

VS_TNF_TREATED_TREG_2H_UP, GSE18893_CTRL_VS_TNF_TREATED_TREG_2H_UP

GSE31082_CD4_VS_CD8_SP_THYMOCYTE_UP, GSE31082_CD4_VS_CD8_SP_THYMOCYTE_UP
GSE40273_GATA1_KO_VS_WT_TREG_DN, GSE40273_GATA1_KO_VS_WT_TREG_DN
GSE18893_TCONV_VS_TREG_2H_CULTURE_UP, GSE18893_TCONV_VS_TREG_2H_CULTURE_UP
GSE32423_MEMORY_VS_NAIVE_CD8_TCELL_DN, GSE32423_MEMORY_VS_NAIVE_CD8_TCELL_DN
GSE26030_UNSTIM_VS_RESTIM_TH1_DAY5_POST_POLARIZATION_UP, GSE26030_UNSTIM_VS_RESTIM_TH1_DAY5_POST_POLARIZATION_UP
GSE37301_PRO_BCELL_VS_RAG2_KO_NK_CELL_DN, GSE37301_PRO_BCELL_VS_RAG2_KO_NK_CELL_DN
GSE22601_DOUBLE_POSITIVE_VS_CD8_SINGLE_POSITIVE_THYMOCYTE_UP, GSE22601_DOUBLE_POSITIVE_VS_CD8_SINGLE_POSITIVE_THYMOCYTE_UP
GSE29618_BCELL_VS_PDC_UP, GSE29618_BCELL_VS_PDC_UP
GSE25088_CTRL_VS_IL4_STIM_MACROPHAGE_UP, GSE25088_CTRL_VS_IL4_STIM_MACROPHAGE_UP
GSE25088_WT_VS_STAT6_KO_MACROPHAGE_ROSIGLITAZONE_STIM_DN, GSE25088_WT_VS_STAT6_KO_MACROPHAGE_ROSIGLITAZONE_STIM_DN
GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP
GSE39916_B_CELL_SPLEEN_VS_PLASMA_CELL_BONE_MARROW_UP, GSE39916_B_CELL_SPLEEN_VS_PLASMA_CELL_BONE_MARROW_UP
MIR548U, MIR548U
LAKE_ADULT_KIDNEY_C26_MESANGIAL_CELLS, LAKE_ADULT_KIDNEY_C26_MESANGIAL_CELLS
GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP, GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP
MIR132_3P, MIR132_3P
MIR212_3P, MIR212_3P
GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_1_DN, GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_1_DN
AACTGAC_MIR223, AACTGAC_MIR223
MIR499B_3P, MIR499B_3P
MIR499A_3P, MIR499A_3P
GSE27241_WT_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_THI7_POLARIZING_CONDITIONS_UP, GSE27241_WT_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_THI7_POLARIZING_CONDITIONS_UP
MIR3064_5P, MIR3064_5P
GSE18893_TCONV_VS_TREG_2H_TNF_STIM_UP, GSE18893_TCONV_VS_TREG_2H_TNF_STIM_UP
GSE32533_WT_VS_MIR17_KO_ACT_CD4_TCELL_UP, GSE32533_WT_VS_MIR17_KO_ACT_CD4_TCELL_UP
GSE7831_1H_VS_4H_INFLUENZA_STIM_PDC_DN, GSE7831_1H_VS_4H_INFLUENZA_STIM_PDC_DN
GSE11961_FOLLICULAR_BCELL_VS_MARGINAL_ZONE_BCELL_UP, GSE11961_FOLLICULAR_BCELL_VS_MARGINAL_ZONE_BCELL_UP
GSE29949_MICROGLIA_VS_DC_BRAIN_UP, GSE29949_MICROGLIA_VS_DC_BRAIN_UP
MIR6720_5P, MIR6720_5P
MIR6882_3P, MIR6882_3P
MIR1251_3P, MIR1251_3P
MIR6512_3P, MIR6512_3P
MIR130B_5P, MIR130B_5P
GSE11924_TH2_VS_THI7_CD4_TCELL_UP, GSE11924_TH2_VS_THI7_CD4_TCELL_UP
MIR6852_3P, MIR6852_3P
MIR6509_3P, MIR6509_3P
MIR499B_5P, MIR499B_5P
MIR4277, MIR4277
GINESTIER_BREAST_CANCER_ZNF217_AMPLIFIED_UP, GINESTIER_BREAST_CANCER_ZNF217_AMPLIFIED_UP
MIR4530, MIR4530
GSE9509_10MIN_VS_30MIN_LPS_STIM_IL10_KO_MACROPHAGE_DN, GSE9509_10MIN_VS_30MIN_LPS_STIM_IL10_KO_MACROPHAGE_DN
MIR7850_5P, MIR7850_5P
MIR7155_5P, MIR7155_5P
ZHAN_MULTIPLE_MYELOMA_UP, ZHAN_MULTIPLE_MYELOMA_UP
MIR4794, MIR4794
ROSS_ACUTE_MYELOID_LEUKEMIA_CBF, ROSS_ACUTE_MYELOID_LEUKEMIA_CBF
MIR149_5P, MIR149_5P
GSE2706_R848_VS_R848_AND_LPS_8H_STIM_DC_DN, GSE2706_R848_VS_R848_AND_LPS_8H_STIM_DC_DN
HP_FOCAL_DYSTONIA, HP_FOCAL_DYSTONIA
MIR4640_5P, MIR4640_5P
GOBP_POSITIVE_REGULATION_OF_PROTEIN_BINDING, GOBP_POSITIVE_REGULATION_OF_PROTEIN_BINDING
MIR643, MIR643
MIR4708_5P, MIR4708_5P
MIR6758_3P, MIR6758_3P
GSE13547_CTRL_VS_ANTI_IJM_STIM_ZFX_KO_BCELL_12H_UP, GSE13547_CTRL_VS_ANTI_IJM_STIM_ZFX_KO_BCELL_12H_UP
MIR1255B_2_3P, MIR1255B_2_3P
HP_SEPSIS, HP_SEPSIS
HP_PARESTHESIA, HP_PARESTHESIA
UBE2L_TARGET_GENES, UBE2L_TARGET_GENES
GOBP_BONE_DEVELOPMENT, GOBP_BONE_DEVELOPMENT
GSE21927_EL4_VS_MCA203_TUMOR_MONOCYTES_DN, GSE21927_EL4_VS_MCA203_TUMOR_MONOCYTES_DN
GOBP_REGULATION_OF_RHO_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_REGULATION_OF_RHO_PROTEIN_SIGNAL_TRANSDUCTION
TRAVAGLINI_LUNG_ARTERY_CELL, TRAVAGLINI_LUNG_ARTERY_CELL
BOYLAN_MULTIPLE_MYELOMA_PCA3_UP, BOYLAN_MULTIPLE_MYELOMA_PCA3_UP
MIR675_3P, MIR675_3P
MIR324_3P, MIR324_3P
PID_CDC42_REG_PATHWAY, PID_CDC42_REG_PATHWAY
HOOL_ST7_TARGETS_DN, HOOL_ST7_TARGETS_DN
HP_ABNORMALITY_OF_PROTHROMBIN, HP_ABNORMALITY_OF_PROTHROMBIN
ZHAN_MULTIPLE_MYELOMA_LB_UP, ZHAN_MULTIPLE_MYELOMA_LB_UP
ALX4_TARGET_GENES, ALX4_TARGET_GENES
MIR18B_3P, MIR18B_3P
GOBP_GLOMERULUS_DEVELOPMENT, GOBP_GLOMERULUS_DEVELOPMENT
MODULE_166, MODULE_166
MIR5000_3P, MIR5000_3P
MIR6888_5P, MIR6888_5P
MIR4687_5P, MIR4687_5P
BALDWIN_PRKCI_TARGETS_UP, BALDWIN_PRKCI_TARGETS_UP
MIR4268, MIR4268
LIM_MAMMARY_LUMINAL_MATURE_DN, LIM_MAMMARY_LUMINAL_MATURE_DN
BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_DN, BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_DN
SHI_SPARC_TARGETS_UP, SHI_SPARC_TARGETS_UP
VANASSE_BCL2_TARGETS_UP, VANASSE_BCL2_TARGETS_UP
WP_PRIMARY_FOCAL_SEGMENTAL_GLOMERULOSCLEROSIS_FSGS, WP_PRIMARY_FOCAL_SEGMENTAL_GLOMERULOSCLEROSIS_FSGS
PID_ERBB2_ERBB3_PATHWAY, PID_ERBB2_ERBB3_PATHWAY
GOBP_CHONDROCYTE_DIFFERENTIATION, GOBP_CHONDROCYTE_DIFFERENTIATION
MIR1298_5P, MIR1298_5P
NADERI_BREAST_CANCER_PROGNOSIS_DN, NADERI_BREAST_CANCER_PROGNOSIS_DN
BEGUM_TARGETS_OF_PAX3_FOXO1_FUSION_UP, BEGUM_TARGETS_OF_PAX3_FOXO1_FUSION_UP
MIR4693_3P, MIR4693_3P
GOBP_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT, GOBP_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT
GOBP_CARTILAGE_DEVELOPMENT, GOBP_CARTILAGE_DEVELOPMENT
GOBP_CGMP_MEDIATED_SIGNALING, GOBP_CGMP_MEDIATED_SIGNALING
CUI_DEVELOPING_HEART_CORONARY_VASCULAR_ENDOTHELIAL_CELL, CUI_DEVELOPING_HEART_CORONARY_VASCULAR_ENDOTHELIAL_CELL
GOBP_SPECIFICATION_OF_ANIMAL_ORGAN_IDENTITY, GOBP_SPECIFICATION_OF_ANIMAL_ORGAN_IDENTITY
GOBP_NEPHRON_EPITHELIUM_DEVELOPMENT, GOBP_NEPHRON_EPITHELIUM_DEVELOPMENT
DISTECHE_ESCAPED_FROM_X_INACTIVATION, DISTECHE_ESCAPED_FROM_X_INACTIVATION
HP_EXTRAPYRAMIDAL_MUSCULAR_RIGIDITY, HP_EXTRAPYRAMIDAL_MUSCULAR_RIGIDITY
HP_MALAR_RASH, HP_MALAR_RASH
KAMIKUBO_MYELOID_MN1_NETWORK, KAMIKUBO_MYELOID_MN1_NETWORK
HP_LARYNGEAL_DYSTONIA, HP_LARYNGEAL_DYSTONIA
GOBP_EXPLORATION_BEHAVIOR, GOBP_EXPLORATION_BEHAVIOR
GOBP_EXOCRINE_SYSTEM_DEVELOPMENT, GOBP_EXOCRINE_SYSTEM_DEVELOPMENT
GOBP_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT, GOBP_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT
GOBP_AROMATIC_AMINO_ACID_TRANSPORT, GOBP_AROMATIC_AMINO_ACID_TRANSPORT
VANDESLUIS_NORMAL_EMBRYOS_DN, VANDESLUIS_NORMAL_EMBRYOS_DN
MIR6874_5P, MIR6874_5P
GOBP_MATERNAL_PLACENTA_DEVELOPMENT, GOBP_MATERNAL_PLACENTA_DEVELOPMENT
MIKKELSEN_NPC_LCP_WITH_H3K4ME3, MIKKELSEN_NPC_LCP_WITH_H3K4ME3
GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_SURFACE, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_SURFACE
GOCC_REGION_OF_CYTOSOL, GOCC_REGION_OF_CYTOSOL
IIZUKA_LIVER_CANCER_EARLY_RECURRENCE, IIZUKA_LIVER_CANCER_EARLY_RECURRENCE
HP_HEMATOCHEZIA, HP_HEMATOCHEZIA
GOCC_POSTSYNAPTIC_CYTOSOL, GOCC_POSTSYNAPTIC_CYTOSOL
GOMF_NADPLUS_NUCLEOSIDASE_ACTIVITY, GOMF_NADPLUS_NUCLEOSIDASE_ACTIVITY
GOMF_SOLUTE_PROTON_SYMPORTER_ACTIVITY, GOMF_SOLUTE_PROTON_SYMPORTER_ACTIVITY
GOBP_REGULATION_OF_NERVOUS_SYSTEM_PROCESS, GOBP_REGULATION_OF_NERVOUS_SYSTEM_PROCESS
GOBP_RENAL_TUBULE_DEVELOPMENT, GOBP_RENAL_TUBULE_DEVELOPMENT
GOBP_REGULATION_OF_CHONDROCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_CHONDROCYTE_DIFFERENTIATION
HP_PERSISTENT_BLEEDING_AFTER_TRAUMA, HP_PERSISTENT_BLEEDING_AFTER_TRAUMA
GOBP_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT, GOBP_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT
HP_ELEVATED_CIRCULATING_PARATHYROID_HORMONE_LEVEL, HP_ELEVATED_CIRCULATING_PARATHYROID_HORMONE_LEVEL
KEGG_TASTE_TRANSDUCTION, KEGG_TASTE_TRANSDUCTION
REACTOME_REGULATION_OF_GENE_EXPRESSION_IN_BETA_CELLS, REACTOME_REGULATION_OF_GENE_EXPRESSION_IN_BETA_CELLS
GOBP_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_DEVELOPMENT, GOBP_REGULATION_OF_SKELETAL_MUSCLE_TISSUE_DEVELOPMENT
DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_OLFACTORY_HORIZONTAL_BASAL_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_OLFACTORY_HORIZONTAL_BASAL_CELLS
HP_MACULAR_THICKENING, HP_MACULAR_THICKENING
GCM_ERCC4, GCM_ERCC4
HP_ODONTOGENIC_NEOPLASM, HP_ODONTOGENIC_NEOPLASM
GOBP_REGULATION_OF_CARTILAGE_DEVELOPMENT, GOBP_REGULATION_OF_CARTILAGE_DEVELOPMENT
WP_SCFA_AND_SKELETAL_MUSCLE_SUBSTRATE_METABOLISM, WP_SCFA_AND_SKELETAL_MUSCLE_SUBSTRATE_METABOLISM
GOBP_RESPONSE_TO_FLUID_SHEAR_STRESS, GOBP_RESPONSE_TO_FLUID_SHEAR_STRESS
GOBP_DIAPHRAGM_DEVELOPMENT, GOBP_DIAPHRAGM_DEVELOPMENT
GOBP_AMPA_GLUTAMATE_RECEPTOR_CLUSTERING, GOBP_AMPA_GLUTAMATE_RECEPTOR_CLUSTERING
GOBP_METANEPHROS_MORPHOGENESIS, GOBP_METANEPHROS_MORPHOGENESIS
REACTOME_ESTROGEN_DEPENDENT_NUCLEAR_EVENTS_DOWNSTREAM_OF_ESR_MEMBRANE_SIGNALING, REACTOME_ESTROGEN_DEPENDENT_NUCLEAR_EVENTS_DOWNSTREAM_OF_ESR_MEMBRANE_SIGNALING
GOBP_REGULATION_OF_KIDNEY_DEVELOPMENT, GOBP_REGULATION_OF_KIDNEY_DEVELOPMENT
GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT
HP_PROLONGED_BLEEDING_FOLLOWING_PROCEDURE, HP_PROLONGED_BLEEDING_FOLLOWING_PROCEDURE
GOBP_REGULATION_OF_MYOBLAST_PROLIFERATION, GOBP_REGULATION_OF_MYOBLAST_PROLIFERATION