

L_MEMBRANES_ACT_MAST_CELL_UP,GSE19888_CTRL_VS_T_CELL_MEMBRANES_ACT_MAST_CELL_UP

GSE25890_CTRL_VS_IL33_IL7_TREATED_NUCOTYES_DN,GSE25890_CTRL_VS_IL33_IL7_TREATED_NUCOTYES_DN
GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_DN,GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_DN
GSE20198_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_UP,GSE20198_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_UP
GSE20198_IL12_IL18_VS_IFNA_TREATED_ACT_CD4_TCELL_DN,GSE20198_IL12_IL18_VS_IFNA_TREATED_ACT_CD4_TCELL_DN
GSE5099_MONOCYTE_VS_ALTERNATIVE_M2_MACROPHAGE_DN,GSE5099_MONOCYTE_VS_ALTERNATIVE_M2_MACROPHAGE_DN
GSE19888_NO_PRETREAT_VS_ADENOSINE_A3R_INHIBITOR_PRETREATED_MAST_CELL_TCELL_MEMBRANES_ACT_UP,GSE19888_NO_PRETREAT_VS_ADENOSINE_A3R_INHIBITOR_PRETREATED_MAST_CELL_TCELL_MEMBRANES_ACT_UP
GSE32901_THI7_EMRICHED_VS_THI7_NEG_CD4_TCELL_DN,GSE32901_THI7_EMRICHED_VS_THI7_NEG_CD4_TCELL_DN
GSE9316_CD4_TCELL_BALBC_VS_THI7_ENR1_CD4_TCELL_SKG_PMA_IJONO_STIM_FR4NEG_DN,GSE9316_CD4_TCELL_BALBC_VS_THI7_ENR1_CD4_TCELL_SKG_PMA_IJONO_STIM_FR4NEG_DN
GSE11924_TFH_VS_TH2_CD4_TCELL_DN,GSE11924_TFH_VS_TH2_CD4_TCELL_DN
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN,GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN
PID_MET_PATHWAY,PID_MET_PATHWAY
GOBP_REGULATION_OF_EXOCYTOSIS,GOBP_REGULATION_OF_EXOCYTOSIS
GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_UP,GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_UP
WP_GASTRIN_SIGNALING_PATHWAY,WP_GASTRIN_SIGNALING_PATHWAY
MIR4520_2_3P,MIR4520_2_3P
GOBP_PROTEIN_HOMOOLOGOMERIZATION,GOBP_PROTEIN_HOMOOLOGOMERIZATION
THEILGAARD_NEUTROPHIL_AT_SKIN_WOUND_UP,THEILGAARD_NEUTROPHIL_AT_SKIN_WOUND_UP
MIR1229_3P,MIR1229_3P
MENON_FETAL_KIDNEY_9_ENDOTHELIAL_CELLS,MENON_FETAL_KIDNEY_9_ENDOTHELIAL_CELLS
GSE26928_EFF_MEM_VS_CENTR_MEM_CD4_TCELL_DN,GSE26928_EFF_MEM_VS_CENTR_MEM_CD4_TCELL_DN
GOBP_REGULATION_OF_REGULATED_SECRETORY_PATHWAY,GOBP_REGULATION_OF_REGULATED_SECRETORY_PATHWAY
GSE2585_AIRE_KO_VS_WT_CD80_LOW_MTEC_UP,GSE2585_AIRE_KO_VS_WT_CD80_LOW_MTEC_UP
AMIT_EGF_RESPONSE_60_HELA,AMIT_EGF_RESPONSE_60_HELA
PTPRA_TARGET_GENES,PTPRA_TARGET_GENES
CAFFAREL_RESPONSE_TO_THC_UP,CAFFAREL_RESPONSE_TO_THC_UP
WP_NEURODEGENERATION_WITH_BRAIN_IRON_ACCUMULATION_NBIA_SUBTYPES_PATHWAY,WP_NEURODEGENERATION_WITH_BRAIN_IRON_ACCUMULATION_NBIA_SUBTYPES_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION,GOBP_NEGATIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION
GOBP_RESPONSE_TO_OSMOTIC_STRESS,GOBP_RESPONSE_TO_OSMOTIC_STRESS
GOBP_FOCAL_ADHESION_ASSEMBLY,GOBP_FOCAL_ADHESION_ASSEMBLY
PID_ECADHERIN_NASCENT_AJ_PATHWAY,PID_ECADHERIN_NASCENT_AJ_PATHWAY
PRAMOONJAGO_SOX4_TARGETS_UP,PRAMOONJAGO_SOX4_TARGETS_UP
PID_RHOA_PATHWAY,PID_RHOA_PATHWAY
GSE17721_POLYIC_VS_CPG_0.5H_BMDC_UP,GSE17721_POLYIC_VS_CPG_0.5H_BMDC_UP
GOBP_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT,GOBP_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT
GOBP_ENDOTHELIAL_CELL_DEVELOPMENT,GOBP_ENDOTHELIAL_CELL_DEVELOPMENT
GOBP_SYNAPTIC_VESICLE_EXOCYTOSIS,GOBP_SYNAPTIC_VESICLE_EXOCYTOSIS
GOBP_REGULATION_OF_SYNAPTIC_PLASTICITY,GOBP_REGULATION_OF_SYNAPTIC_PLASTICITY
KEGG_TGF_BETA_SIGNALING_PATHWAY,KEGG_TGF_BETA_SIGNALING_PATHWAY
PID_THROMBIN_PARI_PATHWAY,PID_THROMBIN_PARI_PATHWAY
GTGTGAG_MIR342,GTGTGAG_MIR342
GOBP_REGULATION_OF_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY,GOBP_REGULATION_OF_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY
GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_48H_UP,GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_48H_UP
KUMAR_AUTOPHAGY_NETWORK,KUMAR_AUTOPHAGY_NETWORK
MIR1226_5P,MIR1226_5P
PID_ECADHERIN_STABILIZATION_PATHWAY,PID_ECADHERIN_STABILIZATION_PATHWAY
HP_FATIGUABLE_WEAKNESS_OF_PROXIMAL_LIMB_MUSCLES,HP_FATIGUABLE_WEAKNESS_OF_PROXIMAL_LIMB_MUSCLES
PID_ER_NONGENOMIC_PATHWAY,PID_ER_NONGENOMIC_PATHWAY
GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADDELTA_TCELL_DN,GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADDELTA_TCELL_DN
GOBP_REGULATION_OF_ESTABLISHMENT_OR_MAINTENANCE_OF_CELL_POLARITY,GOBP_REGULATION_OF_ESTABLISHMENT_OR_MAINTENANCE_OF_CELL_POLARITY
GOMF_ALPHA_ACTININ_BINDING,GOMF_ALPHA_ACTININ_BINDING
MARTIN_VIRAL_GPCR_SIGNALING_UP,MARTIN_VIRAL_GPCR_SIGNALING_UP
GOBP_MITOTIC_CYTOKINETIC_PROCESS,GOBP_MITOTIC_CYTOKINETIC_PROCESS
GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATION,GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATION
WP_MICRORNAS_IN_CARDIOMYOCYTE_HYPERTROPHY,WP_MICRORNAS_IN_CARDIOMYOCYTE_HYPERTROPHY
HP_BREEC_PPRESENTATION,HP_BREEC_PPRESENTATION
MIR200C_5P,MIR200C_5P
GSE27786_BCELL_VS_ERYTHROBLAST_DN,GSE27786_BCELL_VS_ERYTHROBLAST_DN
BECKER_TAMOXIFEN_RESISTANCE_UP,BECKER_TAMOXIFEN_RESISTANCE_UP
PID_IL2_P3K_PATHWAY,PID_IL2_P3K_PATHWAY
