

L_VS_POLYIC_12H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_12H_BMDC_UP

GSE17721_CTRL_VS_POLYIC_6H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_6H_BMDC_UP
GSE17721_0.5H_VS_4H_LPS_BMDC_UP, GSE17721_0.5H_VS_4H_LPS_BMDC_UP
GSE17721_CTRL_VS_CPG_8H_BMDC_UP, GSE17721_CTRL_VS_CPG_8H_BMDC_UP
GSE17721_POLYIC_VS_GARDIQUIMOD_0.5H_BMDC_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_0.5H_BMDC_DN
GSE17721_CTRL_VS_LPS_4H_BMDC_UP, GSE17721_CTRL_VS_LPS_4H_BMDC_UP
GSE17721_CTRL_VS_PAM3CSK4_8H_BMDC_UP, GSE17721_CTRL_VS_PAM3CSK4_8H_BMDC_UP
GSE37605_C57BL6_VS_NOD_FOXP3_FUSION_GFP_TCONV_DN, GSE37605_C57BL6_VS_NOD_FOXP3_FUSION_GFP_TCONV_DN
GSE17721_CTRL_VS_CPG_6H_BMDC_UP, GSE17721_CTRL_VS_CPG_6H_BMDC_UP
GSE17721_CTRL_VS_GARDIQUIMOD_4H_BMDC_UP, GSE17721_CTRL_VS_GARDIQUIMOD_4H_BMDC_UP
GSE17721_CTRL_VS_GARDIQUIMOD_8H_BMDC_UP, GSE17721_CTRL_VS_GARDIQUIMOD_8H_BMDC_UP
GSE17721_CTRL_VS_LPS_8H_BMDC_UP, GSE17721_CTRL_VS_LPS_8H_BMDC_UP
GSE17721_CTRL_VS_PAM3CSK4_4H_BMDC_UP, GSE17721_CTRL_VS_PAM3CSK4_4H_BMDC_UP
GSE17721_CTRL_VS_LPS_6H_BMDC_UP, GSE17721_CTRL_VS_LPS_6H_BMDC_UP
GSE17721_CTRL_VS_GARDIQUIMOD_12H_BMDC_UP, GSE17721_CTRL_VS_GARDIQUIMOD_12H_BMDC_UP
GSE17721_CTRL_VS_GARDIQUIMOD_6H_BMDC_UP, GSE17721_CTRL_VS_GARDIQUIMOD_6H_BMDC_UP
GSE17721_CTRL_VS_LPS_12H_BMDC_UP, GSE17721_CTRL_VS_LPS_12H_BMDC_UP
GSE17721_CTRL_VS_POLYIC_24H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_24H_BMDC_UP
GSE17721_CPG_VS_GARDIQUIMOD_6H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_6H_BMDC_DN
GSE17721_CTRL_VS_LPS_2H_BMDC_UP, GSE17721_CTRL_VS_LPS_2H_BMDC_UP
GSE17721_0.5H_VS_8H_LPS_BMDC_UP, GSE17721_0.5H_VS_8H_LPS_BMDC_UP
GSE17721_CTRL_VS_LPS_1H_BMDC_UP, GSE17721_CTRL_VS_LPS_1H_BMDC_UP
GSE17721_CTRL_VS_PAM3CSK4_0.5H_BMDC_UP, GSE17721_CTRL_VS_PAM3CSK4_0.5H_BMDC_UP
GSE17721_0.5H_VS_24H_LPS_BMDC_UP, GSE17721_0.5H_VS_24H_LPS_BMDC_UP
GSE10273_HIGH_VS_LOW_IL7_TREATED_IRF4_8_NULL_PRE_BCELL_UP, GSE10273_HIGH_VS_LOW_IL7_TREATED_IRF4_8_NULL_PRE_BCELL_UP
GSE17721_CTRL_VS_GARDIQUIMOD_24H_BMDC_UP, GSE17721_CTRL_VS_GARDIQUIMOD_24H_BMDC_UP
GSE17721_CTRL_VS_POLYIC_2H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_2H_BMDC_UP
GSE7831_UNSTIM_VS_INFLUENZA_STIM_PDC_1H_UP, GSE7831_UNSTIM_VS_INFLUENZA_STIM_PDC_1H_UP
GSE17721_CTRL_VS_GARDIQUIMOD_2H_BMDC_UP, GSE17721_CTRL_VS_GARDIQUIMOD_2H_BMDC_UP
GSE17721_LPS_VS_CPG_6H_BMDC_DN, GSE17721_LPS_VS_CPG_6H_BMDC_DN
