

AND_DNFB_ALLERGEN_TREATED_DC_UP, GSE20727_CTRL_VS_ROS_INH_AND_DNFB_ALLERGEN_TREATED_DC_UP

GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_2H_UP, GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_2H_UP
GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_17H_UP, GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_17H_UP
GSE9988_LOW_LPS_VS_ANTI_TREM1_AND_LPS_MONOCYTE_UP, GSE9988_LOW_LPS_VS_ANTI_TREM1_AND_LPS_MONOCYTE_UP
GSE37301_COMMON_LYMPHOID_PROGENITOR_VS_CD4_TCELL_DN, GSE37301_COMMON_LYMPHOID_PROGENITOR_VS_CD4_TCELL_DN
GSE26290_CTRL_VS_AKT_INHIBITOR_TREATED_ANTI_CD3_AND_IL2_STIM_CD8_TCELL_DN, GSE26290_CTRL_VS_AKT_INHIBITOR_TREATED_ANTI_CD3_AND_IL2
IRX2_TARGET_GENES, IRX2_TARGET_GENES
GSE4142_GC_BCELL_VS_MEMORY_BCELL_DN, GSE4142_GC_BCELL_VS_MEMORY_BCELL_DN
GOMF_LYASE_ACTIVITY, GOMF_LYASE_ACTIVITY
GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_2H_DN, GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_2H_DN
GSE21063_WT_VS_NFATC1_KO_3H_ANTI_IGM_STIM_BCELL_UP, GSE21063_WT_VS_NFATC1_KO_3H_ANTI_IGM_STIM_BCELL_UP
GSE17721_PAM3CSK4_VS_CPG_2H_BMDC_DN, GSE17721_PAM3CSK4_VS_CPG_2H_BMDC_DN
GSE4984_LPS_VS_VEHICLE_CTRL_TREATED_DC_UP, GSE4984_LPS_VS_VEHICLE_CTRL_TREATED_DC_UP
HP_HYPOPLASIA_OF_THE_ODONTOID_PROCESS, HP_HYPOPLASIA_OF_THE_ODONTOID_PROCESS
HP_NEPHRONOPHTHISIS, HP_NEPHRONOPHTHISIS
HP_ABNORMALITY_OF_THE_ODONTOID_PROCESS, HP_ABNORMALITY_OF_THE_ODONTOID_PROCESS
ENGELMANN_CANCER_PROGENITORS_DN, ENGELMANN_CANCER_PROGENITORS_DN
CHEN_HOXA5_TARGETS_9HR_DN, CHEN_HOXA5_TARGETS_9HR_DN
GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_UP, GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_UP
HP_SHORT_RIBS, HP_SHORT_RIBS
HP_KETOSIS, HP_KETOSIS
WP_MIR5093P_ALTERATION_OF_YAP1ECM_AXIS, WP_MIR5093P_ALTERATION_OF_YAP1ECM_AXIS
HP_RIGHT_BUNDLE_BRANCH_BLOCK, HP_RIGHT_BUNDLE_BRANCH_BLOCK
HP_SKIN_EROSION, HP_SKIN_EROSION
GOBP_CENTRAL_NERVOUS_SYSTEM_PROJECTION_NEURON_AXONOGENESIS, GOBP_CENTRAL_NERVOUS_SYSTEM_PROJECTION_NEURON_AXONOGENESIS
MODULE_444, MODULE_444
GOBP_MITOTIC_NUCLEAR_MEMBRANE_ORGANIZATION, GOBP_MITOTIC_NUCLEAR_MEMBRANE_ORGANIZATION
GOBP_ALDITOL_PHOSPHATE_METABOLIC_PROCESS, GOBP_ALDITOL_PHOSPHATE_METABOLIC_PROCESS
HP_SPONDYLOMETAPHYSEAL_DYSPLASIA, HP_SPONDYLOMETAPHYSEAL_DYSPLASIA
MODULE_516, MODULE_516
WP_SIMPLIFIED_INTERACTION_MAP_BETWEEN_LOXL4_AND_OXIDATIVE_STRESS_PATHWAY, WP_SIMPLIFIED_INTERACTION_MAP_BETWEEN_LOXL4_AND_OX
GOMF_CCR1_CHEMOKINE_RECEPTOR_BINDING, GOMF_CCR1_CHEMOKINE_RECEPTOR_BINDING
GOBP_MYOTUBE_DIFFERENTIATION_INVOLVED_IN_SKELETAL_MUSCLE_REGENERATION, GOBP_MYOTUBE_DIFFERENTIATION_INVOLVED_IN_SKELETAL_MU
GOBP_PEPTIDE_CROSS_LINKING, GOBP_PEPTIDE_CROSS_LINKING
GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_CHEMOTAXIS, GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_CHEMOTAXIS
HP_BELL_SHAPED_THORAX, HP_BELL_SHAPED_THORAX
GOCC_FIBRINOGEN_COMPLEX, GOCC_FIBRINOGEN_COMPLEX
GOBP_PROTEIN_ACTIVATION_CASCADE, GOBP_PROTEIN_ACTIVATION_CASCADE
MODULE_368, MODULE_368
GOBP_EPOXIDE_METABOLIC_PROCESS, GOBP_EPOXIDE_METABOLIC_PROCESS