMENT
GOBP_AXIS_SPECIFICATION, GOBP_AXIS_SPECIFICATION
GOBP_POSITIVE_REGULATION_OF_EPIDERMAL_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_EPIDERMAL_CELL_DIFFERENTIATION
GOMF_FERROUS_IRON_BINDING, GOMF_FERROUS_IRON_BINDING
GALIE_TUMOR_STEMNESS_GENES, GALIE_TUMOR_STEMNESS_GENES
NEUROD2_TARGET_GENES, NEUROD2_TARGET_GENES
GOBP_POSITIVE_REGULATION_OF_OLIGODENDROCYTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_OLIGODENDROCYTE_DIFFERENTIATION
GOBP_POSITIVE_REGULATION_OF_EPIDERMIS_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_EPIDERMIS_DEVELOPMENT
GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION
GOBP_CHEMOKINE_C_C_MOTIF_LIGAND_5_PRODUCTION, GOBP_CHEMOKINE_C_C_MOTIF_LIGAND_5_PRODUCTION
GOCC_UNCONVENTIONAL_MYOSIN_COMPLEX, GOCC_UNCONVENTIONAL_MYOSIN_COMPLEX
HP_BILATERAL_CLEFT_LIP_AND_PALATE, HP_BILATERAL_CLEFT_LIP_AND_PALATE
GOBP_POSITIVE_REGULATION_OF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_SIGNALING_PATHWAY
GOBP_MONOACYLGLYCEROL_CATABOLIC_PROCESS, GOBP_MONOACYLGLYCEROL_CATABOLIC_PROCESS
GOBP_AMINO_ACID_BETAINE_TRANSPORT, GOBP_AMINO_ACID_BETAINE_TRANSPORT
GOBP_POSITIVE_REGULATION_OF_INTERFERON_ALPHA_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERFERON_ALPHA_PRODUCTION
PID_WNT_CANONICAL_PATHWAY, PID_WNT_CANONICAL_PATHWAY
HP_RECURRENT_HAND_FLAPPING, HP_RECURRENT_HAND_FLAPPING
GOBP_REGULATION_OF_OLIGODENDROCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_OLIGODENDROCYTE_DIFFERENTIATION
HP_CALF_MUSCLE_PSEUDOHYPERTROPHY, HP_CALF_MUSCLE_PSEUDOHYPERTROPHY
GOCC_MUTLALPHA_COMPLEX, GOCC_MUTLALPHA_COMPLEX
GOMF_BETA_N_ACETYLHEXOSAMINIDASE_ACTIVITY, GOMF_BETA_N_ACETYLHEXOSAMINIDASE_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_CELL_MIGRATION_BY_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_CELL_MIGRATION_BY_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR
GOBP_PENETRATION_OF_ZONA_PELLUCIDA, GOBP_PENETRATION_OF_ZONA_PELLUCIDA
HP_LIMB_TREMOR, HP_LIMB_TREMOR
MIR6793_5P, MIR6793_5P
HP_RADIAL_DEVIATION_OF_THE_HAND, HP_RADIAL_DEVIATION_OF_THE_HAND
REACTOME_ORGANIC_CATION_ANION_ZWITTERION_TRANSPORT, REACTOME_ORGANIC_CATION_ANION_ZWITTERION_TRANSPORT
GOMF_GLYCINE_BINDING, GOMF_GLYCINE_BINDING
GOBP_ASTROCYTE_CELL_MIGRATION, GOBP_ASTROCYTE_CELL_MIGRATION
GOBP_SEMICIRCULAR_CANAL_DEVELOPMENT, GOBP_SEMICIRCULAR_CANAL_DEVELOPMENT
GOBP_LEFT_RIGHT_AXIS_SPECIFICATION, GOBP_LEFT_RIGHT_AXIS_SPECIFICATION
PANAPASA_BLOOD_FLUENZ_AGE_03_17YO_3DY_4DY_DN, PANAPASA_BLOOD_FLUENZ_AGE_03_17YO_3DY_4DY_DN
GOCC_BARR_BODY, GOCC_BARR_BODY
HP_PILOMATRIXOMA, HP_PILOMATRIXOMA
HP_APLASIA_HYPOPLASIA_OF_THE_NASAL_SEPTUM, HP_APLASIA_HYPOPLASIA_OF_THE_NASAL_SEPTUM
GOBP_POSITIVE_REGULATION_OF_CELLULAR_PH_REDUCTION, GOBP_POSITIVE_REGULATION_OF_CELLULAR_PH_REDUCTION
GOBP_CHORIONIC_TROPHOBLAST_CELL_PROLIFERATION, GOBP_CHORIONIC_TROPHOBLAST_CELL_PROLIFERATION
GOBP_REGULATION_OF_SNRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II, GOBP_REGULATION_OF_SNRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II
GOBP_PROTEIN_POLY_ADP_RIBOSYLATION, GOBP_PROTEIN_POLY_ADP_RIBOSYLATION
GOMF_RDNA_BINDING, GOMF_RDNA_BINDING
GOMF_GTPASE_INHIBITOR_ACTIVITY, GOMF_GTPASE_INHIBITOR_ACTIVITY
