

CT_IL4_AND_ANTI_IL12_4H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_4H_CD4_TCELL_DN

GSE16451.IMMATURE_VS_MATURE_NEURON_CELL_LINE.WEST_EQUINE_ENC_VIRUS_DN,GSE16451.IMMATURE_VS_MATURE_NEURON_CELL_LINE.WEST_EQUINE_ENC_VIRUS_DN
GSE26727.WT_VS_KLF2_KO.LPS.STIM.MACROPHAGE_DN,GSE26727.WT_VS_KLF2_KO.LPS.STIM.MACROPHAGE_DN
GSE37532.TREG_VS_TCONV_PPARG_KO.CD4_TCELL_FROM_LN_DN,GSE37532.TREG_VS_TCONV_PPARG_KO.CD4_TCELL_FROM_LN_DN
GSE45739.NRAS_KO_VS_WT_ACD3.ACD28_STIM_CD4_TCELL_UP,GSE45739.NRAS_KO_VS_WT_ACD3.ACD28_STIM_CD4_TCELL_UP
GSE23321.EFFECTOR_MEMORY_VS_NAIVE_CD8_TCELL_DN,GSE23321.EFFECTOR_MEMORY_VS_NAIVE_CD8_TCELL_DN
GSE6674.ANTI_IGM_VS_CPG_STIM_BCELL_DN,GSE6674.ANTI_IGM_VS_CPG_STIM_BCELL_DN
GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_6H_CD4_TCELL_DN,GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_6H_CD4_TCELL_DN
GSE17974_0H_VS_2H_IN_VITRO_ACT_CD4_TCELL_DN,GSE17974_0H_VS_2H_IN_VITRO_ACT_CD4_TCELL_DN
GSE26559.TCF1_KO_VS_WT_LIN_NEG_CELL_UP,GSE26559.TCF1_KO_VS_WT_LIN_NEG_CELL_UP
GSE11864.CSF1_VS_CSF1_IFNG_IN_MAC_DN,GSE11864.CSF1_VS_CSF1_IFNG_IN_MAC_DN
GSE11864.UNTREATED_VS_CSF1_PAM3CYS_IN_MAC_DN,GSE11864.UNTREATED_VS_CSF1_PAM3CYS_IN_MAC_DN
MORF_EIF4E,MORF_EIF4E
GOCC.PRERIBOSOME,GOCC.PRERIBOSOME
GSE15330.HSC_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_UP,GSE15330.HSC_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_UP
GSE17721.LPS_VS_POLYIC_1H_BMDC_UP,GSE17721.LPS_VS_POLYIC_1H_BMDC_UP
GSE40274_CTRL_VS_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN,GSE40274_CTRL_VS_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE21670.UNTREATED_VS_TGFB_TREATED_STAT3_KO_CD4_TCELL_DN,GSE21670.UNTREATED_VS_TGFB_TREATED_STAT3_KO_CD4_TCELL_DN
GSE6259_DEC205_POS_DC_VS_CD4_TCELL_UP,GSE6259_DEC205_POS_DC_VS_CD4_TCELL_UP
GSE11864.CSF1_VS_CSF1_PAM3CYS_IN_MAC_DN,GSE11864.CSF1_VS_CSF1_PAM3CYS_IN_MAC_DN
SCHAEFFER_PROSTATE_DEVELOPMENT_6HR_UP,SCHAEFFER_PROSTATE_DEVELOPMENT_6HR_UP
GSE17721_CTRL_VS_PAM3CSK4_8H_BMDC_DN,GSE17721_CTRL_VS_PAM3CSK4_8H_BMDC_DN
HALLMARK_ALLOGRAFT_REJECTION,HALLMARK_ALLOGRAFT_REJECTION
GSE40274_CTRL_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP,GSE40274_CTRL_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
