

TEDCOMMON_DC_PROGENITOR_UP, GSE22432_PDC_VS_TGFB1_TREATEDCOMMON_DC_PROGENITOR_UP

GO_TRNA_METABOLIC_PROCESS, GO_TRNA_METABOLIC_PROCESS
GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_DN, GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_DN
GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_UP, GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_UP
GSE41867_DAY15_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN, GSE41867_DAY15_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN
GSE24671_BAKIMULC_VS_SENDAI_VIRUS_INFECTED_MOUSE_SPLENOCYTES_DN, GSE24671_BAKIMULC_VS_SENDAI_VIRUS_INFECTED_MOUSE_SPLENOCYTES_DN
GSE31082_DN_VS_CD4_SP_THYMOCYTE_UP, GSE31082_DN_VS_CD4_SP_THYMOCYTE_UP
GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_2H_UP
GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TREATED_CD4_TCELL_UP, GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TREATED_CD4_TCELL_UP
GSE39556_CD8A_DC_VS_NK_CELL_MOUSE_3H_POST_POLYIC_INJ_UP, GSE39556_CD8A_DC_VS_NK_CELL_MOUSE_3H_POST_POLYIC_INJ_UP
GSE17721_POLYIC_VS_PAM3CSK4_12H_BMDC_DN, GSE17721_POLYIC_VS_PAM3CSK4_12H_BMDC_DN
GSE27786_CD8_TCELL_VS_NEUTROPHIL_UP, GSE27786_CD8_TCELL_VS_NEUTROPHIL_UP
GO_TRNA_PROCESSING, GO_TRNA_PROCESSING
GSE15330_MEGAKARYOCYTE_ERYTHROID_VS GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_UP, GSE15330_MEGAKARYOCYTE_ERYTHROID_VS GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_UP
GSE20198_IL12_VS_IL12_IL18_TREATED_ACT_CD4_TCELL_UP, GSE20198_IL12_VS_IL12_IL18_TREATED_ACT_CD4_TCELL_UP
GSE14308_TH1_VS_NAIVE_CD4_TCELL_UP, GSE14308_TH1_VS_NAIVE_CD4_TCELL_UP
GSE27786_LIN_NEG_VS_NEUTROPHIL_UP, GSE27786_LIN_NEG_VS_NEUTROPHIL_UP
GO_PROTEIN_IMPORT, GO_PROTEIN_IMPORT
MORF_EIF4E, MORF_EIF4E
GSE37301_HEMATOPOIETIC_STEM_CELL_VS_PRO_BCELL_DN, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_PRO_BCELL_DN
GSE14308_TH2_VS_NATURAL_TREG_UP, GSE14308_TH2_VS_NATURAL_TREG_UP
GSE17721_CPG_VS_GARDIQUIMOD_8H_BMDC_UP, GSE17721_CPG_VS_GARDIQUIMOD_8H_BMDC_UP
GO_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GO_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
GSE18893_TCONV_VS_TREG_24H_TNF_STIM_DN, GSE18893_TCONV_VS_TREG_24H_TNF_STIM_DN
MORF_CCNF, MORF_CCNF
GO_TRNA_MODIFICATION, GO_TRNA_MODIFICATION
GO_TRNA_TRANSPORT, GO_TRNA_TRANSPORT
GSE19888_NO_PRETREAT_VS_ADENOSINE_A3R_INHIBITOR_PRETREATED_MAST_CELL_TCELL_MEMBRANES_ACT_UP, GSE19888_NO_PRETREAT_VS_ADENOSINE_A3R_INHIBITOR_PRETREATED_MAST_CELL_TCELL_MEMBRANES_ACT_UP
GSE21033_3H_VS_12H_POLYIC_STIM_DC_UP, GSE21033_3H_VS_12H_POLYIC_STIM_DC_UP
GO_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_IL_PROMOTER, GO_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_IL_PROMOTER
MODULE_273, MODULE_273
GKCCGNNNNNNNTGAYC_UNKNOWN, GKCCGNNNNNNNTGAYC_UNKNOWN
REACTOME_TRANSPORT_OF_MATURE_MRNA_DERIVED_FROM_AN_INTRONLESS_TRANSCRIPT, REACTOME_TRANSPORT_OF_MATURE_MRNA_DERIVED_FROM_AN_INTRONLESS_TRANSCRIPT
GO_COENZYME_BINDING, GO_COENZYME_BINDING
GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP, GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP
GSE360_DC_VS_MAC_1_DONOVANL_UP, GSE360_DC_VS_MAC_1_DONOVANL_UP
GO_RIBOSOMAL_SMALL_SUBUNIT_BIOGENESIS, GO_RIBOSOMAL_SMALL_SUBUNIT_BIOGENESIS
GSE22282_HYPOXIA_VS_NORMOXIA_MYELOID_DC_UP, GSE22282_HYPOXIA_VS_NORMOXIA_MYELOID_DC_UP
BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_CIS, BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_CIS
GSE360_L_DONOVANL_VS_M_TUBERCULOSIS_MAC_UP, GSE360_L_DONOVANL_VS_M_TUBERCULOSIS_MAC_UP
GO_PROTEIN_TRANSMEMBRANE_TRANSPORT, GO_PROTEIN_TRANSMEMBRANE_TRANSPORT
MODULE_22, MODULE_22
ACTWSNACTNY_UNKNOWN, ACTWSNACTNY_UNKNOWN
MODULE_93, MODULE_93
GO_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_IL_PROMOTER, GO_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_IL_PROMOTER
REACTOME_ELONGATION_ARREST_AND_RECOVERY, REACTOME_ELONGATION_ARREST_AND_RECOVERY
GO_MATURATION_OF_SSU_RRNA, GO_MATURATION_OF_SSU_RRNA
GO_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS, GO_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS
GSE40274_LEF1_VS_FOXP3_AND_LEF1_TRANSDUCEDED_ACTIVATED_CD4_TCELL_DN, GSE40274_LEF1_VS_FOXP3_AND_LEF1_TRANSDUCEDED_ACTIVATED_CD4_TCELL_DN
SETLUR_PROSTATE_CANCER_TMPRSS2_ERG_FUSION_UP, SETLUR_PROSTATE_CANCER_TMPRSS2_ERG_FUSION_UP
GSE41176_UNSTIM_VS_ANTL_IGM_STIM_TAK1_KO_BCELL_24H_DN, GSE41176_UNSTIM_VS_ANTL_IGM_STIM_TAK1_KO_BCELL_24H_DN
MODULE_363, MODULE_363
MODULE_332, MODULE_332
GSE43955_10H_VS_60H_ACT_CD4_TCELL_UP, GSE43955_10H_VS_60H_ACT_CD4_TCELL_UP
GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H
GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN_2_OXOGLUTARATE_AS_ONE_DONOR_AND_INCORPORATION_OF_ONE_ATOM_EACH_OF_OXYGEN_INT0_BOTH_DONORS, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN_2_OXOGLUTARATE_AS_ONE_DONOR_AND_INCORPORATION_OF_ONE_ATOM_EACH_OF_OXYGEN_INT0_BOTH_DONORS
GSE13485_DAY3_VS_DAY7_YF17D_VACCINE_PBM_C_UP, GSE13485_DAY3_VS_DAY7_YF17D_VACCINE_PBM_C_UP
GO_STRUCTURAL_CONSTITUENT_OF_NUCLEAR_PORE, GO_STRUCTURAL_CONSTITUENT_OF_NUCLEAR_PORE
GO_NAD_BINDING, GO_NAD_BINDING
KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_DEGRADATION, KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_DEGRADATION
REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION, REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION
GO_TRANSCRIPTION_FACTOR_ACTIVITY_RNA_POLYMERASE_IL_TRANSCRIPTION_FACTOR_BINDING, GO_TRANSCRIPTION_FACTOR_ACTIVITY_RNA_POLYMERASE_IL_TRANSCRIPTION_FACTOR_BINDING
GO_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS, GO_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS
GSE17721_LPS_VS_CPG_24H_BMDC_UP, GSE17721_LPS_VS_CPG_24H_BMDC_UP
GSE5099_DAY3_VS_DAY7_MCSF_TREATED_MACROPHAGE_DN, GSE5099_DAY3_VS_DAY7_MCSF_TREATED_MACROPHAGE_DN
GO_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_IL_PROMOTER, GO_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_IL_PROMOTER
GSE12963_UNINF_VS_ENV_AND_NEF_DEFICIENT_HIV1_INF_CD4_TCELL_DN, GSE12963_UNINF_VS_ENV_AND_NEF_DEFICIENT_HIV1_INF_CD4_TCELL_DN
GO_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GO_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H_QUINONE_OR_SIMILAR_COMPOUND_AS_ACCEPTOR, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H_QUINONE_OR_SIMILAR_COMPOUND_AS_ACCEPTOR
GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_A_SULFUR_GROUP_OF_DONORS, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_A_SULFUR_GROUP_OF_DONORS
DORSAM_HOXA9_TARGETS_UP, DORSAM_HOXA9_TARGETS_UP
GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
GO_90S_PRRIBOSOME, GO_90S_PRRIBOSOME
GO_TRNA_METHYLATION, GO_TRNA_METHYLATION
GO_TRNA_METHYLTRANSFERASE_ACTIVITY, GO_TRNA_METHYLTRANSFERASE_ACTIVITY
AACYNNNNTTCCS_UNKNOWN, AACYNNNNTTCCS_UNKNOWN
PID_P38_ALPHA_BETA_DOWNSTREAM_PATHWAY, PID_P38_ALPHA_BETA_DOWNSTREAM_PATHWAY
GO_DIOXYGENASE_ACTIVITY, GO_DIOXYGENASE_ACTIVITY
INAMURA_LUNG_CANCER_SCC_SUBTYPES_UP, INAMURA_LUNG_CANCER_SCC_SUBTYPES_UP
REACTOME_VIRAL_MESSENGER_RNA_SYNTHESIS, REACTOME_VIRAL_MESSENGER_RNA_SYNTHESIS
KEGG_SELENOAMINO_ACID_METABOLISM, KEGG_SELENOAMINO_ACID_METABOLISM
MODULE_77, MODULE_77
GSE25147_UNSTIM_VS_HELIOBACTER_PYLORI_LPS_STIM_MKN45_CELL_DN, GSE25147_UNSTIM_VS_HELIOBACTER_PYLORI_LPS_STIM_MKN45_CELL_DN
MODULE_350, MODULE_350
GSE31622_WT_VS_KLF3_KO_BCELL_DN, GSE31622_WT_VS_KLF3_KO_BCELL_DN
KEGG_PROPANOATE_METABOLISM, KEGG_PROPANOATE_METABOLISM
GO_FATTY_ACYL_COA_BINDING, GO_FATTY_ACYL_COA_BINDING
GO_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GO_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
GO_GENETIC_IMPRINTING, GO_GENETIC_IMPRINTING
GO_RESPONSE_TO_WATER, GO_RESPONSE_TO_WATER
GENTILE_UV_RESPONSE_CLUSTER_D5, GENTILE_UV_RESPONSE_CLUSTER_D5
GO_NCRNA_3_END_PROCESSING, GO_NCRNA_3_END_PROCESSING
AMIT_EGF_RESPONSE_40_MCF10A, AMIT_EGF_RESPONSE_40_MCF10A
MAYBURD_RESPONSE_TO_L663536_UP, MAYBURD_RESPONSE_TO_L663536_UP
MODULE_42, MODULE_42
GO_ENDORIBONUCLEASE_ACTIVITY, GO_ENDORIBONUCLEASE_ACTIVITY
GO_DNA_DIRECTED_RNA_POLYMERASE_I_COMPLEX, GO_DNA_DIRECTED_RNA_POLYMERASE_I_COMPLEX
GO_REGULATION_OF_GENE_EXPRESSION_BY_GENETIC_IMPRINTING, GO_REGULATION_OF_GENE_EXPRESSION_BY_GENETIC_IMPRINTING
GO_REGULATION_OF_VASCULOGENESIS, GO_REGULATION_OF_VASCULOGENESIS
GO_PEPTIDYL_PROLINE_MODIFICATION, GO_PEPTIDYL_PROLINE_MODIFICATION
GO_NEGATIVE_REGULATION_OF_HISTONE_METHYLATION, GO_NEGATIVE_REGULATION_OF_HISTONE_METHYLATION
GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_OH_GROUP_OF_DONORS_NAD_OR_NADP_AS_ACCEPTOR, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_OH_GROUP_OF_DONORS_NAD_OR_NADP_AS_ACCEPTOR
KEGG_BASAL_TRANSCRIPTION_FACTORS, KEGG_BASAL_TRANSCRIPTION_FACTORS
GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_IL_PROMOTER, GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_IL_PROMOTER
GO_RESPONSE_TO_WATER_DEPRIVATION, GO_RESPONSE_TO_WATER_DEPRIVATION
GO_CRISTAE_FORMATION, GO_CRISTAE_FORMATION
GO_PURINE_NUCLEOBASE_BIOSYNTHETIC_PROCESS, GO_PURINE_NUCLEOBASE_BIOSYNTHETIC_PROCESS
GO_PEROXISOMAL_TRANSPORT, GO_PEROXISOMAL_TRANSPORT
REACTOME_PURINE_RIBONUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHESIS, REACTOME_PURINE_RIBONUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHESIS
ZHAN_V1_LATE_DIFFERENTIATION_GENES_DN, ZHAN_V1_LATE_DIFFERENTIATION_GENES_DN
GSE2585_THYMIC_DC_VS_THYMIC_MACROPHAGE_UP, GSE2585_THYMIC_DC_VS_THYMIC_MACROPHAGE_UP
GO_CHROMATOID_BODY, GO_CHROMATOID_BODY
WACKER_HYPOXIA_TARGETS_OF_VHL, WACKER_HYPOXIA_TARGETS_OF_VHL
MODULE_286, MODULE_286
GO_RRNA_3_END_PROCESSING, GO_RRNA_3_END_PROCESSING
GO_EXOPEPTIDASE_ACTIVITY, GO_EXOPEPTIDASE_ACTIVITY
GSE32423_CTRL_VS_IL7_MEMORY_CD8_TCELL_UP, GSE32423_CTRL_VS_IL7_MEMORY_CD8_TCELL_UP
KEGG_GLYCINE_SERINE_AND_THREONINE_METABOLISM, KEGG_GLYCINE_SERINE_AND_THREONINE_METABOLISM
REACTOME_REGULATION_OF_HYPOXIA_INDUCIBLE_FACTOR_HIF_BY_OXYGEN, REACTOME_REGULATION_OF_HYPOXIA_INDUCIBLE_FACTOR_HIF_BY_OXYGEN
GO_IMP_METABOLIC_PROCESS, GO_IMP_METABOLIC_PROCESS
GSE34156_NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_DN, GSE34156_NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_DN
GO_ANTIPORTER_ACTIVITY, GO_ANTIPORTER_ACTIVITY
REACTOME_OXYGEN_DEPENDENT_PROLINE_HYDROXYLATION_OF_HYPOXIA_INDUCIBLE_FACTOR_ALPHA, REACTOME_OXYGEN_DEPENDENT_PROLINE_HYDROXYLATION_OF_HYPOXIA_INDUCIBLE_FACTOR_ALPHA
GO_PURINE_NUCLEOBASE_METABOLIC_PROCESS, GO_PURINE_NUCLEOBASE_METABOLIC_PROCESS