

TRANSCRIPTION, REACTOME_TRANSCRIPTION

GO_RNA_LOCALIZATION, GO_RNA_LOCALIZATION
GO_RIBONUCLEOPROTEIN_COMPLEX_LOCALIZATION, GO_RIBONUCLEOPROTEIN_COMPLEX_LOCALIZATION
GO_NUCLEOBASE_CONTAINING_COMPOUND_TRANSPORT, GO_NUCLEOBASE_CONTAINING_COMPOUND_TRANSPORT
GO_DNA_TEMPLATED_TRANSCRIPTION_TERMINATION, GO_DNA_TEMPLATED_TRANSCRIPTION_TERMINATION
GSE14000_4H_VS_16H_LPS_DC_DN, GSE14000_4H_VS_16H_LPS_DC_DN
GO_RNA_3_END_PROCESSING, GO_RNA_3_END_PROCESSING
GO_RNA_POLYMERASE_COMPLEX, GO_RNA_POLYMERASE_COMPLEX
REACTOME_RNA_POL_I_RNA_POL_III_AND_MITOCHONDRIAL_TRANSCRIPTION, REACTOME_RNA_POL_I_RNA_POL_III_AND_MITOCHONDRIAL_TRANSCRIPTION
REACTOME_RNA_POL_I_TRANSCRIPTION, REACTOME_RNA_POL_I_TRANSCRIPTION
GO_NUCLEOTIDYLTRANSFERASE_ACTIVITY, GO_NUCLEOTIDYLTRANSFERASE_ACTIVITY
GO_NCRNA_TRANSCRIPTION, GO_NCRNA_TRANSCRIPTION
GO_GENE_SILENCING_BY_RNA, GO_GENE_SILENCING_BY_RNA
GO_DNA_PACKAGING_COMPLEX, GO_DNA_PACKAGING_COMPLEX
GO_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GO_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
REACTOME_DEPOSITION_OF_NEW_CENPA_CONTAINING_NUCLEOSOMES_AT_THE_CENTROMERE, REACTOME_DEPOSITION_OF_NEW_CENPA_CONTAINING_NUCLEOSOMES_AT_THE_CENTROMERE
REACTOME_RNA_POL_I_PROMOTER_OPENING, REACTOME_RNA_POL_I_PROMOTER_OPENING
GO_DNA_TEMPLATED_TRANSCRIPTION_INITIATION, GO_DNA_TEMPLATED_TRANSCRIPTION_INITIATION
GSE41176_WT_VS_TAK1_KO_ANTL_IGM_STIM_BCELL_24H_DN, GSE41176_WT_VS_TAK1_KO_ANTL_IGM_STIM_BCELL_24H_DN
GO_SNRNA_METABOLIC_PROCESS, GO_SNRNA_METABOLIC_PROCESS
MORF_BMI1, MORF_BMI1
GSE32986_UNSTIM_VS_CURDLAN_LOWDOSE_STIM_DC_DN, GSE32986_UNSTIM_VS_CURDLAN_LOWDOSE_STIM_DC_DN
GO_DNA_DIRECTED_RNA_POLYMERASE_II_HOLOENZYME, GO_DNA_DIRECTED_RNA_POLYMERASE_II_HOLOENZYME
GO_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION, GO_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION
GSE3982_NEUTROPHIL_VS_EFF_MEMORY_CD4_TCELL_DN, GSE3982_NEUTROPHIL_VS_EFF_MEMORY_CD4_TCELL_DN
KRCTCNNNNMANAGC_UNKNOWN, KRCTCNNNNMANAGC_UNKNOWN
GO_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER
REACTOME_PACKAGING_OF_TELOMERE_ENDS, REACTOME_PACKAGING_OF_TELOMERE_ENDS
GO_PROTEIN_KINASE_COMPLEX, GO_PROTEIN_KINASE_COMPLEX
GSE12845_IGD_POS_BLOOD_VS_NAIVE_TONSIL_BCELL_UP, GSE12845_IGD_POS_BLOOD_VS_NAIVE_TONSIL_BCELL_UP
REACTOME_AMYLOIDS, REACTOME_AMYLOIDS
GO_POSITIVE_REGULATION_OF_VIRAL_PROCESS, GO_POSITIVE_REGULATION_OF_VIRAL_PROCESS
KEGG_SYSTEMIC_LUPUS_ERYTHEMATOSUS, KEGG_SYSTEMIC_LUPUS_ERYTHEMATOSUS