GSE17721_0.5H_VS_4H_POLYIC_BMDC_UP, GSE17721_0.5H_VS_4H_POLYIC_BMDC_UP
GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_DN
GSE5542_UNTREATED_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP, GSE5542_UNTREATED_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP
LIU_CMYB_TARGETS_UP, LIU_CMYB_TARGETS_UP
GSE17721_LPS_VS_PAM3CSK4_0.5H_BMDC_UP, GSE17721_LPS_VS_PAM3CSK4_0.5H_BMDC_UP
PASQUALUCCI_LYMPHOMA_BY_GC_STAGE_DN, PASQUALUCCI_LYMPHOMA_BY_GC_STAGE_DN
GSE11057_EFF_MEM_VS_CENT_MEM_CD4_TCELL_DN, GSE11057_EFF_MEM_VS_CENT_MEM_CD4_TCELL_DN
DOANE_RESPONSE_TO_ANDROGEN_UP, DOANE_RESPONSE_TO_ANDROGEN_UP
GSE12003_4D_VS_8D_CULTURE_MIR223_KO_BM_PROGENITOR_DN, GSE12003_4D_VS_8D_CULTURE_MIR223_KO_BM_PROGENITOR_DN
GSE6259_FLT3L_INDUCED_DEC205_POS_DC_VS_CD8_TCELL_UP, GSE6259_FLT3L_INDUCED_DEC205_POS_DC_VS_CD8_TCELL_UP
GSE25677_MPL_VS_MPL_AND_R848_STIM_BCELL_UP, GSE25677_MPL_VS_MPL_AND_R848_STIM_BCELL_UP
RACTNNRTTTNC_UNKNOWN, RACTNNRTTTNC_UNKNOWN
HP_GASTROINTESTINAL_DYSMOTILITY, HP_GASTROINTESTINAL_DYSMOTILITY
AACYNNNNTTCCS_UNKNOWN, AACYNNNNTTCCS_UNKNOWN
GSE12003_MIR223_KO_VS_WT_BM_PROGENITOR_8D_CULTURE_UP, GSE12003_MIR223_KO_VS_WT_BM_PROGENITOR_8D_CULTURE_UP
HP_DELAYED_CRANIAL_SUTURE_CLOSURE, HP_DELAYED_CRANIAL_SUTURE_CLOSURE
IWANAGA_CARCIINOGENESIS_BY_KRAS_UP, IWANAGA_CARCIINOGENESIS_BY_KRAS_UP
GOBP_REGULATION_OF_PROTEIN_TARGETING, GOBP_REGULATION_OF_PROTEIN_TARGETING
HP_ABNORMALITY_OF_THE_DIENCEPHALON, HP_ABNORMALITY_OF_THE_DIENCEPHALON
HP_BRADYKINESIA, HP_BRADYKINESIA
HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_HAND, HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_HAND
HP_ABNORMALITY_OF_THE_SENSE_OF_SMELL, HP_ABNORMALITY_OF_THE_SENSE_OF_SMELL
GOCC_CORTICAL_CYTOSKELETON, GOCC_CORTICAL_CYTOSKELETON
GOBP_RECEPTOR_CLUSTERING, GOBP_RECEPTOR_CLUSTERING
GOBP_LOCALIZATION_WITHIN_MEMBRANE, GOBP_LOCALIZATION_WITHIN_MEMBRANE
GOBP_CARBOHYDRATE_DERIVATIVE_TRANSPORT, GOBP_CARBOHYDRATE_DERIVATIVE_TRANSPORT
HP_HYPOSMIA, HP_HYPOSMIA
GOBP_CELLULAR_RESPONSE_TO_CARBOHYDRATE_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_CARBOHYDRATE_STIMULUS
GOBP_CELLULAR_GLUCOSE_HOMEOSTASIS, GOBP_CELLULAR_GLUCOSE_HOMEOSTASIS
GOBP_CELLULAR_EXTRAVASATION, GOBP_CELLULAR_EXTRAVASATION
HP_PROXIMAL_MUSCLE_WEAKNESS_IN_LOWER_LIMBS, HP_PROXIMAL_MUSCLE_WEAKNESS_IN_LOWER_LIMBS
HP_ABNORMAL_AUTONOMIC_NERVOUS_SYSTEM_PHYSIOLOGY, HP_ABNORMAL_AUTONOMIC_NERVOUS_SYSTEM_PHYSIOLOGY