GSE42724_NAIVE_VS_B1_BCELL_UP,GSE42724_NAIVE_VS_B1_BCELL_UP
BOSCO_ALLERGEN_INDUCED_TH2_ASSOCIATED_MODULE,BOSCO_ALLERGEN_INDUCED_TH2_ASSOCIATED_MODULE
IL15_UP.V1_UP,IL15_UP.V1_UP
GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_48H_ACT_CD4_TCELL_UP,GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_48H_ACT_CD4_TCELL_UP
GSE28130_ACTIVATED_VS_INDUCED_TREG_UP,GSE28130_ACTIVATED_VS_INDUCED_TREG_UP
GSE8515_CTRL_VS_IL1_4H_STIM_MAC_DN,GSE8515_CTRL_VS_IL1_4H_STIM_MAC_DN
ESC_V6.5_UP.LATE.V1_DN,ESC_V6.5_UP.LATE.V1_DN
MIR4709_3P,MIR4709_3P
GSE15330.HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_IKAROS_KO_DN,GSE15330.HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_IKAROS_KO_DN
GSE43863_TFH_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN,GSE43863_TFH_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN
GSE16385.UNTREATED_VS_12H_ROSIGLITAZONE_TREATED_MACROPHAGE_DN,GSE16385.UNTREATED_VS_12H_ROSIGLITAZONE_TREATED_MACROPHAGE_DN
GSE2770_IL12_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_DN,GSE2770_IL12_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_DN
CATRRAGC_UNKNOWN,CATRRAGC_UNKNOWN
HP_IMPAIRED_VIBRATORY_SENSATION,HP_IMPAIRED_VIBRATORY_SENSATION
GSE43863_DAY6_EFF_VS_DAY150_MEM_TFH_CD4_TCELL_DN,GSE43863_DAY6_EFF_VS_DAY150_MEM_TFH_CD4_TCELL_DN
GOBP_CERAMIDE_METABOLIC_PROCESS,GOBP_CERAMIDE_METABOLIC_PROCESS
KIM_MYCN_AMPLIFICATION_TARGETS_UP,KIM_MYCN_AMPLIFICATION_TARGETS_UP
MCAATNNNNNGCG_UNKNOWN,MCAATNNNNNGCG_UNKNOWN
GOBP_CELLULAR_RESPONSE_TO_GLUCOSE_STARVATION,GOBP_CELLULAR_RESPONSE_TO_GLUCOSE_STARVATION
HP_FOOT_DORSIFLEXOR_WEAKNESS,HP_FOOT_DORSIFLEXOR_WEAKNESS
REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION_INITIATION_FROM_TYPE_3_PROMOTER,REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION_INITIATION_FROM_TYPE_3_PROMOTER
GOBP_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT,GOBP_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT
MARSON_FOXP3_TARGETS_DN,MARSON_FOXP3_TARGETS_DN
REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION_INITIATION_FROM_TYPE_1_PROMOTER,REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION_INITIATION_FROM_TYPE_1_PROMOTER
GOBP_CARBOHYDRATE_TRANSMEMBRANE_TRANSPORT,GOBP_CARBOHYDRATE_TRANSMEMBRANE_TRANSPORT
GOBP_LYMPHOCYTE_HOMEOSTASIS,GOBP_LYMPHOCYTE_HOMEOSTASIS
HP_ABNORMALITY_OF_THE_ILIUM,HP_ABNORMALITY_OF_THE_ILIUM
HUANG_FOXA2_TARGETS_UP,HUANG_FOXA2_TARGETS_UP
MIR4694_5P,MIR4694_5P
MIR3914,MIR3914
KEGG_RNA_POLYMERASE,KEGG_RNA_POLYMERASE
GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_24H_ACT_CD4_TCELL_UP,GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_24H_ACT_CD4_TCELL_UP
JAZAG_TGFB1_SIGNALING_VIA_SMAD4_DN,JAZAG_TGFB1_SIGNALING_VIA_SMAD4_DN
GOBP_CERAMIDE_BIOSYNTHETIC_PROCESS,GOBP_CERAMIDE_BIOSYNTHETIC_PROCESS
MIR4741,MIR4741
GOBP_POLYSACCHARIDE_BIOSYNTHETIC_PROCESS,GOBP_POLYSACCHARIDE_BIOSYNTHETIC_PROCESS
GOBP_PROTEIN_LOCALIZATION_TO_NUCLCOLUS,GOBP_PROTEIN_LOCALIZATION_TO_NUCLCOLUS
MIR3117_3P,MIR3117_3P
GOBP_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY,GOBP_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
GOBP_CELLULAR_CARBOHYDRATE_BIOSYNTHETIC_PROCESS,GOBP_CELLULAR_CARBOHYDRATE_BIOSYNTHETIC_PROCESS
HP_NAIL_DYSPLASIA,HP_NAIL_DYSPLASIA
GOBP_PERIPHERAL_NERVOUS_SYSTEM_DEVELOPMENT,GOBP_PERIPHERAL_NERVOUS_SYSTEM_DEVELOPMENT
KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE,KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE
GSE43863_NAIVE_VS_TH1_EFF_CD4_TCELL_D6_1CMV_DN,GSE43863_NAIVE_VS_TH1_EFF_CD4_TCELL_D6_1CMV_DN
GOCC_RNA_POLYMERASE_III_COMPLEX,GOCC_RNA_POLYMERASE_III_COMPLEX
REACTOME_RNA_POLYMERASE_III_CHAIN_ELONGATION,REACTOME_RNA_POLYMERASE_III_CHAIN_ELONGATION
HP_ABNORMALITY_OF_THE_ALVEOLAR_RIDGES,HP_ABNORMALITY_OF_THE_ALVEOLAR_RIDGES
HP_VACUOLATED_LYMPHOCYTES,HP_VACUOLATED_LYMPHOCYTES
HP_ATROPHIC_SCARS,HP_ATROPHIC_SCARS
WP_VITAMIN_D_IN_INFLAMMATORY_DISEASES,WP_VITAMIN_D_IN_INFLAMMATORY_DISEASES
GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN,GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN
GOBP_REGULATION_OF_GLUCOSE_TRANSMEMBRANE_TRANSPORT,GOBP_REGULATION_OF_GLUCOSE_TRANSMEMBRANE_TRANSPORT
GOBP_HEAT_GENERATION,GOBP_HEAT_GENERATION
GOBP_DENDRITIC_SPINE_MAINTENANCE,GOBP_DENDRITIC_SPINE_MAINTENANCE
GOBP_CHONDROITIN_SULFATE_BIOSYNTHETIC_PROCESS,GOBP_CHONDROITIN_SULFATE_BIOSYNTHETIC_PROCESS
GOBP_AMINO_ACID_IMPORT,GOBP_AMINO_ACID_IMPORT
GOBP_REGULATION_OF_POSTSYNAPTIC_MEMBRANE_NEUROTRANSMITTER_RECEPTOR_LEVELS,GOBP_REGULATION_OF_POSTSYNAPTIC_MEMBRANE_NEUROTRANSMITTER_RECEPTOR_LEVELS
GOBP_LYMPH_NODE_DEVELOPMENT,GOBP_LYMPH_NODE_DEVELOPMENT
GOBP_NEUROINFLAMMATORY_RESPONSE,GOBP_NEUROINFLAMMATORY_RESPONSE
GOBP_PURINE_NUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS,GOBP_PURINE_NUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_GLUCOSE_TRANSMEMBRANE_TRANSPORT,GOBP_NEGATIVE_REGULATION_OF_GLUCOSE_TRANSMEMBRANE_TRANSPORT
GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS,GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS
BARRIER_CANCER_RELAPSE_NORMAL_SAMPLE_UP,BARRIER_CANCER_RELAPSE_NORMAL_SAMPLE_UP
FRANCO_BLOOD_SANOFI_PASTEUR_SA_INACTIVATED_INFLUENZA_VACCINE_CORRELATED_WITH_ANTIBODY_RESPONSE_AGE_18_40YO_1DY_NEGATIVE,FRANCO_BLOOD_SANOFI_PASTEUR_SA_INACTIVATED_INFLUENZA_VACCINE_CORRELATED_WITH_ANTIBODY_RESPONSE_AGE_18_40YO_1DY_NEGATIVE
GOBP_NITRIC_OXIDE_MEDIATED_SIGNAL_TRANSDUCTION,GOBP_NITRIC_OXIDE_MEDIATED_SIGNAL_TRANSDUCTION
MIR554,MIR554
GOBP_FEVER_GENERATION,GOBP_FEVER_GENERATION
GOBP_REGULATION_OF_GLUCOSE_IMPORT,GOBP_REGULATION_OF_GLUCOSE_IMPORT
PATEL_SKIN_OF_BODY_ZOSTAVAX_AGE_70_93YO_VZV_CHALLENGE_3DY_DN,PATEL_SKIN_OF_BODY_ZOSTAVAX_AGE_70_93YO_VZV_CHALLENGE_3DY_DN
GOBP_INFLAMMATORY_RESPONSE_TO_ANTIAGENIC_STIMULUS,GOBP_INFLAMMATORY_RESPONSE_TO_ANTIAGENIC_STIMULUS
GOBP_L_ALPHA_AMINO_ACID_TRANSMEMBRANE_TRANSPORT,GOBP_L_ALPHA_AMINO_ACID_TRANSMEMBRANE_TRANSPORT
WP_MONOAMINE_TRANSPORT,WP_MONOAMINE_TRANSPORT
HP_HYPOPLASTIC_IL1A,HP_HYPOPLASTIC_IL1A
GOMF_GUANYLATE_CYCLASE_ACTIVITY,GOMF_GUANYLATE_CYCLASE_ACTIVITY
GOBP_REPLACEMENT_OSSIFICATION,GOBP_REPLACEMENT_OSSIFICATION
HP_HEMOPHAGOCYTOSIS,HP_HEMOPHAGOCYTOSIS
REACTOME_TNF_RECEPTOR_SUPERFAMILY_TNFSF_MEMBERS_MEDIATING_NON_CANONICAL_NF_KB_PATHWAY,REACTOME_TNF_RECEPTOR_SUPERFAMILY_TNFSF_MEMBERS_MEDIATING_NON_CANONICAL_NF_KB_PATHWAY
REACTOME_EGR2_AND_SOX10_MEDIATED_INITIATION_OF_SCHWANN_CELL_MYELINATION,REACTOME_EGR2_AND_SOX10_MEDIATED_INITIATION_OF_SCHWANN_CELL_MYELINATION
WP_NO_METABOLISM_IN_CYSTIC_FIBROSIS,WP_NO_METABOLISM_IN_CYSTIC_FIBROSIS
GOCC_DERLIN_1_RETROTRANSLOCATION_COMPLEX,GOCC_DERLIN_1_RETROTRANSLOCATION_COMPLEX
GOBP_REGULATION_OF_HEAT_GENERATION,GOBP_REGULATION_OF_HEAT_GENERATION
GOBP_ENDOCHONDRAL_BONE_MORPHOGENESIS,GOBP_ENDOCHONDRAL_BONE_MORPHOGENESIS
