

GSE37416_CTRL_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN,GSE37416_CTRL_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN,GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN
GSE16385_MONOCYTE_VS_12H_IL4_TREATED_MACROPHAGE_DN,GSE16385_MONOCYTE_VS_12H_IL4_TREATED_MACROPHAGE_DN
GSE14769_UNSTIM_VS_80MIN_LPS_BMDM_UP,GSE14769_UNSTIM_VS_80MIN_LPS_BMDM_UP
GSE27786_CD8_TCELL_VS_ERYTHROBLAST_UP,GSE27786_CD8_TCELL_VS_ERYTHROBLAST_UP
GSE23925_LIGHT_ZONE_VS_NAIVE_BCELL_DN,GSE23925_LIGHT_ZONE_VS_NAIVE_BCELL_DN
GSE17721_POLYIC_VS_PAM3CSK4_1H_BMDC_UP,GSE17721_POLYIC_VS_PAM3CSK4_1H_BMDC_UP
GSE17721_CPG_VS_GARDIQUIMOD_2H_BMDC_DN,GSE17721_CPG_VS_GARDIQUIMOD_2H_BMDC_DN
GSE23925_DARK_ZONE_VS_NAIVE_BCELL_UP,GSE23925_DARK_ZONE_VS_NAIVE_BCELL_UP
GSE17721_LPS_VS_PAM3CSK4_24H_BMDC_UP,GSE17721_LPS_VS_PAM3CSK4_24H_BMDC_UP
GSE411_UNSTIM_VS_400MIN_IL6_STIM_MACROPHAGE_DN,GSE411_UNSTIM_VS_400MIN_IL6_STIM_MACROPHAGE_DN
GSE17721_CTRL_VS_LPS_1H_BMDC_UP,GSE17721_CTRL_VS_LPS_1H_BMDC_UP
GCM_DFFA,GCM_DFFA
GSE1534_ELF4_KO_VS_WT_ACTIVATED_CD8_TCELL_UP,GSE1534_ELF4_KO_VS_WT_ACTIVATED_CD8_TCELL_UP
GSE32423_IL7_VS_IL7_IL4_NAIVE_CD8_TCELL_DN,GSE32423_IL7_VS_IL7_IL4_NAIVE_CD8_TCELL_DN
GSE36327_CD62L_HIGH_VS_CD62L_LOW_TREG_CD69_NEG_KLRG1_NEG_UP,GSE36327_CD62L_HIGH_VS_CD62L_LOW_TREG_CD69_NEG_KLRG1_NEG_UP
GOMF_RRNA_BINDING,GOMF_RRNA_BINDING
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_UP,GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_UP
MIR513C_5P,MIR513C_5P
GSE40274_E0S_VS_FOXP3_AND_E0S_TRANSDUCED_ACTIVATED_CD4_TCELL_UP,GSE40274_E0S_VS_FOXP3_AND_E0S_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
GSE28737_WT_VS_BCL6_KO_MARGINAL_ZONE_BCELL_DN,GSE28737_WT_VS_BCL6_KO_MARGINAL_ZONE_BCELL_DN
MIR514B_5P,MIR514B_5P
GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_UP,GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_UP
GSE17721_0.5H_VS_12H_PAM3CSK4_BMDC_UP,GSE17721_0.5H_VS_12H_PAM3CSK4_BMDC_UP
MIR603,MIR603
GSE41867_NAIVE_VS_DAY15_LCMV_CONE13_EFFECTOR_CD8_TCELL_DN,GSE41867_NAIVE_VS_DAY15_LCMV_CONE13_EFFECTOR_CD8_TCELL_DN
GSE26669_CD4_VS_CD8_TCELL_IN_MLR_DN,GSE26669_CD4_VS_CD8_TCELL_IN_MLR_DN
GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_DN,GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_DN
GSE7548_NAIVE_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_UP,GSE7548_NAIVE_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_UP
GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP,GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP
GSE15659_RESTING_VS_ACTIVATED_TREG_DN,GSE15659_RESTING_VS_ACTIVATED_TREG_DN
GCM_HBP1,GCM_HBP1
MIR6715A_3P,MIR6715A_3P
GSE2898_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP,GSE2898_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP
GSE42021_TCONV_PLN_VS_CD4INT_TCONV_THYMUS_UP,GSE42021_TCONV_PLN_VS_CD4INT_TCONV_THYMUS_UP
GSE9946_MATURE_STIMULATORY_VS_LISTERIA_INF_MATURE_DC_UP,GSE9946_MATURE_STIMULATORY_VS_LISTERIA_INF_MATURE_DC_UP
GCM_CRKL,GCM_CRKL
MIR190B_3P,MIR190B_3P
MIR150_3P,MIR150_3P
FIGUEROA_AML_METHYLATION_CLUSTER_3_UP,FIGUEROA_AML_METHYLATION_CLUSTER_3_UP
MIR6831_5P,MIR6831_5P
MIR155_3P,MIR155_3P
GOBP_MAINTENANCE_OF_CELL_NUMBER,GOBP_MAINTENANCE_OF_CELL_NUMBER
GSE30153_LUPUS_VS_HEALTHY_DONOR_BCELL_UP,GSE30153_LUPUS_VS_HEALTHY_DONOR_BCELL_UP
GSE9601_UNTREATED_VS_P3K_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_DN,GSE9601_UNTREATED_VS_P3K_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_DN
GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_UP,GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_UP
GSE43955_TGFB_IL6_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_UP,GSE43955_TGFB_IL6_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_UP
MIR3668,MIR3668
MIR3927_3P,MIR3927_3P
FLECHNER_PBL_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN,FLECHNER_PBL_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN
MIR9718,MIR9718
ETF_Q6,ETF_Q6
MIR4673,MIR4673
FIGUEROA_AML_METHYLATION_CLUSTER_6_UP,FIGUEROA_AML_METHYLATION_CLUSTER_6_UP
GOCC_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE,GOCC_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE
MIR3679_5P,MIR3679_5P
MIR5000_5P,MIR5000_5P
GSE16385_UNTREATED_VS_12H_IL4_TREATED_MACROPHAGE_DN,GSE16385_UNTREATED_VS_12H_IL4_TREATED_MACROPHAGE_DN
HP_ABNORMALITY_OF_THE_COAGULATION_CASCADE,HP_ABNORMALITY_OF_THE_COAGULATION_CASCADE
IKEDA_MIR133_TARGETS_UP,IKEDA_MIR133_TARGETS_UP
MIR4647,MIR4662B,MIR4647,MIR4662B
GENTILE_RESPONSE_CLUSTER_D3,GENTILE_RESPONSE_CLUSTER_D3
MIR4642,MIR4642
GSE41978_WT_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN,GSE41978_WT_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN
MIR628_5P,MIR628_5P
GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4MID_SORTED_BCELL_DN,GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4MID_SORTED_BCELL_DN
GOBP_REGULATION_OF_PROTEIN_ACETYLATION,GOBP_REGULATION_OF_PROTEIN_ACETYLATION
GSE1112_OTL_CD8A8_VS_HY_CD8AA_THYMOCTE_RTOC_CULTURE_DN,GSE1112_OTL_CD8A8_VS_HY_CD8AA_THYMOCTE_RTOC_CULTURE_DN
GOBP_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY,GOBP_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY
MIR34A_3P,MIR34A_3P
FIGUEROA_AML_METHYLATION_CLUSTER_7_UP,FIGUEROA_AML_METHYLATION_CLUSTER_7_UP
MIR218_1_3P,MIR218_1_3P
GOCC_PCG_PROTEIN_COMPLEX,GOCC_PCG_PROTEIN_COMPLEX
HP_APLASIA_HYPOPLASIA_OF_TOE,HP_APLASIA_HYPOPLASIA_OF_TOE
chr16q22,chr16q22
GTGTGAG_MIR342,GTGTGAG_MIR342
BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_UP,BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_UP
REACTOME_MICRORNA_MIRNA_BIOGENESIS,REACTOME_MICRORNA_MIRNA_BIOGENESIS
MIR1258,MIR1258
MIR1263,MIR1263
MIR7852_3P,MIR7852_3P
GOBP_REGULATION_OF_MYOBLAST_DIFFERENTIATION,GOBP_REGULATION_OF_MYOBLAST_DIFFERENTIATION
MIR619_3P,MIR619_3P
GOBP_MYOBLAST_DIFFERENTIATION,GOBP_MYOBLAST_DIFFERENTIATION
GOCC_ER_TO_GOLGI_TRANSPORT_VESICLE_MEMBRANE,GOCC_ER_TO_GOLGI_TRANSPORT_VESICLE_MEMBRANE
MIR676_3P,MIR676_3P
MIR3161,MIR3161
GCM_BCL2L1,GCM_BCL2L1
GSE9946_IMMATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP,GSE9946_IMMATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP
REACTOME_ANTIEN_PRESENTATION_FOLDING_ASSEMBLY_AND_PEPTIDE_LOADING_OF_CLASS_I_MHC,REACTOME_ANTIEN_PRESENTATION_FOLDING_ASSEMBLY_AND_PEPTIDE_LOADING_OF_CLASS_I_MHC
GOBP_SOMATIC_STEM_CELL_POPULATION_MAINTENANCE,GOBP_SOMATIC_STEM_CELL_POPULATION_MAINTENANCE
DELASERNA_MYOD_TARGETS_DN,DELASERNA_MYOD_TARGETS_DN
GCM_CSNK1D,GCM_CSNK1D
GOBP_CELLULAR_PROTEIN_CONTAINING_COMPLEX_LOCALIZATION,GOBP_CELLULAR_PROTEIN_CONTAINING_COMPLEX_LOCALIZATION
CCCACAT_MIR2993P,CCCACAT_MIR2993P
MIR3614_5P,MIR3614_5P
MIR5699_3P,MIR5699_3P
FIGUEROA_AML_METHYLATION_CLUSTER_4_UP,FIGUEROA_AML_METHYLATION_CLUSTER_4_UP
GENTILE_UV_RESPONSE_CLUSTER_D8,GENTILE_UV_RESPONSE_CLUSTER_D8
WUNDER_INFLAMMATORY_RESPONSE_AND_CHOLESTEROL_UP,WUNDER_INFLAMMATORY_RESPONSE_AND_CHOLESTEROL_UP
HP_ABNORMALITY_OF_THE_HEPATIC_VASCULATURE,HP_ABNORMALITY_OF_THE_HEPATIC_VASCULATURE
GOBP_RENAL_SYSTEM_VASCULATURE_DEVELOPMENT,GOBP_RENAL_SYSTEM_VASCULATURE_DEVELOPMENT
MIR652_5P,MIR652_5P
GOMF_PALMITOYLTRANSFERASE_ACTIVITY,GOMF_PALMITOYLTRANSFERASE_ACTIVITY
DOANE_BREAST_CANCER_CLASSES_DN,DOANE_BREAST_CANCER_CLASSES_DN
MIR12127,MIR12127
HP_APLASIA_HYPOPLASIA_OF_THE_THYMUS,HP_APLASIA_HYPOPLASIA_OF_THE_THYMUS
HP_DECREASED_LYMPHOCYTE_PROLIFERATION_IN_RESPONSE_TO_MITOGEN,HP_DECREASED_LYMPHOCYTE_PROLIFERATION_IN_RESPONSE_TO_MITOGEN
HP_ABNORMAL_CELL_PROLIFERATION,HP_ABNORMAL_CELL_PROLIFERATION
PLASARL_TGFB1_SIGNALING_VIA_NFIC_1HR_UP,PLASARL_TGFB1_SIGNALING_VIA_NFIC_1HR_UP
ELLWOOD_MYC_TARGETS_DN,ELLWOOD_MYC_TARGETS_DN
MEIS1BHOXA9_01,MEIS1BHOXA9_01
HP_ESOPHAGEAL_VARIX,HP_ESOPHAGEAL_VARIX
GOBP_HISTONE_MONOUBIQUITINATION,GOBP_HISTONE_MONOUBIQUITINATION
HP_ABNORMAL_LACRIMAL_DUCT_MORPHOLOGY,HP_ABNORMAL_LACRIMAL_DUCT_MORPHOLOGY
GOCC_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPARTMENT_MEMBRANE,GOCC_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COMPARTMENT_MEMBRANE
HP_NASOLACRIMAL_DUCT_OBSTRUCTION,HP_NASOLACRIMAL_DUCT_OBSTRUCTION
WP_VITAMIN_D_IN_INFLAMMATORY_DISEASES,WP_VITAMIN_D_IN_INFLAMMATORY_DISEASES
MIR431_5P,MIR431_5P
VALK_AML_CLUSTER_11,VALK_AML_CLUSTER_11
HAY_BONE_MARROW_CD34_POS_HSC,HAY_BONE_MARROW_CD34_POS_HSC
GGARNTKYCCA_UNKNOWN,GGARNTKYCCA_UNKNOWN
MIR4421,MIR4421
GOCC_MITOTIC_SPINDLE_MICROTUBULE,GOCC_MITOTIC_SPINDLE_MICROTUBULE
MIR631,MIR631
GOBP_SKIN_EPIDERMIS_DEVELOPMENT,GOBP_SKIN_EPIDERMIS_DEVELOPMENT
KIM_LRRC3B_TARGETS,KIM_LRRC3B_TARGETS
BRACH1AT_RESPONSE_TO_CISPLATIN,BRACH1AT_RESPONSE_TO_CISPLATIN
HP_TRIGONOCEPHALY,HP_TRIGONOCEPHALY
GOBP_ENDOCARDIAL_CUSHION_MORPHOGENESIS,GOBP_ENDOCARDIAL_CUSHION_MORPHOGENESIS
GOBP_PRE_MIRNA_PROCESSING,GOBP_PRE_MIRNA_PROCESSING
MIR1178_3P,MIR1178_3P
JIANG_HYPOXIA_VIA_VHL,JIANG_HYPOXIA_VIA_VHL
DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_PERICYTES,DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_PERICYTES
HP_ABNORMAL_CIRCULATING_JGA_LEVEL,HP_ABNORMAL_CIRCULATING_JGA_LEVEL
GOBP_RNA_INTERFERENCE,GOBP_RNA_INTERFERENCE
REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_CYTOCHROME_C_RELEASE,REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_CYTOCHROME_C_RELEASE
HP_FOLLICULAR_HYPERKERATOSIS,HP_FOLLICULAR_HYPERKERATOSIS
REACTOME_EGR2_AND_SOX10_MEDIATED_INITIATION_OF_SCHWANN_CELL_MYELINATION,REACTOME_EGR2_AND_SOX10_MEDIATED_INITIATION_OF_SCHWANN_CELL_MYELINATION
GOBP_POSITIVE_REGULATION_OF_CELLULAR_EXTRAVASATION,GOBP_POSITIVE_REGULATION_OF_CELLULAR_EXTRAVASATION
GOBP_NEGATIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR,GOBP_NEGATIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
MIR3661,MIR3661
GSE9946_MATURE_STIMULATORY_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP,GSE9946_MATURE_STIMULATORY_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP
MARCIKIAK_ER_STRESS_RESPONSE_VIA_CHOP,MARCIKIAK_ER_STRESS_RESPONSE_VIA_CHOP
GOBP_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR,GOBP_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR
RUSSELLINGER_GASTRIC_REG3A_POSITIVE_CELLS,RUSSELLINGER_GASTRIC_REG3A_POSITIVE_CELLS
KKGG_SPHINGOLIPID_METABOLISM,KKGG_SPHINGOLIPID_METABOLISM
GOBP_RESPONSE_TO_PLATELET_DERIVED_GROWTH_FACTOR,GOBP_RESPONSE_TO_PLATELET_DERIVED_GROWTH_FACTOR
MIR3189_5P,MIR3189_5P
MIR4683,MIR4683
GOBP_ENDOCARDIAL_CUSHION_FORMATION,GOBP_ENDOCARDIAL_CUSHION_FORMATION
HP_ABNORMAL_B_CELL_MORPHOLOGY,HP_ABNORMAL_B_CELL_MORPHOLOGY
GOBP_POSITIVE_REGULATION_OF_MYOBLAST_DIFFERENTIATION,GOBP_POSITIVE_REGULATION_OF_MYOBLAST_DIFFERENTIATION
BERENJENO_ROCK_SIGNALING_NOT_VIA_RHOA_UP,BERENJENO_ROCK_SIGNALING_NOT_VIA_RHOA_UP
GOBP_MEMBRANE_LIPID_CATABOLIC_PROCESS,GOBP_MEMBRANE_LIPID_CATABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RESPONSE_TO_CHEMICAL_STIMULUS,GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RESPONSE_TO_CHEMICAL_STIMULUS
REACTOME_ALPHA_PROTEIN_KINASE_1_SIGNALING_PATHWAY,REACTOME_ALPHA_PROTEIN_KINASE_1_SIGNALING_PATHWAY
MIR200C_5P,MIR200C_5P
GOBP_PROTECTION_FROM_NON_HOMOLOGOUS_END_JOINING_AT_TELOMERE,GOBP_PROTECTION_FROM_NON_HOMOLOGOUS_END_JOINING_AT_TELOMERE
HP_ORTHOSTATIC_HYPOTENSION,HP_ORTHOSTATIC_HYPOTENSION
GOMF_LACTATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY,GOMF_LACTATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOBP_TRANSFORMING_GROWTH_FACTOR_BETA_PRODUCTION,GOBP_TRANSFORMING_GROWTH_FACTOR_BETA_PRODUCTION
KANG_CISPLATIN_RESISTANCE_DN,KANG_CISPLATIN_RESISTANCE_DN
GOBP_MOLTING_CYCLE,GOBP_MOLTING_CYCLE
MIR3529_5P,MIR3529_5P
GOBP_INNER_EAR_AUDITORY_RECEPTOR_CELL_DIFFERENTIATION,GOBP_INNER_EAR_AUDITORY_RECEPTOR_CELL_DIFFERENTIATION
MIR103B,MIR103B
MIR943,MIR943
MIR10400_3P,MIR4674,MIR10400_3P,MIR4674
CHEN_HOXA5_TARGETS_6HR_UP,CHEN_HOXA5_TARGETS_6HR_UP
GOBP_POSITIVE_REGULATION_OF_CHEMOKINE_C_X_C_MOTIF_LIGAND_2_PRODUCTION,GOBP_POSITIVE_REGULATION_OF_CHEMOKINE_C_X_C_MOTIF_LIGAND_2_PRODUCTION
REACTOME_NEF_MEDIATED_DOWNREGULATION_OF_MHC_CLASS_1_COMPLEX_CELL_SURFACE_EXPRESSION,REACTOME_NEF_MEDIATED_DOWNREGULATION_OF_MHC_CLASS_1_COMPLEX_CELL_SURFACE_EXPRESSION
WP_NEURAL_CREST_DIFFERENTIATION,WP_NEURAL_CREST_DIFFERENTIATION
GOMF_HISTONE_ACETYLTTRANSFERASE_BINDING,GOMF_HISTONE_ACETYLTTRANSFERASE_BINDING
GOBP_PRODUCTION_OF_SIRNA_INVOLVED_IN_RNA_INTERFERENCE,GOBP_PRODUCTION_OF_SIRNA_INVOLVED_IN_RNA_INTERFERENCE
SNIDJERS_AMPIFIED_IN_HEAD_AND_NECK_TUMORS,SNIDJERS_AMPIFIED_IN_HEAD_AND_NECK_TUMORS
PAL_PRMT5_TARGETS_DN,PAL_PRMT5_TARGETS_DN
GOBP_LACTATE_TRANSMEMBRANE_TRANSPORT,GOBP_LACTATE_TRANSMEMBRANE_TRANSPORT
SKP2_TARGET_GENES,SKP2_TARGET_GENES
GOBP_VENTRICULAR_SEPTUM_MORPHOGENESIS,GOBP_VENTRICULAR_SEPTUM_MORPHOGENESIS
WEBER_METHYLATED_LCP_IN_SPERM_DN,WEBER_METHYLATED_LCP_IN_SPERM_DN
GOBP_VESICLE_FUSION_WITH_GOLGI_APPARATUS,GOBP_VESICLE_FUSION_WITH_GOLGI_APPARATUS
GOBP_SECONDARY_PALATE_DEVELOPMENT,GOBP_SECONDARY_PALATE_DEVELOPMENT
PID_SIP_META_PATHWAY,PID_SIP_META_PATHWAY
MCCABE_HOXC6_TARGETS_CANCER_DN,MCCABE_HOXC6_TARGETS_CANCER_DN
GOBP_MODULATION_OF_AGE_RELATED_BEHAVIORAL_DECLINE,GOBP_MODULATION_OF_AGE_RELATED_BEHAVIORAL_DECLINE
GOBP_POSITIVE_REGULATION_OF_VASCULOGENESIS,GOBP_POSITIVE_REGULATION_OF_VASCULOGENESIS
GOBP_CELL_CELL_ADHESION_INVOLVED_IN_GASTRULATION,GOBP_CELL_CELL_ADHESION_INVOLVED_IN_GASTRULATION
GOBP_CANONICAL_WNT_SIGNALING_PATHWAY_INVOLVED_IN_HEART_DEVELOPMENT,GOBP_CANONICAL_WNT_SIGNALING_PATHWAY_INVOLVED_IN_HEART_DEVELOPMENT
HP_LARYNGEAL_DYSTONIA,HP_LARYNGEAL_DYSTONIA
GOBP_EXTRACELLULAR_VESICLE_BIOGENESIS,GOBP_EXTRACELLULAR_VESICLE_BIOGENESIS
GOBP_CARDIAC_LEFT_VENTRICLE_MORPHOGENESIS,GOBP_CARDIAC_LEFT_VENTRICLE_MORPHOGENESIS
INAMURA_LUNG_CANCER_SCC_SUBTYPES_UP,INAMURA_LUNG_CANCER_SCC_SUBTYPES_UP
HP_ERYTHRODERMA,HP_ERYTHRODERMA