_RNA_VS_MRNA_4H_LPS_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_4H_LPS_DC_UP

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GSE14000_TRANSLATED_RNA_VS_MRNA_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_DC_UP
 GSE14000_UNSTIM_VS_4H_LPS_DC_UP, GSE14000_UNSTIM_VS_4H_LPS_DC_UP
GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_6H_DN, GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_6H_DN
GSE14000_4H_VS_16H_LPS_DC_DN, GSE14000_4H_VS_16H_LPS_DC_DN
GSE17721_0.5H_VS_4H_POLYIC_BMDC_DN, GSE17721_0.5H_VS_4H_POLYIC_BMDC_DN
GSE18281_MEDULLARY_THYMOCYTE_VS_WHOLE_MEDULLA_THYMUS_DN, GSE18281_MEDULLARY_THYMOCYTE_VS_WHOLE_MEDULLA_THYMUS_DN
GSE17186 MEMORY VS CD21LOW TRANSITIONAL BCELL DN, GSE17186 MEMORY VS CD21LOW TRANSITIONAL BCELL DN
GSE17721_LPS_VS_POLYIC_6H_BMDC_UP, GSE17721_LPS_VS_POLYIC_6H_BMDC_UP
KEGG_RNA_DEGRADATION, KEGG_RNA_DEGRADATION
SEIDEN_ONCOGENESIS_BY_MET, SEIDEN_ONCOGENESIS_BY_MET
 GSE20366_EX_VIVO_VS_DEC205_CONVERSION_DN, GSE20366_EX_VIVO_VS_DEC205_CONVERSION_DN
GSE45365_WT_VS_IFNAR_KO_BCELL_UP, GSE45365_WT_VS_IFNAR_KO_BCELL_UP
 GO_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER
GO_MATURATION_OF_5_8S_RRNA, GO_MATURATION_OF_5_8S_RRNA
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_2H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_2H_UP
GO_RNA_POLYMERASE_BINDING, GO_RNA_POLYMERASE_BINDING
GO_BASAL_TRANSCRIPTION_MACHINERY_BINDING, GO_BASAL_TRANSCRIPTION_MACHINERY_BINDING
GO_EXOSOME_RNASE_COMPLEX_, GO_EXOSOME_RNASE_COMPLEX_
BRUINS_UVC_RESPONSE_MIDDLE, BRUINS_UVC_RESPONSE_MIDDLE
GO_RNA_CAPPING, GO_RNA_CAPPING
REACTOME_FORMATION_OF_RNA_POL_II_ELONGATION_COMPLEX_, REACTOME_FORMATION_OF_RNA_POL_II_ELONGATION_COMPLEX_
WANG METASTASIS OF BREAST CANCER ESR1_UP, WANG METASTASIS OF BREAST CANCER ESR1_UP
GO_RNA_POLYMERASE_CORE_ENZYME_BINDING, GO_RNA_POLYMERASE_CORE_ENZYME_BINDING
GO_REGULATION_OF_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY, GO_REGULATION_OF_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY
GO NUCLEAR EXOSOME RNASE COMPLEX, GO NUCLEAR EXOSOME RNASE COMPLEX
GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
GSE5589_WT_VS_IL6_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_DN, GSE5589_WT_VS_IL6_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_DN
BIOCARTA_IGF1_PATHWAY, BIOCARTA_IGF1_PATHWAY
YANG_BREAST_CANCER_ESR1_BULK_UP, YANG_BREAST_CANCER_ESR1_BULK_UP
GO_NCRNA_CATABOLIC_PROCESS, GO_NCRNA_CATABOLIC_PROCESS
GO_HISTONE_H4_ACETYLATION, GO_HISTONE_H4_ACETYLATION
 GSE29949_CD8_NEG_DC_SPLEEN_VS_MONOCYTE_BONE_MARROW_DN, GSE29949_CD8_NEG_DC_SPLEEN_VS_MONOCYTE_BONE_MARROW_DN
GO_PEROXISOMAL_TRANSPORT, GO_PEROXISOMAL_TRANSPORT
EHLERS_ANEUPLOIDY_UP, EHLERS_ANEUPLOIDY_UP
 GSE19888_CTRL_VS_T_CELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_CTRL_VS_T_CELL_MEMBRANES_ACT_MAST_CELL_DN
HASLINGER_B_CLL_WITH_6Q21_DELETION, HASLINGER_B_CLL_WITH_6Q21_DELETION
GO_RNA_SURVEILLANCE, GO_RNA_SURVEILLANCE
BIOCARTA_PAR1_PATHWAY, BIOCARTA_PAR1_PATHWAY
GO_RRNA_CATABOLIC_PROCESS, GO_RRNA_CATABOLIC_PROCESS
GSE17721_CTRL_VS_CPG_4H_BMDC_DN, GSE17721_CTRL_VS_CPG_4H_BMDC_DN
MIDORIKAWA_AMPLIFIED_IN_LIVER_CANCER, MIDORIKAWA_AMPLIFIED_IN_LIVER_CANCER
GO_TRANSCRIPTION_FACTOR_ACTIVITY_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_BINDING, GO_TRANSCRIPTION_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTOR_FACTO
GO_SWI_SNF_COMPLEX, GO_SWI_SNF_COMPLEX
GO_REGULATION_OF_STRIATED_MUSCLE_CELL_APOPTOTIC_PROCESS, GO_REGULATION_OF_STRIATED_MUSCLE_CELL_APOPTOTIC_PROCESS
GO_RRNA_TRANSCRIPTION, GO_RRNA_TRANSCRIPTION
GO REGULATION OF INTRACELLULAR ESTROGEN RECEPTOR SIGNALING PATHWAY, GO REGULATION OF INTRACELLULAR ESTROGEN RECEPTOR SIGNALING PATHWAY
GO_ADHERENS_JUNCTION_ASSEMBLY, GO_ADHERENS_JUNCTION_ASSEMBLY
GO_RNA_POLYMERASE_II_TRANSCRIPTION_COFACTOR_ACTIVITY, GO_RNA_POLYMERASE_II_TRANSCRIPTION_COFACTOR_ACTIVITY
MARCINIAK ER STRESS RESPONSE VIA CHOP, MARCINIAK ER STRESS RESPONSE VIA CHOP
WAKABAYASHI_ADIPOGENESIS_PPARG_BOUND_36HR, WAKABAYASHI_ADIPOGENESIS_PPARG_BOUND_36HR
EVI1 01, EVI1 01
GO_NEGATIVE_REGULATION_OF_STRIATED_MUSCLE_CELL_APOPTOTIC_PROCESS, GO_NEGATIVE_REGULATION_OF_STRIATED_MUSCLE_CELL_APOPTOTIC_PROCESS
GO_VENTRICULAR_SYSTEM_DEVELOPMENT, GO_VENTRICULAR_SYSTEM_DEVELOPMENT
BURTON_ADIPOGENESIS_10, BURTON_ADIPOGENESIS_10
DAIRKEE_TERT_TARGETS_DN, DAIRKEE_TERT_TARGETS_DN
WILSON PROTEASES AT TUMOR BONE INTERFACE UP, WILSON PROTEASES AT TUMOR BONE INTERFACE UP
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