\_RNA\_VS\_MRNA\_4H\_LPS\_DC\_UP, GSE14000\_TRANSLATED\_RNA\_VS\_MRNA\_4H\_LPS\_DC\_UP

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GSE33162_HDAC3_KO_VS_HDAC3_KO_4H_LPS_STIM_MACROPHAGE_UP, GSE33162_HDAC3_KO_VS_HDAC3_KO_4H_LPS_STIM_MACROPHAGE_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
GSE38696_LIGHT_ZONE_VS_DARK_ZONE_BCELL_UP, GSE38696_LIGHT_ZONE_VS_DARK_ZONE_BCELL_UP
GSE17186_MEMORY_VS_CD21LOW_TRANSITIONAL_BCELL_DN, GSE17186_MEMORY_VS_CD21LOW_TRANSITIONAL_BCELL_DN
GSE18281_MEDULLARY_THYMOCYTE_VS_WHOLE_MEDULLA_THYMUS_DN, GSE18281_MEDULLARY_THYMOCYTE_VS_WHOLE_MEDULLA_THYMUS_DN
KEGG_RNA_DEGRADATION, KEGG_RNA_DEGRADATION
GSE4142_GC_BCELL_VS_MEMORY_BCELL_DN, GSE4142_GC_BCELL_VS_MEMORY_BCELL_DN
GSE29949_CD8_NEG_DC_SPLEEN_VS_MONOCYTE_BONE_MARROW_DN, GSE29949_CD8_NEG_DC_SPLEEN_VS_MONOCYTE_BONE_MARROW_DN
GSE12484_HEALTHY_VS_PERIDONTITIS_NEUTROPHILS_DN, GSE12484_HEALTHY_VS_PERIDONTITIS_NEUTROPHILS_DN
GSE41867_NAIVE_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP
GSE20366_EX_VIVO_VS_DEC205_CONVERSION_DN, GSE20366_EX_VIVO_VS_DEC205_CONVERSION_DN
GSE17721_CTRL_VS_CPG_4H_BMDC_DN, GSE17721_CTRL_VS_CPG_4H_BMDC_DN
GO REGULATION OF INTRACELLULAR STEROID HORMONE RECEPTOR SIGNALING PATHWAY, GO REGULATION OF INTRACELLULAR STEROID HORMONE R
GO_MATURATION_OF_5_8S_RRNA, GO_MATURATION_OF_5_8S_RRNA
RB_P130_DN.V1_DN, RB_P130_DN.V1_DN
GO_BASAL_TRANSCRIPTION_MACHINERY_BINDING, GO_BASAL_TRANSCRIPTION_MACHINERY_BINDING
GSE15324_ELF4_KO_VS_WT_ACTIVATED_CD8_TCELL_DN, GSE15324_ELF4_KO_VS_WT_ACTIVATED_CD8_TCELL_DN
GSE36891_POLYIC_TLR3_VS_PAM_TLR2_STIM_PERITONEAL_MACROPHAGE_DN, GSE36891_POLYIC_TLR3_VS_PAM_TLR2_STIM_PERITONEAL_MACROPHAGE_DN
GO_EXOSOME_RNASE_COMPLEX_, GO_EXOSOME_RNASE_COMPLEX_
GO_RNA_POLYMERASE_CORE_ENZYME_BINDING, GO_RNA_POLYMERASE_CORE_ENZYME_BINDING
GO_ADA2_GCN5_ADA3_TRANSCRIPTION_ACTIVATOR_COMPLEX, GO_ADA2_GCN5_ADA3_TRANSCRIPTION_ACTIVATOR_COMPLEX
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_MAGENTA_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_MAGENTA_DN
GSE14000_TRANSLATED_RNA_VS_MRNA_16H_LPS_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_16H_LPS_DC_UP
YUAN_ZNF143_PARTNERS, YUAN_ZNF143_PARTNERS
MIZUSHIMA_AUTOPHAGOSOME_FORMATION, MIZUSHIMA_AUTOPHAGOSOME_FORMATION
GO_HISTONE_DEUBIQUITINATION, GO_HISTONE_DEUBIQUITINATION
GO NBAF COMPLEX, GO NBAF COMPLEX
PID_INSULIN_PATHWAY, PID_INSULIN_PATHWAY
