

VS_MONOCYTE_UP, GSE7509_DC_VS_MONOCYTE_UP

GSE26343_UNSTIM_VS_LPS_STIM_MACROPHAGE_DN, GSE26343_UNSTIM_VS_LPS_STIM_MACROPHAGE_DN
KARLSSON_TGFB1_TARGETS_DN, KARLSSON_TGFB1_TARGETS_DN
GSE15930_STIM_VS_STIM_AND_IL12_48H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_IL12_48H_CD8_T_CELL_DN
GSE25123_WT_VS_PPARG_KO_MACROPHAGE_ROSIGLITAZONE_STIM_UP, GSE25123_WT_VS_PPARG_KO_MACROPHAGE_ROSIGLITAZONE_STIM_UP
AIZARANI_LIVER_C21_STELLATE_CELLS_1, AIZARANI_LIVER_C21_STELLATE_CELLS_1
GSE5589_LPS_VS_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_UP, GSE5589_LPS_VS_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_UP
GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_CORTEX_THYMUS_UP, GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_CORTEX_THYMUS_UP
TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_10D_DN, TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_10D_DN
GSE15215_CD2_POS_VS_NEG_PDC_UP, GSE15215_CD2_POS_VS_NEG_PDC_UP
GOCC_AZUROPHIL_Granule_Membrane, GOCC_AZUROPHIL_Granule_Membrane
GSE21063_WT_VS_NFATC1_KO_BCELL_UP, GSE21063_WT_VS_NFATC1_KO_BCELL_UP
PID_P53_DOWNSTREAM_PATHWAY, PID_P53_DOWNSTREAM_PATHWAY
GSE17721_POLYIC_VS_PAM3CSK4_1H_BMDC_DN, GSE17721_POLYIC_VS_PAM3CSK4_1H_BMDC_DN
GOBP_SPHINGOLIPID_METABOLIC_PROCESS, GOBP_SPHINGOLIPID_METABOLIC_PROCESS
ONDER_CDH1_TARGETS_1_UP, ONDER_CDH1_TARGETS_1_UP
GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN, GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN
GSE20715_0H_VS_6H_OZONE_LUNG_DN, GSE20715_0H_VS_6H_OZONE_LUNG_DN
GSE27859_MACROPHAGE_VS_DC_DN, GSE27859_MACROPHAGE_VS_DC_DN
HALLMARK_MYOGENESIS, HALLMARK_MYOGENESIS
GOBP_ORGANOPHOSPHATE_ESTER_TRANSPORT, GOBP_ORGANOPHOSPHATE_ESTER_TRANSPORT
REACTOME_DEGRADATION_OF_THE_EXTRACELLULAR_MATRIX, REACTOME_DEGRADATION_OF_THE_EXTRACELLULAR_MATRIX
GSE43955_1H_VS_42H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN, GSE43955_1H_VS_42H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN
GOMF_LIPID_TRANSPORTER_ACTIVITY, GOMF_LIPID_TRANSPORTER_ACTIVITY
HP_ABNORMALITY_OF_PANCREAS_PHYSIOLOGY, HP_ABNORMALITY_OF_PANCREAS_PHYSIOLOGY
GOBP_LIPOSACCHARIDE_METABOLIC_PROCESS, GOBP_LIPOSACCHARIDE_METABOLIC_PROCESS
GSE19923_WT_VS_HEB_KO_DP_THYMOCYTE_UP, GSE19923_WT_VS_HEB_KO_DP_THYMOCYTE_UP
DACOSTA_UV_RESPONSE_VIA_ERCC3_COMMON_UP, DACOSTA_UV_RESPONSE_VIA_ERCC3_COMMON_UP
LI_WILMS_TUMOR_VS_FETAL_KIDNEY_2_DN, LI_WILMS_TUMOR_VS_FETAL_KIDNEY_2_DN
GOBP_POSITIVE_REGULATION_OF_MACROAUTOPHAGY, GOBP_POSITIVE_REGULATION_OF_MACROAUTOPHAGY
GSE22443_NAIVE_VS_ACT_AND_IL2_TREATED_CD8_TCELL_DN, GSE22443_NAIVE_VS_ACT_AND_IL2_TREATED_CD8_TCELL_DN
GOMF_COLLAGEN_BINDING, GOMF_COLLAGEN_BINDING
WEIGEL_OXIDATIVE_STRESS_BY_HNE_AND_TBH, WEIGEL_OXIDATIVE_STRESS_BY_HNE_AND_TBH
GOBP_ARACHIDONIC_ACID_METABOLIC_PROCESS, GOBP_ARACHIDONIC_ACID_METABOLIC_PROCESS
REACTOME_INTEGRIN_CELL_SURFACE_INTERACTIONS, REACTOME_INTEGRIN_CELL_SURFACE_INTERACTIONS
REACTOME_ECM_PROTEOGLYCANS, REACTOME_ECM_PROTEOGLYCANS
DESCARTES_FETAL_KIDNEY_STROMAL_CELLS, DESCARTES_FETAL_KIDNEY_STROMAL_CELLS
KEGG_DILATED_CARDIOMYOPATHY, KEGG_DILATED_CARDIOMYOPATHY
YANG_BREAST_CANCER_ESR1_LASER_UP, YANG_BREAST_CANCER_ESR1_LASER_UP
GSE45365_BCELL_VS_CD8_TCELL_DN, GSE45365_BCELL_VS_CD8_TCELL_DN
REACTOME_COLLAGEN_DEGRADATION, REACTOME_COLLAGEN_DEGRADATION
VANASSE_BCL2_TARGETS_UP, VANASSE_BCL2_TARGETS_UP
GOMF_STEROL_TRANSPORTER_ACTIVITY, GOMF_STEROL_TRANSPORTER_ACTIVITY
GSE21670_TGFB_VS_IL6_TREATED_CD4_TCELL_UP, GSE21670_TGFB_VS_IL6_TREATED_CD4_TCELL_UP
GSE36476_YOUNG_VS_OLD_DONOR_MEMORY_CD4_TCELL_DN, GSE36476_YOUNG_VS_OLD_DONOR_MEMORY_CD4_TCELL_DN
GOCC_COLLAGEN_TRIMER, GOCC_COLLAGEN_TRIMER
MODULE_410, MODULE_410
GOBP_NEGATIVE_REGULATION_OF_GLYCOPROTEIN_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_GLYCOPROTEIN_METABOLIC_PROCESS
GSE40274_IRF4_VS_FOXP3_AND_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_IRF4_VS_FOXP3_AND_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GOCC_PROTEIN_LIPID_COMPLEX, GOCC_PROTEIN_LIPID_COMPLEX
BIOCARTA_BARRESTIN_PATHWAY, BIOCARTA_BARRESTIN_PATHWAY
GOBP_SYNAPSE_PRUNING, GOBP_SYNAPSE_PRUNING
RICKMAN_HEAD_AND_NECK_CANCER_F, RICKMAN_HEAD_AND_NECK_CANCER_F
GOBP_STEROID_CATABOLIC_PROCESS, GOBP_STEROID_CATABOLIC_PROCESS
MIKKELSEN_MCV6_LCP_WITH_H3K27ME3, MIKKELSEN_MCV6_LCP_WITH_H3K27ME3
HP_XANTHELASMA, HP_XANTHELASMA
LY_AGING_MIDDLE_UP, LY_AGING_MIDDLE_UP
HP_ABNORMALITY_OF_THE_DISTAL_PHALANX_OF_THE_5TH_FINGER, HP_ABNORMALITY_OF_THE_DISTAL_PHALANX_OF_THE_5TH_FINGER
KOBAYASHI_RESPONSE_TO_ROMIDEPSIN, KOBAYASHI_RESPONSE_TO_ROMIDEPSIN
GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_DERIVED_FOAM_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_DERIVED_FOAM_CELL_DIFFERENTIATION
WP_MIRNA_TARGETS_IN_ECM_AND_MEMBRANE_RECEPTORS, WP_MIRNA_TARGETS_IN_ECM_AND_MEMBRANE_RECEPTORS
DESCARTES_FETAL_PANCREAS_CCL19_CCL21_POSITIVE_CELLS, DESCARTES_FETAL_PANCREAS_CCL19_CCL21_POSITIVE_CELLS
GOCC_TRIGLYCERIDE_RICH_PLASMA_LIPOPROTEIN_PARTICLE, GOCC_TRIGLYCERIDE_RICH_PLASMA_LIPOPROTEIN_PARTICLE
GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_STIMULUS, GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_VASCULAR_ENDOTHELIAL_GRO