
GSE339_EX_VIVO_VS_IN_CULTURE_CD8POS_DC_UP, GSE339_EX_VIVO_VS_IN_CULTURE_CD8POS_DC_UP
GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_2H_DN
GSE42088_2H_VS_24H_LEISHMANIA_INF_DC_UP, GSE42088_2H_VS_24H_LEISHMANIA_INF_DC_UP
GSE18893_TCONV_VS_TREG_24H_TNF_STIM_DN, GSE18893_TCONV_VS_TREG_24H_TNF_STIM_DN
GSE17721_CTRL_VS_POLYIC_24H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_24H_BMDC_UP
GSE17721_CTRL_VS_PAM3CSK4_6H_BMDC_UP, GSE17721_CTRL_VS_PAM3CSK4_6H_BMDC_UP
GSE31082_DN_VS_CD4_SP_THYMOCYTE_DN, GSE31082_DN_VS_CD4_SP_THYMOCYTE_DN
GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_UP, GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_UP
GSE5679_RARA_AGONIST_AM580_VS_AM580_AND_ROSIGLITAZONE_TREATED_DC_UP, GSE5679_RARA_AGONIST_AM580_VS_AM580_AND_ROSIGLITAZONE_TREATED_DC_UP
GSE17721_LPS_VS_CPG_4H_BMDC_DN, GSE17721_LPS_VS_CPG_4H_BMDC_DN
GSE12198_CTRL_VS_HIGH_IL2_STIM_NK_CELL_DN, GSE12198_CTRL_VS_HIGH_IL2_STIM_NK_CELL_DN
GSE20715_0H_VS_24H_OZONE_LUNG_UP, GSE20715_0H_VS_24H_OZONE_LUNG_UP
GSE16522_MEMORY_VS_NAIVE_ANTI_CD3CD28_STIM_CD8_TCELL_UP, GSE16522_MEMORY_VS_NAIVE_ANTI_CD3CD28_STIM_CD8_TCELL_UP
GSE3982_NEUTROPHIL_VS_NKCELL_DN, GSE3982_NEUTROPHIL_VS_NKCELL_DN
GSE43863_DAY6_EFF_VS_DAY150_MEM_LY6C_INT_CXCR5POS_CD4_TCELL_UP, GSE43863_DAY6_EFF_VS_DAY150_MEM_LY6C_INT_CXCR5POS_CD4_TCELL_UP
GSE9037_CTRL_VS_LPS_4H_STIM_BMDM_UP, GSE9037_CTRL_VS_LPS_4H_STIM_BMDM_UP
GSE41867_LCMV_ARMSTRONG_VS_CLONE13_DAY8_EFFECTOR_CD8_TCELL_UP, GSE41867_LCMV_ARMSTRONG_VS_CLONE13_DAY8_EFFECTOR_CD8_TCELL_UP
PASQUALUCCI_LYMPHOMA_BY_GC_STAGE_DN, PASQUALUCCI_LYMPHOMA_BY_GC_STAGE_DN
GSE17721_ALL_VS_24H_PAM3CSK4_BMDC_DN, GSE17721_ALL_VS_24H_PAM3CSK4_BMDC_DN
GSE17721_POLYIC_VS_CPG_12H_BMDC_UP, GSE17721_POLYIC_VS_CPG_12H_BMDC_UP
GAUSSMANN_MLL_AF4_FUSION_TARGETS_A_UP, GAUSSMANN_MLL_AF4_FUSION_TARGETS_A_UP
GSE16385_MONOCYTE_VS_MACROPHAGE_UP, GSE16385_MONOCYTE_VS_MACROPHAGE_UP
GOBP_VESICLE_BUDDING_FROM_MEMBRANE, GOBP_VESICLE_BUDDING_FROM_MEMBRANE
MIR124_5P, MIR124_5P
GSE32423_CTRL_VS_IL7_IL4_MEMORY_CD8_TCELL_DN, GSE32423_CTRL_VS_IL7_IL4_MEMORY_CD8_TCELL_DN
GSE13306_TREG_VS_TCONV_LAMINA_PROPRIA_UP, GSE13306_TREG_VS_TCONV_LAMINA_PROPRIA_UP
ROSS_AML_WITH_MLL_FUSIONS, ROSS_AML_WITH_MLL_FUSIONS
GOBP_PROTEIN_LOCALIZATION_TO_VACUOLE, GOBP_PROTEIN_LOCALIZATION_TO_VACUOLE
HP_FLAT_FACE, HP_FLAT_FACE
GSE26928_EFF_MEM_VS_CENTR_MEM_CD4_TCELL_UP, GSE26928_EFF_MEM_VS_CENTR_MEM_CD4_TCELL_UP
TRAVAGLINI_LUNG_NATURAL_KILLER_CELL, TRAVAGLINI_LUNG_NATURAL_KILLER_CELL
MIR8089, MIR8089
GOBP_PEROXISOME_ORGANIZATION, GOBP_PEROXISOME_ORGANIZATION
