L_VS_MONO_MAC_UP, GSE27786_BCELL_VS_MONO_MAC_UP
GSE27786_LIN_NEG_VS_MONO_MAC_UP, GSE27786_LIN_NEG_VS_MONO_MAC_UP
GSE27786_BCELL_VS_NEUTROPHIL_UP, GSE27786_BCELL_VS_NEUTROPHIL_UP
GSE27786_CD8_TCELL_VS_ERYTHROBLAST_UP, GSE27786_CD8_TCELL_VS_ERYTHROBLAST_UP
GSE27786_NKTCELL_VS_MONO_MAC_UP, GSE27786_NKTCELL_VS_MONO_MAC_UP
GSE27786_NKCELL_VS_MONO_MAC_UP, GSE27786_NKCELL_VS_MONO_MAC_UP
GSE27786_BCELL_VS_ERYTHROBLAST_UP, GSE27786_BCELL_VS_ERYTHROBLAST_UP
GSE27786_CD8_TCELL_VS_MONO_MAC_UP, GSE27786_CD8_TCELL_VS_MONO_MAC_UP
GSE27786_LSK_VS_MONO_MAC_UP, GSE27786_LSK_VS_MONO_MAC_UP
GSE46606_IRF4HIGH_VS_IRF4MID_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_UP, GSE46606_IRF4HIGH_VS_IRF4MID_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_UP
GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_DN, GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_DN
GSE27786_CD8_TCELL_VS_NEUTROPHIL_UP, GSE27786_CD8_TCELL_VS_NEUTROPHIL_UP
GSE27786_CD4_TCELL_VS_NEUTROPHIL_UP, GSE27786_CD4_TCELL_VS_NEUTROPHIL_UP
GSE27786_CD4_TCELL_VS_MONO_MAC_UP, GSE27786_CD4_TCELL_VS_MONO_MAC_UP
GSE16385_ROSIGLITAZONE_IL4_VS_ROSIGLITAZONE_ALONE_STIM_MACROPHAGE_DN, GSE16385_ROSIGLITAZONE_IL4_VS_ROSIGLITAZONE_ALONE_STIM_MACROPHAGE_DN
GSE20198_IL12_IL18_VS_IFNA_TREATED_ACT_CD4_TCELL_DN, GSE20198_IL12_IL18_VS_IFNA_TREATED_ACT_CD4_TCELL_DN
GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_DN, GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_DN
GSE27786_BCELL_VS_NKCELL_UP, GSE27786_BCELL_VS_NKCELL_UP
GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_UP, GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_UP
GSE23505_IL6_IL1_VS_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP, GSE23505_IL6_IL1_VS_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP
GSE23568_CTRL_VS_ID3_TRANSDUCED_CD8_TCELL_UP, GSE23568_CTRL_VS_ID3_TRANSDUCED_CD8_TCELL_UP
GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_LPS_STIM_UP, GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_LPS_STIM_UP
GSE14308_TH1_VS_INDUCED_TREG_UP, GSE14308_TH1_VS_INDUCED_TREG_UP
MORF_EIF4E, MORF_EIF4E
GSE27786_LSK_VS_BCELL_DN, GSE27786_LSK_VS_BCELL_DN
GSE14308_TH17_VS_NAIVE_CD4_TCELL_DN, GSE14308_TH17_VS_NAIVE_CD4_TCELL_DN
GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_DN
HP_MACROGLOSSIA, HP_MACROGLOSSIA
GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN, GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN
GSE36527_CD62L_HIGH_CD69_NEG_VS_CD62L_LOW_CD69_POS_TREG_KLRG1_NEG_UP, GSE36527_CD62L_HIGH_CD69_NEG_VS_CD62L_LOW_CD69_POS_TREG_KLRG1_NEG_UP
PHB2_TARGET_GENES, PHB2_TARGET_GENES
HP_SKELETAL_MUSCLE_HYPERTROPHY, HP_SKELETAL_MUSCLE_HYPERTROPHY
MORF_ERCC2, MORF_ERCC2
GOBP_SISTER_CHROMATID_COHESION, GOBP_SISTER_CHROMATID_COHESION
GSE5542_IFNG_VS_IFNA_TREATED_EPITHELIAL_CELLS_6H_DN, GSE5542_IFNG_VS_IFNA_TREATED_EPITHELIAL_CELLS_6H_DN
MORF_RAP1A, MORF_RAP1A
MORF_CDC2L5, MORF_CDC2L5
HP_FASCICULATIONS, HP_FASCICULATIONS
MIR3175, MIR3175
MORF_HEAB, MORF_HEAB
GOBP_DNA_MODIFICATION, GOBP_DNA_MODIFICATION
GOMF_DEMETHYLASE_ACTIVITY, GOMF_DEMETHYLASE_ACTIVITY
REACTOME_FORMATION_OF_THE_EARLY_ELONGATION_COMPLEX, REACTOME_FORMATION_OF_THE_EARLY_ELONGATION_COMPLEX
FAN_EMBRYONIC_CTX_BIG_GROUPS_GLIAL, FAN_EMBRYONIC_CTX_BIG_GROUPS_GLIAL
MORF_RFC5, MORF_RFC5
BURTON_ADIPOGENESIS_PEAK_AT_8HR, BURTON_ADIPOGENESIS_PEAK_AT_8HR
SMITH_TERT_TARGETS_DN, SMITH_TERT_TARGETS_DN
GOCC_CILIARY_TIP, GOCC_CILIARY_TIP
GRE_C, GRE_C
GOCC_MLL1_2_COMPLEX, GOCC_MLL1_2_COMPLEX
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN
GOBP_AUTOPHAGOSOME_MATURATION, GOBP_AUTOPHAGOSOME_MATURATION
MIR6749_3P, MIR6749_3P
NIKOLSKY_BREAST_CANCER_7Q21_Q22_AMPLICON, NIKOLSKY_BREAST_CANCER_7Q21_Q22_AMPLICON
GOBP_INTRACILIARY_TRANSPORT_INVOLVED_IN_CILIUM_ASSEMBLY, GOBP_INTRACILIARY_TRANSPORT_INVOLVED_IN_CILIUM_ASSEMBLY
GOBP_DEMETHYLATION, GOBP_DEMETHYLATION
REACTOME_SIGNALING_BY_FGFR2, REACTOME_SIGNALING_BY_FGFR2
REACTOME_SIGNALING_BY_FGFR, REACTOME_SIGNALING_BY_FGFR
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CONTAINING_COMPLEX_DISASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CONTAINING_COMPLEX_DISASSEMBLY