GSE33292_WT_VS_TCF1_KO_DN3_THYMOCYTE_UP, GSE33292_WT_VS_TCF1_KO_DN3_THYMOCYTE_UP
EGFR_UP.V1_DN, EGFR_UP.V1_DN
MORF_RFC5, MORF_RFC5
GO_DNA_REPLICATION_DEPENDENT_NUCLEOSOME_ORGANIZATION, GO_DNA_REPLICATION_DEPENDENT_NUCLEOSOME_ORGANIZATION
GO_CHROMATIN_SILENCING, GO_CHROMATIN_SILENCING
REACTOME_RNA_POL_II_PRE_TRANSCRIPTION_EVENTS, REACTOME_RNA_POL_II_PRE_TRANSCRIPTION_EVENTS
GO_POSITIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION, GO_POSITIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION
GO_CENTROMERE_COMPLEX_ASSEMBLY, GO_CENTROMERE_COMPLEX_ASSEMBLY
GO_PEPTIDE_N_ACETYLTTRANSFERASE_ACTIVITY, GO_PEPTIDE_N_ACETYLTTRANSFERASE_ACTIVITY
GO_ACETYLTTRANSFERASE_ACTIVITY, GO_ACETYLTTRANSFERASE_ACTIVITY
ACEVEDO_LIVER_CANCER_WITH_H3K9ME3_DN, ACEVEDO_LIVER_CANCER_WITH_H3K9ME3_DN
KOBAYASHI_EGFR_SIGNALING_24HR_UP, KOBAYASHI_EGFR_SIGNALING_24HR_UP
GO_TRANSCRIPTION_FROM_RNA_POLYMERASE_I_PROMOTER, GO_TRANSCRIPTION_FROM_RNA_POLYMERASE_I_PROMOTER
GGGNRMNNYCAT_UNKNOWN, GGGNRMNNYCAT_UNKNOWN
GO_HISTONE_EXCHANGE, GO_HISTONE_EXCHANGE
GO_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER
GSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_DN, GSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_DN
GO_N_ACETYLTTRANSFERASE_ACTIVITY, GO_N_ACETYLTTRANSFERASE_ACTIVITY
GO_TRANSCRIPTION_FROM_RNA_POLYMERASE_III_PROMOTER, GO_TRANSCRIPTION_FROM_RNA_POLYMERASE_III_PROMOTER
GO_POSITIVE_REGULATION_OF_PROTEIN_COMPLEX_ASSEMBLY, GO_POSITIVE_REGULATION_OF_PROTEIN_COMPLEX_ASSEMBLY
chr6p22, chr6p22
GO_CHROMATIN_SILENCING_AT_RDNA, GO_CHROMATIN_SILENCING_AT_RDNA
GO_TRANSCRIPTION_ELONGATION_FACTOR_COMPLEX, GO_TRANSCRIPTION_ELONGATION_FACTOR_COMPLEX
REACTOME_RNA_POL_II_TRANSCRIPTION_PRE_INITIATION_AND_PROMOTER_OPENING, REACTOME_RNA_POL_II_TRANSCRIPTION_PRE_INITIATION_AND_PROMOTER_OPENING
GSE360_CTRL_VS_B_MALAYI_HIGH_DOSE_MAC_UP, GSE360_CTRL_VS_B_MALAYI_HIGH_DOSE_MAC_UP
REACTOME_FORMATION_OF_RNA_POL_II_ELONGATION_COMPLEX, REACTOME_FORMATION_OF_RNA_POL_II_ELONGATION_COMPLEX
GSE25088_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DN, GSE25088_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DN
GO_HISTONE_MRNA_METABOLIC_PROCESS, GO_HISTONE_MRNA_METABOLIC_PROCESS
REACTOME_FORMATION_OF_THE_HIV1_EARLY_ELONGATION_COMPLEX, REACTOME_FORMATION_OF_THE_HIV1_EARLY_ELONGATION_COMPLEX
TTNNANAGCYR_UNKNOWN, TTTNNANAGCYR_UNKNOWN
REACTOME_REGULATORY_RNA_PATHWAYS, REACTOME_REGULATORY_RNA_PATHWAYS