MARTIN_NFKB_TARGETS_UP,MARTIN_NFKB_TARGETS_UP
MIR4732_5P,MIR4732_5P
PID_VEGFR1_PATHWAY,PID_VEGFR1_PATHWAY
CTCTAGA_MIR526C_MIR518F_MIR526A,CTCTAGA_MIR526C_MIR518F_MIR526A
HP_UPPER_LIMB_AMYOTROPHY,HP_UPPER_LIMB_AMYOTROPHY
GOBP_POSITIVE_REGULATION_OF_T_CELL_CYTOKINE_PRODUCTION,GOBP_POSITIVE_REGULATION_OF_T_CELL_CYTOKINE_PRODUCTION
GOBP_POSITIVE_REGULATION_OF_DENDRITIC_SPINE_DEVELOPMENT,GOBP_POSITIVE_REGULATION_OF_DENDRITIC_SPINE_DEVELOPMENT
MORF_DCC,MORF_DCC
GOBP_REGULATION_OF_SYNAPTIC_VESICLE_EXOCYTOSIS,GOBP_REGULATION_OF_SYNAPTIC_VESICLE_EXOCYTOSIS
GOBP_REGULATION_OF_RESPONSE_TO_OSMOTIC_STRESS,GOBP_REGULATION_OF_RESPONSE_TO_OSMOTIC_STRESS
GOBP_REGULATION_OF_VASCULAR_PERMEABILITY,GOBP_REGULATION_OF_VASCULAR_PERMEABILITY
LE_NEURONAL_DIFFERENTIATION_UP,LE_NEURONAL_DIFFERENTIATION_UP
PID_SIP1_PATHWAY,PID_SIP1_PATHWAY
PID_SHP2_PATHWAY,PID_SHP2_PATHWAY
MIR191_5P,MIR191_5P
GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSLATION,GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSLATION
GOMF_TRANSLATION_REPRESSOR_ACTIVITY,GOMF_TRANSLATION_REPRESSOR_ACTIVITY
PID_AVB3_OPN_PATHWAY,PID_AVB3_OPN_PATHWAY
MIR3200_3P,MIR3200_3P
HASINA_NOL7_TARGETS_DN,HASINA_NOL7_TARGETS_DN
GOBP_POSITIVE_REGULATION_OF_GLIAL_CELL_PROLIFERATION,GOBP_POSITIVE_REGULATION_OF_GLIAL_CELL_PROLIFERATION
MIR7845_5P,MIR7845_5P
WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_AUTOPHAGY,WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_AUTOPHAGY
REACTOME_CD28_DEPENDENT_P3K_AKT_SIGNALING,REACTOME_CD28_DEPENDENT_P3K_AKT_SIGNALING
REACTOME_SEMA4D_IN_SEMAPHORIN_SIGNALING,REACTOME_SEMA4D_IN_SEMAPHORIN_SIGNALING
GOBP_REGULATION_OF_PEPITIDYL_SERINE_THIOPHOSPHORYLATION,GOBP_REGULATION_OF_PEPITIDYL_SERINE_THIOPHOSPHORYLATION
GOBP_POSITIVE_REGULATION_OF_REGULATED_SECRETORY_PATHWAY,GOBP_POSITIVE_REGULATION_OF_REGULATED_SECRETORY_PATHWAY
MCMURRAY_TP53_HRAS_COOPERATION_RESPONSE_DN,MCMURRAY_TP53_HRAS_COOPERATION_RESPONSE_DN
GOBP_PROTEIN_LOCALIZATION_TO_EARLY_ENDOSOME,GOBP_PROTEIN_LOCALIZATION_TO_EARLY_ENDOSOME
GOBP_ENDOTHELIUM_DEVELOPMENT,GOBP_ENDOTHELIUM_DEVELOPMENT
GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_ADHESION_TO_VASCULAR_ENDOTHELIAL_CELL,GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_ADHESION_TO_VASCULAR_ENDOTHELIAL_CELL
MIR8058,MIR8058
MIR4290,MIR4290
HP_SCAFULAR_WINGING,HP_SCAFULAR_WINGING
GOBP_REGULATION_OF_AUTOPHAGOSOME_MATURATION,GOBP_REGULATION_OF_AUTOPHAGOSOME_MATURATION