MIR1976, MIR1976
MIR6813_5P, MIR6813_5P
GOBP_DNA_METHYLATION_OR_DEMETHYLATION, GOBP_DNA_METHYLATION_OR_DEMETHYLATION
GOBP_REGULATION_OF_SISTER_CHROMATID_COHESION, GOBP_REGULATION_OF_SISTER_CHROMATID_COHESION
MIR6085, MIR6085
GOLUB_ALL_VS_AML_UP, GOLUB_ALL_VS_AML_UP
GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I
HP_HYPOCHROMIC_ANEMIA, HP_HYPOCHROMIC_ANEMIA
GOBP_PROTEIN_K48_LINKED_DEUBIQUITINATION, GOBP_PROTEIN_K48_LINKED_DEUBIQUITINATION
HP_ABNORMAL_MORPHOLOGY_OF_THE_ABDOMINAL_MUSCULATURE, HP_ABNORMAL_MORPHOLOGY_OF_THE_ABDOMINAL_MUSCULATURE
GOBP_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
MIR675_3P, MIR675_3P
MIR1287_3P, MIR1287_3P
HP_PROTRUDING_TONGUE, HP_PROTRUDING_TONGUE
HP_MUSCLE_FIBRILLATION, HP_MUSCLE_FIBRILLATION
MIR3923, MIR3923
MIR4759, MIR4759
WP_INTRAFLAGELLAR_TRANSPORT_PROTEINS_BINDING_TO_DYNEIN, WP_INTRAFLAGELLAR_TRANSPORT_PROTEINS_BINDING_TO_DYNEIN
GOBP_NEGATIVE_REGULATION_OF_TOR_SIGNALING, GOBP_NEGATIVE_REGULATION_OF_TOR_SIGNALING
PID_ATF2_PATHWAY, PID_ATF2_PATHWAY
HP_ABNORMAL_TONGUE_PHYSIOLOGY, HP_ABNORMAL_TONGUE_PHYSIOLOGY
MIR378E, MIR378E
MIR3681_5P, MIR3681_5P
MIR378A_3P_MIR378C_MIR378D_MIR378I, MIR378A_3P_MIR378C_MIR378D_MIR378I
GOBP_MAINTENANCE_OF_SISTER_CHROMATID_COHESION, GOBP_MAINTENANCE_OF_SISTER_CHROMATID_COHESION
TCTGATC_MIR383, TCTGATC_MIR383
GOBP_NEGATIVE_REGULATION_OF_MACROAUTOPHAGY, GOBP_NEGATIVE_REGULATION_OF_MACROAUTOPHAGY
GOBP_FATTY_ACID_DERIVATIVE_BIOSYNTHETIC_PROCESS, GOBP_FATTY_ACID_DERIVATIVE_BIOSYNTHETIC_PROCESS
REACTOME_CONSTITUTIVE_SIGNALING_BY_AKT1_E17K_IN_CANCER, REACTOME_CONSTITUTIVE_SIGNALING_BY_AKT1_E17K_IN_CANCER
MIR378H, MIR378H
MIR422A, MIR422A
MIR296_5P, MIR296_5P
MIR378F, MIR378F
PID_LYMPH_ANGIOGENESIS_PATHWAY, PID_LYMPH_ANGIOGENESIS_PATHWAY
CAAGGAT_MIR362, CAAGGAT_MIR362
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_CYAN_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_CYAN_UP
GOCC_ENDOPLASMIC_RETICULUM_EXIT_SITE, GOCC_ENDOPLASMIC_RETICULUM_EXIT_SITE
MIR8083, MIR8083
YUAN_ZNF143_PARTNERS, YUAN_ZNF143_PARTNERS
GOBP_REGULATION_OF_ACTIN_NUCLEATION, GOBP_REGULATION_OF_ACTIN_NUCLEATION
GOBP_REGULATION_OF_FILOPODIUM_ASSEMBLY, GOBP_REGULATION_OF_FILOPODIUM_ASSEMBLY
MIR6849_5P, MIR6849_5P
GOCC_PRERIBOSOME_SMALL_SUBUNIT_PRECURSOR, GOCC_PRERIBOSOME_SMALL_SUBUNIT_PRECURSOR
MIR378B, MIR378B
GOBP_GLYCEROL_ETHER_METABOLIC_PROCESS, GOBP_GLYCEROL_ETHER_METABOLIC_PROCESS
WEIGEL_OXIDATIVE_STRESS_RESPONSE, WEIGEL_OXIDATIVE_STRESS_RESPONSE
MIR4663, MIR4663
GOBP_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION, GOBP_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION
HP_MICROCYTIC_ANEMIA, HP_MICROCYTIC_ANEMIA
GOBP_ETHER_METABOLIC_PROCESS, GOBP_ETHER_METABOLIC_PROCESS
HP_TONGUE_FASCICULATIONS, HP_TONGUE_FASCICULATIONS
GOBP_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION
GOMF_SH2_DOMAIN_BINDING, GOMF_SH2_DOMAIN_BINDING
HOUSTIS_ROS, HOUSTIS_ROS
MIR4790_3P, MIR4790_3P
GOBP_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION, GOBP_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION
MIR101_5P, MIR101_5P
MATTIOLI_MULTIPLE_MYELOMA_SUBGROUPS, MATTIOLI_MULTIPLE_MYELOMA_SUBGROUPS
THRAP3_TARGET_GENES, THRAP3_TARGET_GENES
GOBP_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING, GOBP_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING
GOBP_GLANDULAR_EPITHELIAL_CELL_DEVELOPMENT, GOBP_GLANDULAR_EPITHELIAL_CELL_DEVELOPMENT
MIR4659A_5P, MIR4659A_5P
GOMF_LYSINE_ACETYLATED_HISTONE_BINDING, GOMF_LYSINE_ACETYLATED_HISTONE_BINDING
MIR4733_3P, MIR4733_3P
GOBP_RESPONSE_TO_NERVE_GROWTH_FACTOR, GOBP_RESPONSE_TO_NERVE_GROWTH_FACTOR
HP_BOWEL_DIVERTICULOSIS, HP_BOWEL_DIVERTICULOSIS
GOBP_REGULATION_OF_GOLGI_ORGANIZATION, GOBP_REGULATION_OF_GOLGI_ORGANIZATION
HP_LIMB_GIRDLE_MUSCULAR_DYSTROPHY, HP_LIMB_GIRDLE_MUSCULAR_DYSTROPHY
GOMF_LYS63_SPECIFIC_DEUBIQUITINASE_ACTIVITY, GOMF_LYS63_SPECIFIC_DEUBIQUITINASE_ACTIVITY
HP_APLASIA_HYPOPLASIA_INVOLVING_THE_FEMORAL_HEAD_AND_NECK, HP_APLASIA_HYPOPLASIA_INVOLVING_THE_FEMORAL_HEAD_AND_NECK
HP_PHIMOSIS, HP_PHIMOSIS