GOBP_RECEPTOR_LOCALIZATION_TO_SYNAPSE,GOBP_RECEPTOR_LOCALIZATION_TO_SYNAPSE
GOBP_GLYCOPHINGOLIPID_BIOSYNTHETIC_PROCESS,GOBP_GLYCOPHINGOLIPID_BIOSYNTHETIC_PROCESS
GOBP_GANGLIOSIDE_BIOSYNTHETIC_PROCESS,GOBP_GANGLIOSIDE_BIOSYNTHETIC_PROCESS
BILANGES_SERUM_SENSITIVE_VIA_TSC1,BILANGES_SERUM_SENSITIVE_VIA_TSC1
HP_ROTARY_NYSTAGMUS,HP_ROTARY_NYSTAGMUS
HP_SHORT_MIDDLE_PHALANX_OF_FINGER,HP_SHORT_MIDDLE_PHALANX_OF_FINGER
GOBP_GANGLIOSIDE_METABOLIC_PROCESS,GOBP_GANGLIOSIDE_METABOLIC_PROCESS
GOMF_GALACTOSYLTRANSFERASE_ACTIVITY,GOMF_GALACTOSYLTRANSFERASE_ACTIVITY
GOBP_REGULATION_OF_DENDRITIC_SPINE_MAINTENANCE,GOBP_REGULATION_OF_DENDRITIC_SPINE_MAINTENANCE
MIR887_3P,MIR887_3P
GOMF_UDP_GLUCOSYLTRANSFERASE_ACTIVITY,GOMF_UDP_GLUCOSYLTRANSFERASE_ACTIVITY
GOBP_NEGATIVE_REGULATION_OF_REGULATED_SECRETORY_PATHWAY,GOBP_NEGATIVE_REGULATION_OF_REGULATED_SECRETORY_PATHWAY
GOBP_NEUROTRANSMITTER_REUPTAKE,GOBP_NEUROTRANSMITTER_REUPTAKE
GOBP_NEGATIVE_REGULATION_OF_GLUCOSE_IMPORT,GOBP_NEGATIVE_REGULATION_OF_GLUCOSE_IMPORT
GREENBAUM_E2A_TARGETS_DN,GREENBAUM_E2A_TARGETS_DN
GOBP_REGULATION_OF_RESPIRATORY_GASEOUS_EXCHANGE,GOBP_REGULATION_OF_RESPIRATORY_GASEOUS_EXCHANGE
GOBP_INOSITOL_METABOLIC_PROCESS,GOBP_INOSITOL_METABOLIC_PROCESS
STEIN_ESTROGEN_RESPONSE_NOT_VIA_ESRRA,STEIN_ESTROGEN_RESPONSE_NOT_VIA_ESRRA
GOBP_NEUROTRANSMITTER_UPTAKE,GOBP_NEUROTRANSMITTER_UPTAKE
MODULE_65,MODULE_65
GOBP_PROTEIN_O_LINKED_GLYCOSYLATION_VIA_SERINE,GOBP_PROTEIN_O_LINKED_GLYCOSYLATION_VIA_SERINE
GOMF_GALACTOSIDE_BINDING,GOMF_GALACTOSIDE_BINDING
KEGG_O_GLYCAN_BIOSYNTHESIS,KEGG_O_GLYCAN_BIOSYNTHESIS
GOBP_REGULATION_OF_CALCIUM_ION_IMPORT,GOBP_REGULATION_OF_CALCIUM_ION_IMPORT
GOBP_PROSTATE_GLANDULAR_ACINUS_DEVELOPMENT,GOBP_PROSTATE_GLANDULAR_ACINUS_DEVELOPMENT
ZWANG_CLASS_2_TRANSIENTLY_INDUCED_BY_EGF,ZWANG_CLASS_2_TRANSIENTLY_INDUCED_BY_EGF
GOBP_VASCULAR_WOUND_HEALING,GOBP_VASCULAR_WOUND_HEALING
HP_INSIDIOUS_ONSET,HP_INSIDIOUS_ONSET
GOBP_KERATAN_SULFATE_BIOSYNTHETIC_PROCESS,GOBP_KERATAN_SULFATE_BIOSYNTHETIC_PROCESS
REACTOME_KERATAN_SULFATE_BIOSYNTHESIS,REACTOME_KERATAN_SULFATE_BIOSYNTHESIS
GHANDHI_BYSTANDER_IRRADIATION_DN,GHANDHI_BYSTANDER_IRRADIATION_DN
GOBP_REGULATION_OF_PINOCYTOSIS,GOBP_REGULATION_OF_PINOCYTOSIS
