

GSE2589_HEALTHY_VS_HIV_AND_SIV_INFECTED_DN,GSE22589_HEALTHY_VS_HIV_AND_SIV_INFECTED_DN,GSE2601_DOUBLE_NEGATIVE_VS_CD4_SINGLE_POSITIVE_THYMOCYTE_UP,GSE22601_DOUBLE_NEGATIVE_VS_CD4_SINGLE_POSITIVE_THYMOCYTE_UP,GSE32423_MEMORY_VS_NAIVE_CD8_TCELL_IL7_IL4_DN,GSE32423_MEMORY_VS_NAIVE_CD8_TCELL_IL7_IL4_DN,LAKE_ADULT_KIDNEY_C26_MESANGIAL_CELLS,LAKE_ADULT_KIDNEY_C26_MESANGIAL_CELLS,GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL6_KO_MACROPHAGE_45MIN_UP,GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL6_KO_MACROPHAGE_45MIN_UP,WP_EGFEGR_SIGNALING_PATHWAY,WP_EGFEGR_SIGNALING_PATHWAY,GSE31082_CD4_VS_CD8_SP_THYMOCYTE_UP,GSE31082_CD4_VS_CD8_SP_THYMOCYTE_UP,GSE19825_CD24LOW_VS_IL2RA_HIGH_DAY3_EFF_CD8_TCELL_UP,GSE19825_CD24LOW_VS_IL2RA_HIGH_DAY3_EFF_CD8_TCELL_UP,GSE8835_HEALTHY_VS_CLI_CD8_TCELL_UP,GSE8835_HEALTHY_VS_CLI_CD8_TCELL_UP,GSE33424_CD161_INT_VS_NEG_CD8_TCELL_DN,GSE33424_CD161_INT_VS_NEG_CD8_TCELL_DN,GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_48H_UP,GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_48H_UP,GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN,GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN,GSE37301_CD4_TCELL_VS_RAG2_KO_NK_CELL_DN,GSE37301_CD4_TCELL_VS_RAG2_KO_NK_CELL_DN,GSE32255_UNSTIM_VS_4H_LPS_STIM_DC_UP,GSE32255_UNSTIM_VS_4H_LPS_STIM_DC_UP,MIR499B_3P,MIR499B_3P,GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TREG_DN,GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TREG_DN,MIR499A_3P,MIR499A_3P,GSE5503_MLN_DC_VS_PLN_DC_ACTIVATED_ALLOGENIC_TCELL_DN,GSE5503_MLN_DC_VS_PLN_DC_ACTIVATED_ALLOGENIC_TCELL_DN,GSE39916_B_CELL_SPLEEN_VS_PLASMA_CELL_BONE_MARROW_UP,GSE39916_B_CELL_SPLEEN_VS_PLASMA_CELL_BONE_MARROW_UP,MIR302A_5P,MIR302A_5P,MIR7161_5P,MIR7161_5P,GSE24142_ADULT_VS_FETAL_EARLY_THYMIC_PROGENITOR_UP,GSE24142_ADULT_VS_FETAL_EARLY_THYMIC_PROGENITOR_UP,GSE25088_WT_VS_STAT6_KO_MACROPHAGE_ROSIGLITAZONE_STIM_DN,GSE25088_WT_VS_STAT6_KO_MACROPHAGE_ROSIGLITAZONE_STIM_DN,GSE2770_IL12_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_DN,GSE2770_IL12_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_DN,GSE27241_WT_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_THI17_POLARIZING_CONDITIONS_UP,GSE27241_WT_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_THI17_POLARIZING_CONDITIONS_UP,MIR132_3P,MIR132_3P,MIR212_3P,MIR212_3P,GSE15735_2H_VS_12H_HDAC_INHIBITOR_TREATED_CD4_TCELL_UP,GSE15735_2H_VS_12H_HDAC_INHIBITOR_TREATED_CD4_TCELL_UP,GSE21360_NAIVE_VS_PRIMARY_MEMORY_CD8_TCELL_UP,GSE21360_NAIVE_VS_PRIMARY_MEMORY_CD8_TCELL_UP,LAKE_ADULT_KIDNEY_C22_ENDOTHELIAL_CELLS_GLOMERULAR_CAPILLARIES,LAKE_ADULT_KIDNEY_C22_ENDOTHELIAL_CELLS_GLOMERULAR_CAPILLARIES,MIR3064_5P,MIR3064_5P,MIR6720_5P,MIR6720_5P,MIR548U,MIR548U,GSE29949_MICROGLIA_VS_DC_BRAIN_UP,GSE29949_MICROGLIA_VS_DC_BRAIN_UP,GSE17721_CPG_VS_GARDIQUIMOD_1H_BMDC_UP,GSE17721_CPG_VS_GARDIQUIMOD_1H_BMDC_UP,MIR223_3P,MIR223_3P,MIR6504_5P,MIR6504_5P,MIR561_5P,MIR561_5P,GSE17721_LPS_VS_GARDIQUIMOD_16H_BMDC_UP,GSE17721_LPS_VS_GARDIQUIMOD_16H_BMDC_UP,MIR450A_1_3P,MIR450A_1_3P,GSE32533_WT_VS_MIR17_KO_ACT_CD4_TCELL_UP,GSE32533_WT_VS_MIR17_KO_ACT_CD4_TCELL_UP,MIR130B_5P,MIR130B_5P,MIR103A_2_5P,MIR103A_2_5P,MIR499B_5P,MIR499B_5P,GSE12392_IFNAR_KO_VS_IFNB_KO_CD8_NEG_SPLEEN_DC_DN,GSE12392_IFNAR_KO_VS_IFNB_KO_CD8_NEG_SPLEEN_DC_DN,MIR1245B_3P,MIR1245B_3P,TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_8D_UP,TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_8D_UP,MIR103A_1_5P,MIR103A_1_5P,MIR6882_3P,MIR6882_3P,GSE17721_LPS_VS_POLYIC_4H_BMDC_UP,GSE17721_LPS_VS_POLYIC_4H_BMDC_UP,MIR7162_5P,MIR7162_5P,GSE17721_PAM3CSK4_VS_GADIQUIMOD_16H_BMDC_UP,GSE17721_PAM3CSK4_VS_GADIQUIMOD_16H_BMDC_UP,MIR3611,MIR3611,GOBP_REGULATION_OF_CELL_SIZE,GOBP_REGULATION_OF_CELL_SIZE,PID_CXCR4_PATHWAY,PID_CXCR4_PATHWAY,WP_GASTRIN_SIGNALING_PATHWAY,WP_GASTRIN_SIGNALING_PATHWAY,MIR4679,MIR4679,HUTTMANN_B_CELL_POOR_SURVIVAL_DN,HUTTMANN_B_CELL_POOR_SURVIVAL_DN,MIR516A_3P,MIR516B_3P,MIR516A_3P,MIR516B_3P,MIR578,MIR578,MIR8076,MIR8076,MIR149_5P,MIR149_5P,IVANOVSKA_MIR106B_TARGETS,IVANOVSKA_MIR106B_TARGETS,WP_ERBB_SIGNALING_PATHWAY,WP_ERBB_SIGNALING_PATHWAY,GGAMTNNNNNTCCY_UNKNOWN,GGAMTNNNNNTCCY_UNKNOWN,GSE11924_TH2_VS_THI17_CD4_TCELL_UP,GSE11924_TH2_VS_THI17_CD4_TCELL_UP,GSE11961_FOLLICULAR_BCELL_VS_MARGINAL_ZONE_BCELL_UP,GSE11961_FOLLICULAR_BCELL_VS_MARGINAL_ZONE_BCELL_UP,GINESTIER_BREAST_CANCER_20Q13_AMPLIFICATION_UP,GINESTIER_BREAST_CANCER_20Q13_AMPLIFICATION_UP,MIR3744_3P,MIR3744_3P,GSE18893_TCONV_VS_TREG_2H_TNF_STIM_UP,GSE18893_TCONV_VS_TREG_2H_TNF_STIM_UP,WTCNATGCGY_UNKNOWN,WTCNATGCGY_UNKNOWN,AACGTGAC,MIR223,AACGTGAC,MIR223,GSE43955_1H_VS_60H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN,GSE43955_1H_VS_60H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN,MIR6758_3P,MIR6758_3P,HOO1_S17_TARGETS_DN,HOO1_S17_TARGETS_DN,MIR4708_5P,MIR4708_5P,GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP,GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP,MIR4530,MIR4530,GSE7831_1H_VS_4H_INFLUENZA_STIM_PDC_DN,GSE7831_1H_VS_4H_INFLUENZA_STIM_PDC_DN,MIR876_5P,MIR876_5P,MENON_FETAL_KIDNEY_3_STROMAL_CELLS,MENON_FETAL_KIDNEY_3_STROMAL_CELLS,GOBP_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION,GOBP_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION,MIR617,MIR617,MIR6782_3P,MIR6782_3P,KEGG_ERBB_SIGNALING_PATHWAY,KEGG_ERBB_SIGNALING_PATHWAY,WP_RAS_SIGNALING,WP_RAS_SIGNALING,GOLDRATH_IMMUNE_MEMORY,GOLDRATH_IMMUNE_MEMORY,KEGG_RENAL_CELL_CARCINOMA,KEGG_RENAL_CELL_CARCINOMA,MIR4640_5P,MIR4640_5P,WP_CHROMOSOMAL_AND_MICROSATELLITE_INSTABILITY_IN_COLORECTAL_CANCER,WP_CHROMOSOMAL_AND_MICROSATELLITE_INSTABILITY_IN_COLORECTAL_CANCER,MIR3168,MIR3168,GGCKCATGS_UNKNOWN,GGCKCATGS_UNKNOWN,GSE13493_CD4INTCD8POS_VS_CD8POS_THYMOCYTE_DN,GSE13493_CD4INTCD8POS_VS_CD8POS_THYMOCYTE_DN,GOBP_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT,GOBP_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT,MIR7155_5P,MIR7155_5P,BROWNE_HCMV_INFECTION_20HR_DN,BROWNE_HCMV_INFECTION_20HR_DN,GSE23925_LIGHT_ZONE_VS_DARK_ZONE_BCELL_UP,GSE23925_LIGHT_ZONE_VS_DARK_ZONE_BCELL_UP,MIR643,MIR643,LIM_MAMMARY_LUMINAL_MATURE_DN,LIM_MAMMARY_LUMINAL_MATURE_DN,MIR3167,MIR3167,MIR384,MIR384,MIR5000_3P,MIR5000_3P,MIR4527,MIR4527,MIR6748_5P,MIR6748_5P,WWTAAGGC_UNKNOWN,WWTAAGGC_UNKNOWN,MIR324_3P,MIR324_3P,MIR675_3P,MIR675_3P,PID_RET_PATHWAY,PID_RET_PATHWAY,GOBP_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT,GOBP_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT,MIR4445_5P,MIR4445_5P,GOBP_CELLULAR_RESPONSE_TO_CARBOHYDRATE_STIMULUS,GOBP_CELLULAR_RESPONSE_TO_CARBOHYDRATE_STIMULUS,MIR4436A,MIR4436A,GOMF_ACTIVATING_TRANSCRIPTION_FACTOR_BINDING,GOMF_ACTIVATING_TRANSCRIPTION_FACTOR_BINDING,MIR6503_5P,MIR6503_5P,RB_DN.V1_DN,RB_DN.V1_DN,GOBP_RHO_PROTEIN_SIGNAL_TRANSDUCTION,GOBP_RHO_PROTEIN_SIGNAL_TRANSDUCTION,HINATA_NFKB_TARGETS_FIBROBLAST_UP,HINATA_NFKB_TARGETS_FIBROBLAST_UP,MIR6842_3P,MIR6842_3P,MIR592,MIR592,MIR4704_5P,MIR4704_5P,MIR3132,MIR3132,BEIER_GLIOMA_STEM_CELL_DN,BEIER_GLIOMA_STEM_CELL_DN,GOBP_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT,GOBP_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT,PID_P38_ALPHA_BETA_DOWNSTREAM_PATHWAY,PID_P38_ALPHA_BETA_DOWNSTREAM_PATHWAY,GOBP_HEART_VALVE_DEVELOPMENT,GOBP_HEART_VALVE_DEVELOPMENT,HAN_JNK_SINGALING_UP,HAN_JNK_SINGALING_UP,REACTOME_REGULATION_OF_MECP2_EXPRESSION_AND_ACTIVITY,REACTOME_REGULATION_OF_MECP2_EXPRESSION_AND_ACTIVITY,MIR6888_5P,MIR6888_5P,REACTOME_RUNX1_INTERACTS_WITH_CO_FACTORS_WHOSE_PRECISE_EFFECT_ON_RUNX1_TARGETS_IS_NOT_KNOWN,REACTOME_RUNX1_INTERACTS_WITH_CO_FACTORS_WHOSE_PRECISE_EFFECT_ON_RUNX1_TARGETS_IS_NOT_KNOWN,GSE22025_TGFB1_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_DN,GSE22025_TGFB1_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_DN,MIR371B_3P,MIR371B_3P,GOBP_KIDNEY_EPITHELIUM_DEVELOPMENT,GOBP_KIDNEY_EPITHELIUM_DEVELOPMENT,MIR6793_3P,MIR6793_3P,GOBP_CHONDROCYTE_DIFFERENTIATION,GOBP_CHONDROCYTE_DIFFERENTIATION,MIR6762_3P,MIR6762_3P,BOYLAN_MULTIPLE_MYELOMA_C_CLUSTER_DN,BOYLAN_MULTIPLE_MYELOMA_C_CLUSTER_DN,MIR2861,MIR2861,BOYLAN_MULTIPLE_MYELOMA_C_DN,BOYLAN_MULTIPLE_MYELOMA_C_DN,MIR106A_3P,MIR106A_3P,GOBP_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION,GOBP_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION,MIR3689D,MIR3689D,CAGNYGKNAAA_UNKNOWN,CAGNYGKNAAA_UNKNOWN,BALDWIN_PRKCL_TARGETS_UP,BALDWIN_PRKCL_TARGETS_UP,DUNNE_TARGETS_OF_AML1_MTG8_FUSION_DN,DUNNE_TARGETS_OF_AML1_MTG8_FUSION_DN,BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_DN,BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_DN,GOBP_NEPHRON_EPITHELIUM_DEVELOPMENT,GOBP_NEPHRON_EPITHELIUM_DEVELOPMENT,GSE21927_SPLENIC_C26GM_TUMOROUS_VS_BONE_MARROW_MONOCYTES_UP,GSE21927_SPLENIC_C26GM_TUMOROUS_VS_BONE_MARROW_MONOCYTES_UP,GOBP_CARTILAGE_DEVELOPMENT,GOBP_CARTILAGE_DEVELOPMENT,TCGA_GLOBLASTOMA_COPY_NUMBER_UP,TCGA_GLOBLASTOMA_COPY_NUMBER_UP,GOBP_REGULATION_OF_RHO_PROTEIN_SIGNAL_TRANSDUCTION,GOBP_REGULATION_OF_RHO_PROTEIN_SIGNAL_TRANSDUCTION,HP_FOCAL_DYSTONIA,HP_FOCAL_DYSTONIA,MIR3663_5P,MIR3663_5P,WP_COPPER_HOMEOSTASIS,WP_COPPER_HOMEOSTASIS,MIR335_5P,MIR335_5P,HP_PORENCEPHALIC_CYST,HP_PORENCEPHALIC_CYST,MURAKAMI_UV_RESPONSE_6HR_UP,MURAKAMI_UV_RESPONSE_6HR_UP,HP_NEOPLASM_OF_THE_LIVER,HP_NEOPLASM_OF_THE_LIVER,GOBP_CHONDROCYTE_DEVELOPMENT,GOBP_CHONDROCYTE_DEVELOPMENT,MIR412_3P,MIR412_3P,WP_MESODERMAL_COMMITMENT_PATHWAY,WP_MESODERMAL_COMMITMENT_PATHWAY,GOBP_atrial_SePTUM_Development,GOBP_atrial_SePTUM_Development,MIR4687_5P,MIR4687_5P,CHANG_IMMORTALIZED_BY_HPV31_UP,CHANG_IMMORTALIZED_BY_HPV31_UP,MIR12126,MIR12126,MIR135A_3P,MIR135A_3P,HP_HEPATOCYLLULAR_CARCINOMA,HP_HEPATOCYLLULAR_CARCINOMA,GOBP_PERIPHERAL_NERVOUS_SYSTEM_DEVELOPMENT,GOBP_PERIPHERAL_NERVOUS_SYSTEM_DEVELOPMENT,GSE13522_CTRL_VS_T_CRUZLY_STRAIN_INF_SKIN_BALBC_MOUSE_DN,GSE13522_CTRL_VS_T_CRUZLY_STRAIN_INF_SKIN_BALBC_MOUSE_DN,GOBP_POSITIVE_REGULATION_OF_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY,GOBP_POSITIVE_REGULATION_OF_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY,GOBP_RESPONSE_TO_BMP,GOBP_RESPONSE_TO_BMP,HP_ABNORMALITY_OF_THE_METOPIC_SUTURE,HP_ABNORMALITY_OF_THE_METOPIC_SUTURE,SHL_SPARC_TARGETS_UP,SHL_SPARC_TARGETS_UP,YNTTTNNNANGCARM_UNKNOWN,YNTTTNNNANGCARM_UNKNOWN,WP_MELATONIN_METABOLISM_AND_EFFECTS,WP_MELATONIN_METABOLISM_AND_EFFECTS,HP_METAPHYSEAL_IRREGULARITY,HP_METAPHYSEAL_IRREGULARITY,KEGG_VIRAL_MYOCARDITIS,KEGG_VIRAL_MYOCARDITIS,MIR6503_3P,MIR6503_3P,GOBP_CELLULAR_RESPONSE_TO_INTERFERON_BETA,GOBP_CELLULAR_RESPONSE_TO_INTERFERON_BETA,IIZUKA_LIVER_CANCER_EARLY_RECURRENCE,IIZUKA_LIVER_CANCER_EARLY_RECURRENCE,GOBP_RENAL_TUBULE_DEVELOPMENT,GOBP_RENAL_TUBULE_DEVELOPMENT,MIR509_3P,MIR509_3P,GOBP_TOLL_LIKE_RECEPTOR_9_SIGNALING_PATHWAY,GOBP_TOLL_LIKE_RECEPTOR_9_SIGNALING_PATHWAY,GOBP_CARDIAC_ATRIUM_DEVELOPMENT,GOBP_CARDIAC_ATRIUM_DEVELOPMENT,HP_EPISTAXIS,HP_EPISTAXIS,DISTECHE_ESCAPED_FROM_X_INACTIVATION,DISTECHE_ESCAPED_FROM_X_INACTIVATION,GOBP_REGULATION_OF_COLLAGEN_METABOLIC_PROCESS,GOBP_REGULATION_OF_COLLAGEN_METABOLIC_PROCESS,GOMF_SYMPORTER_ACTIVITY,GOMF_SYMPORTER_ACTIVITY,GOBP_COMMISSURAL_NEURON_AXON_GUIDANCE,GOBP_COMMISSURAL_NEURON_AXON_GUIDANCE,MIR4693_3P,MIR4693_3P,GOMF_SOLUTE_CATION_SYMPORTER_ACTIVITY,GOMF_SOLUTE_CATION_SYMPORTER_ACTIVITY,HAN_JNK_SINGALING_DN,HAN_JNK_SINGALING_DN,VANHARANTA_UTERINE_FIBROID_UP,VANHARANTA_UTERINE_FIBROID_UP,TURIJANSKI_MARKH4_TARGETS,TURIJANSKI_MARKH4_TARGETS,GOBP_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION,GOBP_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION,GOBP_SUCKLING_BEHAVIOR,GOBP_SUCKLING_BEHAVIOR,PID_INTEGRIN_A9B1_PATHWAY,PID_INTEGRIN_A9B1_PATHWAY,MIR4268,MIR4268,GOBP_NEUROINFLAMMATORY_RESPONSE,GOBP_NEUROINFLAMMATORY_RESPONSE,CHARAFE_BREAST_CANCER_BASAL_VS_MESENCHYMAL_DN,CHARAFE_BREAST_CANCER_BASAL_VS_MESENCHYMAL_DN,GOBP_ENDOTHELIAL_CELL_PROLIFERATION,GOBP_ENDOTHELIAL_CELL_PROLIFERATION,REACTOME_REGULATION_OF_GENE_EXPRESSION_IN_BETA_CELLS,REACTOME_REGULATION_OF_GENE_EXPRESSION_IN_BETA_CELLS,GOBP_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT,GOBP_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT,REACTOME_ADORA2B_MEDIATED_ANTI_INFLAMMATORY_CYTOKINES_PRODUCTION,REACTOME_ADORA2B_MEDIATED_ANTI_INFLAMMATORY_CYTOKINES_PRODUCTION,MIR4535,MIR4535,HP_ANTINEUTROPHIL_ANTIbody_POSITIVITY,HP_ANTINEUTROPHIL_ANTIbody_POSITIVITY,GOBP_REGULATION_OF_CARTILAGE_DEVELOPMENT,GOBP_REGULATION_OF_CARTILAGE_DEVELOPMENT,HP_TRISMUS,HP_TRISMUS,MIR194_3P,MIR194_3P,MIR3649,MIR3649,GCM_HMGA2,GCM_HMGA2,GOBP_KIDNEY_MORPHOGENESIS,GOBP_KIDNEY_MORPHOGENESIS,HEIDENBLAD_AMPLIFIED_IN_SOFT_TISSUE_CANCER,HEIDENBLAD_AMPLIFIED_IN_SOFT_TISSUE_CANCER,GOBP_AROMATIC_AMINO_ACID_TRANSPORT,GOBP_AROMATIC_AMINO_ACID_TRANSPORT,WP_DEVELOPMENT_OF_URETERIC_COLLECTION_SYSTEM,WP_DEVELOPMENT_OF_URETERIC_COLLECTION_SYSTEM,MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_27,MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_27,HP_SEVERE_EXPRESSIVE_LANGUAGE_DELAY,HP_SEVERE_EXPRESSIVE_LANGUAGE_DELAY,GOBP_LIPOPROTEIN_CATABOLIC_PROCESS,GOBP_LIPOPROTEIN_CATABOLIC_PROCESS,GOCC_POSISYNAPTIC_CYTOSOL,GOCC_POSISYNAPTIC_CYTOSOL,BIOCARTA_DREAM_PATHWAY,BIOCARTA_DREAM_PATHWAY,GOBP_REGULATION_OF_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE,GOBP_REGULATION_OF_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE,GOCC_COSTAMERE,GOCC_COSTAMERE,GOBP_POSITIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY,GOBP_POSITIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY,GOBP_CARDIAC_VENTRICLE_FORMATIO,GOBP_CARDIAC_VENTRICLE_FORMATIO,GOBP_NEUROTRANSMITTER_REUPTAKE,GOBP_NEUROTRANSMITTER_REUPTAKE,MIR6086,MIR6086,BIOCARTA_AGR_PATHWAY,BIOCARTA_AGR_PATHWAY,GOBP_REPLACEMENT_OSSIFICATION,GOBP_REPLACEMENT_OSSIFICATION,GCM_PTPRU,GCM_PTPRU,GOBP_REGULATION_OF_NEUROTRANSMITTER_UPTAKE,GOBP_REGULATION_OF_NEUROTRANSMITTER_UPTAKE,BIOCARTA_MCALPAIN_PATHWAY,BIOCARTA_MCALPAIN_PATHWAY,GOBP_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY,GOBP_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY,HP_ABNORMAL_COCCYX_MORPHOLOGY,HP_ABNORMAL_COCCYX_MORPHOLOGY,GOCC_DYSTROGLYCAN_COMPLEX,GOCC_DYSTROGLYCAN_COMPLEX,GOBP_MESENCHYMAL_TO_EPITHELIAL_TRANSITION,GOBP_MESENCHYMAL_TO_EPITHELIAL_TRANSITION,BIOCARTA_CARDIACEGF_PATHWAY,BIOCARTA_CARDIACEGF_PATHWAY,HP_UNERUPTED_TOOTH,HP_UNERUPTED_TOOTH,GOBP_DEVELOPMENTAL_INDUCATION,GOBP_DEVELOPMENTAL_INDUCATION,VALK_AML_CLUSTER_1,VALK_AML_CLUSTER_1,GOMF_CYCLIC_NUCLEOTIDE_BINDING,GOMF_CYCLIC_NUCLEOTIDE_BINDING,GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_PROLIFERATION,GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_PROLIFERATION,GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_9_SIGNALING_PATHWAY,GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_9_SIGNALING_PATHWAY,MIR377_5P,MIR377_5P,NAKAMURA_CANCER_MICROENVIRONMENT_UP,NAKAMURA_CANCER_MICROENVIRONMENT_UP,MIR3936,MIR3936