

CATABOLIC_PROCESS, GO_TRNA_METABOLIC_PROCESS

GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_10H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_10H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_4H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_4H_UP
GO_RNA_MODIFICATION, GO_RNA_MODIFICATION
GO_NCRNA_TRANSCRIPTION, GO_NCRNA_TRANSCRIPTION
GO_TRNA_PROCESSING, GO_TRNA_PROCESSING
GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_DN
GSE17974_2H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_UP, GSE17974_2H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_UP
GSE4984_GALECTIN1_VS_VEHICLE_CTRL_TREATED_DC_DN, GSE4984_GALECTIN1_VS_VEHICLE_CTRL_TREATED_DC_DN
GO_TRNA_MODIFICATION, GO_TRNA_MODIFICATION
GSE6092_CTRL_VS_BORRELIA_BIRGDOFERI_INF_ENDOTHELIAL_CELL_UP, GSE6092_CTRL_VS_BORRELIA_BIRGDOFERI_INF_ENDOTHELIAL_CELL_UP
GO_S_ADENOSYLMETHIONINE_DEPENDENT_METHYLTRANSFERASE_ACTIVITY, GO_S_ADENOSYLMETHIONINE_DEPENDENT_METHYLTRANSFERASE_ACTIVITY
GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS
REACTOME_UNFOLDED_PROTEIN_RESPONSE, REACTOME_UNFOLDED_PROTEIN_RESPONSE
GO_AMINO_ACID_ACTIVATION, GO_AMINO_ACID_ACTIVATION
GO_RNA_METHYLATION, GO_RNA_METHYLATION
GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_DN, GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_DN
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_PRODUCING_5_P
MYC_UP.V1_UP, MYC_UP.V1_UP
GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP, GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP
KEGG_RNA_DEGRADATION, KEGG_RNA_DEGRADATION
GSE4748_CYANOBACTERIUM_LPSLIKE_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP, GSE4748_CYANOBACTERIUM_LPSLIKE_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP
GO_RNA_METHYLTRANSFERASE_ACTIVITY, GO_RNA_METHYLTRANSFERASE_ACTIVITY
GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_UP, GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_UP
GO_MATURATION_OF_5_8S_RRNA, GO_MATURATION_OF_5_8S_RRNA
GO_ISOMERASE_ACTIVITY, GO_ISOMERASE_ACTIVITY
GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_EXONUCLEOLYTIC, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_EXONUCLEOLYTIC
GSE29617_DAY3_VS_DAY7_TIV_FLU_VACCINE_PBMCDN, GSE29617_DAY3_VS_DAY7_TIV_FLU_VACCINE_PBMCDN
GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_MONOCYTE_UP, GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_MONOCYTE_UP
RB_P107_DN.V1_DN, RB_P107_DN.V1_DN
GSE37301_MULTIPOTENT_PROGENITOR_VS_LYMPHOID_PRIMED_MPP_UP, GSE37301_MULTIPOTENT_PROGENITOR_VS_LYMPHOID_PRIMED_MPP_UP
GO_N_METHYLTRANSFERASE_ACTIVITY, GO_N_METHYLTRANSFERASE_ACTIVITY
GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_BCELL_DN, GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_BCELL_DN
GSE17721_CTRL_VS_GARDIQUIMOD_6H_BMDC_DN, GSE17721_CTRL_VS_GARDIQUIMOD_6H_BMDC_DN
GO_NUCLEAR_TRANSCRIPTION_FACTOR_COMPLEX, GO_NUCLEAR_TRANSCRIPTION_FACTOR_COMPLEX
GO_RNA_POLYADENYLATION, GO_RNA_POLYADENYLATION
GSE17721_CTRL_VS_POLYIC_2H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_2H_BMDC_UP
GO_SNRNA_PROCESSING, GO_SNRNA_PROCESSING
GO_RIBONUCLEASE_ACTIVITY, GO_RIBONUCLEASE_ACTIVITY
