
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_18H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_18H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_14H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_14H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_UP
GSE14000_UNSTIM_VS_4H_LPS_DC_UP, GSE14000_UNSTIM_VS_4H_LPS_DC_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_6H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_6H_UP
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_10H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_10H_UP
GSE40666_UNTREATED_VS_IFNA_STIM_CD8_TCELL_90MIN_DN, GSE40666_UNTREATED_VS_IFNA_STIM_CD8_TCELL_90MIN_DN
GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_DN, GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_DN
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_6H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_6H_UP
GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_6H_DN, GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_6H_DN
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_18H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_18H_UP
GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_DN
GSE2706_UNSTIM_VS_2H_R848_DC_UP, GSE2706_UNSTIM_VS_2H_R848_DC_UP
GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_BCELL_DN, GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_BCELL_DN
GSE46606_IRF4MID_VS_WT_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_UP, GSE46606_IRF4MID_VS_WT_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_UP
GSE7348_UNSTIM_VS_LPS_STIM_MACROPHAGE_UP, GSE7348_UNSTIM_VS_LPS_STIM_MACROPHAGE_UP
GSE2706_UNSTIM_VS_2H_LPS_AND_R848_DC_UP, GSE2706_UNSTIM_VS_2H_LPS_AND_R848_DC_UP
GSE2706_UNSTIM_VS_2H_LPS_DC_UP, GSE2706_UNSTIM_VS_2H_LPS_DC_UP
GO_RNA_METHYLATION, GO_RNA_METHYLATION
BROWNE_HCMV_INFECTION_6HR_DN, BROWNE_HCMV_INFECTION_6HR_DN
GSE40685_NAIVE_CD4_TCELL_VS_TREG_UP, GSE40685_NAIVE_CD4_TCELL_VS_TREG_UP
GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS
GO_RNA_METHYLTRANSFERASE_ACTIVITY, GO_RNA_METHYLTRANSFERASE_ACTIVITY
GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_DC_DN, GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_DC_DN
GO_CULLIN_RING_UBIQUITIN_LIGASE_COMPLEX, GO_CULLIN_RING_UBIQUITIN_LIGASE_COMPLEX
GSE26928_NAIVE_VS_CXCR5_POS_CD4_TCELL_UP, GSE26928_NAIVE_VS_CXCR5_POS_CD4_TCELL_UP
GO_TRANSCRIPTION_FROM_RNA_POLYMERASE_III_PROMOTER, GO_TRANSCRIPTION_FROM_RNA_POLYMERASE_III_PROMOTER
GO_RIBONUCLEASE_ACTIVITY, GO_RIBONUCLEASE_ACTIVITY
GO_EXONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS, GO_EXONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODU
GOTTWEIN_TARGETS_OF_KSHV_MIR_K12_11, GOTTWEIN_TARGETS_OF_KSHV_MIR_K12_11
GSE4748_CYANOBACTERIUM_LPSLIKE_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP, GSE4748_CYANOBACTERIUM_LPSLIKE_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP
ESC_J1_UP_LATE.V1_DN, ESC_J1_UP_LATE.V1_DN
GSE3691_IFN_PRODUCING_KILLER_DC_VS_CONVENTIONAL_DC_SPLEEN_DN, GSE3691_IFN_PRODUCING_KILLER_DC_VS_CONVENTIONAL_DC_SPLEEN_DN
REACTOME_PI_METABOLISM, REACTOME_PI_METABOLISM
