

ND_DNFB_ALLERGEN_TREATED_DC_DN, GSE20727_CTRL_VS_ROS_INH_AND_DNFB_ALLERGEN_TREATED_DC_DN

GSE22886_IGG_IGA_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_DN, GSE22886_IGG_IGA_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_DN
GSE7509_UNSTIM_VS_IFNA_STIM_IMMATURE_DC_UP, GSE7509_UNSTIM_VS_IFNA_STIM_IMMATURE_DC_UP
GSE26030_TH1_VS_TH17_RESTIMULATED_DAY15_POST_POLARIZATION_UP, GSE26030_TH1_VS_TH17_RESTIMULATED_DAY15_POST_POLARIZATION_UP
GO_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN, GO_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN
GO_MITOTIC_CELL_CYCLE_CHECKPOINT, GO_MITOTIC_CELL_CYCLE_CHECKPOINT
GSE17721_CTRL_VS_POLYIC_6H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_6H_BMDC_UP
GO_NEGATIVE_REGULATION_OF_PROTEIN_COMPLEX_DISASSEMBLY, GO_NEGATIVE_REGULATION_OF_PROTEIN_COMPLEX_DISASSEMBLY
GSE17721_12H_VS_24H_POLYIC_BMDC_UP, GSE17721_12H_VS_24H_POLYIC_BMDC_UP
GSE20198_UNTREATED_VS_IL12_IL18_TREATED_ACT_CD4_TCELL_DN, GSE20198_UNTREATED_VS_IL12_IL18_TREATED_ACT_CD4_TCELL_DN
GSE37301 GRANULOCYTE_MONOCYTE_PROGENITOR_VS_RAG2_KO_NK_CELL_UP, GSE37301 GRANULOCYTE_MONOCYTE_PROGENITOR_VS_RAG2_KO_NK_CELL_UP
REACTOME_MITOTIC_G2_G2_M_PHASES, REACTOME_MITOTIC_G2_G2_M_PHASES
GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_48H_UP, GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_48H_UP
GO_POSITIVE_REGULATION_OF_PEPTIDASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_PEPTIDASE_ACTIVITY
GO_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN, GO_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN
GSE27241_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_TH17_POLARIZING_CONDITIONS_UP, GSE27241_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_TH17_POLARIZING_CONDITIONS_UP
REACTOME_LOSS_OF_NLP_FROM_MITOTIC_CENTROSOMES, REACTOME_LOSS_OF_NLP_FROM_MITOTIC_CENTROSOMES
GSE17721_CTRL_VS_GARDIQUIMOD_2H_BMDC_DN, GSE17721_CTRL_VS_GARDIQUIMOD_2H_BMDC_DN
KAECH_NAIVE_VS_MEMORY_CD8_TCELL_UP, KAECH_NAIVE_VS_MEMORY_CD8_TCELL_UP
GO_MEIOTIC_CELL_CYCLE, GO_MEIOTIC_CELL_CYCLE
GO_SPINDLE_ASSEMBLY, GO_SPINDLE_ASSEMBLY
GSE36078_UNTREATED_VS_AD5_INF_IL1R_KO_MOUSE_LUNG_DC_DN, GSE36078_UNTREATED_VS_AD5_INF_IL1R_KO_MOUSE_LUNG_DC_DN
GSE411_WT_VS_SOCS3_KO_MACROPHAGE_IL6_STIM_400MIN_UP, GSE411_WT_VS_SOCS3_KO_MACROPHAGE_IL6_STIM_400MIN_UP
ONDER_CDH1_TARGETS_1_UP, ONDER_CDH1_TARGETS_1_UP
GSE45365_WT_VS_IFNAR_KO_BCELL_DN, GSE45365_WT_VS_IFNAR_KO_BCELL_DN
GSE14308_TH2_VS_TH1_DN, GSE14308_TH2_VS_TH1_DN
TSAI_RESPONSE_TO_IONIZING_RADIATION, TSAI_RESPONSE_TO_IONIZING_RADIATION
ZHANG_BREAST_CANCER_PROGENITORS_DN, ZHANG_BREAST_CANCER_PROGENITORS_DN
PID_P73PATHWAY, PID_P73PATHWAY
PETROVA_ENDOTHELIUM_LYMPHATIC_VS_BLOOD_UP, PETROVA_ENDOTHELIUM_LYMPHATIC_VS_BLOOD_UP
GO_MEIOTIC_CELL_CYCLE_PROCESS, GO_MEIOTIC_CELL_CYCLE_PROCESS
GSE12839_CTRL_VS_IL12_TREATED_PPMC_UP, GSE12839_CTRL_VS_IL12_TREATED_PPMC_UP
GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_30MIN_DN, GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_30MIN_DN
GSE26669_CD4_VS_CD8_TCELL_IN_MLR_COSTIM_BLOCK_DN, GSE26669_CD4_VS_CD8_TCELL_IN_MLR_COSTIM_BLOCK_DN
HOFMANN_CELL_LYMPHOMA_DN, HOFMANN_CELL_LYMPHOMA_DN
GSE45365_WT_VS_IFNAR_KO_BCELL_MCMV_INFECTION_DN, GSE45365_WT_VS_IFNAR_KO_BCELL_MCMV_INFECTION_DN
GO_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION, GO_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION
GO_POSITIVE_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_POSITIVE_REGULATION_OF_CHROMOSOME_SEGREGATION
GO_ENDOPLASMIC_RETICULUM_ORGANIZATION, GO_ENDOPLASMIC_RETICULUM_ORGANIZATION
GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN_2_OXOGLUTARATE_AS_ONE_DONOR_AND_INCORPORATION_OF_ONE_ATOM_EACH_OF_OXYGEN_INTO_BOTH_DONORS, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN_2_OXOGLUTARATE_AS_ONE_DONOR_AND_INCORPORATION_OF_ONE_ATOM_EACH_OF_OXYGEN_INTO_BOTH_DONORS
GSE17721_LPS_VS_PAM3CSK4_4H_BMDC_UP, GSE17721_LPS_VS_PAM3CSK4_4H_BMDC_UP
MODULE_303, MODULE_303
GSE9650_EXHAUSTED_VS_MEMORY_CD8_TCELL_UP, GSE9650_EXHAUSTED_VS_MEMORY_CD8_TCELL_UP
GO_ER_NUCLEUS_SIGNALING_PATHWAY, GO_ER_NUCLEUS_SIGNALING_PATHWAY
REACTOME_BIOSYNTHESIS_OF_THE_N_GLYCAN_PRECURSOR_DOLICHOL_LIPID_LINKED_OLIGOSACCHARIDE_LLO_AND_TRANSFER_TO_A_NASCENT_PROTEIN, REACTOME_BIOSYNTHESIS_OF_THE_N_GLYCAN_PRECURSOR_DOLICHOL_LIPID_LINKED_OLIGOSACCHARIDE_LLO_AND_TRANSFER_TO_A_NASCENT_PROTEIN
GSE13411_SWITCHED_MEMORY_BCELL_VS_PLASMA_CELL_DN, GSE13411_SWITCHED_MEMORY_BCELL_VS_PLASMA_CELL_DN
GO_POSITIVE_REGULATION_OF_NUCLEASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_NUCLEASE_ACTIVITY
GO_N_GLYCAN_PROCESSING, GO_N_GLYCAN_PROCESSING
GSE45365_CD8A_DC_VS_CD11B_DC_IFNAR_KO_MCMV_INFECTION_UP, GSE45365_CD8A_DC_VS_CD11B_DC_IFNAR_KO_MCMV_INFECTION_UP
GSE4142_NAIVE_VS_MEMORY_BCELL_UP, GSE4142_NAIVE_VS_MEMORY_BCELL_UP
GO_ORGAN_REGENERATION, GO_ORGAN_REGENERATION
TERAO_AOX4_TARGETS_HG_UP, TERA0_AOX4_TARGETS_HG_UP
CHUANG_OXIDATIVE_STRESS_RESPONSE_DN, CHUANG_OXIDATIVE_STRESS_RESPONSE_DN
GO_MITOTIC_SPINDLE_ASSEMBLY, GO_MITOTIC_SPINDLE_ASSEMBLY
GO_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEIN_CATABOLIC_PROCESS, GO_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEIN_CATABOLIC_PROCESS
GO_NEGATIVE_REGULATION_OF_T_CELL_PROLIFERATION, GO_NEGATIVE_REGULATION_OF_T_CELL_PROLIFERATION
KYNG_DNA_DAMAGE_BY_GAMMA_AND_UV_RADIATION, KYNG_DNA_DAMAGE_BY_GAMMA_AND_UV_RADIATION
PACHER_TARGETS_OF_IGF1_AND_IGF2_UP, PACHER_TARGETS_OF_IGF1_AND_IGF2_UP
GO_REGULATION_OF_TRANSLATIONAL_FIDELITY, GO_REGULATION_OF_TRANSLATIONAL_FIDELITY
GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN
GO_DEVELOPMENTAL_PROGRAMMED_CELL_DEATH, GO_DEVELOPMENTAL_PROGRAMMED_CELL_DEATH
GO_OLIGOSACCHARIDE_METABOLIC_PROCESS, GO_OLIGOSACCHARIDE_METABOLIC_PROCESS
GO_PROTEIN_O_LINKED_MANNOSYLATION, GO_PROTEIN_O_LINKED_MANNOSYLATION