

S_IN_CULTURE_CD4POS_DC_UP, GSE339_EX_VIVO_VS_IN_CULTURE_CD4POS_DC_UP

- GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_UP, GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_UP
- GSE339_EX_VIVO_VS_IN_CULTURE_CD4CD8DN_DC_UP, GSE339_EX_VIVO_VS_IN_CULTURE_CD4CD8DN_DC_UP
- GSE17721_LPS_VS_PAM3CSK4_1H_BMDC_UP, GSE17721_LPS_VS_PAM3CSK4_1H_BMDC_UP
- GSE43955_1H_VS_42H_ACT_CD4_TCELL_WITH_TGFB_IL6_UP, GSE43955_1H_VS_42H_ACT_CD4_TCELL_WITH_TGFB_IL6_UP
- GSE35825_UNTREATED_VS_IFNG_STIM_MACROPHAGE_DN, GSE35825_UNTREATED_VS_IFNG_STIM_MACROPHAGE_DN
- GSE339_CD4POS_VS_CD8POS_DC_UP, GSE339_CD4POS_VS_CD8POS_DC_UP
- GSE12392_WT_VS_IFNB_KO_CD8A_NEG_SPLEEN_DC_DN, GSE12392_WT_VS_IFNB_KO_CD8A_NEG_SPLEEN_DC_DN
- GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_BCL6_HET_UP, GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_BCL6_HET_UP
- GSE45365_NK_CELL_VS_CD8A_DC_UP, GSE45365_NK_CELL_VS_CD8A_DC_UP
- GSE27786_LSK_VS_LIN_NEG_CELL_UP, GSE27786_LSK_VS_LIN_NEG_CELL_UP
- GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_DN, GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_DN
- GSE26488_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMOCYTE_DN, GSE26488_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMOCYTE_DN
- GSE7831_UNSTIM_VS_INFLUENZA_STIM_PDC_4H_DN, GSE7831_UNSTIM_VS_INFLUENZA_STIM_PDC_4H_DN
- GSE43955_1H_VS_42H_ACT_CD4_TCELL_UP, GSE43955_1H_VS_42H_ACT_CD4_TCELL_UP
- GSE339_CD8POS_VS_CD4CD8DN_DC_DN, GSE339_CD8POS_VS_CD4CD8DN_DC_DN
- GSE43955_10H_VS_30H_ACT_CD4_TCELL_WITH_TGFB_IL6_UP, GSE43955_10H_VS_30H_ACT_CD4_TCELL_WITH_TGFB_IL6_UP
- HP_ABNORMAL_FEMORAL_NECK_HEAD_MORPHOLOGY, HP_ABNORMAL_FEMORAL_NECK_HEAD_MORPHOLOGY
- HP_SLOW_PROGRESSION, HP_SLOW_PROGRESSION
- GSE28783_ANTI_MIR33_VS_UNTREATED_ATHEROSCLEROSIS_MACROPHAGE_UP, GSE28783_ANTI_MIR33_VS_UNTREATED_ATHEROSCLEROSIS_MACROPHAGE_UP
- GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_DN, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_DN
- HP_CIRRHOSIS, HP_CIRRHOSIS
- GOBP_NEGATIVE_REGULATION_OF_NEURON_PROJECTION_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_NEURON_PROJECTION_DEVELOPMENT
- GOBP_INOSITOL_LIPID_MEDIATED_SIGNALING, GOBP_INOSITOL_LIPID_MEDIATED_SIGNALING
- GOBP_BONE_DEVELOPMENT, GOBP_BONE_DEVELOPMENT
- MSX1_01, MSX1_01
- REACTOME_PI3K_AKT_SIGNALING_IN_CANCER, REACTOME_PI3K_AKT_SIGNALING_IN_CANCER
- LEE_CALORIE_RESTRICTION_MUSCLE_DN, LEE_CALORIE_RESTRICTION_MUSCLE_DN
- DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_MATURE_NEURONS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_MATURE_NEURONS
- HP_SPASTIC_GAIT, HP_SPASTIC_GAIT
- MIR4715_3P, MIR4715_3P
- GSE22443_NAIVE_VS_ACT_AND_IL2_TREATED_CD8_TCELL_UP, GSE22443_NAIVE_VS_ACT_AND_IL2_TREATED_CD8_TCELL_UP
- GOBP_ARTERY_DEVELOPMENT, GOBP_ARTERY_DEVELOPMENT
- REACTOME_SPHINGOLIPID_DE_NOVO_BIOSYNTHESIS, REACTOME_SPHINGOLIPID_DE_NOVO_BIOSYNTHESIS