GOBP_ESTABLISHMENT_OF_EPITHELIAL_CELL_POLARITY, GOBP_ESTABLISHMENT_OF_EPITHELIAL_CELL_POLARITY
STAT1_01, STAT1_01
HP_HYPOSMIA, HP_HYPOSMIA
GOMF_DYNEIN_INTERMEDIATE_CHAIN_BINDING, GOMF_DYNEIN_INTERMEDIATE_CHAIN_BINDING
MIR4710, MIR4710
WP_MECP2_AND_ASSOCIATED_RETT_SYNDROME, WP_MECP2_AND_ASSOCIATED_RETT_SYNDROME
GOBP_OXIDATIVE_DEMETHYLATION, GOBP_OXIDATIVE_DEMETHYLATION
HP_OVERGROWTH, HP_OVERGROWTH
HP_DIASTASIS_RECTI, HP_DIASTASIS_RECTI
MIR4474_3P, MIR4474_3P
REACTOME_NGF_STIMULATED_TRANSCRIPTION, REACTOME_NGF_STIMULATED_TRANSCRIPTION
GOBP_POLARIZED_EPITHELIAL_CELL_DIFFERENTIATION, GOBP_POLARIZED_EPITHELIAL_CELL_DIFFERENTIATION
GOBP_POSITIVE_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY
GOCC_RNA_POLYMERASE_II_CORE_COMPLEX, GOCC_RNA_POLYMERASE_II_CORE_COMPLEX
GOMF_RNA_POLYMERASE_I_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING, GOMF_RNA_POLYMERASE_I_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING
GOBP_INTRACILIARY_RETROGRADE_TRANSPORT, GOBP_INTRACILIARY_RETROGRADE_TRANSPORT
HP_HAND_TREMOR, HP_HAND_TREMOR
GOBP_PEPTIDYL_LYSINE_MONOMETHYLATION, GOBP_PEPTIDYL_LYSINE_MONOMETHYLATION
GOMF_ARMADILLO_REPEAT_DOMAIN_BINDING, GOMF_ARMADILLO_REPEAT_DOMAIN_BINDING
GOCC_TORC2_COMPLEX, GOCC_TORC2_COMPLEX
REACTOME_PIWI_INTERACTING_RNA_PIRNA_BIOGENESIS, REACTOME_PIWI_INTERACTING_RNA_PIRNA_BIOGENESIS
MIR4535, MIR4535
HP_COLONIC_DIVERTICULA, HP_COLONIC_DIVERTICULA
HP_LIMB_GIRDLE_MUSCLE_ATROPHY, HP_LIMB_GIRDLE_MUSCLE_ATROPHY
GOCC_TOR_COMPLEX, GOCC_TOR_COMPLEX
GOBP_NEGATIVE_REGULATION_OF_CELL_AGING, GOBP_NEGATIVE_REGULATION_OF_CELL_AGING
GOBP_NEGATIVE_REGULATION_OF_CELL_MORPHOGENESIS_INVOLVED_IN_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_CELL_MORPHOGENESIS_INVOLVED_IN_DIFFERENTIATION
REACTOME_EGR2_AND_SOX10_MEDIATED_INITIATION_OF_SCHWANN_CELL_MYELINATION, REACTOME_EGR2_AND_SOX10_MEDIATED_INITIATION_OF_SCHWANN_CELL_MYELINATION
MIR6800_5P, MIR6800_5P
WP_OXIDATIVE_STRESS, WP_OXIDATIVE_STRESS
REACTOME_CYTOSOLIC_IRON_SULFUR_CLUSTER_ASSEMBLY, REACTOME_CYTOSOLIC_IRON_SULFUR_CLUSTER_ASSEMBLY
GOBP_NEGATIVE_REGULATION_OF_SECRETION, GOBP_NEGATIVE_REGULATION_OF_SECRETION
GOBP_SCHWANN_CELL_DIFFERENTIATION, GOBP_SCHWANN_CELL_DIFFERENTIATION
FARMER_BREAST_CANCER_CLUSTER_7, FARMER_BREAST_CANCER_CLUSTER_7
GOBP_ETHER_BIOSYNTHETIC_PROCESS, GOBP_ETHER_BIOSYNTHETIC_PROCESS
GOBP_INTERFERON_ALPHA_PRODUCTION, GOBP_INTERFERON_ALPHA_PRODUCTION
ATF2_UP.V1_UP, ATF2_UP.V1_UP
HP_TALIPES_CALCANEOVALGUS, HP_TALIPES_CALCANEOVALGUS
GOCC_INTRACILIARY_TRANSPORT_PARTICLE_A, GOCC_INTRACILIARY_TRANSPORT_PARTICLE_A
WP_CHOLESTEROL_BIOSYNTHESIS_WITH_SKELETAL_DYSPLASIAS, WP_CHOLESTEROL_BIOSYNTHESIS_WITH_SKELETAL_DYSPLASIAS
REACTOME_CREB_PHOSPHORYLATION, REACTOME_CREB_PHOSPHORYLATION
KORKOLA_TERATOMA_UP, KORKOLA_TERATOMA_UP
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BROWN_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BROWN_UP
GOCC_DNA_HELICASE_COMPLEX, GOCC_DNA_HELICASE_COMPLEX
MIR6724_5P, MIR6724_5P
SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX1_DN, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX1_DN
GOMF_ARRESTIN_FAMILY_PROTEIN_BINDING, GOMF_ARRESTIN_FAMILY_PROTEIN_BINDING
GOMF_LYS48_SPECIFIC_DEUBIQUITINASE_ACTIVITY, GOMF_LYS48_SPECIFIC_DEUBIQUITINASE_ACTIVITY
XU_RESPONSE_TO_TRETINOIN_DN, XU_RESPONSE_TO_TRETINOIN_DN
AR_02, AR_02
HP_SYNOVITIS, HP_SYNOVITIS
ROY_WOUND_BLOOD_VESSEL_DN, ROY_WOUND_BLOOD_VESSEL_DN
GOBP_REGULATION_OF_MAINTENANCE_OF_SISTER_CHROMATID_COHESION, GOBP_REGULATION_OF_MAINTENANCE_OF_SISTER_CHROMATID_COHESION
GOCC_TORC1_COMPLEX, GOCC_TORC1_COMPLEX
HP_GERM_CELL_NEOPLASIA, HP_GERM_CELL_NEOPLASIA
HP_DECREASED_MEAN_CORPUSCULAR_VOLUME, HP_DECREASED_MEAN_CORPUSCULAR_VOLUME
GOBP_CELLULAR_RESPONSE_TO_LEUCINE, GOBP_CELLULAR_RESPONSE_TO_LEUCINE
REACTOME_FGFR3B_LIGAND_BINDING_AND_ACTIVATION, REACTOME_FGFR3B_LIGAND_BINDING_AND_ACTIVATION
HP_THORACOLUMBAR_SCOLIOSIS, HP_THORACOLUMBAR_SCOLIOSIS
HP_SCLERODERMA, HP_SCLERODERMA
GOBP_LUNG_SACCULE_DEVELOPMENT, GOBP_LUNG_SACCULE_DEVELOPMENT
GOBP_METANEPHRIC_NEPHRON_TUBULE_MORPHOGENESIS, GOBP_METANEPHRIC_NEPHRON_TUBULE_MORPHOGENESIS
GOBP_RESPONSE_TO_PLATELET_DERIVED_GROWTH_FACTOR, GOBP_RESPONSE_TO_PLATELET_DERIVED_GROWTH_FACTOR