REACTOME_SYNTHESIS_OF_ACTIVE_UBIQUITIN_ROLES_OF_E1_AND_E2_ENZYMES, REACTOME_SYNTHESIS_OF_ACTIVE_UBIQUITIN_ROLES_OF_E1_AND_E2_ENZYMES
GSE32255_WT_VS_JMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_UP, GSE32255_WT_VS_JMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_UP
GENTILE_UV_RESPONSE_CLUSTER_D5, GENTILE_UV_RESPONSE_CLUSTER_D5
HAHTOLA_MYCOSIS_FUNGOIDES_CD4_UP, HAHTOLA_MYCOSIS_FUNGOIDES_CD4_UP
GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_UP, GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_UP
MIR936, MIR936
ZHOU_TNF_SIGNALING_4HR, ZHOU_TNF_SIGNALING_4HR
REACTOME_SPHINGOLIPID_METABOLISM, REACTOME_SPHINGOLIPID_METABOLISM
SETLUR_PROSTATE_CANCER_TMPRSS2_ERG_FUSION_UP, SETLUR_PROSTATE_CANCER_TMPRSS2_ERG_FUSION_UP
HP_DEMYELINATING_PERIPHERAL_NEUROPATHY, HP_DEMYELINATING_PERIPHERAL_NEUROPATHY
HP_APLASIA_HYPOPLASIA_INVOLVING_THE_VERTEBRAL_COLUMN, HP_APLASIA_HYPOPLASIA_INVOLVING_THE_VERTEBRAL_COLUMN
MIR4700_3P, MIR4700_3P
GSE22886_NAIVE_BCELL_VS_MONOCYTE_UP, GSE22886_NAIVE_BCELL_VS_MONOCYTE_UP
ROSS_AML_WITH_AML1_ETO_FUSION, ROSS_AML_WITH_AML1_ETO_FUSION
KEGG_VEGF_SIGNALING_PATHWAY, KEGG_VEGF_SIGNALING_PATHWAY
JAZAERI_BREAST_CANCER_BRCA1_VS_BRCA2_UP, JAZAERI_BREAST_CANCER_BRCA1_VS_BRCA2_UP
REACTOME_SYNTHESIS_OF_SUBSTRATES_IN_N_GLYCAN_BIOSYTHESIS, REACTOME_SYNTHESIS_OF_SUBSTRATES_IN_N_GLYCAN_BIOSYTHESIS
HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_56D_TOP_100_DEG_AFTER_IN_VITRO_RE_STIMULATION_UP, HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_56D_TOP_100_DEG_AFTER_IN_VITRO_R
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CONTAINING_COMPLEX_ASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CONTAINING_COMPLEX_ASSEMBLY
GOBP_NUCLEOTIDE_SUGAR_METABOLIC_PROCESS, GOBP_NUCLEOTIDE_SUGAR_METABOLIC_PROCESS
ZHU_CMV_8_HR_DN, ZHU_CMV_8_HR_DN
MARSON_FOXP3_TARGETS_UP, MARSON_FOXP3_TARGETS_UP
GOBP_ENDOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_ENDOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY
KEGG_LEUKOCYTE_TRANSENDOTHELIAL_MIGRATION, KEGG_LEUKOCYTE_TRANSENDOTHELIAL_MIGRATION
REACTOME_SIGNALING_BY_BRAF_AND_RAF_FUSIONS, REACTOME_SIGNALING_BY_BRAF_AND_RAF_FUSIONS
GSE34156_UNTREATED_VS_6H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP, GSE34156_UNTREATED_VS_6H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP
GCM_MAPK10, GCM_MAPK10
MIR3668, MIR3668
GOBP_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_MULTIVESICULAR_BODY_SORTING_PATHWAY
MIR7155_5P, MIR7155_5P
KEGG_AMINO_SUGAR_AND_NUCLEOTIDE_SUGAR_METABOLISM, KEGG_AMINO_SUGAR_AND_NUCLEOTIDE_SUGAR_METABOLISM
KEGG_LEISHMANIA_INFECTION, KEGG_LEISHMANIA_INFECTION
MIR664A_5P, MIR664A_5P