GOCC_DYSTROGLYCAN_COMPLEX, GOCC_DYSTROGLYCAN_COMPLEX
PETRETTO_LEFT_VENTRICLE_MASS_QTL_CIS_UP, PETRETTO_LEFT_VENTRICLE_MASS_QTL_CIS_UP
HP_ENDOCARDIAL_FIBROELASTOSIS, HP_ENDOCARDIAL_FIBROELASTOSIS
GOBP_DETECTION_OF_VIRUS, GOBP_DETECTION_OF_VIRUS
GOBP_HISTONE_H3_K36_TRIMETHYLATION, GOBP_HISTONE_H3_K36_TRIMETHYLATION
GOBP_REGULATION_OF_FOCAL_ADHESION_DISASSEMBLY, GOBP_REGULATION_OF_FOCAL_ADHESION_DISASSEMBLY
GOBP_VITAMIN_CATABOLIC_PROCESS, GOBP_VITAMIN_CATABOLIC_PROCESS
PID_ARF6_DOWNSTREAM_PATHWAY, PID_ARF6_DOWNSTREAM_PATHWAY
REACTOME_TRAF3_DEPENDENT_IRF_ACTIVATION_PATHWAY, REACTOME_TRAF3_DEPENDENT_IRF_ACTIVATION_PATHWAY
GOBP_POSITIVE_REGULATION_OF_SKELETAL_MUSCLE_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_SKELETAL_MUSCLE_CELL_DIFFERENTIATION
HP_IMPAIRED_THERMAL_SENSITIVITY, HP_IMPAIRED_THERMAL_SENSITIVITY
GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_CYSTEINE_S_NITROSYLATION, GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_CYSTEINE_S_NITROSYLATION
GOBP_NEURAL_PLATE_DEVELOPMENT, GOBP_NEURAL_PLATE_DEVELOPMENT
GOBP_GANGLIOSIDE_CATABOLIC_PROCESS, GOBP_GANGLIOSIDE_CATABOLIC_PROCESS
GOBP_INTESTINAL_EPITHELIAL_STRUCTURE_MAINTENANCE, GOBP_INTESTINAL_EPITHELIAL_STRUCTURE_MAINTENANCE
GOBP_EXTRAEMBRYONIC_MEMBRANE_DEVELOPMENT, GOBP_EXTRAEMBRYONIC_MEMBRANE_DEVELOPMENT
HP_FOOT_OLIGODACTYLY, HP_FOOT_OLIGODACTYLY
GOBP_GLIAL_CELL_DERIVED_NEUROTROPHIC_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_GLIAL_CELL_DERIVED_NEUROTROPHIC_FACTOR_RECEPTOR_SIGNALING_PATHWAY
HP_ESOPHAGEAL_CARCINOMA, HP_ESOPHAGEAL_CARCINOMA
GOMF_QUATERNARY_AMMONIUM_GROUP_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_QUATERNARY_AMMONIUM_GROUP_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOMF_PHOSPHATIDYLINOSITOL_3_KINASE_REGULATORY_SUBUNIT_BINDING, GOMF_PHOSPHATIDYLINOSITOL_3_KINASE_REGULATORY_SUBUNIT_BINDING
HP_CUTANEOUS_CYST, HP_CUTANEOUS_CYST
HP_THORACIC_DYSPLASIA, HP_THORACIC_DYSPLASIA
BIOCARTA_WNT_PATHWAY, BIOCARTA_WNT_PATHWAY
HP_ORBITAL_CYST, HP_ORBITAL_CYST
HP_SACROCOCCYGEAL_TERATOMA, HP_SACROCOCCYGEAL_TERATOMA
GOBP_DELAMINATION, GOBP_DELAMINATION
GOBP_REGULATION_OF_FOREBRAIN_NEURON_DIFFERENTIATION, GOBP_REGULATION_OF_FOREBRAIN_NEURON_DIFFERENTIATION
GOBP_POSITIVE_REGULATION_OF_CHEMOKINE_C_C_MOTIF_LIGAND_5_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_CHEMOKINE_C_C_MOTIF_LIGAND_5_PRODUCTION
HP_HYPERPIGMENTATION_OF_THE_FUNDUS, HP_HYPERPIGMENTATION_OF_THE_FUNDUS
GOBP_CARNITINE_TRANSPORT, GOBP_CARNITINE_TRANSPORT
GOBP_CHORION_DEVELOPMENT, GOBP_CHORION_DEVELOPMENT
GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATIONAL_INITIATION, GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATIONAL_INITIATION
HP_ADRENOCORTICAL_ADENOMA, HP_ADRENOCORTICAL_ADENOMA
WP_OSTEOPONTIN_SIGNALING, WP_OSTEOPONTIN_SIGNALING
GOBP_NEGATIVE_REGULATION_OF_MULTICELLULAR_ORGANISM_GROWTH, GOBP_NEGATIVE_REGULATION_OF_MULTICELLULAR_ORGANISM_GROWTH
GOBP_GROWTH_PLATE_CARTILAGE_CHONDROCYTE_DIFFERENTIATION, GOBP_GROWTH_PLATE_CARTILAGE_CHONDROCYTE_DIFFERENTIATION
GOBP_CELL_FATE_DETERMINATION, GOBP_CELL_FATE_DETERMINATION
GOBP_PEPTIDOGLYCAN_METABOLIC_PROCESS, GOBP_PEPTIDOGLYCAN_METABOLIC_PROCESS
