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GSE23321_EFFECTOR_MEMORY_VS_NAIVE_CD8_TCELL_DN, GSE23321_EFFECTOR_MEMORY_VS_NAIVE_CD8_TCELL_DN
 GSE17974_1H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_UP, GSE17974_1H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_UP
 GSE32533_WT_VS_MIR17_OVEREXPRESS_ACT_CD4_TCELL_UP, GSE32533_WT_VS_MIR17_OVEREXPRESS_ACT_CD4_TCELL_UP
 GSE17974_0.5H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_UP, GSE17974_0.5H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_UP
 GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_DN, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_DN
 GSE24634_NAIVE_CD4_TCELL_VS_DAY7_IL4_CONV_TREG_UP, GSE24634_NAIVE_CD4_TCELL_VS_DAY7_IL4_CONV_TREG_UP
 GSE19888 ADENOSINE A3R INH PRETREAT AND ACT BY A3R VS A3R INH AND TCELL MEMBRANES ACT MAST CELL UP, GSE19888 ADENOSINE A3R INH PRETREAT AND ACT BY A3R VS A3R INH AND TCELL MEMBRANES ACT MAST CELL UP
 GSE17974_2H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_UP, GSE17974_2H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_UP
 GSE17974_1.5H_VS_72H_IL4_AND_ANTI_IL12_ACT_CD4_TCELL_UP, GSE17974_1.5H_VS_72H_IL4_AND_ANTI_IL12_ACT_CD4_TCELL_UP
 GSE28408_LY6G_POS_VS_NEG_DC_UP, GSE28408_LY6G_POS_VS_NEG_DC_UP
 GSE16697_CD4_TCELL_VS_TFH_CD4_TCELL_UP, GSE16697_CD4_TCELL_VS_TFH_CD4_TCELL_UP
 GSE45365_NK_CELL_VS_BCELL_UP, GSE45365_NK_CELL_VS_BCELL_UP
 GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_2H_DN
 GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_NFATC1_KO_3H_DN, GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_NFATC1_KO_3H_DN
 GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_17H_UP, GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_17H_UP
 GSE25088_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_DN, GSE25088_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_DN
 GSE44649_WT_VS_MIR155_KO_ACTIVATED_CD8_TCELL_UP, GSE44649_WT_VS_MIR155_KO_ACTIVATED_CD8_TCELL_UP
 GSE13411_NAIVE_BCELL_VS_PLASMA_CELL_UP, GSE13411_NAIVE_BCELL_VS_PLASMA_CELL_UP
 GSE17974_0H_VS_12H_IN_VITRO_ACT_CD4_TCELL_UP, GSE17974_0H_VS_12H_IN_VITRO_ACT_CD4_TCELL_UP
 GSE17974_0H_VS_48H_IN_VITRO_ACT_CD4_TCELL_UP, GSE17974_0H_VS_48H_IN_VITRO_ACT_CD4_TCELL_UP
 GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_24H_CD4_TCELL_UP, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_24H_CD4_TCELL_UP
GSE36476_YOUNG_VS_OLD_DONOR_MEMORY_CD4_TCELL_72H_TSST_ACT_DN, GSE36476_YOUNG_VS_OLD_DONOR_MEMORY_CD4_TCELL_72H_TSST_ACT_DN
 GSE22033_UNTREATED_VS_ROSIGLITAZONE_TREATED_MEF_DN, GSE22033_UNTREATED_VS_ROSIGLITAZONE_TREATED_MEF_DN
 GSE22886_NAIVE_BCELL_VS_BLOOD_PLASMA_CELL_UP, GSE22886_NAIVE_BCELL_VS_BLOOD_PLASMA_CELL_UP
 GSE7460_WT_VS_FOXP3_HET_ACT_WITH_TGFB_TCONV_UP, GSE7460_WT_VS_FOXP3_HET_ACT_WITH_TGFB_TCONV_UP
 GSE17721_PAM3CSK4_VS_GADIQUIMOD_6H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_6H_BMDC_DN
 GSE43863 LY6C INT CXCR5POS VS LY6C LOW CXCR5NEG EFFECTOR CD4 TCELL DN, GSE43863 LY6C INT CXCR5POS VS LY6C LOW CXCR5NEG EFFECTOR CD4 TCELL DN
 GSE5589_WT_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_DN, GSE5589_WT_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_DN
 GSE14769_UNSTIM_VS_120MIN_LPS_BMDM_DN, GSE14769_UNSTIM_VS_120MIN_LPS_BMDM_DN
 GSE369_SOCS3_KO_VS_WT_LIVER_POST_IL6_INJECTION_DN, GSE369_SOCS3_KO_VS_WT_LIVER_POST_IL6_INJECTION_DN
GSE17721_PAM3CSK4_VS_CPG_2H_BMDC_DN, GSE17721_PAM3CSK4_VS_CPG_2H_BMDC_DN
GSE1460_INTRATHYMIC_T_PROGENITOR_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_DN, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_DN
GSE21360_PRIMARY_VS_SECONDARY_MEMORY_CD8_TCELL_DN, GSE21360_PRIMARY_VS_SECONDARY_MEMORY_CD8_TCELL_DN
GSE1460_INTRATHYMIC_T_PROGENITOR_VS_DP_THYMOCYTE_DN, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_DP_THYMOCYTE_DN
GOMF_RNA_METHYLTRANSFERASE_ACTIVITY, GOMF_RNA_METHYLTRANSFERASE_ACTIVITY
GSE41867_DAY15_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_UP, GSE41867_DAY15_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_UP
GSE28783_CTRL_ANTI_MIR_VS_UNTREATED_ATHEROSCLEROSIS_MACROPHAGE_DN, GSE28783_CTRL_ANTI_MIR_VS_UNTREATED_ATHEROSCLEROSIS_MACROPHAGE_DN
MIR6128, MIR6128
GSE14415_FOXP3_KO_NATURAL_TREG_VS_TCONV_DN, GSE14415_FOXP3_KO_NATURAL_TREG_VS_TCONV_DN
GSE3982_DC_VS_CENT_MEMORY_CD4_TCELL_DN, GSE3982_DC_VS_CENT_MEMORY_CD4_TCELL_DN
MIR758 3P, MIR758 3P
GSE12845_NAIVE_VS_DARKZONE_GC_TONSIL_BCELL_UP, GSE12845_NAIVE_VS_DARKZONE_GC_TONSIL_BCELL_UP
MIR103A_1_5P, MIR103A_1_5P
MIR370 5P, MIR370 5P
GSE27241 WT CTRL VS DIGOXIN TREATED RORGT KO CD4 TCELL IN TH17 POLARIZING CONDITIONS UP, GSE27241 WT CTRL VS DIGOXIN TREATED RORGT KO CD4 TCELL IN TH17 POLARIZING CONDITIONS UP
GSE40441 NRP1 POS INDUCED TREG VS NRP1 NEG NATURAL TREG DN, GSE40441 NRP1 POS INDUCED TREG VS NRP1 NEG NATURAL TREG DN
GOBP_T_CELL_PROLIFERATION, GOBP_T_CELL_PROLIFERATION
GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_UP, GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_UP
GSE40274_CTRL_VS_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_CTRL_VS_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GOBP ALPHA BETA T CELL ACTIVATION, GOBP ALPHA BETA T CELL ACTIVATION
GSE1460_INTRATHYMIC_T_PROGENITOR_VS_NAIVE_CD4_TCELL_CORD_BLOOD_DN, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_NAIVE_CD4_TCELL_CORD_BLOOD_DN
GAGCCTG_MIR484, GAGCCTG_MIR484
GOBP_LEUKOCYTE_APOPTOTIC_PROCESS, GOBP_LEUKOCYTE_APOPTOTIC_PROCESS
 MIR376C 3P, MIR376C 3P
 GRADE COLON AND RECTAL CANCER DN, GRADE COLON AND RECTAL CANCER DN
 MIR4468, MIR4468
 MIR182 3P, MIR182 3P
 AMIT_EGF_RESPONSE_240_HELA, AMIT_EGF_RESPONSE_240_HELA
ZHENG_FOXP3_TARGETS_IN_T_LYMPHOCYTE_DN, ZHENG_FOXP3_TARGETS_IN_T_LYMPHOCYTE_DN
 GOBP_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
 MIR500B 3P, MIR500B 3P
 GOBP_REGULATION_OF_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_REGULATION_OF_ALPHA_BETA_T_CELL_ACTIVATION
 GSE37319_WT_VS_RC3H1_KO_CD44LOW_CD8_TCELL_UP, GSE37319_WT_VS_RC3H1_KO_CD44LOW_CD8_TCELL_UP
 TRAVAGLINI_LUNG_ARTERY_CELL, TRAVAGLINI_LUNG_ARTERY_CELL
 GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_PROLIFERATION
 GOBP_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION
 GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_PDC_DN, GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_PDC_DN
 BORCZUK_MALIGNANT_MESOTHELIOMA_DN, BORCZUK_MALIGNANT_MESOTHELIOMA_DN
 TARTE_PLASMA_CELL_VS_B_LYMPHOCYTE_DN, TARTE_PLASMA_CELL_VS_B_LYMPHOCYTE_DN
 GOBP_REGULATION_OF_LEUKOCYTE_APOPTOTIC_PROCESS, GOBP_REGULATION_OF_LEUKOCYTE_APOPTOTIC_PROCESS
 KEGG_SELENOAMINO_ACID_METABOLISM, KEGG_SELENOAMINO_ACID_METABOLISM
MIR3152_3P, MIR3152_3P
 MODULE 145, MODULE 145
DESCARTES_MAIN_FETAL_LYMPHOID_CELLS, DESCARTES_MAIN_FETAL_LYMPHOID_CELLS
GOBP_CARDIAC_CHAMBER_MORPHOGENESIS, GOBP_CARDIAC_CHAMBER_MORPHOGENESIS
GOBP_REGULATION_OF_HISTONE_DEACETYLATION, GOBP_REGULATION_OF_HISTONE_DEACETYLATION
GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I
GOBP REGULATION OF OSSIFICATION, GOBP REGULATION OF OSSIFICATION
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_CYAN_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_CYAN_UP
MIR3155A_MIR3155B, MIR3155A_MIR3155B
EN1_01, EN1_01
GSE2706_R848_VS_LPS_2H_STIM_DC_DN, GSE2706_R848_VS_LPS_2H_STIM_DC_DN
 DESCARTES_FETAL_ADRENAL_LYMPHOID_CELLS, DESCARTES_FETAL_ADRENAL_LYMPHOID_CELLS
YANG_BREAST_CANCER_ESR1_BULK_DN, YANG_BREAST_CANCER_ESR1_BULK_DN
GOBP POSITIVE T CELL SELECTION, GOBP POSITIVE T CELL SELECTION
GOBP_REGULATION_OF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
GOMF I SMAD BINDING, GOMF I SMAD BINDING
GOBP_POSITIVE_REGULATION_OF_PHOSPHATASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_PHOSPHATASE_ACTIVITY
MIR484, MIR484
GSE2706_R848_VS_LPS_8H_STIM_DC_UP, GSE2706_R848_VS_LPS_8H_STIM_DC_UP
 DESCARTES_FETAL_PANCREAS_LYMPHOID_CELLS, DESCARTES_FETAL_PANCREAS_LYMPHOID_CELLS
 GOBP_CARDIAC_VENTRICLE_DEVELOPMENT, GOBP_CARDIAC_VENTRICLE_DEVELOPMENT
 GOBP_VENTRICULAR_SEPTUM_DEVELOPMENT, GOBP_VENTRICULAR_SEPTUM_DEVELOPMENT
 TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_LOBULAR_NORMAL_DN, TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_LOBULAR_NORMAL_DN
GOBP_T_CELL_SELECTION, GOBP_T_CELL_SELECTION
 GOBP MODIFIED AMINO ACID TRANSPORT, GOBP MODIFIED AMINO ACID TRANSPORT
 HARALAMBIEVA_PBMC_FLUARIX_AGE_50_74YO_CORR_WITH_28D_MEM_B_CELL_RESPONSE_AT_28DY_LATE_GENE_EXPR_INDIVID_GENE_MODELS_PRED_PEAK_B
 GOMF_SIGNALING_RECEPTOR_COMPLEX_ADAPTOR_ACTIVITY, GOMF_SIGNALING_RECEPTOR_COMPLEX_ADAPTOR_ACTIVITY
 GOBP_REGULATION_OF_HISTONE_H4_ACETYLATION, GOBP_REGULATION_OF_HISTONE_H4_ACETYLATION
 GOBP_REGULATION_OF_T_CELL_DIFFERENTIATION_IN_THYMUS, GOBP_REGULATION_OF_T_CELL_DIFFERENTIATION_IN_THYMUS
REACTOME SIGNALING BY BMP, REACTOME SIGNALING BY BMP
HAHTOLA_CTCL_PATHOGENESIS, HAHTOLA_CTCL_PATHOGENESIS
 GOBP HEART TRABECULA MORPHOGENESIS, GOBP HEART TRABECULA MORPHOGENESIS
 MIR6759_5P, MIR6759_5P
 PEDERSEN METASTASIS BY ERBB2 ISOFORM 5, PEDERSEN METASTASIS BY ERBB2 ISOFORM 5
 GOBP_CARDIAC_MUSCLE_TISSUE_MORPHOGENESIS, GOBP_CARDIAC_MUSCLE_TISSUE_MORPHOGENESIS
GOBP_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE, GOBP_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE
 MIZUSHIMA_AUTOPHAGOSOME_FORMATION, MIZUSHIMA_AUTOPHAGOSOME_FORMATION
GOBP_CD4_POSITIVE_OR_CD8_POSITIVE_ALPHA_BETA_T_CELL_LINEAGE_COMMITMENT, GOBP_CD4_POSITIVE_OR_CD8_POSITIVE_ALPHA_BETA_T_CELL_LINEAGE_COMMITMENT
 GOBP_METANEPHROS_DEVELOPMENT, GOBP_METANEPHROS_DEVELOPMENT
GOBP_BODY_MORPHOGENESIS, GOBP_BODY_MORPHOGENESIS
 GOBP CARDIAC EPITHELIAL TO MESENCHYMAL TRANSITION, GOBP CARDIAC EPITHELIAL TO MESENCHYMAL TRANSITION
 GOBP_MUSCLE_ORGAN_MORPHOGENESIS, GOBP_MUSCLE_ORGAN_MORPHOGENESIS
 KLEIN_PRIMARY_EFFUSION_LYMPHOMA_DN, KLEIN_PRIMARY_EFFUSION_LYMPHOMA_DN
 KEGG_PRIMARY_IMMUNODEFICIENCY, KEGG_PRIMARY_IMMUNODEFICIENCY
 ROETH_TERT_TARGETS_UP, ROETH_TERT_TARGETS_UP
 HP_FAILURE_TO_THRIVE_SECONDARY_TO_RECURRENT_INFECTIONS, HP_FAILURE_TO_THRIVE_SECONDARY_TO_RECURRENT_INFECTIONS
 MIR7848 3P, MIR7848 3P
 NIKOLSKY_BREAST_CANCER_15Q26_AMPLICON, NIKOLSKY_BREAST_CANCER_15Q26_AMPLICON
 GOBP_POSITIVE_REGULATION_OF_TISSUE_REMODELING, GOBP_POSITIVE_REGULATION_OF_TISSUE_REMODELING
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WANG_IMMORTALIZED_BY_HOXA9_AND_MEIS1_DN, WANG_IMMORTALIZED_BY_HOXA9_AND_MEIS1_DN

GOMF_ACTIVIN_BINDING, GOMF_ACTIVIN_BINDING

_72H_IL4_AND_ANTI_IL12_ACT_CD4_TCELL_UP, GSE17974_2.5H_VS_72H_IL4_AND_ANTI_IL12_ACT_CD4_TCELL_UF