

L_MEMBRANES_ACT_MAST_CELL_UP, GSE19888_CTRL_VS_T_CELL_MEMBRANES_ACT_MAST_CELL_UP

GSE25890_CTRL_VS_IL33_IL7_TREATED_NUOCYTES_DN, GSE25890_CTRL_VS_IL33_IL7_TREATED_NUOCYTES_DN
GSE20198_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_UP, GSE20198_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_UP
GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_DN, GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_DN
GSE20198_IL12_IL18_VS_IFNA_TREATED_ACT_CD4_TCELL_DN, GSE20198_IL12_IL18_VS_IFNA_TREATED_ACT_CD4_TCELL_DN
GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_DN, GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_DN
GSE5099_MONOCYTE_VS_ALTERNATIVE_M2_MACROPHAGE_DN, GSE5099_MONOCYTE_VS_ALTERNATIVE_M2_MACROPHAGE_DN
GSE11924_TFH_VS_TH2_CD4_TCELL_DN, GSE11924_TFH_VS_TH2_CD4_TCELL_DN
GSE17721_PAM3CSK4_VS_CPG_1H_BMDC_DN, GSE17721_PAM3CSK4_VS_CPG_1H_BMDC_DN
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN
PID_MET_PATHWAY, PID_MET_PATHWAY
WP_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY, WP_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY
BHATI_G2M_ARREST_BY_2METHOXYESTRADIOL_DN, BHATI_G2M_ARREST_BY_2METHOXYESTRADIOL_DN
GOBP_REGULATION_OF_REGULATED_SECRETORY_PATHWAY, GOBP_REGULATION_OF_REGULATED_SECRETORY_PATHWAY
GOBP_REGULATION_OF_PROTEIN_TARGETING, GOBP_REGULATION_OF_PROTEIN_TARGETING
GOBP_REGULATION_OF_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_INTRACELLULAR_STEROID_HORMONE
GNF2_CBBF, GNF2_CBBF
GOBP_PROTEIN_HOMOOLIGOMERIZATION, GOBP_PROTEIN_HOMOOLIGOMERIZATION
PTPRA_TARGET_GENES, PTPRA_TARGET_GENES
GOBP_FOCAL_ADHESION_ASSEMBLY, GOBP_FOCAL_ADHESION_ASSEMBLY
MIR1229_3P, MIR1229_3P
MILI_PSEUDOPODIA, MILI_PSEUDOPODIA
MIR4520_2_3P, MIR4520_2_3P
MIR875_5P, MIR875_5P
GOBP_POSITIVE_REGULATION_OF_ACTIN_FILAMENT_BUNDLE_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_ACTIN_FILAMENT_BUNDLE_ASSEMBLY
MIR4475, MIR4475
GOBP_POSITIVE_REGULATION_OF_EXOCYTOSIS, GOBP_POSITIVE_REGULATION_OF_EXOCYTOSIS
GOBP_POSITIVE_REGULATION_OF_STRESS_FIBER_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_STRESS_FIBER_ASSEMBLY
MIR4790_3P, MIR4790_3P
GOBP_PROTEIN_HOMOTETRAMERIZATION, GOBP_PROTEIN_HOMOTETRAMERIZATION
GSE45365_HEALTHY_VS_MCMV_INFECTION_CD8_TCELL_IFNAR_KO_DN, GSE45365_HEALTHY_VS_MCMV_INFECTION_CD8_TCELL_IFNAR_KO_DN
HP_BULBAR_SIGNS, HP_BULBAR_SIGNS
GOMF_TAU_PROTEIN_BINDING, GOMF_TAU_PROTEIN_BINDING
GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4_KO_BCELL_UP, GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4_KO_BCELL_UP
HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_HAND, HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_HAND
GOBP_SYNAPTIC_VESICLE_EXOCYTOSIS, GOBP_SYNAPTIC_VESICLE_EXOCYTOSIS
MORF_DCC, MORF_DCC
BECKER_TAMOXIFEN_RESISTANCE_UP, BECKER_TAMOXIFEN_RESISTANCE_UP