HP_ABNORMAL_MUSCLE_FIBER_ALPHA_DYSTROGLYCAN, HP_ABNORMAL_MUSCLE_FIBER_ALPHA_DYSTROGLYCAN
GOBP_POSTSYNAPTIC_SPECIALIZATION_ORGANIZATION, GOBP_POSTSYNAPTIC_SPECIALIZATION_ORGANIZATION
GOBP_METANEPHRIC_TUBULE_MORPHOGENESIS, GOBP_METANEPHRIC_TUBULE_MORPHOGENESIS
HP_SEPTATE_VAGINA, HP_SEPTATE_VAGINA
REACTOME_AKT_PHOSPHORYLATES_TARGETS_IN_THE_NUCLEUS, REACTOME_AKT_PHOSPHORYLATES_TARGETS_IN_THE_NUCLEUS
GAO_STOMACH_24W_C6_PIT_CELL, GAO_STOMACH_24W_C6_PIT_CELL
HP_FAIR_HAIR, HP_FAIR_HAIR
HP_ENLARGED_KIDNEY, HP_ENLARGED_KIDNEY
GOBP_NEGATIVE_REGULATION_OF_CELL_SUBSTRATE_ADHESION, GOBP_NEGATIVE_REGULATION_OF_CELL_SUBSTRATE_ADHESION
GSE29615_CTRL_VS_LAIV_FLU_VACCINE_PBMC_DN, GSE29615_CTRL_VS_LAIV_FLU_VACCINE_PBMC_DN
GOBP_NEGATIVE_REGULATION_OF_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY
GOBP_HISTONE_H3_K4_DEMETHYLATION, GOBP_HISTONE_H3_K4_DEMETHYLATION
GOBP_COMMISSURAL_NEURON_AXON_GUIDANCE, GOBP_COMMISSURAL_NEURON_AXON_GUIDANCE
GOBP_REGULATION_OF_DEACETYLASE_ACTIVITY, GOBP_REGULATION_OF_DEACETYLASE_ACTIVITY
GOMF_TYPE_1_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_BINDING, GOMF_TYPE_1_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_BINDING
WP_MITOCHONDRIAL_FATTY_ACID_SYNTHESIS_PATHWAY, WP_MITOCHONDRIAL_FATTY_ACID_SYNTHESIS_PATHWAY
GOMF_CHROMATIN_INSULATOR_SEQUENCE_BINDING, GOMF_CHROMATIN_INSULATOR_SEQUENCE_BINDING
GOBP_POSTSYNAPTIC_DENSITY_PROTEIN_95_CLUSTERING, GOBP_POSTSYNAPTIC_DENSITY_PROTEIN_95_CLUSTERING
GOBP_REGULATION_OF_HISTONE_H3_K9_TRIMETHYLATION, GOBP_REGULATION_OF_HISTONE_H3_K9_TRIMETHYLATION
GOBP_GENETIC_IMPRINTING, GOBP_GENETIC_IMPRINTING
REACTOME_ACTIVATION_OF_THE_AP_1_FAMILY_OF_TRANSCRIPTION_FACTORS, REACTOME_ACTIVATION_OF_THE_AP_1_FAMILY_OF_TRANSCRIPTION_FACTORS
GOBP_PROTEIN_K11_LINKED_DEUBIQUITINATION, GOBP_PROTEIN_K11_LINKED_DEUBIQUITINATION
HP_ABNORMAL_SPACED_INCISORS, HP_ABNORMAL_SPACED_INCISORS
GOMF_STRUCTURAL_CONSTITUENT_OF_TOOTH_ENAMEL, GOMF_STRUCTURAL_CONSTITUENT_OF_TOOTH_ENAMEL
GOBP_POSITIVE_REGULATION_OF_NUCLEAR_CELL_CYCLE_DNA_REPLICATION, GOBP_POSITIVE_REGULATION_OF_NUCLEAR_CELL_CYCLE_DNA_REPLICATION
HP_CARCINOMA, HP_CARCINOMA
MCCABE_HOXC6_TARGETS_CANCER_UP, MCCABE_HOXC6_TARGETS_CANCER_UP
TACGGGT_MIR99A_MIR100_MIR99B, TACGGGT_MIR99A_MIR100_MIR99B
GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_EARLY_ENDOSOME, GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_EARLY_ENDOSOME
GOMF_HISTONE_DEMETHYLASE_ACTIVITY_H3_K4_SPECIFIC, GOMF_HISTONE_DEMETHYLASE_ACTIVITY_H3_K4_SPECIFIC
REACTOME_FGFR3_LIGAND_BINDING_AND_ACTIVATION, REACTOME_FGFR3_LIGAND_BINDING_AND_ACTIVATION
FAN_EMBRYONIC_CTX_IN_1_INTERNEURON, FAN_EMBRYONIC_CTX_IN_1_INTERNEURON
GOBP_RESPONSE_TO_MUSCLE_INACTIVITY, GOBP_RESPONSE_TO_MUSCLE_INACTIVITY
GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H3_K36_SPECIFIC, GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H3_K36_SPECIFIC
GOBP_CENTROMERIC_SISTER_CHROMATID_COHESION, GOBP_CENTROMERIC_SISTER_CHROMATID_COHESION
GOCC_COSTAMERE, GOCC_COSTAMERE
GOBP_HEPATOCYTE_APOPTOTIC_PROCESS, GOBP_HEPATOCYTE_APOPTOTIC_PROCESS
HP_ASYMMETRY_OF_THE_THORAX, HP_ASYMMETRY_OF_THE_THORAX
HP_CONGENITAL_MUSCULAR_TORTICOLLIS, HP_CONGENITAL_MUSCULAR_TORTICOLLIS
FULLER_PBMC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_18HR_TO_336HR_SUSTAINED_UP, FULLER_PBMC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_18HR_TO_336HR_SUSTAINED_UP
MATZUK_STEROIDOGENESIS, MATZUK_STEROIDOGENESIS
GOBP_NEGATIVE_REGULATION_OF_INSULIN_SECRETION_INVOLVED_IN_CELLULAR_RESPONSE_TO_GLUCOSE_STIMULUS, GOBP_NEGATIVE_REGULATION_OF_INSULIN_SECRETION_INVOLVED_IN_CELLULAR_RESPONSE_TO_GLUCOSE_STIMULUS
HP_POSTERIOR_HELIX_PIT, HP_POSTERIOR_HELIX_PIT
BIOCARTA_MELANOCYTE_PATHWAY, BIOCARTA_MELANOCYTE_PATHWAY
BIOCARTA_HBX_PATHWAY, BIOCARTA_HBX_PATHWAY
HP_TALL_CHIN, HP_TALL_CHIN
GOBP_MAINTENANCE_OF_DNA_METHYLATION, GOBP_MAINTENANCE_OF_DNA_METHYLATION
GOBP_METANEPHRIC_DISTAL_TUBULE_DEVELOPMENT, GOBP_METANEPHRIC_DISTAL_TUBULE_DEVELOPMENT
HP_APLASIA_CUTIS_CONGENITA_OF_SCALP, HP_APLASIA_CUTIS_CONGENITA_OF_SCALP
GOMF_PEROXISOME_TARGETING_SEQUENCE_BINDING, GOMF_PEROXISOME_TARGETING_SEQUENCE_BINDING
GOBP_NEGATIVE_REGULATION_OF_PEPTIDE_SECRETION, GOBP_NEGATIVE_REGULATION_OF_PEPTIDE_SECRETION
HP_ENLARGED_EPIPHYSES, HP_ENLARGED_EPIPHYSES
REACTOME_REGULATION_OF_TP53_ACTIVITY_THROUGH_ASSOCIATION_WITH_CO_FACTORS, REACTOME_REGULATION_OF_TP53_ACTIVITY_THROUGH_ASSOCIATION_WITH_CO_FACTORS
SARTIPY_NORMAL_AT_INSULIN_RESISTANCE_DN, SARTIPY_NORMAL_AT_INSULIN_RESISTANCE_DN
REACTOME_CREB3_FACTORS_ACTIVATE_GENES, REACTOME_CREB3_FACTORS_ACTIVATE_GENES
WP_FOLATEALCOHOL_AND_CANCER_PATHWAY_HYPOTHESES, WP_FOLATEALCOHOL_AND_CANCER_PATHWAY_HYPOTHESES
MIR1537_3P, MIR1537_3P
REACTOME_FGFR1C_LIGAND_BINDING_AND_ACTIVATION, REACTOME_FGFR1C_LIGAND_BINDING_AND_ACTIVATION
REACTOME_GRB7_EVENTS_IN_ERBB2_SIGNALING, REACTOME_GRB7_EVENTS_IN_ERBB2_SIGNALING
HP_HYPOPLASIA_OF_THE_PRIMARY_TEETH, HP_HYPOPLASIA_OF_THE_PRIMARY_TEETH
HP_CLINODACTYLY_OF_THE_2ND_FINGER, HP_CLINODACTYLY_OF_THE_2ND_FINGER
GOBP_PROTEIN_K29_LINKED_DEUBIQUITINATION, GOBP_PROTEIN_K29_LINKED_DEUBIQUITINATION
GOBP_NEGATIVE_REGULATION_OF_SISTER_CHROMATID_COHESION, GOBP_NEGATIVE_REGULATION_OF_SISTER_CHROMATID_COHESION
GOCC_KICSTOR_COMPLEX, GOCC_KICSTOR_COMPLEX
HP_ABNORMAL_SARCOMERE_MORPHOLOGY, HP_ABNORMAL_SARCOMERE_MORPHOLOGY
HP_JUVENILE_CATARACT, HP_JUVENILE_CATARACT
GOMF_5_3_DNA_HELICASE_ACTIVITY, GOMF_5_3_DNA_HELICASE_ACTIVITY
MATZUK_EARLY_ANTRAL_FOLLICLE, MATZUK_EARLY_ANTRAL_FOLLICLE
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_SURFACE, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_SURFACE
GOBP_REGULATION_OF_HISTONE_DEACETYLASE_ACTIVITY, GOBP_REGULATION_OF_HISTONE_DEACETYLASE_ACTIVITY
MCCABE_HOXC6_TARGETS_UP, MCCABE_HOXC6_TARGETS_UP
GOBP_COLUMNAR_CUBOIDAL_EPITHELIAL_CELL_DEVELOPMENT, GOBP_COLUMNAR_CUBOIDAL_EPITHELIAL_CELL_DEVELOPMENT
HP_CEREBRAL_BERRY_ANEURYSM, HP_CEREBRAL_BERRY_ANEURYSM
GOMF_RNA_POLYMERASE_II_TRANSCRIPTION_COACTIVATOR_BINDING, GOMF_RNA_POLYMERASE_II_TRANSCRIPTION_COACTIVATOR_BINDING
SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX5_UP, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX5_UP
HP_TERATOMA, HP_TERATOMA
GOMF_RNA_POLYMERASE_I_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY, GOMF_RNA_POLYMERASE_I_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY
MIR100_5P_MIR99A_5P_MIR99B_5P, MIR100_5P_MIR99A_5P_MIR99B_5P
STAT3_01, STAT3_01
GOBP_LYMPHOCYTE_AGGREGATION, GOBP_LYMPHOCYTE_AGGREGATION
GOMF_TYPE_2_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_BINDING, GOMF_TYPE_2_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_BINDING
BIOCARTA_FOSB_PATHWAY, BIOCARTA_FOSB_PATHWAY
MIR1247_5P, MIR1247_5P
HP_ABNORMAL_PULMONARY_VALVE_CUSP_MORPHOLOGY, HP_ABNORMAL_PULMONARY_VALVE_CUSP_MORPHOLOGY
GOBP_POSITIVE_REGULATION_OF_MAINTENANCE_OF_SISTER_CHROMATID_COHESION, GOBP_POSITIVE_REGULATION_OF_MAINTENANCE_OF_SISTER_CHROMATID_COHESION
GOBP_REGULATION_OF_NEUTROPHIL_EXTRAVASATION, GOBP_REGULATION_OF_NEUTROPHIL_EXTRAVASATION
HP_HEMIHYPERTROPHY, HP_HEMIHYPERTROPHY
REACTOME_P38MAPK_EVENTS, REACTOME_P38MAPK_EVENTS
CHO_NR4A1_TARGETS, CHO_NR4A1_TARGETS
GOBP_CELLULAR_RESPONSE_TO_BRAIN_DERIVED_NEUROTROPHIC_FACTOR_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_BRAIN_DERIVED_NEUROTROPHIC_FACTOR_STIMULUS
GOBP_POSITIVE_REGULATION_OF_PODOSOME_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_PODOSOME_ASSEMBLY
HP_MACRODONTIA_OF_PERMANENT_MAXILLARY_CENTRAL_INCISOR, HP_MACRODONTIA_OF_PERMANENT_MAXILLARY_CENTRAL_INCISOR
GOMF_CYCLIN_DEPENDENT_PROTEIN_SERINE_THREONINE_KINASE_ACTIVATOR_ACTIVITY, GOMF_CYCLIN_DEPENDENT_PROTEIN_SERINE_THREONINE_KINASE_ACTIVATOR_ACTIVITY
HP_CONGENITAL_POSTERIOR_URETHRAL_VALVE, HP_CONGENITAL_POSTERIOR_URETHRAL_VALVE
GOBP_AMINO_ACID_SALVAGE, GOBP_AMINO_ACID_SALVAGE
GOBP_PREPULSE_INHIBITION, GOBP_PREPULSE_INHIBITION
REACTOME_NOTCH2_INTRACELLULAR_DOMAIN_REGULATES_TRANSCRIPTION, REACTOME_NOTCH2_INTRACELLULAR_DOMAIN_REGULATES_TRANSCRIPTION
GOBP_REGULATION_OF_INFLAMMATORY_RESPONSE_TO_WOUNDING, GOBP_REGULATION_OF_INFLAMMATORY_RESPONSE_TO_WOUNDING
GOMF_ARYLESTERASE_ACTIVITY, GOMF_ARYLESTERASE_ACTIVITY
FRASOR_TAMOXIFEN_RESPONSE_DN, FRASOR_TAMOXIFEN_RESPONSE_DN