GOBP_NEGATIVE_REGULATION_OF_SERINE_TYPE_PEPTIDASE_ACTIVITY,GOBP_NEGATIVE_REGULATION_OF_SERINE_TYPE_PEPTIDASE_ACTIVITY
HP_LOBULATED_TONGUE,HP_LOBULATED_TONGUE
GOBP_REGULATION_OF_SERINE_TYPE_PEPTIDASE_ACTIVITY,GOBP_REGULATION_OF_SERINE_TYPE_PEPTIDASE_ACTIVITY
GOBP_ANGIOGENESIS_INVOLVED_IN_WOUND_HEALING,GOBP_ANGIOGENESIS_INVOLVED_IN_WOUND_HEALING
chr12p11,chr12p11
HP_OVERLAPPING_FINGERS,HP_OVERLAPPING_FINGERS
HP_DECREASED_LYMPHOCYTE_PROLIFERATION_IN_RESPONSE_TO_ANTI_CD3,HP_DECREASED_LYMPHOCYTE_PROLIFERATION_IN_RESPONSE_TO_ANTI_CD3
GAUSSMANN_MLL_A4_FUSION_TARGETS_B_DN,GAUSSMANN_MLL_A4_FUSION_TARGETS_B_DN
GOBP_REGULATION_OF_STORE_OPERATED_CALCIUM_ENTRY,GOBP_REGULATION_OF_STORE_OPERATED_CALCIUM_ENTRY
HOEK_MYELOID_DENDRITIC_CELL_2011_2012_TIV_ADULT_7DY_DN,HOEK_MYELOID_DENDRITIC_CELL_2011_2012_TIV_ADULT_7DY_DN
GOBP_POSITIVE_REGULATION_OF_T_CELL_DIFFERENTIATION_IN_THYMUS,GOBP_POSITIVE_REGULATION_OF_T_CELL_DIFFERENTIATION_IN_THYMUS
REACTOME_MULTIFUNCTIONAL_ANION_EXCHANGERS,REACTOME_MULTIFUNCTIONAL_ANION_EXCHANGERS
GOBP_POSITIVE_REGULATION_OF_AUTOPHAGOSOME_MATURATION,GOBP_POSITIVE_REGULATION_OF_AUTOPHAGOSOME_MATURATION
GOBP_NEGATIVE_REGULATION_OF_T_CELL_MIGRATION,GOBP_NEGATIVE_REGULATION_OF_T_CELL_MIGRATION
PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_6,PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_6
GOBP_TRIPARTITE_REGIONAL_SUBDIVISION,GOBP_TRIPARTITE_REGIONAL_SUBDIVISION
YAMASHITA_METHYLATED_IN_PROSTATE_CANCER,YAMASHITA_METHYLATED_IN_PROSTATE_CANCER
HP_PANCREATIC_CALCIFICATION,HP_PANCREATIC_CALCIFICATION
GOMF_SOLUTE_BICARBONATE_SYMPORTER_ACTIVITY,GOMF_SOLUTE_BICARBONATE_SYMPORTER_ACTIVITY
GOBP_LUNG_GROWTH,GOBP_LUNG_GROWTH
GOMF_GLUCOSE_SODIUM_SYMPORTER_ACTIVITY,GOMF_GLUCOSE_SODIUM_SYMPORTER_ACTIVITY
GOBP_NEGATIVE_REGULATION_OF_MACROPHAGE_ACTIVATION,GOBP_NEGATIVE_REGULATION_OF_MACROPHAGE_ACTIVATION
HP_CENTRAL_HYPOTHYROIDISM,HP_CENTRAL_HYPOTHYROIDISM
GOBP_POSITIVE_REGULATION_OF_ACTIVATED_T_CELL_PROLIFERATION,GOBP_POSITIVE_REGULATION_OF_ACTIVATED_T_CELL_PROLIFERATION
GOBP_PROTEIN_LOCALIZATION_TO_POSTSYNAPSE,GOBP_PROTEIN_LOCALIZATION_TO_POSTSYNAPSE
GOBP_NEGATIVE_REGULATION_OF_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION,GOBP_NEGATIVE_REGULATION_OF_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION