

1_KO_8H_ANTI_IGM_STIM_BCELL_DN, GSE21063_WT_VS_NFATC1_KO_8H_ANTI_IGM_STIM_BCELL_DN

GSE12003_MIR223_KO_VS_WT_BM_PROGENITOR_4D_CULTURE_UP
GOLDRATH_EFF_VS_MEMORY_CD8_TCELL_DN, GOLDRATH_EFF_VS_MEMORY_CD8_TCELL_DN
GSE43863_TH1_VS_1Y6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN, GSE43863_TH1_VS_1Y6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN
GSE7852_LN_VS_THYMUS_TREG_UP, GSE7852_LN_VS_THYMUS_TREG_UP
MIR603, MIR603
GSE17186_NAIVE_VS_CD21HIGH_TRANSITIONAL_BCELL_CORD_BLOOD_DN, GSE17186_NAIVE_VS_CD21HIGH_TRANSITIONAL_BCELL_CORD_BLOOD_DN
GSE43955_TGFB_IL6_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_60H_UP, GSE43955_TGFB_IL6_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_60H_UP
GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_6H_UP, GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_6H_UP
GSE21033_1H_VS_24H_POLYIC_STIM_DC_UP, GSE21033_1H_VS_24H_POLYIC_STIM_DC_UP
GSE23321_CENTRAL_VS_EFFECTOR_MEMORY_CD8_TCELL_UP, GSE23321_CENTRAL_VS_EFFECTOR_MEMORY_CD8_TCELL_UP
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_DN, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_DN
GSE7460_FOXP3_MUT_VS_WT_ACT_TCONV_DN, GSE7460_FOXP3_MUT_VS_WT_ACT_TCONV_DN
GSE18203_CTRL_VS_INTRATUMORAL_CPG_INJ_MC38_TUMOR_UP, GSE18203_CTRL_VS_INTRATUMORAL_CPG_INJ_MC38_TUMOR_UP
GSE17721_0.5H_VS_4H_LPS_BMDC_DN, GSE17721_0.5H_VS_4H_LPS_BMDC_DN
MCBRYAN_PUBERTAL_BREAST_5_6WK_DN, MCBRYAN_PUBERTAL_BREAST_5_6WK_DN
GSE3982_DC_VS_CENT_MEMORY_CD4_TCELL_DN, GSE3982_DC_VS_CENT_MEMORY_CD4_TCELL_DN
HP_SELF_INJURIOUS_BEHAVIOR, HP_SELF_INJURIOUS_BEHAVIOR
THRA_TARGET_GENES, THRA_TARGET_GENES
GSE43955_10H_VS_30H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN, GSE43955_10H_VS_30H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN
GSE36476_YOUNG_VS_OLD_DONOR_MEMORY_CD4_TCELL_16H_TSST_ACT_DN, GSE36476_YOUNG_VS_OLD_DONOR_MEMORY_CD4_TCELL_16H_TSST_ACT_DN
GSE22886_IGM_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_UP, GSE22886_IGM_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_UP
GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_DN, GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_DN
GSE22229_UNTREATED_VS_IMMUNOSUPP_THERAPY_RENAL_TRANSPLANT_PATIENT_PBMIC_UP, GSE22229_UNTREATED_VS_IMMUNOSUPP_THERAPY_RENAL_TRANSPLANT_PATIENT_PBMIC_UP
GSE30971_2H_VS_4H_LPS_STIM_MACROPHAGE_WBP7_KO_UP, GSE30971_2H_VS_4H_LPS_STIM_MACROPHAGE_WBP7_KO_UP
YGCANTGCR_UNKNOWN, YGCANTGCR_UNKNOWN
GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG2_AND_FOXP3 TRASDUCED_UP, GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG2_AND_FOXP3 TRASDUCED_UP
MIR6809_5P, MIR6809_5P
MIR324_3P, MIR324_3P
GOBP_HISTONE_DEACETYLATION, GOBP_HISTONE_DEACETYLATION
GAO_ESOPHAGUS_25W_C3_FGFR1LOW_EPITHELIAL_CELLS, GAO_ESOPHAGUS_25W_C3_FGFR1LOW_EPITHELIAL_CELLS
GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_DN, GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_DN
WANG_PROSTATE_CANCER_ANDROGEN_INDEPENDENT, WANG_PROSTATE_CANCER_ANDROGEN_INDEPENDENT
GSE17721_PAM3CSK4_VS_GADIQUIMOD_1H_BMDC_UP, GSE17721_PAM3CSK4_VS_GADIQUIMOD_1H_BMDC_UP
MIR3613_5P, MIR3613_5P
GOBP_FOCAL_ADHESION_ASSEMBLY, GOBP_FOCAL_ADHESION_ASSEMBLY
GSE17721_POLYIC_VS_CPG_0.5H_BMDC_UP, GSE17721_POLYIC_VS_CPG_0.5H_BMDC_UP
REACTOME_GLYCEROPHOSPHOLIPID_BIOSYNTHESIS, REACTOME_GLYCEROPHOSPHOLIPID_BIOSYNTHESIS
GOCC_POLYSOMAL_RIBOSOME, GOCC_POLYSOMAL_RIBOSOME
MIR3194_5P, MIR3194_5P
MIR4289, MIR4289
AR_03, AR_03
HP_MACRODONTIA, HP_MACRODONTIA
HP_FRONTAL_CORTICAL_ATROPHY, HP_FRONTAL_CORTICAL_ATROPHY
GOBP_POSITIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
HP_APLASIA_HYPOPLASIA_OF_THE_HALLUX, HP_APLASIA_HYPOPLASIA_OF_THE_HALLUX
GOBP_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS, GOBP_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS
KEGG_GLYCEROLIPID_METABOLISM, KEGG_GLYCEROLIPID_METABOLISM
HMEF2_Q6, HMEF2_Q6
GOBP_NEGATIVE_REGULATION_OF_GTPASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_GTPASE_ACTIVITY
GGGGCCC_MIR296, GGGGCC_MIR296
HOWLIN_CITED1_TARGETS_1_UP, HOWLIN_CITED1_TARGETS_1_UP
HAY_BONE_MARROW_CD34_POS_PRE_PC, HAY_BONE_MARROW_CD34_POS_PRE_PC
REACTOME_GLYCOGEN_BREAKDOWN_GLYCOGENOLYSIS, REACTOME_GLYCOGEN_BREAKDOWN_GLYCOGENOLYSIS
HP_INCREASED_MUSCLE_GLYCOGEN_CONTENT, HP_INCREASED_MUSCLE_GLYCOGEN_CONTENT
HP_ABNORMAL_MUSCLE_GLYCOGEN_CONTENT, HP_ABNORMAL_MUSCLE_GLYCOGEN_CONTENT
GAO_LARGE_INTESTINE_24W_C6_SECRETORY_PROGENITOR, GAO_LARGE_INTESTINE_24W_C6_SECRETORY_PROGENITOR
GOBP_REGULATION_OF_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN, GOBP_REGULATION_OF_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN
GOBP_REGULATION_OF_KETONE_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_KETONE_BIOSYNTHETIC_PROCESS
GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATION, GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATION
HEDENFALK_BREAST_CANCER_BRACX_UP, HEDENFALK_BREAST_CANCER_BRACX_UP
GOBP_C21_STEROID_HORMONE_BIOSYNTHETIC_PROCESS, GOBP_C21_STEROID_HORMONE_BIOSYNTHETIC_PROCESS
CLASPER_LYMPHATIC_VESSELS_DURING_METASTASIS_UP, CLASPER_LYMPHATIC_VESSELS_DURING_METASTASIS_UP
REACTOME_CALCITONIN_LIKE_LIGAND_RECEPTORS, REACTOME_CALCITONIN_LIKE_LIGAND_RECEPTORS
GOCC_SPECTRIN_ASSOCIATED_CYTOSKELETON, GOCC_SPECTRIN_ASSOCIATED_CYTOSKELETON
HARALAMBIEVA_PBMIC_FLUARIX_AGE_50_74YO_CORR_WITH_28D_MEM_B_CELL_RESPONSE_AT_28DY_LATE_GENE_EXPR_INDIVID.GENE_MODELS_PRED_PEAK_B_CELL_ELISPOT_RESP_POSITIVE, HARALAMBIEVA_PBMIC_FLUARIX_AGE_50_74YO_CORR_WITH_28D_MEM_B_CELL_RESPONSE_AT_28DY_LATE_GENE_EXPR_INDIVID.GENE_MODELS_PRED_PEAK_B_CELL_ELISPOT_RESP_POSITIVE
GOBP_ISOPENTENYL_DIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_ISOPENTENYL_DIPHOSPHATE_BIOSYNTHETIC_PROCESS
GOBP_SIGNALING_RECEPTOR_LIGAND_PRECURSOR_PROCESSING, GOBP_SIGNALING_RECEPTOR_LIGAND_PRECURSOR_PROCESSING
GOBP_DEFENSE_RESPONSE_TO_TUMOR_CELL, GOBP_DEFENSE_RESPONSE_TO_TUMOR_CELL
GOBP_CALCITONIN_FAMILY_RECEPTOR_SIGNALING_PATHWAY, GOBP_CALCITONIN_FAMILY_RECEPTOR_SIGNALING_PATHWAY
HP_ABNORMAL_ATRIOVENTRICULAR_CONNECTION, HP_ABNORMAL_ATRIOVENTRICULAR_CONNECTION
GOCC_AXOLEMMA, GOCC_AXOLEMMA
HP_ROUND_EAR, HP_ROUND_EAR
GOBP_ANDROGEN_METABOLIC_PROCESS, GOBP_ANDROGEN_METABOLIC_PROCESS
GOBP_OLEFINIC_COMPOUND_BIOSYNTHETIC_PROCESS, GOBP_OLEFINIC_COMPOUND_BIOSYNTHETIC_PROCESS
GOMF_LIM_DOMAIN_BINDING, GOMF_LIM_DOMAIN_BINDING
DESCARTES_FETAL_EYE_AMACRINE_CELLS, DESCARTES_FETAL_EYE_AMACRINE_CELLS
ZNF416_TARGET_GENES, ZNF416_TARGET_GENES
HP_LOP_EAR, HP_LOP_EAR
HP_INFLEXIBLE_ADHERENCE_TO_ROUTINES_OR_RITUALS, HP_INFLEXIBLE_ADHERENCE_TO_ROUTINES_OR_RITUALS
GOBP_POSITIVE_REGULATION_OF_BLOOD_CIRCULATION, GOBP_POSITIVE_REGULATION_OF_BLOOD_CIRCULATION
ZNF491_TARGET_GENES, ZNF491_TARGET_GENES
GOMF_BENZODIAZEPINE_RECEPTOR_ACTIVITY, GOMF_BENZODIAZEPINE_RECEPTOR_ACTIVITY
GOBP_INTERLEUKIN_21_PRODUCTION, GOBP_INTERLEUKIN_21_PRODUCTION
PASTURAL_RIZ1_TARGETS_DN, PASTURAL_RIZ1_TARGETS_DN
GOMF_G_PROTEIN_COUPLED_ADENOSINE_RECEPTOR_ACTIVITY, GOMF_G_PROTEIN_COUPLED_ADENOSINE_RECEPTOR_ACTIVITY
DESCARTES_MAIN_FETAL_LIMBIC_SYSTEM_NEURONS, DESCARTES_MAIN_FETAL_LIMBIC_SYSTEM_NEURONS
GOBP_PROGESTERONE_BIOSYNTHETIC_PROCESS, GOBP_PROGESTERONE_BIOSYNTHETIC_PROCESS