GOBP_AMP_METABOLIC_PROCESS,GOBP_AMP_METABOLIC_PROCESS
REACTOME_SURFACTANT_METABOLISM,REACTOME_SURFACTANT_METABOLISM
AMIT_EGF_RESPONSE_480_MCF10A,AMIT_EGF_RESPONSE_480_MCF10A
HP_NEUROMUSCULAR_DYSPHAGIA,HP_NEUROMUSCULAR_DYSPHAGIA
REACTOME_SEMA4D_INDUCED_CELL_MIGRATION_AND_GROWTH_CONE_COLLAPSE,REACTOME_SEMA4D_INDUCED_CELL_MIGRATION_AND_GROWTH_CONE_COLLAPSE
REACTOME_RHO_GTPASES_ACTIVATE_ROCKS,REACTOME_RHO_GTPASES_ACTIVATE_ROCKS
REACTOME_INTEGRIN_SIGNALING,REACTOME_INTEGRIN_SIGNALING
GOBP_MODIFICATION_OF_POSTSYNAPTIC_STRUCTURE,GOBP_MODIFICATION_OF_POSTSYNAPTIC_STRUCTURE
REACTOME_REGULATION_OF_GENE_EXPRESSION_BY_HYPOXIA_INDUCIBLE_FACTOR,REACTOME_REGULATION_OF_GENE_EXPRESSION_BY_HYPOXIA_INDUCIBLE_FACTOR
GOBP_CATECHOLAMINE_SECRETION,GOBP_CATECHOLAMINE_SECRETION
OVSYANNIKOVA_PBMCM_FLUARIX_AGE_50_74YO_COMMON_WITH_BOTH_HAI_AND_VNA_28DY_VS_0DY_USED_IN_HAI_AND_VNA_RESPONSE_MODELS_DN,OVSYANNIKOVA_PBMCM_FLUARIX_AGE_50_74YO_COMMON_WITH_BOTH_HAI_AND_VNA_28DY_VS_0DY_USED_IN_HAI_AND_VNA_RESPONSE_MODELS_DN
REACTOME_SIGNALING_BY_TYPE_1_INSULIN_LIKE_GROWTH_FACTOR_1_RECEPTOR_IGF1R,REACTOME_SIGNALING_BY_TYPE_1_INSULIN_LIKE_GROWTH_FACTOR_1_RECEPTOR_IGF1R
HP_SEVERE_COMBINED_IMMUNODEFICIENCY,HP_SEVERE_COMBINED_IMMUNODEFICIENCY
GOBP_RESPONSE_TO_IRON_IJON,GOBP_RESPONSE_TO_IRON_IJON
HP_REDUCED_VITAL_CAPACITY,HP_REDUCED_VITAL_CAPACITY
GOBP_CARDIAC_CELL_DEVELOPMENT,GOBP_CARDIAC_CELL_DEVELOPMENT
HP_HYPOPROTEINEMIA,HP_HYPOPROTEINEMIA
GOBP_NEGATIVE_REGULATION_OF_VASCULAR_PERMEABILITY,GOBP_NEGATIVE_REGULATION_OF_VASCULAR_PERMEABILITY
GOMF_G_PROTEIN_COUPLED_ADENOSINE_RECEPTOR_ACTIVITY,GOMF_G_PROTEIN_COUPLED_ADENOSINE_RECEPTOR_ACTIVITY
GOBP_REGULATION_OF_CALCIIUM_IJON_DEPENDENT_EXOCYTOSIS,GOBP_REGULATION_OF_CALCIIUM_IJON_DEPENDENT_EXOCYTOSIS
GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_ENDOSOME,GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_ENDOSOME
GOBP_VASODILATION,GOBP_VASODILATION
GOBP_NEGATIVE_REGULATION_OF_VIRAL_TRANSCRIPTION,GOBP_NEGATIVE_REGULATION_OF_VIRAL_TRANSCRIPTION
GOBP_MONOCYTE_DIFFERENTIATION,GOBP_MONOCYTE_DIFFERENTIATION
WP_GLIAL_CELL_DIFFERENTIATION,WP_GLIAL_CELL_DIFFERENTIATION
GAUSSMANN_MLL_AF4_FUSION_TARGETS_F_DN,GAUSSMANN_MLL_AF4_FUSION_TARGETS_F_DN
GOBP_NEGATIVE_REGULATION_OF_CELL_CELL_ADHESION_MEDIATED_BY_CADHERIN,GOBP_NEGATIVE_REGULATION_OF_CELL_CELL_ADHESION_MEDIATED_BY_CADHERIN