HP_DISPROPORTIONATE_SHORT_STATURE, HP_DISPROPORTIONATE_SHORT_STATURE
LIM_MAMMARY_LUMINAL_MATURE_UP, LIM_MAMMARY_LUMINAL_MATURE_UP
HP_THICK_UPPER_LIP_VERMILION, HP_THICK_UPPER_LIP_VERMILION
GOBP_GLYCOSPHINGOLIPID_METABOLIC_PROCESS, GOBP_GLYCOSPHINGOLIPID_METABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_8_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_8_PRODUCTION
HP_DYSOSTOSIS_MULTIPLEX, HP_DYSOSTOSIS_MULTIPLEX
TOMLINS_PROSTATE_CANCER_UP, TOMLINS_PROSTATE_CANCER_UP
BIOCARTA_ARENRF2_PATHWAY, BIOCARTA_ARENRF2_PATHWAY
GSE22025_UNTREATED_VS_PROGESTERONE_TREATED_CD4_TCELL_UP, GSE22025_UNTREATED_VS_PROGESTERONE_TREATED_CD4_TCELL_UP
GOBP_REGULATION_OF_TRANSLATIONAL_INITIATION_IN_RESPONSE_TO_STRESS, GOBP_REGULATION_OF_TRANSLATIONAL_INITIATION_IN_RESPONSE_TO_STRESS
KEGG_GNRH_SIGNALING_PATHWAY, KEGG_GNRH_SIGNALING_PATHWAY
HP_ACETABULAR_DYSPLASIA, HP_ACETABULAR_DYSPLASIA
BIOCARTA_BCR_PATHWAY, BIOCARTA_BCR_PATHWAY
GOBP_STEROID_BIOSYNTHETIC_PROCESS, GOBP_STEROID_BIOSYNTHETIC_PROCESS
WP_NONGENOMIC_ACTIONS_OF_125_DIHYDROXYVITAMIN_D3, WP_NONGENOMIC_ACTIONS_OF_125_DIHYDROXYVITAMIN_D3
REACTOME_ENDOSOMAL_SORTING_COMPLEX_REQUIRED_FOR_TRANSPORT_ESCRT, REACTOME_ENDOSOMAL_SORTING_COMPLEX_REQUIRED_FOR_TRANSPORT_ESCRT
BIOCARTA_CALCINEURIN_PATHWAY, BIOCARTA_CALCINEURIN_PATHWAY
MIR380_5P, MIR380_5P
GRABARCZYK_BCL11B_TARGETS_DN, GRABARCZYK_BCL11B_TARGETS_DN
REACTOME_GLYCOSPHINGOLIPID_METABOLISM, REACTOME_GLYCOSPHINGOLIPID_METABOLISM
REACTOME_RHO_GTPASES_ACTIVATE_NADPH_OXIDASES, REACTOME_RHO_GTPASES_ACTIVATE_NADPH_OXIDASES
MIR637, MIR637
GOBP_NUCLEOTIDE_SUGAR_BIOSYNTHETIC_PROCESS, GOBP_NUCLEOTIDE_SUGAR_BIOSYNTHETIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_INTERFERON_GAMMA_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERFERON_GAMMA_PRODUCTION
HAHTOLA_SEZARY_SYNDROM_DN, HAHTOLA_SEZARY_SYNDROM_DN
GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT, GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT
GOBP_MITOCHONDRIAL_FISSION, GOBP_MITOCHONDRIAL_FISSION
BIOCARTA_GPCR_PATHWAY, BIOCARTA_GPCR_PATHWAY
WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_MAPK_SIGNALING, WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_MAPK_SIGNALING
BIOCARTA_SPPA_PATHWAY, BIOCARTA_SPPA_PATHWAY
BIOCARTA_HDAC_PATHWAY, BIOCARTA_HDAC_PATHWAY
BIOCARTA_MEF2D_PATHWAY, BIOCARTA_MEF2D_PATHWAY
PID_NFKAPPAB_ATYPICAL_PATHWAY, PID_NFKAPPAB_ATYPICAL_PATHWAY
GOCC_CLATHRIN_ADAPTOR_COMPLEX, GOCC_CLATHRIN_ADAPTOR_COMPLEX
BILBAN_B_CLL_LPL_UP, BILBAN_B_CLL_LPL_UP
GOBP_PIGMENT_GRANULE_ORGANIZATION, GOBP_PIGMENT_GRANULE_ORGANIZATION