- GSE40274_CTRL_VS_FOXP3_AND_SATB1_TRANSDUCEd_ACTIVATED_CD4_TCELL_DN, GSE40274_CTRL_VS_FOXP3_AND_SATB1_TRANSDUCEd_ACTIVATED_CD4_TCELL_DN
- MIR663B, MIR663B
- GSE33292_DN3_THYMOCYTE_VS_TCF1_KO_TCELL_LYMPHOMA_DN, GSE33292_DN3_THYMOCYTE_VS_TCF1_KO_TCELL_LYMPHOMA_DN
- GOBP_EXTRACELLULAR_MATRIX_DISASSEMBLY, GOBP_EXTRACELLULAR_MATRIX_DISASSEMBLY
- GOBP_POSITIVE_REGULATION_OF_PRI_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II, GOBP_POSITIVE_REGULATION_OF_PRI_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II
- GOBP_SUBSTRATE_DEPENDENT_CELL_MIGRATION, GOBP_SUBSTRATE_DEPENDENT_CELL_MIGRATION
- GOBP_ENTEROENDOCRINE_CELL_DIFFERENTIATION, GOBP_ENTEROENDOCRINE_CELL_DIFFERENTIATION
- HP_RECURRENT_OPPORTUNISTIC_INFECTIONS, HP_RECURRENT_OPPORTUNISTIC_INFECTIONS
- HP_OPPORTUNISTIC_INFECTION, HP_OPPORTUNISTIC_INFECTION
- GOBP_NEGATIVE_REGULATION_OF_LIPID_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_LIPID_CATABOLIC_PROCESS
- GOBP_CORPUS_CALLOSUM_DEVELOPMENT, GOBP_CORPUS_CALLOSUM_DEVELOPMENT
- GOBP_SINGLE_STRANDED_VIRAL_RNA_REPLICATION_VIA_DOUBLE_STRANDED_DNA_INTERMEDIATE, GOBP_SINGLE_STRANDED_VIRAL_RNA_REPLICATION_VIA_DOUBLE_STRANDED_DNA_INTERMEDIATE
- GOBP_NEGATIVE_REGULATION_OF_SINGLE_STRANDED_VIRAL_RNA_REPLICATION_VIA_DOUBLE_STRANDED_DNA_INTERMEDIATE, GOBP_NEGATIVE_REGULATION_OF_SINGLE_STRANDED_VIRAL_RNA_REPLICATION_VIA_DOUBLE_STRANDED_DNA_INTERMEDIATE
- HP_OVERGROWTH_OF_EXTERNAL_GENITALIA, HP_OVERGROWTH_OF_EXTERNAL_GENITALIA
- CERIBELLI_PROMOTERS_INACTIVE_AND_BOUND_BY_NFY, CERIBELLI_PROMOTERS_INACTIVE_AND_BOUND_BY_NFY
- HP_PROPORTIONATE_TALL_STATUTE, HP_PROPORTIONATE_TALL_STATUTE
- GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_REGULATORY_REGION_DNA_BINDING, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_REGULATORY_REGION_DNA_BINDING
- LIU_CDX2_TARGETS_UP, LIU_CDX2_TARGETS_UP
- GOBP_ASSOCIATIVE_LEARNING, GOBP_ASSOCIATIVE_LEARNING
- HP_POSTERIOR_PITUITARY_DYSGENESIS, HP_POSTERIOR_PITUITARY_DYSGENESIS
- GOBP_HIGH_DENSITY_LIPOPROTEIN_PARTICLE_ASSEMBLY, GOBP_HIGH_DENSITY_LIPOPROTEIN_PARTICLE_ASSEMBLY
- GOMF_TRIGLYCERIDE_LIPASE_ACTIVITY, GOMF_TRIGLYCERIDE_LIPASE_ACTIVITY
- KEGG_ASCORBATE_AND_ALDARATE_METABOLISM, KEGG_ASCORBATE_AND_ALDARATE_METABOLISM
- HP_LIMITED_NECK_RANGE_OF_MOTION, HP_LIMITED_NECK_RANGE_OF_MOTION
- GOBP_ASCENDING_AORTA_DEVELOPMENT, GOBP_ASCENDING_AORTA_DEVELOPMENT
- LIU_CDX2_TARGETS_DN, LIU_CDX2_TARGETS_DN
- GOBP_CARDIAC_NEURAL_CREST_CELL_DEVELOPMENT_INVOLVED_IN_OUTFLOW_TRACT_MORPHOGENESIS, GOBP_CARDIAC_NEURAL_CREST_CELL_DEVELOPMENT_INVOLVED_IN_OUTFLOW_TRACT_MORPHOGENESIS
- HP_YELLOW_WHITE_LESIONS_OF_THE_MACULA, HP_YELLOW_WHITE_LESIONS_OF_THE_MACULA
- HP_PHOTSENSITIVE_MYOCLONIC_SEIZURE, HP_PHOTSENSITIVE_MYOCLONIC_SEIZURE
- GOBP_COMMA_SHAPED_BODY_MORPHOGENESIS, GOBP_COMMA_SHAPED_BODY_MORPHOGENESIS
- HP_CYANOTIC_EPISODE, HP_CYANOTIC_EPISODE