GO_REGULATION_OF_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALING_PATHWAY, GO_REGULATION_OF_INTRACELLULAR_ESTROGEN_RECEPTOR_SIGNALIN
TURASHVILI_BREAST_NORMAL_DUCTAL_VS_LOBULAR_UP, TURASHVILI_BREAST_NORMAL_DUCTAL_VS_LOBULAR_UP
BIOCARTA_IGF1MTOR_PATHWAY, BIOCARTA_IGF1MTOR_PATHWAY
MODULE_192, MODULE_192
GSE5589_WT_VS_IL6_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_DN, GSE5589_WT_VS_IL6_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_DN
MODULE_279, MODULE_279
GO_RIBONUCLEOPROTEIN_COMPLEX_DISASSEMBLY, GO_RIBONUCLEOPROTEIN_COMPLEX_DISASSEMBLY
SHETH_LIVER_CANCER_VS_TXNIP_LOSS_PAM6, SHETH_LIVER_CANCER_VS_TXNIP_LOSS_PAM6
REACTOME_SYNTHESIS_OF_PIPS_AT_THE_EARLY_ENDOSOME_MEMBRANE, REACTOME_SYNTHESIS_OF_PIPS_AT_THE_EARLY_ENDOSOME_MEMBRANE
YANG_BREAST_CANCER_ESR1_BULK_UP, YANG_BREAST_CANCER_ESR1_BULK_UP
GSE19888_CTRL_VS_T_CELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_CTRL_VS_T_CELL_MEMBRANES_ACT_MAST_CELL_DN
GO_RRNA_CATABOLIC_PROCESS, GO_RRNA_CATABOLIC_PROCESS
MODULE_334, MODULE_334
PID_IGF1_PATHWAY, PID_IGF1_PATHWAY
SCHRAMM_INHBA_TARGETS_DN, SCHRAMM_INHBA_TARGETS_DN
PID_VEGFR1_PATHWAY, PID_VEGFR1_PATHWAY
SASSON_RESPONSE_TO_FORSKOLIN_UP, SASSON_RESPONSE_TO_FORSKOLIN_UP
BIOCARTA_INSULIN_PATHWAY, BIOCARTA_INSULIN_PATHWAY
CAGTCAC_MIR134, CAGTCAC_MIR134
MIDORIKAWA AMPLIFIED IN LIVER CANCER, MIDORIKAWA AMPLIFIED IN LIVER CANCER
EVI1_01, EVI1_01
GO_CELL_SUBSTRATE_ADHERENS_JUNCTION_ASSEMBLY, GO_CELL_SUBSTRATE_ADHERENS_JUNCTION_ASSEMBLY
GSE17721_4_VS_24H_CPG_BMDC_UP, GSE17721_4_VS_24H_CPG_BMDC_UP
GO CENTROSOME LOCALIZATION, GO CENTROSOME LOCALIZATION
BIOCARTA_PAR1_PATHWAY, BIOCARTA_PAR1_PATHWAY
GO_NEGATIVE_REGULATION_OF_STRIATED_MUSCLE_CELL_APOPTOTIC_PROCESS, GO_NEGATIVE_REGULATION_OF_STRIATED_MUSCLE_CELL_APOPTOTIC_PROCI
GO_RNA_SURVEILLANCE, GO_RNA_SURVEILLANCE
GO_POSITIVE_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_DEVELOPMENT, GO_POSITIVE_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_DEVELOPMENT
AGTTCTC MIR146A MIR146B, AGTTCTC MIR146A MIR146B
ZHAN_MULTIPLE_MYELOMA_MF_DN, ZHAN_MULTIPLE_MYELOMA_MF_DN
BIOCARTA_MYOSIN_PATHWAY, BIOCARTA_MYOSIN_PATHWAY
REACTOME_DOWNSTREAM_TCR_SIGNALING, REACTOME_DOWNSTREAM_TCR_SIGNALING
GO_MITOCHONDRION_DISTRIBUTION, GO_MITOCHONDRION_DISTRIBUTION
BIOCARTA_GSK3_PATHWAY, BIOCARTA_GSK3_PATHWAY
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