GSE12845_IGD_POS_VS_NEG_BLOOD_BCELL_UP, GSE12845_IGD_POS_VS_NEG_BLOOD_BCELL_UP
GO_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY, GO_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY
GO_MLL1_2_COMPLEX, GO_MLL1_2_COMPLEX
GO_PRE_MRNA_BINDING, GO_PRE_MRNA_BINDING
GO_HISTONE_H3_ACETYLTATION, GO_HISTONE_H3_ACETYLTATION
REACTOME_MRNA_CAPPING, REACTOME_MRNA_CAPPING
REACTOME_RNA_POL_I_TRANSCRIPTION_TERMINATION, REACTOME_RNA_POL_I_TRANSCRIPTION_TERMINATION
GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_DUPLEX_UNWINDING, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_DUPLEX_UNWINDING
GO_EXON_EXON_JUNCTION_COMPLEX, GO_EXON_EXON_JUNCTION_COMPLEX
GO_RNA_POLYMERASE_BINDING, GO_RNA_POLYMERASE_BINDING
GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_LARGE_VS_TINY_DN, LIANG_HEMATOPOIESIS_STEM_CELL_NUMBER_LARGE_VS_TINY_DN
REACTOME_RNA_POL_I_TRANSCRIPTION_INITIATION, REACTOME_RNA_POL_I_TRANSCRIPTION_INITIATION
REACTOME_RNA_POL_III_TRANSCRIPTION, REACTOME_RNA_POL_III_TRANSCRIPTION
GO_NUCLEAR_NUCLEOSOME, GO_NUCLEAR_NUCLEOSOME
REACTOME_RNA_POL_III_TRANSCRIPTION_INITIATION_FROM_TYPE_2_PROMOTER, REACTOME_RNA_POL_III_TRANSCRIPTION_INITIATION_FROM_TYPE_2_PROMOTER
REACTOME_MICRORNA_MIRNA_BIOGENESIS, REACTOME_MICRORNA_MIRNA_BIOGENESIS
REACTOME_RNA_POL_III_TRANSCRIPTION_INITIATION_FROM_TYPE_3_PROMOTER, REACTOME_RNA_POL_III_TRANSCRIPTION_INITIATION_FROM_TYPE_3_PROMOTER
GO_SAGA_TYPE_COMPLEX, GO_SAGA_TYPE_COMPLEX
ROSS_AML_WITH_AML1_ETO_FUSION, ROSS_AML_WITH_AML1_ETO_FUSION
GO_POSITIVE_REGULATION_OF_MULTI_ORGANISM_PROCESS, GO_POSITIVE_REGULATION_OF_MULTI_ORGANISM_PROCESS
GEORGANTAS_HSC_MARKERS, GEORGANTAS_HSC_MARKERS
GO_PROTEIN_HETEROTETRAMERIZATION, GO_PROTEIN_HETEROTETRAMERIZATION
GO_RNA_POLYADENYLATION, GO_RNA_POLYADENYLATION
GO_HUMORAL_IMMUNE_RESPONSE, GO_HUMORAL_IMMUNE_RESPONSE
REACTOME_FORMATION_OF_TRANSCRIPTION_COUPLED_NER_TC_NER_REPAIR_COMPLEX, REACTOME_FORMATION_OF_TRANSCRIPTION_COUPLED_NER_TC_NER_REPAIR_COMPLEX
GO_POSITIVE_REGULATION_OF_VIRAL_TRANSCRIPTION, GO_POSITIVE_REGULATION_OF_VIRAL_TRANSCRIPTION
GO_RRNA_TRANSCRIPTION, GO_RRNA_TRANSCRIPTION
BANDRES_RESPONSE_TO_CARMUSTIN_MGMT_48HR_UP, BANDRES_RESPONSE_TO_CARMUSTIN_MGMT_48HR_UP
REACTOME_ABORTIVE_ELONGATION_OF_HIV1_TRANSCRIPT_IN_THE_ABSENCE_OF_TAT, REACTOME_ABORTIVE_ELONGATION_OF_HIV1_TRANSCRIPT_IN_THE_ABSENCE_OF_TAT
GO_N_ACYLTRANSFERASE_ACTIVITY, GO_N_ACYLTRANSFERASE_ACTIVITY