MIR4520_5P, MIR4520_5P
HP_FRACTURES_OF_THE_LONG_BONES, HP_FRACTURES_OF_THE_LONG_BONES
GOBP_ASYMMETRIC_NEUROBLAST_DIVISION, GOBP_ASYMMETRIC_NEUROBLAST_DIVISION
HP_INCREASED_ARM_SPAN, HP_INCREASED_ARM_SPAN
GOBP_NEGATIVE_REGULATION_OF_DENDRITIC_SPINE_MORPHOGENESIS, GOBP_NEGATIVE_REGULATION_OF_DENDRITIC_SPINE_MORPHOGENESIS
GOBP_HISTONE_H3_K36_TRIMETHYLATION, GOBP_HISTONE_H3_K36_TRIMETHYLATION
GOBP_REGULATION_OF_FOCAL_ADHESION_DISASSEMBLY, GOBP_REGULATION_OF_FOCAL_ADHESION_DISASSEMBLY
GOBP_REGULATION_OF_RIBONUCLEOPROTEIN_COMPLEX_LOCALIZATION, GOBP_REGULATION_OF_RIBONUCLEOPROTEIN_COMPLEX_LOCALIZATION
GOMF_GLUCURONOSYL_N_ACETYLGALACTOSAMINYL_PROTEOGLYCAN_4_BETA_N_ACETYLGALACTOSAMINYLTRANSFERASE_ACTIVITY, GOMF_GLUCURONOSYL_N_ACETYLGALACTOSAMINYL_PROTEOGLYCAN_4_BETA_N_ACETYLGALACTOSAMINYLTRANSFERASE_ACTIVITY
HP_APLASIA_HYPOPLASIA_OF_THE_CAPITAL_FEMORAL_EPIPHYSIS, HP_APLASIA_HYPOPLASIA_OF_THE_CAPITAL_FEMORAL_EPIPHYSIS
GOBP_REGULATION_OF_POSTSYNAPTIC_DENSITY_ORGANIZATION, GOBP_REGULATION_OF_POSTSYNAPTIC_DENSITY_ORGANIZATION
HP_ABNORMALITY_OF_FRONTAL_SINUS, HP_ABNORMALITY_OF_FRONTAL_SINUS
HP_CALF_MUSCLE_PSEUDOHYPERTROPHY, HP_CALF_MUSCLE_PSEUDOHYPERTROPHY
HP_PROMINENT_NASAL_TIP, HP_PROMINENT_NASAL_TIP
GOBP_MAINTENANCE_OF_SYNAPSE_STRUCTURE, GOBP_MAINTENANCE_OF_SYNAPSE_STRUCTURE
HP_HYPOPROTEINEMIA, HP_HYPOPROTEINEMIA
HARALAMBIEVA_PBMC_FLUARIX_AGE_50_74YO_CORR_WITH_28D_MEM_B_CELL_RESPONSE_AT_28DY_LEUK_MIGR_MAPK_ACT_CYTOK_SIG_DIAB_OF_THE_YNG_NEGATIVE, HARALAMBIEVA_PBMC_FLUARIX_AGE_50_74YO_CORR_WITH_28D_MEM_B_CELL_RESPONSE_AT_28DY_LEUK_MIGR_MAPK_ACT_CYTOK_SIG_DIA
GOBP_METANEPHRIC_COLLECTING_DUCT_DEVELOPMENT, GOBP_METANEPHRIC_COLLECTING_DUCT_DEVELOPMENT
GOBP_NEGATIVE_REGULATION_OF_AMYLOID_BETA_CLEARANCE, GOBP_NEGATIVE_REGULATION_OF_AMYLOID_BETA_CLEARANCE
MONTERO_THYROID_CANCER_POOR_SURVIVAL_DN, MONTERO_THYROID_CANCER_POOR_SURVIVAL_DN
GOBP_RESPONSE_TO_LIGHT_INTENSITY, GOBP_RESPONSE_TO_LIGHT_INTENSITY
GOBP_KERATINOCYTE_MIGRATION, GOBP_KERATINOCYTE_MIGRATION
REACTOME_GASTRIN_CREB_SIGNALLING_PATHWAY_VIA_PKC_AND_MAPK, REACTOME_GASTRIN_CREB_SIGNALLING_PATHWAY_VIA_PKC_AND_MAPK
GOBP_RESPONSE_TO_INACTIVITY, GOBP_RESPONSE_TO_INACTIVITY
GOBP_NEGATIVE_REGULATION_OF_INTERFERON_BETA_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_INTERFERON_BETA_PRODUCTION
HP_RECURRENT_HAND_FLAPPING, HP_RECURRENT_HAND_FLAPPING
MIR184, MIR184
HP_METAPHYSEAL_STRIATIONS, HP_METAPHYSEAL_STRIATIONS
GOBP_MESONEPHRIC_DUCT_DEVELOPMENT, GOBP_MESONEPHRIC_DUCT_DEVELOPMENT
HP_LUMBAR_SCOLIOSIS, HP_LUMBAR_SCOLIOSIS
GOBP_ZYMOGEN_INHIBITION, GOBP_ZYMOGEN_INHIBITION
GOBP_POSITIVE_REGULATION_OF_SKELETAL_MUSCLE_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_SKELETAL_MUSCLE_CELL_DIFFERENTIATION
HP_WIDELY_SPACED_MAXILLARY_CENTRAL_INCISORS, HP_WIDELY_SPACED_MAXILLARY_CENTRAL_INCISORS
HP_ADVANCED_OSSIFICATION_OF_CARPAL_BONES, HP_ADVANCED_OSSIFICATION_OF_CARPAL_BONES
HP_INTRAALVEOLAR_PHOSPHOLIPID_ACCUMULATION, HP_INTRAALVEOLAR_PHOSPHOLIPID_ACCUMULATION
GLINSKY_CANCER_DEATH_UP, GLINSKY_CANCER_DEATH_UP
REACTOME_WAX_AND_PLASMALOGEN_BIOSYNTHESIS, REACTOME_WAX_AND_PLASMALOGEN_BIOSYNTHESIS
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_KINASE_B_SIGNALING, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_KINASE_B_SIGNALING
GOBP_NEGATIVE_REGULATION_OF_PEPTIDE_HORMONE_SECRETION, GOBP_NEGATIVE_REGULATION_OF_PEPTIDE_HORMONE_SECRETION
NEBEN_AML_WITH_FLT3_OR_NRAS_UP, NEBEN_AML_WITH_FLT3_OR_NRAS_UP
WP_PHASE_I_BIOTRANSFORMATIONS_NON_P450, WP_PHASE_I_BIOTRANSFORMATIONS_NON_P450
GOBP_NEGATIVE_REGULATION_OF_FIBROBLAST_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_FIBROBLAST_APOPTOTIC_PROCESS
WP_GLIAL_CELL_DIFFERENTIATION, WP_GLIAL_CELL_DIFFERENTIATION
REACTOME_FGFR2C_LIGAND_BINDING_AND_ACTIVATION, REACTOME_FGFR2C_LIGAND_BINDING_AND_ACTIVATION
HP_ABNORMAL_BRONCHOALVEOLAR_LAVAGE_FLUID_MORPHOLOGY, HP_ABNORMAL_BRONCHOALVEOLAR_LAVAGE_FLUID_MORPHOLOGY
MIR656_5P, MIR656_5P
HP_INCREASED_DENSITY_OF_LONG_BONES, HP_INCREASED_DENSITY_OF_LONG_BONES
GOCC_INSULIN_RESPONSIVE_COMPARTMENT, GOCC_INSULIN_RESPONSIVE_COMPARTMENT