GSE2585_THYMIC_DC_VS_MTEC_DN, GSE2585_THYMIC_DC_VS_MTEC_DN
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN
GAL_LEUKEMIC_STEM_CELL_UP, GAL_LEUKEMIC_STEM_CELL_UP
REACTOME_RNA_POL_III_TRANSCRIPTION, REACTOME_RNA_POL_III_TRANSCRIPTION
GO_NCRNA_CATABOLIC_PROCESS, GO_NCRNA_CATABOLIC_PROCESS
BIOCARTA_BCELLSURVIVAL_PATHWAY, BIOCARTA_BCELLSURVIVAL_PATHWAY
GO_MEDIATOR_COMPLEX, GO_MEDIATOR_COMPLEX
GO_CUL4_RING_E3_UBIQUITIN_LIGASE_COMPLEX, GO_CUL4_RING_E3_UBIQUITIN_LIGASE_COMPLEX
STAT5A_02, STAT5A_02
REACTOME_INTERFERON_GAMMA_SIGNALING, REACTOME_INTERFERON_GAMMA_SIGNALING
GO_DNA_MODIFICATION, GO_DNA_MODIFICATION
GSE45365_NK_CELL_VS_BCELL_DN, GSE45365_NK_CELL_VS_BCELL_DN
REACTOME_DESTABILIZATION_OF_MRNA_BY_KSRP, REACTOME_DESTABILIZATION_OF_MRNA_BY_KSRP
GO_RRNA_CATABOLIC_PROCESS, GO_RRNA_CATABOLIC_PROCESS
REACTOME_DESTABILIZATION_OF_MRNA_BY_BRF1, REACTOME_DESTABILIZATION_OF_MRNA_BY_BRF1
REACTOME_DESTABILIZATION_OF_MRNA_BY_TRISTETRAPROLIN_TTP, REACTOME_DESTABILIZATION_OF_MRNA_BY_TRISTETRAPROLIN_TTP
GSE20366_EX_VIVO_VS_DEC205_CONVERSION_NAIVE_CD4_TCELL_DN, GSE20366_EX_VIVO_VS_DEC205_CONVERSION_NAIVE_CD4_TCELL_DN
LIU_IL13_PRIMING_MODEL, LIU_IL13_PRIMING_MODEL
REACTOME_RNA_POL_III_TRANSCRIPTION_INITIATION_FROM_TYPE_2_PROMOTER, REACTOME_RNA_POL_III_TRANSCRIPTION_INITIATION_FROM_TYPE_2_PROMOTER
GO_TRANSFERASE_ACTIVITY_TRANSFERRING_PENTOSYL_GROUPS, GO_TRANSFERASE_ACTIVITY_TRANSFERRING_PENTOSYL_GROUPS
REACTOME_MRNA_DECAY_BY_3_TO_5_EXORIBONUCLEASE, REACTOME_MRNA_DECAY_BY_3_TO_5_EXORIBONUCLEASE
GSE369_PRE_VS_POST_IL6_INJECTION_IFNG_WT_LIVER_UP, GSE369_PRE_VS_POST_IL6_INJECTION_IFNG_WT_LIVER_UP
GO_RNA_SPLICING_VIA_ENDONUCLEOLYTIC_CLEAVAGE_AND_LIGATION, GO_RNA_SPLICING_VIA_ENDONUCLEOLYTIC_CLEAVAGE_AND_LIGATION
GO_RIBOSOMAL_LARGE_SUBUNIT_ASSEMBLY, GO_RIBOSOMAL_LARGE_SUBUNIT_ASSEMBLY
GO_REGULATION_OF_INTERFERON_BETA_PRODUCTION, GO_REGULATION_OF_INTERFERON_BETA_PRODUCTION
PID_PDGFRA_PATHWAY, PID_PDGFRA_PATHWAY
REACTOME_RNA_POL_III_CHAIN_ELONGATION, REACTOME_RNA_POL_III_CHAIN_ELONGATION
GO_DNA_DIRECTED_RNA_POLYMERASE_III_COMPLEX, GO_DNA_DIRECTED_RNA_POLYMERASE_III_COMPLEX
REACTOME_RNA_POL_III_TRANSCRIPTION_TERMINATION, REACTOME_RNA_POL_III_TRANSCRIPTION_TERMINATION
GO_RRNA_TRANSCRIPTION, GO_RRNA_TRANSCRIPTION
CYCLIN_D1_UP.V1_DN, CYCLIN_D1_UP.V1_DN
GO_PROTEIN_SERINE_THREONINE_TYROSINE_KINASE_ACTIVITY, GO_PROTEIN_SERINE_THREONINE_TYROSINE_KINASE_ACTIVITY
GO_PHOSPHOLIPID_CATABOLIC_PROCESS, GO_PHOSPHOLIPID_CATABOLIC_PROCESS
chr6q23, chr6q23
GO_O_METHYLTRANSFERASE_ACTIVITY, GO_O_METHYLTRANSFERASE_ACTIVITY
KEGG_CYTOSOLIC_DNA_SENSING_PATHWAY, KEGG_CYTOSOLIC_DNA_SENSING_PATHWAY
BIOCARTA_NGF_PATHWAY, BIOCARTA_NGF_PATHWAY
GSE17721_CTRL_VS_CPG_24H_BMDC_UP, GSE17721_CTRL_VS_CPG_24H_BMDC_UP
WORSCHECH_TUMOR_EVASION_AND_TOLEROGENICITY_DN, WORSCHECH_TUMOR_EVASION_AND_TOLEROGENICITY_DN
GO_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS_VIA_SOMATIC_MUTATION, GO_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS_VIA_SOMATIC_MUTATION
GO_HETEROCHROMATIN_ORGANIZATION, GO_HETEROCHROMATIN_ORGANIZATION
_NEWCASTLE_VIRUS_DC_10H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_10H_UP