REACTOME_SYNTHESIS_OF_BILE_ACIDS_AND_BILE_SALTS_VIA_7ALPHA_HYDROXYCHOLESTEROL, REACTOME_SYNTHESIS_OF_BILE_ACIDS_AND_BILE_SALTS_VIA_7ALPHA_HYDROXYCHOLESTEROL
HP_LARGE_FOREHEAD, HP_LARGE_FOREHEAD
KEGG_LONG_TERM_POTENTIATION, KEGG_LONG_TERM_POTENTIATION
GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY
GOBP_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION, GOBP_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION
GOMF_CADHERIN_BINDING_INVOLVED_IN_CELL_CELL_ADHESION, GOMF_CADHERIN_BINDING_INVOLVED_IN_CELL_CELL_ADHESION
HP_DELAYED_ABILITY_TO_SIT, HP_DELAYED_ABILITY_TO_SIT
BIOCARTA_BIOPEPTIDES_PATHWAY, BIOCARTA_BIOPEPTIDES_PATHWAY
HP_HYPOPLASTIC_VERTEBRAL_BODIES, HP_HYPOPLASTIC_VERTEBRAL_BODIES
WP_WNT_SIGNALING, WP_WNT_SIGNALING
CAGGGTC_MIR504, CAGGGTC_MIR504
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_POLYMERIZATION, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_POLYMERIZATION
ZNF502_TARGET_GENES, ZNF502_TARGET_GENES
WP_MYOMETRIAL_RELAXATION_AND_CONTRACTION_PATHWAYS, WP_MYOMETRIAL_RELAXATION_AND_CONTRACTION_PATHWAYS
FARDIN_HYPOXIA_11, FARDIN_HYPOXIA_11
WP_CHOLESTEROL_BIOSYNTHESIS_WITH_SKELETAL_DYSPLASIAS, WP_CHOLESTEROL_BIOSYNTHESIS_WITH_SKELETAL_DYSPLASIAS
HP_DISPROPORTIONATE_SHORT_LIMB_SHORT_STATURE, HP_DISPROPORTIONATE_SHORT_LIMB_SHORT_STATURE
GOBP_INTERLEUKIN_8_PRODUCTION, GOBP_INTERLEUKIN_8_PRODUCTION
SCHRAETS_MLL_TARGETS_UP, SCHRAETS_MLL_TARGETS_UP
LMNB2_TARGET_GENES, LMNB2_TARGET_GENES
MIR563, MIR563
BIOCARTA_PAR1_PATHWAY, BIOCARTA_PAR1_PATHWAY
GOBP_SIGNAL_TRANSDUCTION_IN_ABSENCE_OF_LIGAND, GOBP_SIGNAL_TRANSDUCTION_IN_ABSENCE_OF_LIGAND
WP_NANOPARTICLEMEDIATED_ACTIVATION_OF_RECEPTOR_SIGNALING, WP_NANOPARTICLEMEDIATED_ACTIVATION_OF_RECEPTOR_SIGNALING
WP_EICOSANOID_METABOLISM_VIA_CYCLO_OXYGENASES_COX, WP_EICOSANOID_METABOLISM_VIA_CYCLO_OXYGENASES_COX
MODULE_530, MODULE_530
HP_ABNORMALITY_OF_CARPAL_BONE_OSSIFICATION, HP_ABNORMALITY_OF_CARPAL_BONE_OSSIFICATION
HP_BARREL_SHAPED_CHEST, HP_BARREL_SHAPED_CHEST
HP_ABNORMAL_THALAMIC_MRI_SIGNAL_INTENSITY, HP_ABNORMAL_THALAMIC_MRI_SIGNAL_INTENSITY
HP_HYPOPLASIA_OF_THE_ODONTOID_PROCESS, HP_HYPOPLASIA_OF_THE_ODONTOID_PROCESS
BMI1_DN_MEL18_DN.V1_DN, BMI1_DN_MEL18_DN.V1_DN
MIR432_5P, MIR432_5P
KLEIN_TARGETS_OF_BCR_ABL1_FUSION, KLEIN_TARGETS_OF_BCR_ABL1_FUSION
REACTOME_WNT5A_DEPENDENT_INTERNALIZATION_OF_FZD4, REACTOME_WNT5A_DEPENDENT_INTERNALIZATION_OF_FZD4
PID_THROMBIN_PAR1_PATHWAY, PID_THROMBIN_PAR1_PATHWAY
DOANE_BREAST_CANCER_CLASSES_UP, DOANE_BREAST_CANCER_CLASSES_UP
PID_ALK1_PATHWAY, PID_ALK1_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_ACTIN_NUCLEATION, GOBP_NEGATIVE_REGULATION_OF_ACTIN_NUCLEATION
GOCC_TRAPP_COMPLEX, GOCC_TRAPP_COMPLEX
