

GSE5589_LPS_AND_IL10_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_DN, GSE5589_LPS_AND_IL10_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_DN
GSE21774_CD62L_POS_CD56_BRIGHT_VS_CD62L_NEG_CD56_DIM_NK_CELL_DN, GSE21774_CD62L_POS_CD56_BRIGHT_VS_CD62L_NEG_CD56_DIM_NK_CELL_DN
GSE21380_NON_TFH_VS_TFH_CD4_TCELL_UP, GSE21380_NON_TFH_VS_TFH_CD4_TCELL_UP
GSE21546_WT_VS_SAP1A_KO_AND_ELK1_KO_DP_THYMOCYTES_DN, GSE21546_WT_VS_SAP1A_KO_AND_ELK1_KO_DP_THYMOCYTES_DN
GSE36392_TYPE_2_MYELOID_VS_NEUTROPHIL_IL25_TREATED_LUNG_UP, GSE36392_TYPE_2_MYELOID_VS_NEUTROPHIL_IL25_TREATED_LUNG_UP
TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_10D_DN, TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_10D_DN
GSE37605_FOXP3_FUSION_GFP_VS_IRES_GFP_TREG_C57BL6_DN, GSE37605_FOXP3_FUSION_GFP_VS_IRES_GFP_TREG_C57BL6_DN
GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_IRF8_KO_UP, GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_IRF8_KO_UP
GSE5589_WT_VS_IL6_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN, GSE5589_WT_VS_IL6_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN
GSE4748_CTRL_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP, GSE4748_CTRL_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP
GSE21670_STAT3_KO_VS_WT_CD4_TCELL_IL6_TREATED_DN, GSE21670_STAT3_KO_VS_WT_CD4_TCELL_IL6_TREATED_DN
GSE24142_DN2_VS_DN3_THYMOCYTE_ADULT_UP, GSE24142_DN2_VS_DN3_THYMOCYTE_ADULT_UP
MODULE_122, MODULE_122
GSE7831_1H_VS_4H_INFLUENZA_STIM_PDC_UP, GSE7831_1H_VS_4H_INFLUENZA_STIM_PDC_UP
GSE22443_NAIVE_VS_ACT_AND_IL2_TREATED_CD8_TCELL_DN, GSE22443_NAIVE_VS_ACT_AND_IL2_TREATED_CD8_TCELL_DN
SWEET_KRAS_TARGETS_UP, SWEET_KRAS_TARGETS_UP
SNF5_DN.V1_DN, SNF5_DN.V1_DN
HALLMARK_MYOGENESIS, HALLMARK_MYOGENESIS
GSE16385_ROSIGLITAZONE_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_DN, GSE16385_ROSIGLITAZONE_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_DN
BOYLAN_MULTIPLE_MYELOMA_PCA1_UP, BOYLAN_MULTIPLE_MYELOMA_PCA1_UP
MCDOWELL_ACUTE_LUNG_INJURY_UP, MCDOWELL_ACUTE_LUNG_INJURY_UP
GO_CELL_ADHESION_MOLECULE_BINDING, GO_CELL_ADHESION_MOLECULE_BINDING
GSE25502_WT_VS_KLF13_KO_THYMIC_MEMORY_LIKE_CD8_TCELL_DN, GSE25502_WT_VS_KLF13_KO_THYMIC_MEMORY_LIKE_CD8_TCELL_DN
NOTCH_DN.V1_DN, NOTCH_DN.V1_DN
GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP, GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP
GO_PEPTIDYL_TYROSINE_MODIFICATION, GO_PEPTIDYL_TYROSINE_MODIFICATION
GSE22229_UNTREATED_VS_IMMUNOSUPP_THERAPY_RENAL_TRANSPLANT_PATIENT_PBMC_DN, GSE22229_UNTREATED_VS_IMMUNOSUPP_THERAPY_RENAL_TRANSPLANT_PATIENT_PBMC_DN
GSE21380_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_UP, GSE21380_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_UP
GSE24210_IL35_TREATED_VS_RESTING_TREG_DN, GSE24210_IL35_TREATED_VS_RESTING_TREG_DN
BMI1_DN.V1_UP, BMI1_DN.V1_UP
HALLMARK_APICAL_JUNCTION, HALLMARK_APICAL_JUNCTION
BERENJENO_ROCK_SIGNALING_NOT_VIA_RHOA_DN, BERENJENO_ROCK_SIGNALING_NOT_VIA_RHOA_DN
GSE17721_CTRL_VS_POLYIC_1H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_1H_BMDC_DN
GSE21380_NON_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_UP, GSE21380_NON_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_UP
GO_REGULATION_OF_ENDOTHELIAL_CELL_PROLIFERATION, GO_REGULATION_OF_ENDOTHELIAL_CELL_PROLIFERATION
GSE3982_BASOPHIL_VS_CENT_MEMORY_CD4_TCELL_DN, GSE3982_BASOPHIL_VS_CENT_MEMORY_CD4_TCELL_DN
GO_EPITHELIAL_CELL_DEVELOPMENT, GO_EPITHELIAL_CELL_DEVELOPMENT
GO_ACTOMYOSIN, GO_ACTOMYOSIN
MODULE_199, MODULE_199
MODULE_259, MODULE_259
GSE21670_UNTREATED_VS_TGFB_TREATED_CD4_TCELL_DN, GSE21670_UNTREATED_VS_TGFB_TREATED_CD4_TCELL_DN
GO_ACTIN_FILAMENT_BUNDLE, GO_ACTIN_FILAMENT_BUNDLE
RADMACHER_AML_PROGNOSIS, RADMACHER_AML_PROGNOSIS
GO_DETOXIFICATION, GO_DETOXIFICATION
GSE15324_ELF4_KO_VS_WT_NAIVE_CD8_TCELL_DN, GSE15324_ELF4_KO_VS_WT_NAIVE_CD8_TCELL_DN
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_6, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_6
RAF_UP.V1_DN, RAF_UP.V1_DN
MODULE_85, MODULE_85
GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_TYROSINE_KINASE_ACTIVITY, GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_TYROSINE_KINASE_ACTIVITY
TONKS_TARGETS_OF_RUNX1_RUNXIT1_FUSION GRANULOCYTE_UP, TONKS_TARGETS_OF_RUNX1_RUNXIT1_FUSION GRANULOCYTE_UP
GO_MULTICELLULAR_ORGANISM_METABOLIC_PROCESS, GO_MULTICELLULAR_ORGANISM_METABOLIC_PROCESS
GO_SECOND_MESSENGER_MEDIATED_SIGNALING, GO_SECOND_MESSENGER_MEDIATED_SIGNALING
GO_POSITIVE_REGULATION_OF_CALCIUM_ION_TRANSPORT, GO_POSITIVE_REGULATION_OF_CALCIUM_ION_TRANSPORT
GSE22601_IMMATURE_CD4_SINGLE_POSITIVE_VS_DOUBLE_POSITIVE_THYMOCYTE_UP, GSE22601_IMMATURE_CD4_SINGLE_POSITIVE_VS_DOUBLE_POSITIVE_THYMOCYTE_UP
GO_REGULATION_OF_CALCIUM_ION_TRANSMEMBRANE_TRANSPORT, GO_REGULATION_OF_CALCIUM_ION_TRANSMEMBRANE_TRANSPORT
GO_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_PROLIFERATION, GO_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_PROLIFERATION
GO_GLOMERULUS_DEVELOPMENT, GO_GLOMERULUS_DEVELOPMENT
GO_CELL_SUBSTRATE_ADHERENS_JUNCTION_ASSEMBLY, GO_CELL_SUBSTRATE_ADHERENS_JUNCTION_ASSEMBLY
GO_ADHERENS_JUNCTION_ORGANIZATION, GO_ADHERENS_JUNCTION_ORGANIZATION
GSE7568_CTRL_VS_24H_TGFB_TREATED_MACROPHAGES_WITH_IL4_AND_DEXAMETHASONE_DN, GSE7568_CTRL_VS_24H_TGFB_TREATED_MACROPHAGES_WITH_IL4_AND_DEXAMETHASONE_DN
GO_PROTEIN_ACTIVATION_CASCADE, GO_PROTEIN_ACTIVATION_CASCADE
GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_DN, GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_DN
LINDGREN_BLADDER_CANCER_HIGH_RECURRENCE, LINDGREN_BLADDER_CANCER_HIGH_RECURRENCE
GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_UP, GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_UP
GO_EXTRACELLULAR_MATRIX_STRUCTURAL_CONSTITUENT, GO_EXTRACELLULAR_MATRIX_STRUCTURAL_CONSTITUENT
TONKS_TARGETS_OF_RUNX1_RUNXIT1_FUSION_SUSTAINED_IN_MONOCYTE_UP, TONKS_TARGETS_OF_RUNX1_RUNXIT1_FUSION_SUSTAINED_IN_MONOCYTE_UP
ZHENG_GLIOBLASTOMA_PLASTICITY_DN, ZHENG_GLIOBLASTOMA_PLASTICITY_DN
GO_ACTIN_FILAMENT_BINDING, GO_ACTIN_FILAMENT_BINDING
GO_NEGATIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT, GO_NEGATIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT
GO_REGULATION_OF_RELEASE_OF_SEQUESTERED_CALCIUM_ION_INTO_CYTOSOL, GO_REGULATION_OF_RELEASE_OF_SEQUESTERED_CALCIUM_ION_INTO_CYTOSOL
GNF2_CDH11, GNF2_CDH11
LINDVALL_IMMORTALIZED_BY_TERT_UP, LINDVALL_IMMORTALIZED_BY_TERT_UP
LL_WILMS_TUMOR_VS_FETAL_KIDNEY_2_DN, LL_WILMS_TUMOR_VS_FETAL_KIDNEY_2_DN
VART_KSHV_INFECTION_ANGIOGENIC_MARKERS_DN, VART_KSHV_INFECTION_ANGIOGENIC_MARKERS_DN
REACTOME_COLLAGEN_FORMATION, REACTOME_COLLAGEN_FORMATION
GO_FILOPODIUM, GO_FILOPODIUM
TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_DUCTAL_NORMAL_UP, TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_DUCTAL_NORMAL_UP
GO_CHANNEL_REGULATOR_ACTIVITY, GO_CHANNEL_REGULATOR_ACTIVITY
BIOCARTA_INTRINSIC_PATHWAY, BIOCARTA_INTRINSIC_PATHWAY
GSE19772_HCMV_INFL_VS_HCMV_INF_MONOCYTES_AND_PI3K_INHIBITION_UP, GSE19772_HCMV_INFL_VS_HCMV_INF_MONOCYTES_AND_PI3K_INHIBITION_UP
GO_LYMPH_VESSEL_DEVELOPMENT, GO_LYMPH_VESSEL_DEVELOPMENT
GSE29949_MICROGLIA_VS_DC_BRAIN_DN, GSE29949_MICROGLIA_VS_DC_BRAIN_DN
WANG_BARRETTS_ESOPHAGUS_AND_ESOPHAGUS_CANCER_UP, WANG_BARRETTS_ESOPHAGUS_AND_ESOPHAGUS_CANCER_UP
GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_BCL6_HET_DN, GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_BCL6_HET_DN
LEE_LIVER_CANCER_ACOX1_UP, LEE_LIVER_CANCER_ACOX1_UP
GO_GROWTH_FACTOR_BINDING, GO_GROWTH_FACTOR_BINDING
GO_CALCIUM_MEDIATED_SIGNALING, GO_CALCIUM_MEDIATED_SIGNALING
GSE29618_PRE_VS_DAY7_POST_LAIV_FLU_VACCINE_PDC_UP, GSE29618_PRE_VS_DAY7_POST_LAIV_FLU_VACCINE_PDC_UP
GSE37319_WT_VS_RC3H1_KO_CD44LOW_CD8_TCELL_DN, GSE37319_WT_VS_RC3H1_KO_CD44LOW_CD8_TCELL_DN
GO_ADHERENS_JUNCTION_ASSEMBLY, GO_ADHERENS_JUNCTION_ASSEMBLY
MODULE_51, MODULE_51
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_16, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_16
GO_IGG_BINDING, GO_IGG_BINDING
GO_REGULATION_OF_SMOOTH_MUSCLE_CONTRACTION, GO_REGULATION_OF_SMOOTH_MUSCLE_CONTRACTION
IL21_UP.V1_UP, IL21_UP.V1_UP
GO_ANCHORED_COMPONENT_OF_EXTERNAL_SIDE_OF_PLASMA_MEMBRANE, GO_ANCHORED_COMPONENT_OF_EXTERNAL_SIDE_OF_PLASMA_MEMBRANE
BIOCARTA_PLATELETAPP_PATHWAY, BIOCARTA_PLATELETAPP_PATHWAY
GO_REGULATION_OF_CALCIUM_ION_TRANSPORT_INTO_CYTOSOL, GO_REGULATION_OF_CALCIUM_ION_TRANSPORT_INTO_CYTOSOL
GSE37532_TREG_VS_TCONV_CD4_TCELL_FROM_VISCERAL_ADIPOSE_TISSUE_UP, GSE37532_TREG_VS_TCONV_CD4_TCELL_FROM_VISCERAL_ADIPOSE_TISSUE_UP
ESC_V6.5_UP_LATE.V1_UP, ESC_V6.5_UP_LATE.V1_UP
GSE6681_DELETED_FOXP3_VS_WT_TREG_UP, GSE6681_DELETED_FOXP3_VS_WT_TREG_UP
GO_AUTONOMIC_NERVOUS_SYSTEM_DEVELOPMENT, GO_AUTONOMIC_NERVOUS_SYSTEM_DEVELOPMENT
GO_NEGATIVE_REGULATION_OF_ENDOTHELIAL_CELL_PROLIFERATION, GO_NEGATIVE_REGULATION_OF_ENDOTHELIAL_CELL_PROLIFERATION
GO_DEVELOPMENTAL_GROWTH_INVOLVED_IN_MORPHOGENESIS, GO_DEVELOPMENTAL_GROWTH_INVOLVED_IN_MORPHOGENESIS
WESTON_VEGFA_TARGETS, WESTON_VEGFA_TARGETS
TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_LOBULAR_NORMAL_DN, TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_LOBULAR_NORMAL_DN
TING_SILENCED_BY_DICER, TING_SILENCED_BY_DICER
GO_COMPLEX_OF_COLLAGEN_TRIMERS, GO_COMPLEX_OF_COLLAGEN_TRIMERS
NABA_BASEMENT_MEMBRANES, NABA_BASEMENT_MEMBRANES
HALLMARK_HEDGEHOG_SIGNALING, HALLMARK_HEDGEHOG_SIGNALING
GO_PHOTORECEPTOR_OUTER_SEGMENT, GO_PHOTORECEPTOR_OUTER_SEGMENT
GO_BLOOD_VESSEL_REMODELING, GO_BLOOD_VESSEL_REMODELING
MODULE_80, MODULE_80
BIOCARTA_VITCB_PATHWAY, BIOCARTA_VITCB_PATHWAY
GO_REGULATION_OF_CHEMOTAXIS, GO_REGULATION_OF_CHEMOTAXIS
GO_POSITIVE_REGULATION_OF_CATION_TRANSMEMBRANE_TRANSPORT, GO_POSITIVE_REGULATION_OF_CATION_TRANSMEMBRANE_TRANSPORT
GNF2_PTX3, GNF2_PTX3
GO_SECRETORY_GRANULE_MEMBRANE, GO_SECRETORY_GRANULE_MEMBRANE
RICKMAN_HEAD_AND_NECK_CANCER_D, RICKMAN_HEAD_AND_NECK_CANCER_D
GNF2_HPX, GNF2_HPX
GSE13522_WT_VS_IFNG_KO_SKIN_UP, GSE13522_WT_VS_IFNG_KO_SKIN_UP
GO_NITRIC_OXIDE_METABOLIC_PROCESS, GO_NITRIC_OXIDE_METABOLIC_PROCESS
GO_CRANIAL_NERVE_MORPHOGENESIS, GO_CRANIAL_NERVE_MORPHOGENESIS
GO_ANATOMICAL_STRUCTURE_ARRANGEMENT, GO_ANATOMICAL_STRUCTURE_ARRANGEMENT
GO_GANGLION_DEVELOPMENT, GO_GANGLION_DEVELOPMENT
GNF2_HPN, GNF2_HPN
GO_FOREBRAIN_CELL_MIGRATION, GO_FOREBRAIN_CELL_MIGRATION
GO_CALCIUM_MEDIATED_SIGNALING_USING_INTRACELLULAR_CALCIUM_SOURCE, GO_CALCIUM_MEDIATED_SIGNALING_USING_INTRACELLULAR_CALCIUM_SOURCE
YAMASHITA_LIVER_CANCER_STEM_CELL_DN, YAMASHITA_LIVER_CANCER_STEM_CELL_DN
GSE21379_TFH_VS_NON_TFH_CD4_TCELL_DN, GSE21379_TFH_VS_NON_TFH_CD4_TCELL_DN
GSE3203_HEALTHY_VS_INFLUENZA_INFECTED_LN_BCELL_DN, GSE3203_HEALTHY_VS_INFLUENZA_INFECTED_LN_BCELL_DN
NIELSEN_SCHWANNOMA_DN, NIELSEN_SCHWANNOMA_DN