GOBP_FOREBRAIN_RADIAL_GLIAL_CELL_DIFFERENTIATION, GOBP_FOREBRAIN_RADIAL_GLIAL_CELL_DIFFERENTIATION
WP_AUTOSOMAL_RECESSIVE_OSTEOPETROSIS_PATHWAYS, WP_AUTOSOMAL_RECESSIVE_OSTEOPETROSIS_PATHWAYS
MIR6788_3P, MIR6788_3P
HOEK_MONOCYTE_2011_2012_TIV_ADULT_3DY_UP, HOEK_MONOCYTE_2011_2012_TIV_ADULT_3DY_UP
GOBP_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_OXIDATIVE_STRESS, GOBP_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_OXIDATIVE_STRESS
GOBP_NEGATIVE_REGULATION_OF_DENDRITE_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_DENDRITE_DEVELOPMENT
LIU_CMYB_TARGETS_DN, LIU_CMYB_TARGETS_DN
GOMF_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOBP_LATE_ENDOSOME_TO_LYSOSOME_TRANSPORT, GOBP_LATE_ENDOSOME_TO_LYSOSOME_TRANSPORT
GOMF_PYRIMIDINE_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_PYRIMIDINE_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOBP_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORT, GOBP_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORT
GOCC_CILIARY_BASE, GOCC_CILIARY_BASE
GOBP_SMOOTHENED_SIGNALING_PATHWAY_INVOLVED_IN_DORSAL_VENTRAL_NEURAL_TUBE_PATTERNING, GOBP_SMOOTHENED_SIGNALING_PATHWAY_INVOLVED_IN_DORSAL_VENTRAL_NEURAL_TUBE_PATTERNING
GOBP_NEGATIVE_REGULATION_OF_PRI_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II, GOBP_NEGATIVE_REGULATION_OF_PRI_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II
GOCC_STEREOCILIUM_MEMBRANE, GOCC_STEREOCILIUM_MEMBRANE
HP_ALTERNATING_ESOTROPIA, HP_ALTERNATING_ESOTROPIA
GOBP_PYRIMIDINE_NUCLEOSIDE_TRANSPORT, GOBP_PYRIMIDINE_NUCLEOSIDE_TRANSPORT
GOBP_REGULATION_OF_ENDODERMAL_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_ENDODERMAL_CELL_DIFFERENTIATION
GOMF_URIDINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_URIDINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOBP_CHROMOSOME_LOCALIZATION_TO_NUCLEAR_ENVELOPE_INVOLVED_IN_HOMOLOGOUS_CHROMOSOME_SEGREGATION, GOBP_CHROMOSOME_LOCALIZATION_TO_NUCLEAR_ENVELOPE_INVOLVED_IN_HOMOLOGOUS_CHRO
GOBP_XENOBIOTIC_DETOXIFICATION_BY_TRANSMEMBRANE_EXPORT_ACROSS_THE_PLASMA_MEMBRANE, GOBP_XENOBIOTIC_DETOXIFICATION_BY_TRANSMEMBRANE_EXPORT_ACROSS_THE_PLASMA_MEMBRANE
HP_ODONTOGENIC_NEOPLASM, HP_ODONTOGENIC_NEOPLASM
HP_CALCIFICATION_OF_FALX_CEREBRI, HP_CALCIFICATION_OF_FALX_CEREBRI
HP_INCREASED_ARM_SPAN, HP_INCREASED_ARM_SPAN
GOBP_NEGATIVE_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GOBP_NEGATIVE_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I
GOBP_POST_EMBRYONIC_CAMERA_TYPE_EYE_DEVELOPMENT, GOBP_POST_EMBRYONIC_CAMERA_TYPE_EYE_DEVELOPMENT
GOBP_PEPTIDE_MODIFICATION, GOBP_PEPTIDE_MODIFICATION
GOBP_NEGATIVE_REGULATION_OF_DENDRITIC_SPINE_MORPHOGENESIS, GOBP_NEGATIVE_REGULATION_OF_DENDRITIC_SPINE_MORPHOGENESIS
GOMF_METALLOENDOPEPTIDASE_INHIBITOR_ACTIVITY, GOMF_METALLOENDOPEPTIDASE_INHIBITOR_ACTIVITY
GOBP_GLUCOSAMINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS, GOBP_GLUCOSAMINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS
HP_TALL_CHIN, HP_TALL_CHIN
REACTOME_TRAF6_MEDIATED_IRF7_ACTIVATION, REACTOME_TRAF6_MEDIATED_IRF7_ACTIVATION
GOBP_CELL_DIFFERENTIATION_IN_HINDBRAIN, GOBP_CELL_DIFFERENTIATION_IN_HINDBRAIN