

VS_TGFBETA3_IL6_IL23A_CD4_TCELL_UP, GSE39820_CTRL_VS_TGFBETA3_IL6_IL23A_CD4_TCELL_UP

GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_DN, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_DN
GSE39820_CTRL_VS_TGFBETA1_IL6_IL23A_CD4_TCELL_UP, GSE39820_CTRL_VS_TGFBETA1_IL6_IL23A_CD4_TCELL_UP
GSE39820_CTRL_VS_TGFBETA3_IL6_CD4_TCELL_UP, GSE39820_CTRL_VS_TGFBETA3_IL6_CD4_TCELL_UP
GSE39820_CTRL_VS_TGFBETA1_IL6_CD4_TCELL_UP, GSE39820_CTRL_VS_TGFBETA1_IL6_CD4_TCELL_UP
GSE37532_TREG_VS_TCONV_PPARG_KO_CD4_TCELL_FROM_VISCERAL_ADIPOSE_TISSUE_DN, GSE37532_TREG_VS_TCONV_PPARG_KO_CD4_TCELL_FROM_VISCERAL_ADIPOSE_TISSUE_DN
GSE19941_UNSTIM_VS_LPS_STIM_IL10_KO_MACROPHAGE_DN, GSE19941_UNSTIM_VS_LPS_STIM_IL10_KO_MACROPHAGE_DN
GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_6H_DN, GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_6H_DN
GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_MATURE_NEURON_CELL_LINE_UP, GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_MATURE_NEURON_CELL_LINE_UP
GSE13493_CD4INTCD8POS_VS_CD8POS_THYMOCYTE_UP, GSE13493_CD4INTCD8POS_VS_CD8POS_THYMOCYTE_UP
GSE39820_CTRL_VS_IL1B_IL6_CD4_TCELL_UP, GSE39820_CTRL_VS_IL1B_IL6_CD4_TCELL_UP
GSE40277_GATA1_AND_SATB1_TRANSDUCED_VS_CTRL_CD4_TCELL_DN, GSE40277_GATA1_AND_SATB1_TRANSDUCED_VS_CTRL_CD4_TCELL_DN
BENPORATH_NOS_TARGETS, BENPORATH_NOS_TARGETS
GSE40274_FOXP3_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_FOXP3_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE20366_TREG_VS_TCONV_DN, GSE20366_TREG_VS_TCONV_DN
HP_ABNORMALITY_OF_BONE_MARROW_CELL_MORPHOLOGY, HP_ABNORMALITY_OF_BONE_MARROW_CELL_MORPHOLOGY
MIR548BA, MIR548BA
GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_6H_DN, GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_6H_DN
MIR548AG, MIR548AG
MIR548AI_MIR570_5P, MIR548AI_MIR570_5P
MIR103A_1_5P, MIR103A_1_5P
MIR5579_3P, MIR5579_3P
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_UP, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_UP
MIR593_5P, MIR593_5P
MIR12131, MIR12131
LU_AGING_BRAIN_DN, LU_AGING_BRAIN_DN
MIR548S, MIR548S
YYCATTCAWW_UNKNOWN, YYCATTCAWW_UNKNOWN
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_CONJUGATION_OR_REMOVAL, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_CONJUGATION_OR_REMOVAL
MIR4711_5P, MIR4711_5P
MIR6883_3P, MIR6883_3P
MIR606, MIR606
MIR4659B_5P, MIR4659B_5P
MODULE_525, MODULE_525
MIR3913_5P, MIR3913_5P
MIR8070, MIR8070
MIR6750_3P, MIR6750_3P
MIR190B_5P, MIR190B_5P
MIR190A_5P, MIR190A_5P
MIR1244, MIR1244
KIM_GERMINAL_CENTER_T_HELPER_UP, KIM_GERMINAL_CENTER_T_HELPER_UP
HP_UNUSUAL_INFECTION_BY_ANATOMICAL_SITE, HP_UNUSUAL_INFECTION_BY_ANATOMICAL_SITE
MARTORIATI_MDM4_TARGETS_NEUROEPITHELIUM_DN, MARTORIATI_MDM4_TARGETS_NEUROEPITHELIUM_DN
PIAS4_TARGET_GENES, PIAS4_TARGET_GENES
MIR6751_3P, MIR6751_3P
MIR1199_5P, MIR1199_5P
STREICHER_LSM1_TARGETS_UP, STREICHER_LSM1_TARGETS_UP
HP_MENINGITIS, HP_MENINGITIS
GOBP_POSITIVE_REGULATION_OF_RNA_SPLICING, GOBP_POSITIVE_REGULATION_OF_RNA_SPLICING
MIR875_5P, MIR875_5P
GOMF_PHOSPHOTRANSFERASE_ACTIVITY_FOR_OTHER_SUBSTITUTED_PHOSPHATE_GROUPS, GOMF_PHOSPHOTRANSFERASE_ACTIVITY_FOR_OTHER_SUBSTITUTED_PHOSPHATE_GROUPS
BRUINS_UVC_RESPONSE_VIA_TP53_GROUP_C, BRUINS_UVC_RESPONSE_VIA_TP53_GROUP_C
GOBP_INTERFERON_BETA_PRODUCTION, GOBP_INTERFERON_BETA_PRODUCTION
MIR7112_3P, MIR7112_3P
MIR4317, MIR4317
MIR574_5P, MIR574_5P
MIR2276_5P, MIR2276_5P
MIR6884_3P, MIR6884_3P
MIR6740_5P, MIR6740_5P
GOMF_CIS_TRANS_ISOMERASE_ACTIVITY, GOMF_CIS_TRANS_ISOMERASE_ACTIVITY
MIR15B_3P, MIR15B_3P
HP_BROAD_RIBS, HP_BROAD_RIBS
MIR6823_5P, MIR6823_5P
MIR6826_5P, MIR6826_5P
GOBP_PROTEIN_PEPTIDYL_PROLYL_ISOMERIZATION, GOBP_PROTEIN_PEPTIDYL_PROLYL_ISOMERIZATION
MIR454_5P, MIR454_5P
GOMF_TAU_PROTEIN_KINASE_ACTIVITY, GOMF_TAU_PROTEIN_KINASE_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_INTERFERON_BETA_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERFERON_BETA_PRODUCTION
RNCTGNYNRNCTGNY_UNKNOWN, RNCTGNYNRNCTGNY_UNKNOWN
GOCC_POSTSYNAPTIC_SPECIALIZATION_INTRACELLULAR_COMPONENT, GOCC_POSTSYNAPTIC_SPECIALIZATION_INTRACELLULAR_COMPONENT
BROWNE_HCMV_INFECTION_4HR_UP, BROWNE_HCMV_INFECTION_4HR_UP
HP_MATERNAL_DIABETES, HP_MATERNAL_DIABETES
GOBP_REGULATION_OF_MITOCHONDRIAL_FUSION, GOBP_REGULATION_OF_MITOCHONDRIAL_FUSION
HP_FEMORAL_HERNIA, HP_FEMORAL_HERNIA
GOCC_POSTSYNAPTIC_DENSITY_INTRACELLULAR_COMPONENT, GOCC_POSTSYNAPTIC_DENSITY_INTRACELLULAR_COMPONENT
BUSSLINGER_GASTRIC_D_CELLS, BUSSLINGER_GASTRIC_D_CELLS
HP_APLASIA_HYPOPLASIA_OF_THE_OVARY, HP_APLASIA_HYPOPLASIA_OF_THE_OVARY
HP_ANODONTIA, HP_ANODONTIA
GOBP_MICROVILLUS_ASSEMBLY, GOBP_MICROVILLUS_ASSEMBLY
LU_TUMOR_ANGIOGENESIS_UP, LU_TUMOR_ANGIOGENESIS_UP
GOBP_CHITIN_METABOLIC_PROCESS, GOBP_CHITIN_METABOLIC_PROCESS
GOBP_REGULATION_OF_NUCLEOTIDE_BINDING_OLIGOMERIZATION_DOMAIN_CONTAINING_SIGNALING_PATHWAY, GOBP_REGULATION_OF_NUCLEOTIDE_BINDING_OLIGOMERIZATION_DOMAIN_CONTAINING_SIGNALING_PATHWAY
chr6q24, chr6q24
chrXq25, chrXq25
CAO_BLOOD_FLUMIST_AGE_05_14YO_CORRELATED_WITH_H3N2_VN_TITTER_7DY_POSITIVE, CAO_BLOOD_FLUMIST_AGE_05_14YO_CORRELATED_WITH_H3N2_VN_TITTER_7DY_POSITIVE
TCGA_GLIOMASTOMA_COPY_NUMBER_DN, TCGA_GLIOMASTOMA_COPY_NUMBER_DN
GOBP_POSITIVE_REGULATION_OF_SYNAPTIC_VESICLE_EXOCYTOSIS, GOBP_POSITIVE_REGULATION_OF_SYNAPTIC_VESICLE_EXOCYTOSIS
chrXq13, chrXq13
BIOCARTA_EPHA4_PATHWAY, BIOCARTA_EPHA4_PATHWAY
REACTOME_RUNX1_REGULATES_EXPRESSION_OF_COMPONENTS_OF_TIGHT_JUNCTIONS, REACTOME_RUNX1_REGULATES_EXPRESSION_OF_COMPONENTS_OF_TIGHT_JUNCTIONS
HP_IRIS_ATROPHY, HP_IRIS_ATROPHY
GOBP_SEQUESTERING_OF_EXTRACELLULAR_LIGAND_FROM_RECEPTOR, GOBP_SEQUESTERING_OF_EXTRACELLULAR_LIGAND_FROM_RECEPTOR
GOBP_EXTRACELLULAR_REGULATION_OF_SIGNAL_TRANSDUCTION, GOBP_EXTRACELLULAR_REGULATION_OF_SIGNAL_TRANSDUCTION
ABBUD_LIF_SIGNALING_2_DN, ABBUD_LIF_SIGNALING_2_DN
REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_WNT_SIGNALING, REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_WNT_SIGNALING
GOCC_BLOOD_MICROPARTICLE, GOCC_BLOOD_MICROPARTICLE
HP_RIDGED_FINGERNAIL, HP_RIDGED_FINGERNAIL
GOBP_MIDBRAIN_HINDBRAIN_BOUNDARY_DEVELOPMENT, GOBP_MIDBRAIN_HINDBRAIN_BOUNDARY_DEVELOPMENT
chr12q21, chr12q21
HP_APLASIA_HYPOPLASIA_OF_THE_FALLOPIAN_TUBE, HP_APLASIA_HYPOPLASIA_OF_THE_FALLOPIAN_TUBE
MIR652_3P, MIR652_3P
GOBP_REGULATION_OF_BINDING_OF_SPERM_TO_ZONA_PELLUCIDA, GOBP_REGULATION_OF_BINDING_OF_SPERM_TO_ZONA_PELLUCIDA
GOCC_EGG_COAT, GOCC_EGG_COAT
REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_INTERLEUKIN_SIGNALING, REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_INTERLEUKIN_SIGNALING
LEI_HOXC8_TARGETS_DN, LEI_HOXC8_TARGETS_DN
chr3q22, chr3q22
HP_ONYCHOGRYPOSIS_OF_FINGERNAIL, HP_ONYCHOGRYPOSIS_OF_FINGERNAIL
GOBP_MAINTENANCE_OF_PROTEIN_LOCATION_IN_EXTRACELLULAR_REGION, GOBP_MAINTENANCE_OF_PROTEIN_LOCATION_IN_EXTRACELLULAR_REGION
GOMF_BILE_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_BILE_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
FUJIWARA_PARK2_IN_LIVER_CANCER_DN, FUJIWARA_PARK2_IN_LIVER_CANCER_DN