

ABOLIC_PROCESS, GO_TRNA_METABOLIC_PROCESS

GSE43863_NAIVE_VS_MEMORY_LY6C_INT_CXCR5POS_CD4_TCELL_D150_LCMV_DN, GSE43863_NAIVE_VS_MEMORY_LY6C_INT_CXCR5POS_CD4_TCELL_D150_LCMV_DN
GSE37532_TREG_VS_TCONV_PPARG_KO_CD4_TCELL_FROM_LN_DN, GSE37532_TREG_VS_TCONV_PPARG_KO_CD4_TCELL_FROM_LN_DN
GSE6674_ANTI_IGM_VS_PL2_3_STIM_BCELL_DN, GSE6674_ANTI_IGM_VS_PL2_3_STIM_BCELL_DN
GSE46606_UNSTIM_VS_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_DN, GSE46606_UNSTIM_VS_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_DN
GSE6674_UNSTIM_VS_ANTI_IGM_AND_CPG_STIM_BCELL_DN, GSE6674_UNSTIM_VS_ANTI_IGM_AND_CPG_STIM_BCELL_DN
GO_NUCLEASE_ACTIVITY, GO_NUCLEASE_ACTIVITY
GO_TRNA_PROCESSING, GO_TRNA_PROCESSING
GSE27786_LIN_NEG_VS_MONO_MAC_UP, GSE27786_LIN_NEG_VS_MONO_MAC_UP
GO_RNA_MODIFICATION, GO_RNA_MODIFICATION
GSE4984_GALECTIN1_VS_VEHICLE_CTRL_TREATED_DC_DN, GSE4984_GALECTIN1_VS_VEHICLE_CTRL_TREATED_DC_DN
GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS
GSE17974_2H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_UP, GSE17974_2H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_UP
GO_S_ADENOSYLMETHIONINE_DEPENDENT_METHYLTRANSFERASE_ACTIVITY, GO_S_ADENOSYLMETHIONINE_DEPENDENT_METHYLTRANSFERASE_ACTIVITY
MYC_UP.V1_UP, MYC_UP.V1_UP
RB_P107_DN.V1_DN, RB_P107_DN.V1_DN
GO_NUCLEOLAR_PART, GO_NUCLEOLAR_PART
GSE25087_TREG_VS_TCONV_FETUS_DN, GSE25087_TREG_VS_TCONV_FETUS_DN
GO_ENDONUCLEASE_ACTIVITY, GO_ENDONUCLEASE_ACTIVITY
GO_RIBONUCLEASE_ACTIVITY, GO_RIBONUCLEASE_ACTIVITY
GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_DN, GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_DN
GSE37301_MULTIPOTENT_PROGENITOR_VS_LYMPHOID_PRIMED_MPP_UP, GSE37301_MULTIPOTENT_PROGENITOR_VS_LYMPHOID_PRIMED_MPP_UP
GO_RNA_METHYLTRANSFERASE_ACTIVITY, GO_RNA_METHYLTRANSFERASE_ACTIVITY
GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_EXONUCLEOLYTIC, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_EXONUCLEOLYTIC
REACTOME_UNFOLDED_PROTEIN_RESPONSE, REACTOME_UNFOLDED_PROTEIN_RESPONSE
GSE34217_MIR17_92_OVEREXPRESS_VS_WT_ACT_CD8_TCELL_DN, GSE34217_MIR17_92_OVEREXPRESS_VS_WT_ACT_CD8_TCELL_DN
GSE3982_DC_VS_MAC_LPS_STIM_UP, GSE3982_DC_VS_MAC_LPS_STIM_UP
GSE4748_CYANOBACTERIUM_LPSLIKE_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP, GSE4748_CYANOBACTERIUM_LPSLIKE_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_S
GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD4_TCELL_UP, GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD4_TCELL_UP
GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_UP, GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_UP
GO_N_METHYLTRANSFERASE_ACTIVITY, GO_N_METHYLTRANSFERASE_ACTIVITY
GO_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS, GO_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS
GO_NUCLEAR_TRANSCRIPTION_FACTOR_COMPLEX, GO_NUCLEAR_TRANSCRIPTION_FACTOR_COMPLEX
GO_TRNA_BINDING, GO_TRNA_BINDING
GO_TRANSCRIPTION_FROM_RNA_POLYMERASE_III_PROMOTER, GO_TRANSCRIPTION_FROM_RNA_POLYMERASE_III_PROMOTER
GO_INTRAMOLECULAR_TRANSFERASE_ACTIVITY, GO_INTRAMOLECULAR_TRANSFERASE_ACTIVITY
GO_CLEAVAGE_INVOLVED_IN_RRNA_PROCESSING, GO_CLEAVAGE_INVOLVED_IN_RRNA_PROCESSING
GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_ENDONUCLEOLYTIC, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_ENDONUCLEOLYTIC
REACTOME_PERK_REGULATED_GENE_EXPRESSION, REACTOME_PERK_REGULATED_GENE_EXPRESSION
REACTOME_ACTIVATION_OF_GENES_BY_ATF4, REACTOME_ACTIVATION_OF_GENES_BY_ATF4
GSE22886_IGA_VS_IGM_MEMORY_BCELL_UP, GSE22886_IGA_VS_IGM_MEMORY_BCELL_UP
GO_SNRNA_PROCESSING, GO_SNRNA_PROCESSING
GSE17721_0.5H_VS_4H_POLYIC_BMDC_UP, GSE17721_0.5H_VS_4H_POLYIC_BMDC_UP
GO_RNA_POLYADENYLATION, GO_RNA_POLYADENYLATION
REACTOME_DESTABILIZATION_OF_MRNA_BY_KSRP, REACTOME_DESTABILIZATION_OF_MRNA_BY_KSRP
GO_RRNA_MODIFICATION, GO_RRNA_MODIFICATION
GO_MITOCHONDRIAL_GENOME_MAINTENANCE, GO_MITOCHONDRIAL_GENOME_MAINTENANCE
GO_CARBON_NITROGEN_LIGASE_ACTIVITY_WITH_Glutamine_AS_AMIDO_N_DONOR, GO_CARBON_NITROGEN_LIGASE_ACTIVITY_WITH_Glutamine_AS_AMIDO_N_DONOR
GO_ADENYLYLTRANSFERASE_ACTIVITY, GO_ADENYLYLTRANSFERASE_ACTIVITY
GO_PSEUDOURIDINE_SYNTHASE_ACTIVITY, GO_PSEUDOURIDINE_SYNTHASE_ACTIVITY
IRF2_01, IRF2_01
GO_MRNA_CLEAVAGE_FACTOR_COMPLEX, GO_MRNA_CLEAVAGE_FACTOR_COMPLEX
chr14q12, chr14q12
KRIGE_AMINO_ACID_DEPRIVATION, KRIGE_AMINO_ACID_DEPRIVATION
CAFFAREL_RESPONSE_TO_THC_8HR_5_UP, CAFFAREL_RESPONSE_TO_THC_8HR_5_UP
GO_RNA_SPLICING_VIA_ENDONUCLEOLYTIC_CLEAVAGE_AND_LIGATION, GO_RNA_SPLICING_VIA_ENDONUCLEOLYTIC_CLEAVAGE_AND_LIGATION
GO_AMINOACYL_TRNA_EDITING_ACTIVITY, GO_AMINOACYL_TRNA_EDITING_ACTIVITY
GO_MRNA_CLEAVAGE_AND_POLYADENYLATION_SPECIFICITY_FACTOR_COMPLEX, GO_MRNA_CLEAVAGE_AND_POLYADENYLATION_SPECIFICITY_FACTOR_COMPLEX
GO_ENDORIBONUCLEASE_COMPLEX, GO_ENDORIBONUCLEASE_COMPLEX
REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION, REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BLACK_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BLACK_DN