GOBP_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT, GOBP_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT
GOBP_POSITIVE_REGULATION_OF_REGULATED_SECRETORY_PATHWAY, GOBP_POSITIVE_REGULATION_OF_REGULATED_SECRETORY_PATHWAY
GOBP_REGULATION_OF_MITOCHONDRIAL_FISSION, GOBP_REGULATION_OF_MITOCHONDRIAL_FISSION
GOBP_REGULATION_OF_SYNAPTIC_VESICLE_EXOCYTOSIS, GOBP_REGULATION_OF_SYNAPTIC_VESICLE_EXOCYTOSIS
GOBP_ACIDIC_AMINO_ACID_TRANSPORT, GOBP_ACIDIC_AMINO_ACID_TRANSPORT
REACTOME_SEMA4D_IN_SEMAPHORIN_SIGNALING, REACTOME_SEMA4D_IN_SEMAPHORIN_SIGNALING
REACTOME_SEMA4D_INDUCED_CELL_MIGRATION_AND_GROWTH_CONE_COLLAPSE, REACTOME_SEMA4D_INDUCED_CELL_MIGRATION_AND_GROWTH_CONE_COLLAPSE
HP_UPPER_LIMB_AMYOTROPHY, HP_UPPER_LIMB_AMYOTROPHY
WP_MICRORNAS_IN_CARDIOMYOCYTE_HYPERTROPHY, WP_MICRORNAS_IN_CARDIOMYOCYTE_HYPERTROPHY
MIR627_5P, MIR627_5P
ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_6HR_DN, ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_6HR_DN
HP_HAND_MUSCLE_WEAKNESS, HP_HAND_MUSCLE_WEAKNESS
PBX1_02, PBX1_02
GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT
GOMF_ALPHA_ACTININ_BINDING, GOMF_ALPHA_ACTININ_BINDING
GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_48H_UP
GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_SECRETION, GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_SECRETION
GOBP_REGULATION_OF_PROTEIN_TARGETING_TO_MEMBRANE, GOBP_REGULATION_OF_PROTEIN_TARGETING_TO_MEMBRANE
GOBP_MITOTIC_CYTOKINETIC_PROCESS, GOBP_MITOTIC_CYTOKINETIC_PROCESS
GOBP_RETICULOPHAGY, GOBP_RETICULOPHAGY
GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSLATION, GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSLATION
GOBP_GOLGI_TO_LYSOSOME_TRANSPORT, GOBP_GOLGI_TO_LYSOSOME_TRANSPORT
GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_OSMOTIC_STRESS, GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_OSMOTIC_STRESS
MIR1226_5P, MIR1226_5P
HP_BREECH_PRESENTATION, HP_BREECH_PRESENTATION
GOBP_MODIFICATION_OF_SYNAPTIC_STRUCTURE, GOBP_MODIFICATION_OF_SYNAPTIC_STRUCTURE
LE_NEURONAL_DIFFERENTIATION_UP, LE_NEURONAL_DIFFERENTIATION_UP
GOBP_VASODILATION, GOBP_VASODILATION
GOBP_POSITIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MEMBRANE, GOBP_POSITIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MEMBRANE
GOBP_PEPTIDYL_TYROSINE_AUTOPHOSPHORYLATION, GOBP_PEPTIDYL_TYROSINE_AUTOPHOSPHORYLATION
chr17p11, chr17p11
GOBP_EPITHELIAL_CELL_APOPTOTIC_PROCESS, GOBP_EPITHELIAL_CELL_APOPTOTIC_PROCESS
GOBP_SLEEP, GOBP_SLEEP
GAUSSMANN_MLL_AF4_FUSION_TARGETS_F_DN, GAUSSMANN_MLL_AF4_FUSION_TARGETS_F_DN
MARTIN_NFKB_TARGETS_UP, MARTIN_NFKB_TARGETS_UP
ZNF354A_TARGET_GENES, ZNF354A_TARGET_GENES
GOBP_GLUTAMATE_SECRETION, GOBP_GLUTAMATE_SECRETION
GOBP_REGULATION_OF_RESPONSE_TO_OSMOTIC_STRESS, GOBP_REGULATION_OF_RESPONSE_TO_OSMOTIC_STRESS