GOBP_RESPONSE_TO_STIMULUS_INVOLVED_IN_REGULATION_OF_MUSCLE_ADAPTATION, GOBP_RESPONSE_TO_STIMULUS_INVOLVED_IN_REGULATION_OF_MUSCLE_ADAPTATION
GOBP_CORONARY_VASCULATURE_MORPHOGENESIS, GOBP_CORONARY_VASCULATURE_MORPHOGENESIS
GOBP_COLLECTING_DUCT_DEVELOPMENT, GOBP_COLLECTING_DUCT_DEVELOPMENT
GOBP_REGULATION_OF_GENE_EXPRESSION_BY_GENETIC_IMPRINTING, GOBP_REGULATION_OF_GENE_EXPRESSION_BY_GENETIC_IMPRINTING
ZHONG_PFC_C6_DLX5_GAD1_GAD2_POS_INTERNEURON, ZHONG_PFC_C6_DLX5_GAD1_GAD2_POS_INTERNEURON
HP_BONE_FRACTURE, HP_BONE_FRACTURE
GOCC_NODE_OF_RANVIER, GOCC_NODE_OF_RANVIER
REACTOME_TRIGLYCERIDE_BIOSYNTHESIS, REACTOME_TRIGLYCERIDE_BIOSYNTHESIS
HP_URETHROVAGINAL_FISTULA, HP_URETHROVAGINAL_FISTULA
chr2p23, chr2p23
HP_ABSENT_LACRIMAL_PUNCTUM, HP_ABSENT_LACRIMAL_PUNCTUM
GOBP_BRANCHIOMOTOR_NEURON_AXON_GUIDANCE, GOBP_BRANCHIOMOTOR_NEURON_AXON_GUIDANCE
GOBP_APOPTOTIC_CHROMOSOME_CONDENSATION, GOBP_APOPTOTIC_CHROMOSOME_CONDENSATION
GOBP_RESPONSE_TO_PHENYLPROPANOID, GOBP_RESPONSE_TO_PHENYLPROPANOID
GOMF_NEUREGULIN_BINDING, GOMF_NEUREGULIN_BINDING
GOBP_PROXIMAL_TUBULE_DEVELOPMENT, GOBP_PROXIMAL_TUBULE_DEVELOPMENT
HP_SACROCOCCYGEAL_TERATOMA, HP_SACROCOCCYGEAL_TERATOMA
HP_ABNORMALITY_OF_MESENTERY_MORPHOLOGY, HP_ABNORMALITY_OF_MESENTERY_MORPHOLOGY
GOBP_REGULATION_OF_MRNA_EXPORT_FROM_NUCLEUS, GOBP_REGULATION_OF_MRNA_EXPORT_FROM_NUCLEUS
GOBP_HISTONE_H3_K4_DEMETHYLATION_TRIMETHYL_H3_K4_SPECIFIC, GOBP_HISTONE_H3_K4_DEMETHYLATION_TRIMETHYL_H3_K4_SPECIFIC
GOBP_METANEPHRIC_LOOP_OF_HENLE_DEVELOPMENT, GOBP_METANEPHRIC_LOOP_OF_HENLE_DEVELOPMENT
GOCC_CIA_COMPLEX, GOCC_CIA_COMPLEX
GOMF_ERBB_3_CLASS_RECEPTOR_BINDING, GOMF_ERBB_3_CLASS_RECEPTOR_BINDING
REACTOME_MECP2_REGULATES_TRANSCRIPTION_FACTORS, REACTOME_MECP2_REGULATES_TRANSCRIPTION_FACTORS
GOBP_PROTEIN_DEGLUTAMYLATION, GOBP_PROTEIN_DEGLUTAMYLATION
GOBP_HISTONE_H2A_K119_MONOUBIQUITINATION, GOBP_HISTONE_H2A_K119_MONOUBIQUITINATION
GOBP_PROTEIN_LOCALIZATION_TO_BASOLATERAL_PLASMA_MEMBRANE, GOBP_PROTEIN_LOCALIZATION_TO_BASOLATERAL_PLASMA_MEMBRANE
TURJANSKI_MAPK8_AND_MAPK9_TARGETS, TURJANSKI_MAPK8_AND_MAPK9_TARGETS
HP_ABSENT_FRONTAL_SINUSES, HP_ABSENT_FRONTAL_SINUSES
HP_EEG_WITH_FOCAL_SPIKE_WAVES, HP_EEG_WITH_FOCAL_SPIKE_WAVES
HP_FIBROSARCOMA, HP_FIBROSARCOMA
GOBP_SERINE_FAMILY_AMINO_ACID_CATABOLIC_PROCESS, GOBP_SERINE_FAMILY_AMINO_ACID_CATABOLIC_PROCESS
GOMF_1_PHOSPHATIDYLINOSITOL_3_KINASE_REGULATOR_ACTIVITY, GOMF_1_PHOSPHATIDYLINOSITOL_3_KINASE_REGULATOR_ACTIVITY
GOBP_FATTY_ACID_HOMEOSTASIS, GOBP_FATTY_ACID_HOMEOSTASIS
GOBP_LUNG_EPITHELIUM_DEVELOPMENT, GOBP_LUNG_EPITHELIUM_DEVELOPMENT
HP_ERYTHROID_HYPERPLASIA, HP_ERYTHROID_HYPERPLASIA
REACTOME_PHOSPHOLIPASE_C_MEDIATED_CASCADE_FGFR4, REACTOME_PHOSPHOLIPASE_C_MEDIATED_CASCADE_FGFR4
HP_PATCHY_CHANGES_OF_BONE_MINERAL_DENSITY, HP_PATCHY_CHANGES_OF_BONE_MINERAL_DENSITY
HP_NEOPLASM_OF_THE_MIDDLE_EAR, HP_NEOPLASM_OF_THE_MIDDLE_EAR
GOBP_NEGATIVE_REGULATION_OF_INTERFERON_ALPHA_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_INTERFERON_ALPHA_PRODUCTION
MIR1973, MIR1973
HP_GASTRIC_ULCER, HP_GASTRIC_ULCER
FAN_EMBRYONIC_CTX_IN_3_INTERNEURON, FAN_EMBRYONIC_CTX_IN_3_INTERNEURON
ZHONG_PFC_C2_ASCL1_POS_ASTROCYTE, ZHONG_PFC_C2_ASCL1_POS_ASTROCYTE
GOBP_GERMINAL_CENTER_B_CELL_DIFFERENTIATION, GOBP_GERMINAL_CENTER_B_CELL_DIFFERENTIATION
GOBP_PENTOSE_PHOSPHATE_SHUNT_NON_OXIDATIVE_BRANCH, GOBP_PENTOSE_PHOSPHATE_SHUNT_NON_OXIDATIVE_BRANCH
GOBP_NEGATIVE_REGULATION_OF_AUTOPHAGOSOME_MATURATION, GOBP_NEGATIVE_REGULATION_OF_AUTOPHAGOSOME_MATURATION
GOBP_NEGATIVE_REGULATION_OF_DENDRITE_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_DENDRITE_DEVELOPMENT
GOMF_1_4_ALPHA_OLIGOGLUCAN_PHOSPHORYLASE_ACTIVITY, GOMF_1_4_ALPHA_OLIGOGLUCAN_PHOSPHORYLASE_ACTIVITY
REACTOME_METHIONINE_SALVAGE_PATHWAY, REACTOME_METHIONINE_SALVAGE_PATHWAY
REACTOME_FGFR2_LIGAND_BINDING_AND_ACTIVATION, REACTOME_FGFR2_LIGAND_BINDING_AND_ACTIVATION
chr7q21, chr7q21
GOMF_PHOSPHATIDATE_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDATE_PHOSPHATASE_ACTIVITY
GOBP_HEMOGLOBIN_METABOLIC_PROCESS, GOBP_HEMOGLOBIN_METABOLIC_PROCESS
MIR3649, MIR3649
HOEK_MYELOID_DENDRITIC_CELL_2011_2012_TIV_ADULT_7DY_DN, HOEK_MYELOID_DENDRITIC_CELL_2011_2012_TIV_ADULT_7DY_DN
GOCC_DYSTROGLYCAN_COMPLEX, GOCC_DYSTROGLYCAN_COMPLEX
GOBP_RESPONSE_TO_L_ASCORBIC_ACID, GOBP_RESPONSE_TO_L_ASCORBIC_ACID
GOMF_SPHINGOSINE_1_PHOSPHATE_PHOSPHATASE_ACTIVITY, GOMF_SPHINGOSINE_1_PHOSPHATE_PHOSPHATASE_ACTIVITY
HP_U_SHAPED_UPPER_LIP_VERMILION, HP_U_SHAPED_UPPER_LIP_VERMILION
HP_SHOCK, HP_SHOCK
GOBP_REGULATION_OF_CELLULAR_PH_REDUCTION, GOBP_REGULATION_OF_CELLULAR_PH_REDUCTION
GOCC_MICROTUBULE_MINUS_END, GOCC_MICROTUBULE_MINUS_END
HP_AKINESIA, HP_AKINESIA
GOBP_SERTOLI_CELL_DIFFERENTIATION, GOBP_SERTOLI_CELL_DIFFERENTIATION
GOMF_WNT_ACTIVATED_RECEPTOR_ACTIVITY, GOMF_WNT_ACTIVATED_RECEPTOR_ACTIVITY
GOBP_REGULATION_OF_MUSCLE_ORGAN_DEVELOPMENT, GOBP_REGULATION_OF_MUSCLE_ORGAN_DEVELOPMENT
HP_NEONATAL_HYPOGLYCEMIA, HP_NEONATAL_HYPOGLYCEMIA
ZNF697_TARGET_GENES, ZNF697_TARGET_GENES
KOHOUTEK_CCNT2_TARGETS, KOHOUTEK_CCNT2_TARGETS
GOMF_MAP_KINASE_ACTIVITY, GOMF_MAP_KINASE_ACTIVITY
PLASARI_TGFB1_TARGETS_1HR_DN, PLASARI_TGFB1_TARGETS_1HR_DN
HP_INCREASED_VLDL_CHOLESTEROL_CONCENTRATION, HP_INCREASED_VLDL_CHOLESTEROL_CONCENTRATION
HP_PREMATURE_LOSS_OF_PERMANENT_TEETH, HP_PREMATURE_LOSS_OF_PERMANENT_TEETH
GOBP_MEMBRANE_TO_MEMBRANE_DOCKING, GOBP_MEMBRANE_TO_MEMBRANE_DOCKING
GOBP_NEGATIVE_REGULATION_OF_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY
GOBP_REGULATION_OF_CARDIAC_MUSCLE_TISSUE_DEVELOPMENT, GOBP_REGULATION_OF_CARDIAC_MUSCLE_TISSUE_DEVELOPMENT
GOBP_SYNAPTIC_TRANSMISSION_GLYCINERGIC, GOBP_SYNAPTIC_TRANSMISSION_GLYCINERGIC
BIOCARTA_ERBB3_PATHWAY, BIOCARTA_ERBB3_PATHWAY
GOBP_L_METHIONINE_SALVAGE_FROM_METHYLTHIOADENOSINE, GOBP_L_METHIONINE_SALVAGE_FROM_METHYLTHIOADENOSINE
GOBP_POSITIVE_REGULATION_OF_CELLULAR_PH_REDUCTION, GOBP_POSITIVE_REGULATION_OF_CELLULAR_PH_REDUCTION
GOBP_TRANSFORMING_GROWTH_FACTOR_BETA3_PRODUCTION, GOBP_TRANSFORMING_GROWTH_FACTOR_BETA3_PRODUCTION
GOBP_PROTEIN_LOCALIZATION_TO_SITE_OF_DOUBLE_STRAND_BREAK, GOBP_PROTEIN_LOCALIZATION_TO_SITE_OF_DOUBLE_STRAND_BREAK
GOBP_STRESS_INDUCED_PREMATURE_SENESCENCE, GOBP_STRESS_INDUCED_PREMATURE_SENESCENCE
HP_ZOLLINGER_ELLISON_SYNDROME, HP_ZOLLINGER_ELLISON_SYNDROME
HP_LIMITATION_OF_MOVEMENT_AT_ANKLES, HP_LIMITATION_OF_MOVEMENT_AT_ANKLES
GOBP_TRANS_SYNAPTIC_SIGNALING_BY_TRANS_SYNAPTIC_COMPLEX, GOBP_TRANS_SYNAPTIC_SIGNALING_BY_TRANS_SYNAPTIC_COMPLEX
HP_ABNORMALITY_OF_THE_MASTOID, HP_ABNORMALITY_OF_THE_MASTOID
WP_BILE_ACIDS_SYNTHESIS_AND_ENTEROHEPATIC_CIRCULATION, WP_BILE_ACIDS_SYNTHESIS_AND_ENTEROHEPATIC_CIRCULATION
GOBP_IMMUNOLOGICAL_SYNAPSE_FORMATION, GOBP_IMMUNOLOGICAL_SYNAPSE_FORMATION
GOBP_PHOTORECEPTOR_CELL_OUTER_SEGMENT_ORGANIZATION, GOBP_PHOTORECEPTOR_CELL_OUTER_SEGMENT_ORGANIZATION
HP_NEOPLASM_OF_THE_LARYNX, HP_NEOPLASM_OF_THE_LARYNX
FAN_EMBRYONIC_CTX_IN_4_INTERNEURON, FAN_EMBRYONIC_CTX_IN_4_INTERNEURON
GOBP_PROSTATE_GLAND_GROWTH, GOBP_PROSTATE_GLAND_GROWTH
GOBP_OSTEOBLAST_DEVELOPMENT, GOBP_OSTEOBLAST_DEVELOPMENT
GOBP_CELLULAR_RESPONSE_TO_CHOLESTEROL, GOBP_CELLULAR_RESPONSE_TO_CHOLESTEROL
BIOCARTA_CIRCADIAN_PATHWAY, BIOCARTA_CIRCADIAN_PATHWAY
GOMF_AMINOACYLASE_ACTIVITY, GOMF_AMINOACYLASE_ACTIVITY
GOMF_LAMININ_1_BINDING, GOMF_LAMININ_1_BINDING
GOBP_NEGATIVE_REGULATION_OF_NITRIC_OXIDE_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_NITRIC_OXIDE_METABOLIC_PROCESS
GOBP_CELL_AGGREGATION, GOBP_CELL_AGGREGATION
GOBP_NEGATIVE_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS
GOBP_REGULATION_OF_APPETITE, GOBP_REGULATION_OF_APPETITE
HP_POSTPRANDIAL_HYPERGLYCEMIA, HP_POSTPRANDIAL_HYPERGLYCEMIA
CGCAAAA_MIR450, CGCAAAA_MIR450
GOBP_SERTOLI_CELL_DEVELOPMENT, GOBP_SERTOLI_CELL_DEVELOPMENT
chrXq12, chrXq12
GOMF_LIPID_PHOSPHATASE_ACTIVITY, GOMF_LIPID_PHOSPHATASE_ACTIVITY
GOBP_CELLULAR_RESPONSE_TO_ZINC_ION, GOBP_CELLULAR_RESPONSE_TO_ZINC_ION
GOBP_SPINAL_CORD_MOTOR_NEURON_DIFFERENTIATION, GOBP_SPINAL_CORD_MOTOR_NEURON_DIFFERENTIATION
WENG_POR_TARGETS_GLOBAL_UP, WENG_POR_TARGETS_GLOBAL_UP
GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_FOOD, GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_FOOD
GOBP_RESPONSE_TO_FOOD, GOBP_RESPONSE_TO_FOOD
GOBP_REGULATION_OF_RESPONSE_TO_FOOD, GOBP_REGULATION_OF_RESPONSE_TO_FOOD