GOBP_ORGANISM_EMERGENCE_FROM_PROTECTIVE_STRUCTURE, GOBP_ORGANISM_EMERGENCE_FROM_PROTECTIVE_STRUCTURE
GOBP_CAMP_METABOLIC_PROCESS, GOBP_CAMP_METABOLIC_PROCESS
ZFP28_TARGET_GENES, ZFP28_TARGET_GENES
GOBP_FUCOSE_METABOLIC_PROCESS, GOBP_FUCOSE_METABOLIC_PROCESS
REACTOME_INTERLEUKIN_6_SIGNALING, REACTOME_INTERLEUKIN_6_SIGNALING
GOMF_OPIOID_RECEPTOR_BINDING, GOMF_OPIOID_RECEPTOR_BINDING
REACTOME_ACYL_CHAIN_REMODELING_OF_DAG_AND_TAG, REACTOME_ACYL_CHAIN_REMODELING_OF_DAG_AND_TAG
GOBP_NEGATIVE_REGULATION_OF_LYASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_LYASE_ACTIVITY
GOCC_APICAL_DENDRITE, GOCC_APICAL_DENDRITE
GOBP_SYNAPTIC_TRANSMISSION_DOPAMINERGIC, GOBP_SYNAPTIC_TRANSMISSION_DOPAMINERGIC
GOBP_ACYLGLYCEROL_ACYL_CHAIN_REMODELING, GOBP_ACYLGLYCEROL_ACYL_CHAIN_REMODELING
GOBP_MONOACYLGLYCEROL_BIOSYNTHETIC_PROCESS, GOBP_MONOACYLGLYCEROL_BIOSYNTHETIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_NEURON_MIGRATION, GOBP_POSITIVE_REGULATION_OF_NEURON_MIGRATION
HP_PAROXYSMAL_DYSKINESIA, HP_PAROXYSMAL_DYSKINESIA
HP_STRIDOR, HP_STRIDOR
GOBP_NEUTROPHIL_EXTRAVASATION, GOBP_NEUTROPHIL_EXTRAVASATION
PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_6, PEDERSEN_METASTASIS_BY_ERBB2_ISOFORM_6
GOBP_GLYCOSIDE_TRANSPORT, GOBP_GLYCOSIDE_TRANSPORT
GOBP_TESTOSTERONE_BIOSYNTHETIC_PROCESS, GOBP_TESTOSTERONE_BIOSYNTHETIC_PROCESS
GOMF_VOLTAGE_GATED_CALCIUM_CHANNEL_ACTIVITY_INVOLVED_IN_REGULATION_OF_CYTOSOLIC_CALCIUM_LEVELS, GOMF_VOLTAGE_GATED_CALCIUM_CHANNEL_ACTIVITY_INVOLVED_IN_REGULATION_OF.C
FIGUEROA_AML_METHYLATION_CLUSTER_2_DN, FIGUEROA_AML_METHYLATION_CLUSTER_2_DN
BIOCARTA_FOSB_PATHWAY, BIOCARTA_FOSB_PATHWAY
GOMF_DIACYLGLYCEROL_O_ACYLTRANSFERASE_ACTIVITY, GOMF_DIACYLGLYCEROL_O_ACYLTRANSFERASE_ACTIVITY
GOBP_REGULATION_OF_ATRIAL_CARDIAC_MUSCLE_CELL_MEMBRANE_REPOLARIZATION, GOBP_REGULATION_OF_ATRIAL_CARDIAC_MUSCLE_CELL_MEMBRANE_REPOLARIZATION
REACTOME_TERMINAL_PATHWAY_OF_COMPLEMENT, REACTOME_TERMINAL_PATHWAY_OF_COMPLEMENT
GOCC_MEMBRANE_ATTACK_COMPLEX, GOCC_MEMBRANE_ATTACK_COMPLEX
GOBP_MEMBRANE_REPOLARIZATION_DURING_ATRIAL_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL, GOBP_MEMBRANE_REPOLARIZATION_DURING_ATRIAL_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL
HP_ANISOSPONDYLY, HP_ANISOSPONDYLY
GOCC_CALCITONIN_FAMILY_RECEPTOR_COMPLEX, GOCC_CALCITONIN_FAMILY_RECEPTOR_COMPLEX
GOBP_ATRIAL_CARDIAC_MUSCLE_CELL_MEMBRANE_REPOLARIZATION, GOBP_ATRIAL_CARDIAC_MUSCLE_CELL_MEMBRANE_REPOLARIZATION