HP_PROMINENT_SUPRAORBITAL_RIDGES, HP_PROMINENT_SUPRAORBITAL_RIDGES
PID_S1P_S1P1_PATHWAY, PID_S1P_S1P1_PATHWAY
GOBP_REGULATION_OF_AUTOPHAGOSOME_MATURATION, GOBP_REGULATION_OF_AUTOPHAGOSOME_MATURATION
HP_RHEUMATOID_ARTHRITIS, HP_RHEUMATOID_ARTHRITIS
GOBP_POSITIVE_REGULATION_OF_T_CELL_CYTOKINE_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_T_CELL_CYTOKINE_PRODUCTION
KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_HEPARAN_SULFATE, KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_HEPARAN_SULFATE
GOMF_PHOSPHOLIPASE_C_ACTIVITY, GOMF_PHOSPHOLIPASE_C_ACTIVITY
GOBP_MODIFICATION_OF_POSTSYNAPTIC_STRUCTURE, GOBP_MODIFICATION_OF_POSTSYNAPTIC_STRUCTURE
MIR6800_3P, MIR6800_3P
GOBP_PRIMARY_MIRNA_PROCESSING, GOBP_PRIMARY_MIRNA_PROCESSING
GOBP_GLUTAMATE_RECEPTOR_SIGNALING_PATHWAY, GOBP_GLUTAMATE_RECEPTOR_SIGNALING_PATHWAY
DACOSTA_UV_RESPONSE_VIA_ERCC3_XPCS_UP, DACOSTA_UV_RESPONSE_VIA_ERCC3_XPCS_UP
MAINA_VHL_TARGETS_UP, MAINA_VHL_TARGETS_UP
GOBP_REGULATION_OF_CALCIUM_ION_DEPENDENT_EXOCYTOSIS, GOBP_REGULATION_OF_CALCIUM_ION_DEPENDENT_EXOCYTOSIS
GOBP_POSITIVE_REGULATION_OF_CIRCADIAN_SLEEP_WAKE_CYCLE_SLEEP, GOBP_POSITIVE_REGULATION_OF_CIRCADIAN_SLEEP_WAKE_CYCLE_SLEEP
GOBP_SYNAPTIC_TRANSMISSION_GABAERGIC, GOBP_SYNAPTIC_TRANSMISSION_GABAERGIC
GOBP_AMP_METABOLIC_PROCESS, GOBP_AMP_METABOLIC_PROCESS
WEST_ADRENOCORTICAL_CARCINOMA_VS_ADENOMA_DN, WEST_ADRENOCORTICAL_CARCINOMA_VS_ADENOMA_DN
WP_SLEEP_REGULATION, WP_SLEEP_REGULATION
GOBP_POSITIVE_REGULATION_OF_AMINE_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_AMINE_TRANSPORT
REACTOME_ERYTHROCYTES_TAKE_UP_OXYGEN_AND_RELEASE CARBON_DIOXIDE, REACTOME_ERYTHROCYTES_TAKE_UP_OXYGEN_AND_RELEASE CARBON_DIOXIDE
GOBP_REGULATION_OF_SYNAPTIC_TRANSMISSION_GABAERGIC, GOBP_REGULATION_OF_SYNAPTIC_TRANSMISSION_GABAERGIC
MODULE_512, MODULE_512
REACTOME_EPHA_MEDIATED_GROWTH_CONE_COLLAPSE, REACTOME_EPHA_MEDIATED_GROWTH_CONE_COLLAPSE
GOBP_ONE CARBON_COMPOUND_TRANSPORT, GOBP_ONE CARBON_COMPOUND_TRANSPORT
HP_REDUCED_VITAL_CAPACITY, HP_REDUCED_VITAL_CAPACITY
HP_PROTEIN_LOSING_ENTEROPATHY, HP_PROTEIN_LOSING_ENTEROPATHY
GOBP_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION, GOBP_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION
GOBP_REGULATION_OF_PEPTIDYL_TYROSINE_AUTOPHOSPHORYLATION, GOBP_REGULATION_OF_PEPTIDYL_TYROSINE_AUTOPHOSPHORYLATION
GOBP CARBON_DIOXIDE_TRANSPORT, GOBP CARBON_DIOXIDE_TRANSPORT
GOCC_ER_MEMBRANE_INSERTION_COMPLEX, GOCC_ER_MEMBRANE_INSERTION_COMPLEX
WP_HYPOTHETICAL_CRANIOFACIAL_DEVELOPMENT_PATHWAY, WP_HYPOTHETICAL_CRANIOFACIAL_DEVELOPMENT_PATHWAY
GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_ADHESION_TO_VASCULAR_ENDOTHELIAL_CELL, GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_ADHESION_TO_VASCULAR_ENDOTHELIAL_CELL