GO_REGULATION_OF_MRNA_3_END_PROCESSING, GO_REGULATION_OF_MRNA_3_END_PROCESSING
PID_RETINOIC_ACID_PATHWAY, PID_RETINOIC_ACID_PATHWAY
GO_DEFENSE_RESPONSE_TO_VIRUS, GO_DEFENSE_RESPONSE_TO_VIRUS
HASLINGER_B_CLL_WITH_6Q21_DELETION, HASLINGER_B_CLL_WITH_6Q21_DELETION
GNATENKO_PLATELET_SIGNATURE, GNATENKO_PLATELET_SIGNATURE
REACTOME_ELONGATION_ARREST_AND_RECOVERY, REACTOME_ELONGATION_ARREST_AND_RECOVERY
REACTOME_TRANSCRIPTIONAL_ACTIVITY_OF_SMAD2_SMAD3_SMAD4_HETEROTRIMER, REACTOME_TRANSCRIPTIONAL_ACTIVITY_OF_SMAD2_SMAD3_SMAD4_HETEROTRIMER
GSE12963_UNINF_VS_ENV_AND_NEF_AND_VPR_DEFICIENT_HIV1_INF_CD4_TCELL_UP, GSE12963_UNINF_VS_ENV_AND_NEF_AND_VPR_DEFICIENT_HIV1_INF_CD4_TCELL_UP
GO_NEGATIVE_REGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION
GO_MAINTENANCE_OF_CELL_NUMBER, GO_MAINTENANCE_OF_CELL_NUMBER
GO_REGULATION_OF_INTERFERON_BETA_PRODUCTION, GO_REGULATION_OF_INTERFERON_BETA_PRODUCTION
GO_DNA_DIRECTED_RNA_POLYMERASE_I_COMPLEX, GO_DNA_DIRECTED_RNA_POLYMERASE_I_COMPLEX
GO_REGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION, GO_REGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION
CHUNG_BLISTER_CYTOTOXICITY_DN, CHUNG_BLISTER_CYTOTOXICITY_DN
NADLER_HYPERGLYCEMIA_AT_OBESITY, NADLER_HYPERGLYCEMIA_AT_OBESITY
GTACAGG_MIR486, GTACAGG_MIR486
RAGHAVACHARI_PLATELET_SPECIFIC_GENES, RAGHAVACHARI_PLATELET_SPECIFIC_GENES
CHIARADONNA_NEOPLASTIC_TRANSFORMATION_CDC25_UP, CHIARADONNA_NEOPLASTIC_TRANSFORMATION_CDC25_UP
BIOCARTA_RARRXR_PATHWAY, BIOCARTA_RARRXR_PATHWAY
RIZKI_TUMOR_INVASIVENESS_2D_UP, RIZKI_TUMOR_INVASIVENESS_2D_UP
GO_NEGATIVE_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION
GO_PROTEIN_HETEROOLIGOMERIZATION, GO_PROTEIN_HETEROOLIGOMERIZATION
REACTOME_PROCESSING_OF_INTRONLESS_PRE_MRNAS, REACTOME_PROCESSING_OF_INTRONLESS_PRE_MRNAS
BIOCARTA_CARM_ER_PATHWAY, BIOCARTA_CARM_ER_PATHWAY
AHR_01, AHR_01
GO_POSITIVE_REGULATION_OF_G1_S_TRANSITION_OF_MITOTIC_CELL_CYCLE, GO_POSITIVE_REGULATION_OF_G1_S_TRANSITION_OF_MITOTIC_CELL_CYCLE
GO_TBP_CLASS_PROTEIN_BINDING, GO_TBP_CLASS_PROTEIN_BINDING
GSE29615_CTRL_VS_DAY3_LAIV_IFLU_VACCINE_PPMC_DN, GSE29615_CTRL_VS_DAY3_LAIV_IFLU_VACCINE_PPMC_DN
GATA2_01, GATA2_01
LIANG_SILENCED_BY_METHYLATION_UP, LIANG_SILENCED_BY_METHYLATION_UP
GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_MONOCYTE_UP, GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_MONOCYTE_UP
CTCTAGA_MIR526C_MIR518F_MIR526A, CTCTAGA_MIR526C_MIR518F_MIR526A
GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_DN, GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_DN
GSE3982_MAC_VS_NEUTROPHIL_LPS_STIM_DN, GSE3982_MAC_VS_NEUTROPHIL_LPS_STIM_DN
GO_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION, GO_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION
GO_ALTERNATIVE_MRNA_SPLICING_VIA_SPLICEOSOME, GO_ALTERNATIVE_MRNA_SPLICING_VIA_SPLICEOSOME
GRAHAM_NORMAL QUIESCENT_VS_NORMAL_DIVIDING_UP, GRAHAM_NORMAL QUIESCENT_VS_NORMAL_DIVIDING_UP
GSE37533_PPARG2_FOXP3_VS_FOXP3_TRANSDUCED_CD4_TCELL_PIOGLITAZONE_TREATED_UP, GSE37533_PPARG2_FOXP3_VS_FOXP3_TRANSDUCED_CD4_TCELL_PIOGLITAZONE_TREATED_UP
GO_REGULATION_OF_MRNA_POLYADENYLATION, GO_REGULATION_OF_MRNA_POLYADENYLATION
GO_CORE_PROMOTER_SEQUENCE_SPECIFIC_DNA_BINDING, GO_CORE_PROMOTER_SEQUENCE_SPECIFIC_DNA_BINDING
CAFFAREL_RESPONSE_TO_THC_8HR_5_DN, CAFFAREL_RESPONSE_TO_THC_8HR_5_DN
GO_STAGA_COMPLEX, GO_STAGA_COMPLEX
NAGY_TFTC_COMPONENTS_HUMAN, NAGY_TFTC_COMPONENTS_HUMAN
GAZDA_DIAMOND_BLACKFAN_ANEMIA_PROGENITOR_UP, GAZDA_DIAMOND_BLACKFAN_ANEMIA_PROGENITOR_UP
REACTOME_VIRAL_MESSENGER_RNA_SYNTHESIS, REACTOME_VIRAL_MESSENGER_RNA_SYNTHESIS
GO_RNA_CAP_BINDING, GO_RNA_CAP_BINDING
PAX2_01, PAX2_01
GO_SOMATIC_STEM_CELL_POPULATION_MAINTENANCE, GO_SOMATIC_STEM_CELL_POPULATION_MAINTENANCE
SUBTIL_PROGESTIN_TARGETS, SUBTIL_PROGESTIN_TARGETS
GO_DNA_DIRECTED_RNA_POLYMERASE_II_CORE_COMPLEX, GO_DNA_DIRECTED_RNA_POLYMERASE_II_CORE_COMPLEX
CTCTATG_MIR368, CTCTATG_MIR368
GO_MRNA_CLEAVAGE_AND_POLYADENYLATION_SPECIFICITY_FACTOR_COMPLEX, GO_MRNA_CLEAVAGE_AND_POLYADENYLATION_SPECIFICITY_FACTOR_COMPLEX
GO_TRANSCRIPTION_FACTOR_TFTC_COMPLEX, GO_TRANSCRIPTION_FACTOR_TFTC_COMPLEX
GO_LIVER_REGENERATION, GO_LIVER_REGENERATION
KYNG_WERNER_SYNDROM_DN, KYNG_WERNER_SYNDROM_DN
NAGY_STAGA_COMPONENTS_HUMAN, NAGY_STAGA_COMPONENTS_HUMAN
GO_RNA_POLYMERASE_II_CARBOXY_TERMINAL_DOMAIN_KINASE_ACTIVITY, GO_RNA_POLYMERASE_II_CARBOXY_TERMINAL_DOMAIN_KINASE_ACTIVITY
chr1q42, chr1q42
GO_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR, GO_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_C
GO_RNA_CAP_BINDING_COMPLEX, GO_RNA_CAP_BINDING_COMPLEX
GO_NEGATIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_NEGATIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
DAZARD_UV_RESPONSE_CLUSTER_G2_DAZARD_UV_RESPONSE_CLUSTER_G2
GO_NEGATIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION