

AY15_LCMV_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY15_LCMV_EFFECTOR_CD8_TCELL_UP

GSE41867_NAIVE_VS_DAY30_LCMV_ARMSTRONG_MEMORY_CD8_TCELL_DN, GSE41867_NAIVE_VS_DAY30_LCMV_ARMSTRONG_MEMORY_CD8_TCELL_DN
GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_DN, GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_DN
GSE17721_CTRL_VS_PAM3CSK4_1H_BMDC_UP, GSE17721_CTRL_VS_PAM3CSK4_1H_BMDC_UP
GSE36009_WT_VS_NLRP10_KO_DC_UP, GSE36009_WT_VS_NLRP10_KO_DC_UP
GSE40274_HELIOS_VS_FOXP3_AND_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_HELIOS_VS_FOXP3_AND_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE14308_TH1_VS_NATURAL_TREG_UP, GSE14308_TH1_VS_NATURAL_TREG_UP
GSE16522_MEMORY_VS_NAIVE_ANTI_CD3CD28_STIM_CD8_TCELL_DN, GSE16522_MEMORY_VS_NAIVE_ANTI_CD3CD28_STIM_CD8_TCELL_DN
GSE21670_TGFB_VS_TGFB_AND_IL6_TREATED_STAT3_KO_CD4_TCELL_UP, GSE21670_TGFB_VS_TGFB_AND_IL6_TREATED_STAT3_KO_CD4_TCELL_UP
GSE32423_CTRL_VS_IL4_MEMORY_CD8_TCELL_UP, GSE32423_CTRL_VS_IL4_MEMORY_CD8_TCELL_UP
GSE42724_B1_BCELL_VS_PLASMABLAST_UP, GSE42724_B1_BCELL_VS_PLASMABLAST_UP
GSE1448_ANTI_VALPHA2_VS_VBETA5_DP_THYMOCYTE_DN, GSE1448_ANTI_VALPHA2_VS_VBETA5_DP_THYMOCYTE_DN
GSE41867_NAIVE_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP
GSE6092_UNSTIM_VS_IFNG_STIM_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_UP, GSE6092_UNSTIM_VS_IFNG_STIM_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_UP
GO_SOMATIC_DIVERSIFICATION_OF_IMMUNOGLOBULINS, GO_SOMATIC_DIVERSIFICATION_OF_IMMUNOGLOBULINS
GO_MISMATCH_REPAIR, GO_MISMATCH_REPAIR
HALLMARK_CHOLESTEROL_HOMEOSTASIS, HALLMARK_CHOLESTEROL_HOMEOSTASIS
GO_PROTEIN_N_LINKED_GLYCOSYLATION, GO_PROTEIN_N_LINKED_GLYCOSYLATION
KEGG_AMINO_SUGAR_AND_NUCLEOTIDE_SUGAR_METABOLISM, KEGG_AMINO_SUGAR_AND_NUCLEOTIDE_SUGAR_METABOLISM
GO_HISTONE_DEACETYLASE_COMPLEX, GO_HISTONE_DEACETYLASE_COMPLEX
GO_MICROBODY_PART, GO_MICROBODY_PART
GSE360_DC_VS_MAC_L_MAJOR_DN, GSE360_DC_VS_MAC_L_MAJOR_DN
RIZKI_TUMOR_INVASIVENESS_2D_DN, RIZKI_TUMOR_INVASIVENESS_2D_DN
GO_PROTEIN_DEACETYLASE_ACTIVITY, GO_PROTEIN_DEACETYLASE_ACTIVITY
BIOCARTA_CARM_ER_PATHWAY, BIOCARTA_CARM_ER_PATHWAY
GO_MISMATCH_REPAIR_COMPLEX, GO_MISMATCH_REPAIR_COMPLEX
BIOCARTA_INTEGRIN_PATHWAY, BIOCARTA_INTEGRIN_PATHWAY
GO_NUCLEAR_UBIQUITIN_LIGASE_COMPLEX, GO_NUCLEAR_UBIQUITIN_LIGASE_COMPLEX
GO_DEACETYLASE_ACTIVITY, GO_DEACETYLASE_ACTIVITY
GO_PROTEIN_K11_LINKED_UBIQUITINATION, GO_PROTEIN_K11_LINKED_UBIQUITINATION
GO_ANAPHASE_PROMOTING_COMPLEX, GO_ANAPHASE_PROMOTING_COMPLEX
GO_PEROXISOMAL_TRANSPORT, GO_PEROXISOMAL_TRANSPORT
GO_INTRACILIARY_TRANSPORT_PARTICLE_B, GO_INTRACILIARY_TRANSPORT_PARTICLE_B
GO_BETA_TUBULIN_BINDING, GO_BETA_TUBULIN_BINDING
GO_MICROBODY_MEMBRANE, GO_MICROBODY_MEMBRANE
GO_ALCOHOL_BIOSYNTHETIC_PROCESS, GO_ALCOHOL_BIOSYNTHETIC_PROCESS
BIOCARTA_MET_PATHWAY, BIOCARTA_MET_PATHWAY
BUCKANOVICH_T_LYMPHOCYTE_HOMING_ON_TUMOR_DN, BUCKANOVICH_T_LYMPHOCYTE_HOMING_ON_TUMOR_DN
REACTOME_PHOSPHORYLATION_OF_THE_APC_C, REACTOME_PHOSPHORYLATION_OF_THE_APC_C
GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_UP, GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_UP
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
SILIGAN_TARGETS_OF_EWS_FLI1_FUSION_DN, SILIGAN_TARGETS_OF_EWS_FLI1_FUSION_DN
GSE19888_ADENOSINE_A3R_INH_VS_ACT_WITH_INHIBITOR_PRETREATMENT_IN_MAST_CELL_UP, GSE19888_ADENOSINE_A3R_INH_VS_ACT_WITH_INHIBITOR_PRETREATMENT_IN_MAST_CELL_UP
GO_NAD_DEPENDENT_PROTEIN_DEACETYLASE_ACTIVITY, GO_NAD_DEPENDENT_PROTEIN_DEACETYLASE_ACTIVITY
WAESCH_ANAPHASE_PROMOTING_COMPLEX, WAESCH_ANAPHASE_PROMOTING_COMPLEX
GO_MISMATCHED_DNA_BINDING, GO_MISMATCHED_DNA_BINDING
NADLER_HYPERGLYCEMIA_AT_OBESITY, NADLER_HYPERGLYCEMIA_AT_OBESITY
TUOMISTO_TUMOR_SUPPRESSION_BY_COL13A1_DN, TUOMISTO_TUMOR_SUPPRESSION_BY_COL13A1_DN
CHEN_LIVER_METABOLISM_QTL_CIS, CHEN_LIVER_METABOLISM_QTL_CIS
REACTOME_PPARA_ACTIVATES_GENE_EXPRESSION, REACTOME_PPARA_ACTIVATES_GENE_EXPRESSION
GO_REGULATION_OF_MITOCHONDRIAL_TRANSLATION, GO_REGULATION_OF_MITOCHONDRIAL_TRANSLATION
GO_RESPONSE_TO_MISFOLDED_PROTEIN, GO_RESPONSE_TO_MISFOLDED_PROTEIN
GO_PHOSPHATIDYLCHOLINE_BIOSYNTHETIC_PROCESS, GO_PHOSPHATIDYLCHOLINE_BIOSYNTHETIC_PROCESS
GO_HISTONE_DEACETYLASE_ACTIVITY_H3_K14_SPECIFIC_, GO_HISTONE_DEACETYLASE_ACTIVITY_H3_K14_SPECIFIC_
GO_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEIN_CATABOLIC_PROCESS, GO_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEIN_CATABOLIC_PROCESS
GO_HISTONE_H3_DEACETYLATION, GO_HISTONE_H3_DEACETYLATION
GO_REGULATION_OF_HORMONE_METABOLIC_PROCESS, GO_REGULATION_OF_HORMONE_METABOLIC_PROCESS
GO_REGULATION_OF_HORMONE_BIOSYNTHETIC_PROCESS, GO_REGULATION_OF_HORMONE_BIOSYNTHETIC_PROCESS
GO_NUCLEOSIDE_BISPHOSPHATE_METABOLIC_PROCESS, GO_NUCLEOSIDE_BISPHOSPHATE_METABOLIC_PROCESS
GO_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY, GO_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY
KAMIKUBO_MYELOID_MN1_NETWORK, KAMIKUBO_MYELOID_MN1_NETWORK
GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_PDC_DN, GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_PDC_DN