

GSE9988_LPS_VS_CTRL_TREATED_MONOCYTE_DN,GSE9988_LPS_VS_CTRL_TREATED_MONOCYTE_DN
GSE2706_UNSTIM_VS_8H_LPS_AND_R848_DC_UP,GSE2706_UNSTIM_VS_8H_LPS_AND_R848_DC_UP
GSE9988_ANTL_TREM1_AND_LPS_VS_CTRL_TREATED_MONOCYTES_DN,GSE9988_ANTL_TREM1_AND_LPS_VS_CTRL_TREATED_MONOCYTES_DN
GSE22886_NAIVE_CD4_TCELL_VS_MEMORY_TCELL_DN,GSE22886_NAIVE_CD4_TCELL_VS_MEMORY_TCELL_DN
GCM_MYST2,GCM_MYST2
GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP,GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP
GSE17721_CTRL_VS_LPS_4H_BMDC_UP,GSE17721_CTRL_VS_LPS_4H_BMDC_UP
GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_2H_UP,GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_2H_UP
GSE360_CTRL_VS_M_TUBERCULOSIS_DC_UP,GSE360_CTRL_VS_M_TUBERCULOSIS_DC_UP
GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TCONV_DN,GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TCONV_DN
GSE9960_GRAM_NEG_VS_GRAM_POS_SEPSIS_PBMCDN,GSE9960_GRAM_NEG_VS_GRAM_POS_SEPSIS_PBMCDN
GSE13484_UNSTIM_VS_3H_YF17D_VACCINE_STIM_PBMCDN,GSE13484_UNSTIM_VS_3H_YF17D_VACCINE_STIM_PBMCDN
GCM_UBE2N,GCM_UBE2N
GSE22140_HEALTHY_VS_ARTHRITIC_MOUSE_CD4_TCELL_DN,GSE22140_HEALTHY_VS_ARTHRITIC_MOUSE_CD4_TCELL_DN
GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CVS_IN_MAC_UP,GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CVS_IN_MAC_UP
GSE37416_CTRL_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_UP,GSE37416_CTRL_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_UP
GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN,GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN
GSE360_LOW_DOSE_B_MALAYL_VS_M_TUBERCULOSIS_DC_UP,GSE360_LOW_DOSE_B_MALAYL_VS_M_TUBERCULOSIS_DC_UP
GSE9960_HEALTHY_VS_GRAM_NEG_SEPSIS_PBMCDN,GSE9960_HEALTHY_VS_GRAM_NEG_SEPSIS_PBMCDN
GSE41867_NAIVE_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN,GSE41867_NAIVE_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN
GSE37533_PPARG1_FOXP3_VS_FOXP3_TRANSDUCED_CD4_TCELL_PIOGLITAZONE_TREATED_DN,GSE37533_PPARG1_FOXP3_VS_FOXP3_TRANSDUCED_CD4_TCELL_PIOGLITAZONE_TREATED_DN
THAKAR_PBMCDN_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_7DY_UP,THAKAR_PBMCDN_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_7DY_UP
GSE13411_IGM_MEMORY_BCELL_VS_PLASMA_CELL_UP,GSE13411_IGM_MEMORY_BCELL_VS_PLASMA_CELL_UP
GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_DN,GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_DN
GSE7568_IL4_VS_IL4_AND_TGFB_TREATED_MACROPHAGE_24H_UP,GSE7568_IL4_VS_IL4_AND_TGFB_TREATED_MACROPHAGE_24H_UP
GSE13522_CTRL_VS_T_CRUZL_Y_STRAIN_INF_SKIN_IFNG_KO_UP,GSE13522_CTRL_VS_T_CRUZL_Y_STRAIN_INF_SKIN_IFNG_KO_UP
GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_UP,GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_UP
GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_48H_DN,GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_48H_DN
GSE15930_STIM_VS_STIM_AND_IFNAB_72H_CD8_T_CELL_DN,GSE15930_STIM_VS_STIM_AND_IFNAB_72H_CD8_T_CELL_DN
GSE15930_STIM_VS_STIM_AND_IL12_72H_CD8_T_CELL_UP,GSE15930_STIM_VS_STIM_AND_IL12_72H_CD8_T_CELL_UP
GSE37605_C57BL6_VS_NOD_FOXP3_FUSION_GFP_TCONV_UP,GSE37605_C57BL6_VS_NOD_FOXP3_FUSION_GFP_TCONV_UP
GSE41978_ID2_KO_AND_BIM_KO_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN,GSE41978_ID2_KO_AND_BIM_KO_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN
GSE27859_CD11C_INT_F480_HI_MACROPHAGE_VS_CD11C_ING_F480_INT_DC_DN,GSE27859_CD11C_INT_F480_HI_MACROPHAGE_VS_CD11C_ING_F480_INT_DC_DN
GSE42021_TREG_VS_TCONV_PLN_DN,GSE42021_TREG_VS_TCONV_PLN_DN
GSE37301_MULTIPOTENT_PROGENITOR_VS_RAG2_KO_NK_CELL_UP,GSE37301_MULTIPOTENT_PROGENITOR_VS_RAG2_KO_NK_CELL_UP
GSE45739_UNSTIM_VS_ACD3_ACD28_STIM_NRA5_KO_CD4_TCELL_UP,GSE45739_UNSTIM_VS_ACD3_ACD28_STIM_NRA5_KO_CD4_TCELL_UP
LINDGREN_BLADDER_CANCER_WITH_LOH_IN_CHR9Q,LINDGREN_BLADDER_CANCER_WITH_LOH_IN_CHR9Q
GCM_NCAMI,GCM_NCAMI
MORF_PP5SC,MORF_PP5SC
GSE1448_CTRL_VS_ANTL_VALPHA2_DP_THYMOCYTE_DN,GSE1448_CTRL_VS_ANTL_VALPHA2_DP_THYMOCYTE_DN
GSE41867_DAY15_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN,GSE41867_DAY15_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN
LINDGREN_BLADDER_CANCER_CLUSTER_1_UP,LINDGREN_BLADDER_CANCER_CLUSTER_1_UP
GSE3203_HEALTHY_VS_INFLUENZA_INFECTED_I_N_BCELL_UP,GSE3203_HEALTHY_VS_INFLUENZA_INFECTED_I_N_BCELL_UP
GSE19401_PLN_VS_PeyERS_PATCH_FOLLICULAR_DC_UP,GSE19401_PLN_VS_PeyERS_PATCH_FOLLICULAR_DC_UP
GCM_BMPR2,GCM_BMPR2
GNF2_STAT6,GNF2_STAT6
GOMF_NUCLEAR_RECEPTOR_BINDING,GOMF_NUCLEAR_RECEPTOR_BINDING
GSE42021_TREG_PLN_VS_CD24LO_TREG_THYMUS_UP,GSE42021_TREG_PLN_VS_CD24LO_TREG_THYMUS_UP
GAGCTGG_MIR337,GAGCTGG_MIR337
MIR5087,MIR5087
GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_UP,GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_UP
MIR329_5P,MIR329_5P
YAGL_AML_WITH_T_9_11_TRANSLOCATION,YAGL_AML_WITH_T_9_11_TRANSLOCATION
MIR935,MIR935
GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_WITH_EGR2_DELETED_DN,GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_WITH_EGR2_DELETED_DN
GOMF_HORMONE_RECEPTOR_BINDING,GOMF_HORMONE_RECEPTOR_BINDING
MIR708_3P,MIR708_3P
GSE37532_VISCERAL_ADIPOSE_TISSUE_VS_I_N_DERIVED_PPARG_KO_TREG_CD4_TCELL_DN,GSE37532_VISCERAL_ADIPOSE_TISSUE_VS_I_N_DERIVED_PPARG_KO_TREG_CD4_TCELL_DN
GSE42021_CD24INT_VS_CD24LOW_TREG_THYMUS_UP,GSE42021_CD24INT_VS_CD24LOW_TREG_THYMUS_UP
SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5,SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5
TOOKER_GEMCITABINE_RESISTANCE_DN,TOOKER_GEMCITABINE_RESISTANCE_DN
GOBP_CELLULAR_RESPONSE_TO_STEROID_HORMONE_STIMULUS,GOBP_CELLULAR_RESPONSE_TO_STEROID_HORMONE_STIMULUS
MIR6757_3P,MIR6757_3P
MIR3692_5P,MIR3692_5P
GCM_BAG5,GCM_BAG5
MIR3129_3P,MIR3129_3P
NGUYEN_NOTCH1_TARGETS_DN,NGUYEN_NOTCH1_TARGETS_DN
GSE27786_BCELL_VS_MONO_MAC_DN,GSE27786_BCELL_VS_MONO_MAC_DN
GSE22611_UNSTIM_VS_2H_MDP_STIM_NOD2_TRANSDUCED_HEK293T_CELL_DN,GSE22611_UNSTIM_VS_2H_MDP_STIM_NOD2_TRANSDUCED_HEK293T_CELL_DN
MIR5583_5P,MIR5583_5P
GOBP_STEROID_HORMONE_MEDIATED_SIGNALING_PATHWAY,GOBP_STEROID_HORMONE_MEDIATED_SIGNALING_PATHWAY
MIR1224_3P,MIR1224_3P
CTCCAAG_MIR432,CTCCAAG_MIR432
GOBP_HISTONE_H3_ACETYLTATION,GOBP_HISTONE_H3_ACETYLTATION
DARWICHE_PAPILLOMA_RISK_LOW_DN,DARWICHE_PAPILLOMA_RISK_LOW_DN
MIR7843_5P,MIR7843_5P
HP_BONE_MARROW_HYPOCELLULARITY,HP_BONE_MARROW_HYPOCELLULARITY
GCM_RAP2A,GCM_RAP2A
MIR769_5P,MIR769_5P
GNF2_YAV1,GNF2_YAV1
MIR4709_5P,MIR4709_5P
BOYAULT_LIVER_CANCER_SUBCLASS_G12_UP,BOYAULT_LIVER_CANCER_SUBCLASS_G12_UP
ZSCAN5C_TARGET_GENES,ZSCAN5C_TARGET_GENES
SIRNA_EIF4GL_DN,SIRNA_EIF4GL_DN
REACTOME_NRIH3_NRIH2_REGULATE_GENE_EXPRESSION_LINKED_TO_CHOLESTEROL_TRANSPORT_AND_EFFLUX,REACTOME_NRIH3_NRIH2_REGULATE_GENE_EXPRESSION_LINKED_TO_CHOLESTEROL_TRANSPORT_AND_EFFLUX
GCM_DEAF1,GCM_DEAF1
HP_CARIOUS_TEETH,HP_CARIOUS_TEETH
SIG_CHEMOTAXIS,SIG_CHEMOTAXIS
GOBP_REGULATION_OF_HISTONE_METHYLATION,GOBP_REGULATION_OF_HISTONE_METHYLATION
GOBP_PRODUCTION_OF_SMALL_RNA_INVOLVED_IN_GENE_SILENCING_BY_RNA,GOBP_PRODUCTION_OF_SMALL_RNA_INVOLVED_IN_GENE_SILENCING_BY_RNA
GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_24H_CD8_T_CELL_UP,GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_24H_CD8_T_CELL_UP
MIR541_3P,MIR541_3P
MIR654_5P,MIR654_5P
FONTAINE_FOLLICULAR_THYROID_ADENOMA_UP,FONTAINE_FOLLICULAR_THYROID_ADENOMA_UP
HP_ABNORMAL_LARYNX_MORPHOLOGY,HP_ABNORMAL_LARYNX_MORPHOLOGY
MEISSNER_NPC_JCP_WITH_H3K4ME3,MEISSNER_NPC_JCP_WITH_H3K4ME3
CASTELLANO_NRAS_TARGETS_UP,CASTELLANO_NRAS_TARGETS_UP
MODULE_284,MODULE_284
MIR6889_3P,MIR6889_3P
HP_CRANIAL_HYPEROSTOSIS,HP_CRANIAL_HYPEROSTOSIS
MIR6866_5P,MIR6866_5P
ASTON_MAJOR_DEPRESSIVE_DISORDER_UP,ASTON_MAJOR_DEPRESSIVE_DISORDER_UP
GOBP_PROTEIN_LOCALIZATION_TO_CELL_CELL_JUNCTION,GOBP_PROTEIN_LOCALIZATION_TO_CELL_CELL_JUNCTION
BIOCARTA_VDR_PATHWAY,BIOCARTA_VDR_PATHWAY
WP_ENDODERM_DIFFERENTIATION,WP_ENDODERM_DIFFERENTIATION
GOMF_METALLOAMINOPEPTIDASE_ACTIVITY,GOMF_METALLOAMINOPEPTIDASE_ACTIVITY
HP_CRANIOFACIAL_HYPEROSTOSIS,HP_CRANIOFACIAL_HYPEROSTOSIS
RUAN_RESPONSE_TO_TROGLITAZONE_DN,RUAN_RESPONSE_TO_TROGLITAZONE_DN
AR_Q2,AR_Q2
GOBP_PROTEIN_PALMITOYLATION,GOBP_PROTEIN_PALMITOYLATION
REACTOME_NRIH2_AND_NRIH3_MEDIATED_SIGNALING,REACTOME_NRIH2_AND_NRIH3_MEDIATED_SIGNALING
HP_VARICOSE_VEINS,HP_VARICOSE_VEINS
GOBP_PPTIDYL_L_CYSSTEINE_S_PALMITOYLATION,GOBP_PPTIDYL_L_CYSSTEINE_S_PALMITOYLATION
MODULE_362,MODULE_362
GOMF_PHOSPHATIDIC_ACID_BINDING,GOMF_PHOSPHATIDIC_ACID_BINDING
MODULE_166,MODULE_166
GOBP_REGULATION_OF_STEM_CELL_POPULATION_MAINTENANCE,GOBP_REGULATION_OF_STEM_CELL_POPULATION_MAINTENANCE
REACTOME_TRANSCRIPTIONAL_REGULATION_BY_MECP2,REACTOME_TRANSCRIPTIONAL_REGULATION_BY_MECP2
MIR642A_5P,MIR642A_5P
GOBP_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS,GOBP_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS
GNF2_MAP2K3,GNF2_MAP2K3
MIR6881_5P,MIR6881_5P
WP_EBV_LMP1_SIGNALING,WP_EBV_LMP1_SIGNALING
MIR2277_3P,MIR2277_3P
GOBP_NEGATIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT,GOBP_NEGATIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT
CNF2_BNIP3L,GNF2_BNIP3L
MIR645,MIR645
HP_ECTROPION,HP_ECTROPION
GOBP_REGULATION_OF_MRNA_POLYADENYLATION,GOBP_REGULATION_OF_MRNA_POLYADENYLATION
GOBP_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION,GOBP_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION
MCCLUNG_DELTA_FOSB_TARGETS_8WK,MCCLUNG_DELTA_FOSB_TARGETS_8WK
WP_WNT_SIGNALING_PATHWAY_AND_PLURIPOTENCY,WP_WNT_SIGNALING_PATHWAY_AND_PLURIPOTENCY
GOBP_POSITIVE_REGULATION_OF_DENDRITE_MORPHOGENESIS,GOBP_POSITIVE_REGULATION_OF_DENDRITE_MORPHOGENESIS
GOBP_REGULATION_OF_HISTONE_H3_K4_METHYLATION,GOBP_REGULATION_OF_HISTONE_H3_K4_METHYLATION
chr5q22,chr5q22
GOBP_POSITIVE_REGULATION_OF_NON_MOTILE_CILIUM_ASSEMBLY,GOBP_POSITIVE_REGULATION_OF_NON_MOTILE_CILIUM_ASSEMBLY
MIR3174,MIR3174
WIKMAN_ASBESTOS_LUNG_CANCER_UP,WIKMAN_ASBESTOS_LUNG_CANCER_UP
GOBP_MEMBRANE_FISSION,GOBP_MEMBRANE_FISSION
MIR937_5P,MIR937_5P
BOGNI_TREATMENT_RELATED_MYELOID_LEUKEMIA_DN,BOGNI_TREATMENT_RELATED_MYELOID_LEUKEMIA_DN
GOBP_GLOMERULUS_MORPHOGENESIS,GOBP_GLOMERULUS_MORPHOGENESIS
GOBP_INTERLEUKIN_2_PRODUCTION,GOBP_INTERLEUKIN_2_PRODUCTION
HP_MACULAR_THICKENING,HP_MACULAR_THICKENING
HP_HYPOPLASTIC_ILIA,HP_HYPOPLASTIC_ILIA
HP_ULNAR_DEVIATION_OF_THE_HAND,HP_ULNAR_DEVIATION_OF_THE_HAND
HP_MIXED_HEARING_IMPAIRMENT,HP_MIXED_HEARING_IMPAIRMENT
HERNANDEZ_MITOTIC_ARREST_BY_DOCETAXEL_2_UP,HERNANDEZ_MITOTIC_ARREST_BY_DOCETAXEL_2_UP
GOBP_HISTONE_H3_K9_METHYLATION,GOBP_HISTONE_H3_K9_METHYLATION
MIR1193,MIR1193
GOBP_POSITIVE_REGULATION_OF_PROTEIN_CONTAINING_COMPLEX_DISASSEMBLY,GOBP_POSITIVE_REGULATION_OF_PROTEIN_CONTAINING_COMPLEX_DISASSEMBLY
HP_THICKENED_CALVARIA,HP_THICKENED_CALVARIA
PARK_HSC_MARKERS,PARK_HSC_MARKERS
MORF_MYC,MORF_MYC
FIRESTEIN_CTNNB1_PATHWAY,FIRESTEIN_CTNNB1_PATHWAY
GOCC_RNAI_EFFECTOR_COMPLEX,GOCC_RNAI_EFFECTOR_COMPLEX
GOBP_FRUCTOSE_2_6_BISPHOSPHATE_METABOLIC_PROCESS,GOBP_FRUCTOSE_2_6_BISPHOSPHATE_METABOLIC_PROCESS
GOMF_FRUCTOSE_2_6_BISPHOSPHATE_2_PHOSPHATASE_ACTIVITY,GOMF_FRUCTOSE_2_6_BISPHOSPHATE_2_PHOSPHATASE_ACTIVITY
GOCC_DNA_REPLICATION_FACTOR_A_COMPLEX,GOCC_DNA_REPLICATION_FACTOR_A_COMPLEX
GOMF_HISTONE_DEMETHYLASE_ACTIVITY_H3_K9_SPECIFIC,GOMF_HISTONE_DEMETHYLASE_ACTIVITY_H3_K9_SPECIFIC
GSE12707_AT16L1_HYPOMORPH_VS_WT_THYMUS_DN,GSE12707_AT16L1_HYPOMORPH_VS_WT_THYMUS_DN
KYNQ_RESPONSE_TO_H2O2_VIA_ERCC6,KYNQ_RESPONSE_TO_H2O2_VIA_ERCC6
GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION,GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION
PAX8_01,PAX8_01
GOBP_REGULATION_OF_INCLUSION_BODY_ASSEMBLY,GOBP_REGULATION_OF_INCLUSION_BODY_ASSEMBLY
GOBP_NEGATIVE_REGULATION_OF_CHROMATIN_ORGANIZATION,GOBP_NEGATIVE_REGULATION_OF_CHROMATIN_ORGANIZATION
GOMF_UREA_TRANSMEMBRANE_TRANSPORTER_ACTIVITY,GOMF_UREA_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
TOMLINS_METASTASIS_DN,TOMLINS_METASTASIS_DN
WP_LNCRNA_INVOLVEMENT_IN_CANONICAL_WNT_SIGNALING_AND_COLORECTAL_CANCER,WP_LNCRNA_INVOLVEMENT_IN_CANONICAL_WNT_SIGNALING_AND_COLORECTAL_CANCER
chr10q24,chr10q24
GOBP_REGULATION_OF_ER_TO_GOLGI_VESICLE_MEDIATED_TRANSPORT,GOBP_REGULATION_OF_ER_TO_GOLGI_VESICLE_MEDIATED_TRANSPORT
OVSYANNIKOVA_PBMCDN_FLUARIX_AGE_50_74YO_COMMON_WITH_BOTH_HAI_AND_VNA_28DY_VS_0DY_USED_IN_HAI_AND_VNA_RESPONSE_MODELS_UP,OVSYANNIKOVA_PBMCDN_FLUARIX_AGE_50_74YO_COMMON_WITH_BOTH_HAI_AND_VNA_28DY_VS_0DY_USED_IN_HAI_AND_VNA_RESPONSE_MODELS_UP
GOBP_UREA_TRANSPORT,GOBP_UREA_TRANSPORT
REACTOME_SYNTHESIS_OF_PI,REACTOME_SYNTHESIS_OF_PI
GOMF_ROUNDABOUT_BINDING,GOMF_ROUNDABOUT_BINDING
HP_ABNORMALITY_OF_AMINO_ACID_METABOLISM,HP_ABNORMALITY_OF_AMINO_ACID_METABOLISM
HP_ABNORMAL_PLATELET_VOLUME,HP_ABNORMAL_PLATELET_VOLUME
GOBP_POSITIVE_REGULATION_OF_T_CELL_CYTOKINE_PRODUCTION,GOBP_POSITIVE_REGULATION_OF_T_CELL_CYTOKINE_PRODUCTION
MIR5586_3P,MIR5586_3P
GOMF_MAP_KINASE_KINASE_KINASE_ACTIVITY,GOMF_MAP_KINASE_KINASE_KINASE_ACTIVITY
GOBP_PULMONARY_ARTERY_MORPHOGENESIS,GOBP_PULMONARY_ARTERY_MORPHOGENESIS
REACTOME_TICAM1_TRAF6_DEPENDENT_INDUCTION_OF_TAK1_COMPLEX,REACTOME_TICAM1_TRAF6_DEPENDENT_INDUCTION_OF_TAK1_COMPLEX
GOBP_HISTONE_H3_K9_DEMETHYLATION,GOBP_HISTONE_H3_K9_DEMETHYLATION
GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY,GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY