GSE7548_DAY7_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_UP, GSE7548_DAY7_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_UP MODULE_332, MODULE_332 GAUSSMANN_MLL_AF4_FUSION_TARGETS_A_UP, GAUSSMANN_MLL_AF4_FUSION_TARGETS_A_UP GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN GRAHAM_CML_DIVIDING_VS_NORMAL_QUIESCENT_DN, GRAHAM_CML_DIVIDING_VS_NORMAL_QUIESCENT_DN GO_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS, GO_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS GSE1460_INTRATHYMIC_T_PROGENITOR_VS_THYMIC_STROMAL_CELL_DN, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_THYMIC_STROMAL_CELL_DN GSE7852_TREG_VS_TCONV_THYMUS_DN, GSE7852_TREG_VS_TCONV_THYMUS_DN HALLMARK_COAGULATION, HALLMARK_COAGULATION TGAGATT_MIR216, TGAGATT_MIR216 ZHU_CMV_8_HR_DN, ZHU_CMV_8_HR_DN GO_CILIARY_BASAL_BODY, GO_CILIARY_BASAL_BODY GO_INTEGRIN_MEDIATED_SIGNALING_PATHWAY, GO_INTEGRIN_MEDIATED_SIGNALING_PATHWAY ROMEDINU_IN_T_CELL_LINE_6H_DN, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_DN GO_HORMONE_RECEPTOR_BINDING, GO_HORMONE_RECEPTOR_BINDING PID_S1P_S1P3_PATHWAY, PID_S1P_S1P3_PATHWAY GO_CYTOKINE_BINDING, GO_CYTOKINE_BINDING GO POSITIVE REGULATION OF EPITHELIAL CELL MIGRATION, GO POSITIVE REGULATION OF EPITHELIAL CELL MIGRATION GO_RENAL_SYSTEM_VASCULATURE_DEVELOPMENT, GO_RENAL_SYSTEM_VASCULATURE_DEVELOPMENT SANA_TNF_SIGNALING_DN, SANA_TNF_SIGNALING_DN ZHU_CMV_ALL_DN, ZHU_CMV_ALL_DN GO_ESTROGEN_RECEPTOR_BINDING, GO_ESTROGEN_RECEPTOR_BINDING REACTOME_L1CAM_INTERACTIONS, REACTOME_L1CAM_INTERACTIONS PID LYMPH ANGIOGENESIS PATHWAY, PID LYMPH ANGIOGENESIS PATHWAY GO_EXTRACELLULAR_MATRIX_BINDING, GO_EXTRACELLULAR_MATRIX_BINDING SCHOEN_NFKB_SIGNALING, SCHOEN_NFKB_SIGNALING YORDY_RECIPROCAL_REGULATION_BY_ETS1_AND_SP100_UP, YORDY_RECIPROCAL_REGULATION_BY_ETS1_AND_SP100_UP GO_REGULATION_OF_EPITHELIAL_CELL_MIGRATION, GO_REGULATION_OF_EPITHELIAL_CELL_MIGRATION GO_CORECEPTOR_ACTIVITY, GO_CORECEPTOR_ACTIVITY GO_FIBRONECTIN_BINDING, GO_FIBRONECTIN_BINDING GO_CONTRACTILE_ACTIN_FILAMENT_BUNDLE_ASSEMBLY, GO_CONTRACTILE_ACTIN_FILAMENT_BUNDLE_ASSEMBLY GO_MESODERMAL_CELL_DIFFERENTIATION, GO_MESODERMAL_CELL_DIFFERENTIATION REACTOME CHONDROITIN SULFATE DERMATAN SULFATE METABOLISM, REACTOME CHONDROITIN SULFATE DERMATAN SULFATE METABOLISM LABBE_TARGETS_OF_TGFB1_AND_WNT3A_UP, LABBE_TARGETS_OF_TGFB1_AND_WNT3A_UP GO_CYTOKINE_PRODUCTION, GO_CYTOKINE_PRODUCTION COATES_MACROPHAGE_M1_VS_M2_DN, COATES_MACROPHAGE_M1_VS_M2_DN GO_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_MIGRATION, GO_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_MIGRATION GO_NEGATIVE_REGULATION_OF_LIPID_STORAGE, GO_NEGATIVE_REGULATION_OF_LIPID_STORAGE GO_COLLAGEN_BINDING, GO_COLLAGEN_BINDING GO_WNT_SIGNALING_PATHWAY_CALCIUM_MODULATING_PATHWAY, GO_WNT_SIGNALING_PATHWAY_CALCIUM_MODULATING_PATHWAY RELA_DN.V1_UP, RELA_DN.V1_UP GO_POSITIVE_REGULATION_OF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GO_POSITIVE_REGULATION_OF_VASCULAR_ENDOTHELIAL_GROWTH_FACTOI GO_GABA_RECEPTOR_BINDING, GO_GABA_RECEPTOR_BINDING GO_CHONDROITIN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS, GO_CHONDROITIN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS HENDRICKS_SMARCA4_TARGETS_DN, HENDRICKS_SMARCA4_TARGETS_DN PID_S1P_S1P1_PATHWAY, PID_S1P_S1P1_PATHWAY SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX5_UP, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX5_UP

GO_PROTEOGLYCAN_METABOLIC_PROCESS, GO_PROTEOGLYCAN_METABOLIC_PROCESS

GSE34205_RSV_VS_FLU_INF_INFANT_PBMC_UP, GSE34205_RSV_VS_FLU_INF_INFANT_PBMC_UP

GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN

GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP

GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN, GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN

GSE22025_PROGESTERONE_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_UP, GSE22025_PROGESTERONE_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_UP

GSE40666_UNTREATED_VS_IFNA_STIM_STAT4_KO_EFFECTOR_CD8_TCELL_90MIN_DN, GSE40666_UNTREATED_VS_IFNA_STIM_STAT4_KO_EFFECTOR_CD8_TCELL_90MIN_DN

GSE23321_CD8_STEM_CELL_MEMORY_VS_EFFECTOR_MEMORY_CD8_TCELL_UP, GSE23321_CD8_STEM_CELL_MEMORY_VS_EFFECTOR_MEMORY_CD8_TCELL_UP

KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE, KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE

GO_POSITIVE_REGULATION_OF_PEPTIDYL_TYROSINE_PHOSPHORYLATION, GO_POSITIVE_REGULATION_OF_PEPTIDYL_TYROSINE_PHOSPHORYLATION

GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_DN

GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP

GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_3H_DN, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_3H_DN

GSE2770 IL12 AND TGFB VS IL4 TREATED ACT CD4 TCELL 2H DN, GSE2770 IL12 AND TGFB VS IL4 TREATED ACT CD4 TCELL 2H DN

GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP

GSE16385_ROSIGLITAZONE_IL4_VS_IFNG_TNF_STIM_MACROPHAGE_UP, GSE16385_ROSIGLITAZONE_IL4_VS_IFNG_TNF_STIM_MACROPHAGE_UP

GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP, GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP

GSE32533_WT_VS_MIR17_KO_ACT_CD4_TCELL_DN, GSE32533_WT_VS_MIR17_KO_ACT_CD4_TCELL_DN

STEARMAN_LUNG_CANCER_EARLY_VS_LATE_UP, STEARMAN_LUNG_CANCER_EARLY_VS_LATE_UP

GSE24142_DN2_VS_DN3_THYMOCYTE_ADULT_DN, GSE24142_DN2_VS_DN3_THYMOCYTE_ADULT_DN

GSE25123_WT_VS_PPARG_KO_MACROPHAGE_UP, GSE25123_WT_VS_PPARG_KO_MACROPHAGE_UP

GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_UP

GSE41176_UNSTIM_VS_ANTI_IGM_STIM_BCELL_6H_UP, GSE41176_UNSTIM_VS_ANTI_IGM_STIM_BCELL_6H_UP

GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP, GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP

GSE2770 IL12 ACT VS ACT CD4 TCELL 2H DN, GSE2770 IL12 ACT VS ACT CD4 TCELL 2H DN

GSE14308_TH17_VS_INDUCED_TREG_UP, GSE14308_TH17_VS_INDUCED_TREG_UP

GSE25087_TREG_VS_TCONV_FETUS_UP, GSE25087_TREG_VS_TCONV_FETUS_UP

GO CHROMATIN SILENCING, GO CHROMATIN SILENCING