KEGG_AMINOACYL_TRNA_BIOSYNTHESIS, KEGG_AMINOACYL_TRNA_BIOSYNTHESIS
GO_TRNA_METHYLATION, GO_TRNA_METHYLATION
GO_TRANSCRIPTION_FROM_RNA_POLYMERASE_III_PROMOTER, GO_TRANSCRIPTION_FROM_RNA_POLYMERASE_III_PROMOTER
BHATL_G2M_ARREST_BY_2METHOXYESTRADIOL_DN, BHATL_G2M_ARREST_BY_2METHOXYESTRADIOL_DN
GSE22886_IGA_VS_IGM_MEMORY_BCELL_UP, GSE22886_IGA_VS_IGM_MEMORY_BCELL_UP
GO_MATURATION_OF_5_8S_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA_, GO_MATURATION_OF_5_8S_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA_
GSE5542_IFNA_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_24H_UP, GSE5542_IFNA_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_24H_UP
GSE34217_MIR17_92_OVEREXPRESS_VS_WT_ACT_CD8_TCELL_DN, GSE34217_MIR17_92_OVEREXPRESS_VS_WT_ACT_CD8_TCELL_DN
GO_CLEAVAGE_INVOLVED_IN_RRNA_PROCESSING, GO_CLEAVAGE_INVOLVED_IN_RRNA_PROCESSING
REACTOME_DESTABILIZATION_OF_MRNA_BY_KSRP, REACTOME_DESTABILIZATION_OF_MRNA_BY_KSRP
GO_TRNA_METHYLTRANSFERASE_ACTIVITY, GO_TRNA_METHYLTRANSFERASE_ACTIVITY
GO_NCRNA_3_END_PROCESSING, GO_NCRNA_3_END_PROCESSING
GO_RRNA_TRANSCRIPTION, GO_RRNA_TRANSCRIPTION
GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_ENDONUCLEOLYTIC, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_ENDONUCLEOLYTIC
GO_RNA_SPLICING_VIA_ENDONUCLEOLYTIC_CLEAVAGE_AND_LIGATION, GO_RNA_SPLICING_VIA_ENDONUCLEOLYTIC_CLEAVAGE_AND_LIGATION
REACTOME_RNA_POL_III_TRANSCRIPTION, REACTOME_RNA_POL_III_TRANSCRIPTION
GO_TRNA_THREONYLCARBAMOYLADENOSINE_METABOLIC_PROCESS, GO_TRNA_THREONYLCARBAMOYLADENOSINE_METABOLIC_PROCESS
REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION, REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION
GO_RRNA_3_END_PROCESSING, GO_RRNA_3_END_PROCESSING
KEGG_SELENOAMINO_ACID_METABOLISM, KEGG_SELENOAMINO_ACID_METABOLISM
GO_ADENYLYLTRANSFERASE_ACTIVITY, GO_ADENYLYLTRANSFERASE_ACTIVITY
GO_MRNA_CLEAVAGE_FACTOR_COMPLEX, GO_MRNA_CLEAVAGE_FACTOR_COMPLEX
KRIGE_AMINO_ACID_DEPRIVATION, KRIGE_AMINO_ACID_DEPRIVATION
MODULE_322, MODULE_322
REACTOME_ACTIVATION_OF_GENES_BY_ATF4, REACTOME_ACTIVATION_OF_GENES_BY_ATF4
REACTOME_RNA_POL_III_TRANSCRIPTION_INITIATION_FROM_TYPE_2_PROMOTER, REACTOME_RNA_POL_III_TRANSCRIPTION_INITIATION_FROM_TYPE_2_PROMOTER
GO_RRNA_CATABOLIC_PROCESS, GO_RRNA_CATABOLIC_PROCESS
GO_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_3_PHOSPHOMONOESTERS, GO_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_3_P
KEGG_HISTIDINE_METABOLISM, KEGG_HISTIDINE_METABOLISM
KEGG_TYROSINE_METABOLISM, KEGG_TYROSINE_METABOLISM
WAKABAYASHI_ADIPOGENESIS_PPARG_BOUND_36HR, WAKABAYASHI_ADIPOGENESIS_PPARG_BOUND_36HR
chr1q25, chr1q25
GSE3691_IFN_PRODUCING_KILLER_DC_VS_CONVENTIONAL_DC_SPLEEN_DN, GSE3691_IFN_PRODUCING_KILLER_DC_VS_CONVENTIONAL_DC_SPLEEN_DN
GSE2585_THYMIC_MACROPHAGE_VS_MTEC_DN, GSE2585_THYMIC_MACROPHAGE_VS_MTEC_DN
CAFFAREL_RESPONSE_TO_THC_8HR_5_UP, CAFFAREL_RESPONSE_TO_THC_8HR_5_UP
GO_PSEUDOURIDINE_SYNTHASE_ACTIVITY, GO_PSEUDOURIDINE_SYNTHASE_ACTIVITY
REACTOME_MRNA_DECAY_BY_3_TO_5_EXORIBONUCLEASE, REACTOME_MRNA_DECAY_BY_3_TO_5_EXORIBONUCLEASE
LASTOWSKA_COAMPLIFIED_WITH_MYCN, LASTOWSKA_COAMPLIFIED_WITH_MYCN
chr1q41, chr1q41
ZWANG_EGF_PERSISTENTLY_DN, ZWANG_EGF_PERSISTENTLY_DN