GOBP_REGULATION_OF_NITRIC_OXIDE_MEDIATED_SIGNAL_TRANSDUCTION,GOBP_REGULATION_OF_NITRIC_OXIDE_MEDIATED_SIGNAL_TRANSDUCTION
GOBP_CARDIAC_MUSCLE_FIBER_DEVELOPMENT,GOBP_CARDIAC_MUSCLE_FIBER_DEVELOPMENT
GOBP_DIPEPTIDE_TRANSMEMBRANE_TRANSPORT,GOBP_DIPEPTIDE_TRANSMEMBRANE_TRANSPORT
HP_MALAR_PROMINENCE,HP_MALAR_PROMINENCE
GOBP_SYNAPTIC_TRANSMISSION_GABAERGIC,GOBP_SYNAPTIC_TRANSMISSION_GABAERGIC
GOBP_MODIFICATION_OF_SYNAPTIC_STRUCTURE,GOBP_MODIFICATION_OF_SYNAPTIC_STRUCTURE
GOMF_PHOSPHOLIPASE_C_ACTIVITY,GOMF_PHOSPHOLIPASE_C_ACTIVITY
WP_SLEEP_REGULATION,WP_SLEEP_REGULATION
WEST_ADRENOCORTICAL_CARINOMA_VS_ADENOMA_DN,WEST_ADRENOCORTICAL_CARINOMA_VS_ADENOMA_DN
GOBP_RADIAL_GLIAL_CELL_DIFFERENTIATION,GOBP_RADIAL_GLIAL_CELL_DIFFERENTIATION
HP_ABNORMAL_LEFT_ATRIUM_MORPHOLOGY,HP_ABNORMAL_LEFT_ATRIUM_MORPHOLOGY
GOBP_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION,GOBP_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION
GOCC_ER_MEMBRANE_INSERTION_COMPLEX,GOCC_ER_MEMBRANE_INSERTION_COMPLEX
GOBP_VENOUS_BLOOD_VESSEL_MORPHOGENESIS,GOBP_VENOUS_BLOOD_VESSEL_MORPHOGENESIS
GOBP_CORONARY_ARTERY_MORPHOGENESIS,GOBP_CORONARY_ARTERY_MORPHOGENESIS
GOBP_POSITIVE_REGULATION_OF_URINE_VOLUME,GOBP_POSITIVE_REGULATION_OF_URINE_VOLUME
DESCARTES_FETAL_HEART_CARDIOMYOCYTES,DESCARTES_FETAL_HEART_CARDIOMYOCYTES
GOBP_NEUROMUSCULAR_PROCESS,GOBP_NEUROMUSCULAR_PROCESS
GOBP_REGULATION_OF_TISSUE_REMODELING,GOBP_REGULATION_OF_TISSUE_REMODELING
GOBP_RESPONSE_TO_ANGIOTENSIN,GOBP_RESPONSE_TO_ANGIOTENSIN
GOBP_NOREPINEPHRINE_SECRETION,GOBP_NOREPINEPHRINE_SECRETION
chr2q23,chr2q23
GOBP_DICARBOXYLIC_ACID_TRANSPORT,GOBP_DICARBOXYLIC_ACID_TRANSPORT
GOMF_GLUCURONOSYL_N_ACETYLGALACTOSAMINYL_PROTEOGLYCAN_4_BETA_N_ACETYLGALACTOSAMINYLTRANSFERASE_ACTIVITY,GOMF_GLUCURONOSYL_N_ACETYLGALACTOSAMINYL_PROTEOGLYCAN_4_BETA_N_ACETYLGALACTOSAMINYLTRANSFERASE_ACTIVITY
ZNFS34A_TARGET_GENES,ZNFS34A_TARGET_GENES
GOBP_ESTABLISHMENT_OR_MAINTENANCE_OF_MONOPOLAR_CELL_POLARITY,GOBP_ESTABLISHMENT_OR_MAINTENANCE_OF_MONOPOLAR_CELL_POLARITY
chr5q22,chr5q22
HP_ASTERIXIS,HP_ASTERIXIS
GOMF_TYPE_5_METABOTROPIC_GLUTAMATE_RECEPTOR_BINDING,GOMF_TYPE_5_METABOTROPIC_GLUTAMATE_RECEPTOR_BINDING
GOBP_DELAMINATION,GOBP_DELAMINATION
GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_SECRETION,GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_SECRETION
MAINA_VHL_TARGETS_UP,MAINA_VHL_TARGETS_UP