GOMF_FATTY_ACID_BINDING, GOMF_FATTY_ACID_BINDING
WP_SYNAPTIC_VESICLE_PATHWAY, WP_SYNAPTIC_VESICLE_PATHWAY
GOBP_SYNAPTIC_VESICLE_CYTOSKELETAL_TRANSPORT, GOBP_SYNAPTIC_VESICLE_CYTOSKELETAL_TRANSPORT
REACTOME_CALCINEURIN_ACTIVATES_NFAT, REACTOME_CALCINEURIN_ACTIVATES_NFAT
SQSTM1_TARGET_GENES, SQSTM1_TARGET_GENES
GOBP_NEGATIVE_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION, GOBP_NEGATIVE_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION
REACTOME_P38MAPK_EVENTS, REACTOME_P38MAPK_EVENTS
REACTOME_ASSEMBLY_AND_CELL_SURFACE_PRESENTATION_OF_NMDA_RECEPTORS, REACTOME_ASSEMBLY_AND_CELL_SURFACE_PRESENTATION_OF_NMDA_RECEPTORS
GOBP_ALPHA_LINOLENIC_ACID_METABOLIC_PROCESS, GOBP_ALPHA_LINOLENIC_ACID_METABOLIC_PROCESS
REACTOME_ALPHA_LINOLENIC_OMEGA3_AND_LINOLEIC_OMEGA6_ACID_METABOLISM, REACTOME_ALPHA_LINOLENIC_OMEGA3_AND_LINOLEIC_OMEGA6_ACID_METABOLISM
HP_LUMBAR_HYPERLORDOSIS, HP_LUMBAR_HYPERLORDOSIS
BIOCARTA_CCR5_PATHWAY, BIOCARTA_CCR5_PATHWAY
ZHONG_PFC_C1_NEUROD1_POS_EXCITATORY_NEURON, ZHONG_PFC_C1_NEUROD1_POS_EXCITATORY_NEURON
HP_PERIVENTRICULAR_WHITE_MATTER_HYPERDENSITIES, HP_PERIVENTRICULAR_WHITE_MATTER_HYPERDENSITIES
GOBP_PURINE_DEOXYRIBONUCLEOTIDE_METABOLIC_PROCESS, GOBP_PURINE_DEOXYRIBONUCLEOTIDE_METABOLIC_PROCESS
HP_ABNORMALITY_OF_THE_ODONTOID_PROCESS, HP_ABNORMALITY_OF_THE_ODONTOID_PROCESS
GOBP_REGULATION_OF_MITOCHONDRIAL_FISSION, GOBP_REGULATION_OF_MITOCHONDRIAL_FISSION
MAGRANGEAS_MULTIPLE_MYELOMA_IGLL_VS_IGLK_DN, MAGRANGEAS_MULTIPLE_MYELOMA_IGLL_VS_IGLK_DN
BIOCARTA_CREB_PATHWAY, BIOCARTA_CREB_PATHWAY
GOBP_POSITIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
GOBP_HEPARAN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_HEPARAN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS
HP_CHERRY_RED_SPOT_OF_THE_MACULA, HP_CHERRY_RED_SPOT_OF_THE_MACULA
GOBP_SYNAPTIC_VESICLE_RECYCLING_VIA_ENDOSOME, GOBP_SYNAPTIC_VESICLE_RECYCLING_VIA_ENDOSOME
MODULE_327, MODULE_327
GOBP_RESPONSE_TO_REDOX_STATE, GOBP_RESPONSE_TO_REDOX_STATE
REACTOME_TRIGLYCERIDE_CATABOLISM, REACTOME_TRIGLYCERIDE_CATABOLISM
GOCC_STEREOCILIUM_BASE, GOCC_STEREOCILIUM_BASE
HP_HYPOPLASTIC_HIPPOCAMPUS, HP_HYPOPLASTIC_HIPPOCAMPUS
GOBP_PURINE_DEOXYRIBONUCLEOTIDE_CATABOLIC_PROCESS, GOBP_PURINE_DEOXYRIBONUCLEOTIDE_CATABOLIC_PROCESS
GOBP_PURINE_DEOXYRIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS, GOBP_PURINE_DEOXYRIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS
HP_TYPE_II_LISSENCEPHALY, HP_TYPE_II_LISSENCEPHALY
REACTOME_CYTOSOLIC_IRON_SULFUR_CLUSTER_ASSEMBLY, REACTOME_CYTOSOLIC_IRON_SULFUR_CLUSTER_ASSEMBLY
ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_2H_UP