

G_TREATED_ENDOTHELIAL_CELL_DN, GSE3920_UNTREATED_VS_IFNG_TREATED_ENDOTHELIAL_CELL_DN

- GSE3920_IFNB_VS_IFNG_TREATED_ENDOTHELIAL_CELL_DN, GSE3920_IFNB_VS_IFNG_TREATED_ENDOTHELIAL_CELL_DN
- GSE27859_MACROPHAGE_VS_CD11C_INT_F480_HI_MACROPHAGE_DN, GSE27859_MACROPHAGE_VS_CD11C_INT_F480_HI_MACROPHAGE_DN
- GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4MID_BCELL_DN, GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4MID_BCELL_DN
- GSE17721_POLYIC_VS_CPG_1H_BMDC_UP, GSE17721_POLYIC_VS_CPG_1H_BMDC_UP
- GSE6092_CTRL_VS_BORRELIA_BIRGDOFERI_INF_ENDOTHELIAL_CELL_UP, GSE6092_CTRL_VS_BORRELIA_BIRGDOFERI_INF_ENDOTHELIAL_CELL_UP
- GSE21678_WT_VS_FOXO1_FOXO3_KO_TREG_DN, GSE21678_WT_VS_FOXO1_FOXO3_KO_TREG_DN
- MODULE_257, MODULE_257
- GSE27434_WT_VS_DNMT1_KO_TREG_UP, GSE27434_WT_VS_DNMT1_KO_TREG_UP
- GSE40274_CTRL_VS_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_CTRL_VS_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
- GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_UP
- GSE22611_UNSTIM_VS_6H_MDP_STIM_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP, GSE22611_UNSTIM_VS_6H_MDP_STIM_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP
- REACTOME_PHOSPHOLIPID_METABOLISM, REACTOME_PHOSPHOLIPID_METABOLISM
- GSE5589_WT_VS_IL6_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN, GSE5589_WT_VS_IL6_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN
- GO_POSTTRANSCRIPTIONAL_GENE_SILENCING, GO_POSTTRANSCRIPTIONAL_GENE_SILENCING
- GSE8685_IL15_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP, GSE8685_IL15_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP
- REACTOME_REGULATORY_RNA_PATHWAYS, REACTOME_REGULATORY_RNA_PATHWAYS
- ACAWNRNSRCGG_UNKNOWN, ACAWNRNSRCGG_UNKNOWN
- GGCNNMSMYNTTG_UNKNOWN, GGCNNMSMYNTTG_UNKNOWN
- GSE10240_IL17_VS_IL17_AND_IL22_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN, GSE10240_IL17_VS_IL17_AND_IL22_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN
- GO_PHOSPHATIDYLINOSITOL_BIOSYNTHETIC_PROCESS, GO_PHOSPHATIDYLINOSITOL_BIOSYNTHETIC_PROCESS
- GO_PHOSPHATIDYLINOSITOL_METABOLIC_PROCESS, GO_PHOSPHATIDYLINOSITOL_METABOLIC_PROCESS
- GCM_TINF2, GCM_TINF2
- REACTOME_MICRORNA_MIRNA_BIOGENESIS, REACTOME_MICRORNA_MIRNA_BIOGENESIS
- STARK_HYPPOCAMPUS_22Q11_DELETION_UP, STARK_HYPPOCAMPUS_22Q11_DELETION_UP
- GO_CELLULAR_RESPONSE_TO_DSRNA, GO_CELLULAR_RESPONSE_TO_DSRNA
- PAPASPYRIDONOS_UNSTABLE_ATEROSCLEROTIC_PLAQUE_UP, PAPASPYRIDONOS_UNSTABLE_ATEROSCLEROTIC_PLAQUE_UP
- REACTOME_PPARG_ACTIVATES_GENE_EXPRESSION, REACTOME_PPARG_ACTIVATES_GENE_EXPRESSION
- GSE25123_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_DAY10_UP, GSE25123_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_DAY10_UP
- GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_TREATED_MACROPHAGE_DN, GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_TREATED_MACROPHAGE_DN
- REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION, REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION
- GO_NEGATIVE_REGULATION_OF_AUTOPHAGY, GO_NEGATIVE_REGULATION_OF_AUTOPHAGY
- REACTOME_SYNTHESIS_OF_GLYCOSYLPHOSPHATIDYLINOSITOL_GPI, REACTOME_SYNTHESIS_OF_GLYCOSYLPHOSPHATIDYLINOSITOL_GPI
- GSE3982_EFF_MEMORY_VS_CENT_MEMORY_CD4_TCELL_UP, GSE3982_EFF_MEMORY_VS_CENT_MEMORY_CD4_TCELL_UP
- GO_PREASSEMBLY_OF_GPI_ANCHOR_IN_ER_MEMBRANE, GO_PREASSEMBLY_OF_GPI_ANCHOR_IN_ER_MEMBRANE
- PID_TXA2PATHWAY, PID_TXA2PATHWAY
- GO_MOVEMENT_IN_ENVIRONMENT_OF_OTHER_ORGANISM_INVOLVED_IN_SYMBIOTIC_INTERACTION, GO_MOVEMENT_IN_ENVIRONMENT_OF_OTHER_ORGANISM_INVOLVED_IN_SYMBIOTIC_INTERACTION
- GO_PURINE_DEOXYRIBONUCLEOTIDE_METABOLIC_PROCESS, GO_PURINE_DEOXYRIBONUCLEOTIDE_METABOLIC_PROCESS
- GO_PHOTORECEPTOR_OUTER_SEGMENT_MEMBRANE, GO_PHOTORECEPTOR_OUTER_SEGMENT_MEMBRANE
- FONTAINE_PAPILLARY_THYROID_CARCINOMA_UP, FONTAINE_PAPILLARY_THYROID_CARCINOMA_UP
- HEIDENBLAD_AMPLICON_12P11_12_UP, HEIDENBLAD_AMPLICON_12P11_12_UP
- GO_NEGATIVE_REGULATION_OF_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE, GO_NEGATIVE_REGULATION_OF_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE
- LEE_METASTASIS_AND_ALTERNATIVE_SPLICING_DN, LEE_METASTASIS_AND_ALTERNATIVE_SPLICING_DN