GOBP_CARDIAC_VASCULAR_SMOOTH_MUSCLE_CELL_DIFFERENTIATION,GOBP_CARDIAC_VASCULAR_SMOOTH_MUSCLE_CELL_DIFFERENTIATION
REACTOME_RHO_GTPASES_ACTIVATE_RHOTEKIN_AND_RHOPHILINS,REACTOME_RHO_GTPASES_ACTIVATE_RHOTEKIN_AND_RHOPHILINS
GOBP_POSITIVE_REGULATION_OF_CGMP_MEDIATED_SIGNALING,GOBP_POSITIVE_REGULATION_OF_CGMP_MEDIATED_SIGNALING
MIZUKAMI_HYPOXIA_UP,MIZUKAMI_HYPOXIA_UP
WALLACE_PROSTATE_CANCER_DN,WALLACE_PROSTATE_CANCER_DN
GOBP_ACETATE_ETER_TRANSPORT,GOBP_ACETATE_ETER_TRANSPORT
GOBP_MONOACYLGLYCEROL_METABOLIC_PROCESS,GOBP_MONOACYLGLYCEROL_METABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT,GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT
HP_RHEUMATOID_ARTHRITIS,HP_RHEUMATOID_ARTHRITIS
GOMF_INSULIN_LIKE_GROWTH_FACTOR_1_BINDING,GOMF_INSULIN_LIKE_GROWTH_FACTOR_1_BINDING
chr6p23,chr6p23
HP_REDUCED_SYSTOLIC_FUNCTION,HP_REDUCED_SYSTOLIC_FUNCTION
GOBP_NEGATIVE_REGULATION_OF_CALCIIUM_IJON_DEPENDENT_EXOCYTOSIS,GOBP_NEGATIVE_REGULATION_OF_CALCIIUM_IJON_DEPENDENT_EXOCYTOSIS
WP_MICRORNA_FOR_TARGETING_CANCER_GROWTH_AND_VASCULARIZATION_IN_GLIOBLASTOMA,WP_MICRORNA_FOR_TARGETING_CANCER_GROWTH_AND_VASCULARIZATION_IN_GLIOBLASTOMA
REACTOME_P3K_AKT_ACTIVATION,REACTOME_P3K_AKT_ACTIVATION
REACTOME_EPHA_MEDIATED_GROWTH_CONE_COLLAPSE,REACTOME_EPHA_MEDIATED_GROWTH_CONE_COLLAPSE
RBMX_TARGET_GENES,RBMX_TARGET_GENES
GOBP_RESPONSE_TO_GONADOTROPIN_RELEASING_HORMONE,GOBP_RESPONSE_TO_GONADOTROPIN_RELEASING_HORMONE
GOBP_REGULATION_OF_ASTROCYTE_ACTIVATION,GOBP_REGULATION_OF_ASTROCYTE_ACTIVATION
HP_ELEVATED_GAMMA_GLUTAMYLTRANSFERASE_LEVEL,HP_ELEVATED_GAMMA_GLUTAMYLTRANSFERASE_LEVEL
GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_1B,GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_1B
MODULE_201,MODULE_201
GOBP_POSITIVE_REGULATION_OF_GLUTAMATE_SECRETION,GOBP_POSITIVE_REGULATION_OF_GLUTAMATE_SECRETION
GOBP_REGULATION_OF_GLUTAMATE_SECRETION,GOBP_REGULATION_OF_GLUTAMATE_SECRETION
GOBP_REGULATION_OF_SYNAPTIC_TRANSMISSION_GABAERGIC,GOBP_REGULATION_OF_SYNAPTIC_TRANSMISSION_GABAERGIC
HP_FETAL_AKINESIA_SEQUENCE,HP_FETAL_AKINESIA_SEQUENCE
GOBP_NEGATIVE_REGULATION_OF_SYNAPTIC_TRANSMISSION,GOBP_NEGATIVE_REGULATION_OF_SYNAPTIC_TRANSMISSION
GOBP_TRYPTOPHAN_CATABOLIC_PROCESS,GOBP_TRYPTOPHAN_CATABOLIC_PROCESS
MIR4259,MIR4259
GOBP_POSITIVE_REGULATION_OF_TISSUE_REMODELING,GOBP_POSITIVE_REGULATION_OF_TISSUE_REMODELING
ROVERSI_GLIOMA_LOH_REGIONS,ROVERSI_GLIOMA_LOH_REGIONS