

WT_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_UP, GSE46606_IRF4HIGH_VS_WT_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_UP

- GSE24726_WT_VS_E2_2_KO_PDC_UP, GSE24726_WT_VS_E2_2_KO_PDC_UP
- GSE24726_WT_VS_E2_2_KO_PDC_DAY6_POST_DELETION_UP, GSE24726_WT_VS_E2_2_KO_PDC_DAY6_POST_DELETION_UP
- GSE6259_FLT3L_INDUCED_33D1_POS_DC_VS_BCELL_DN, GSE6259_FLT3L_INDUCED_33D1_POS_DC_VS_BCELL_DN
- GSE42724_MEMORY_BCELL_VS_PLASMABLAST_DN, GSE42724_MEMORY_BCELL_VS_PLASMABLAST_DN
- GSE10240_IL17_VS_IL17_AND_IL22_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_UP, GSE10240_IL17_VS_IL17_AND_IL22_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_UP
- GSE33424_CD161_HIGH_VS_NEG_CD8_TCELL_UP, GSE33424_CD161_HIGH_VS_NEG_CD8_TCELL_UP
- GSE22229_RENAL_TRANSPLANT_VS_HEALTHY_PBMCDN, GSE22229_RENAL_TRANSPLANT_VS_HEALTHY_PBMCDN
- GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_UP, GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_UP
- HP_APLASIA_HYPOPLASIA_OF_THE_EAR, HP_APLASIA_HYPOPLASIA_OF_THE_EAR
- GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_UP, GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_UP
- GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP
- GOBP_NEGATIVE_REGULATION_OF_DNA_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_DNA_METABOLIC_PROCESS
- MIR1285_3P, MIR1285_3P
- GSE28737_WT_VS_BCL6_HET_MARGINAL_ZONE_BCELL_UP, GSE28737_WT_VS_BCL6_HET_MARGINAL_ZONE_BCELL_UP
- MIR5189_5P, MIR5189_5P
- MIR1915_3P, MIR1915_3P
- MIR1260A_MIR1260B, MIR1260A_MIR1260B
- MIR3190_3P, MIR3190_3P
- GOMF_TRANSFERASE_ACTIVITY_TRANSFERRING_HEXOSYL_GROUPS, GOMF_TRANSFERASE_ACTIVITY_TRANSFERRING_HEXOSYL_GROUPS
- GOMF_REPRESSING_TRANSCRIPTION_FACTOR_BINDING, GOMF_REPRESSING_TRANSCRIPTION_FACTOR_BINDING
- CHIANG_LIVER_CANCER_SUBCLASS_CTNNB1_UP, CHIANG_LIVER_CANCER_SUBCLASS_CTNNB1_UP
- HP_SHORT_FOOT, HP_SHORT_FOOT
- GOBP_PROTEIN_O_LINKED_GLYCOSYLATION, GOBP_PROTEIN_O_LINKED_GLYCOSYLATION
- GOBP_NEGATIVE_REGULATION_OF_DNA_REPAIR, GOBP_NEGATIVE_REGULATION_OF_DNA_REPAIR
- HP_POSTAXIAL_FOOT_POLYDACTYLY, HP_POSTAXIAL_FOOT_POLYDACTYLY
- GOBP_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS
- GOCC_GOLGI_CISTERNA_MEMBRANE, GOCC_GOLGI_CISTERNA_MEMBRANE
- GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS, GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS
- GOBP_SPERM_AXONEME_ASSEMBLY, GOBP_SPERM_AXONEME_ASSEMBLY
- GOMF_UDP_GLYCOSYLTRANSFERASE_ACTIVITY, GOMF_UDP_GLYCOSYLTRANSFERASE_ACTIVITY
- GSE29618_PRE_VS_DAY7_FLU_VACCINE_PDC_UP, GSE29618_PRE_VS_DAY7_FLU_VACCINE_PDC_UP
- GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS
- WP_LNCRNA_INVOLVEMENT_IN_CANONICAL_WNT_SIGNALING_AND_COLORECTAL_CANCER, WP_LNCRNA_INVOLVEMENT_IN_CANONICAL_WNT_SIGNALING_AND_COLORECTAL_CANCER
- GOMF_MANNOSYLTRANSFERASE_ACTIVITY, GOMF_MANNOSYLTRANSFERASE_ACTIVITY
- GOBP_DERMATAN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS, GOBP_DERMATAN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS
- HP_PETERS_ANOMALY, HP_PETERS_ANOMALY
- MIR6763_5P, MIR6763_5P
- GOBP_POSITIVE_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLATION
- HP_APLASIA_HYPOPLASIA_OF_THE_TIBIA, HP_APLASIA_HYPOPLASIA_OF_THE_TIBIA
- HOEK_MONOCYTE_2011_2012_TIV_ADULT_1DY_DN, HOEK_MONOCYTE_2011_2012_TIV_ADULT_1DY_DN
- REACTOME_SYNTHESIS_OF_PIPS_AT_THE_GOLGI_MEMBRANE, REACTOME_SYNTHESIS_OF_PIPS_AT_THE_GOLGI_MEMBRANE
- HP_ABNORMALITY_OF_THE_PHALANGES_OF_THE_HALLUX, HP_ABNORMALITY_OF_THE_PHALANGES_OF_THE_HALLUX
- GOBP_RESPONSE_TO_ISCHEMIA, GOBP_RESPONSE_TO_ISCHEMIA
- GOBP_CHONDROITIN_SULFATE_BIOSYNTHETIC_PROCESS, GOBP_CHONDROITIN_SULFATE_BIOSYNTHETIC_PROCESS
- GSE13762_CTRL_VS_125_VITAMIND_DAY5_DC_DN, GSE13762_CTRL_VS_125_VITAMIND_DAY5_DC_DN
- GOBP_VENOUS_BLOOD_VESSEL_DEVELOPMENT, GOBP_VENOUS_BLOOD_VESSEL_DEVELOPMENT
- GOMF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY, GOMF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY
- GOBP_VENOUS_BLOOD_VESSEL_MORPHOGENESIS, GOBP_VENOUS_BLOOD_VESSEL_MORPHOGENESIS
- REACTOME_PLATELET_CALCIIUM_HOMEOSTASIS, REACTOME_PLATELET_CALCIIUM_HOMEOSTASIS
- GOMF_GDP DISSOCIATION INHIBITOR ACTIVITY, GOMF_GDP DISSOCIATION INHIBITOR ACTIVITY
- HP_BILIARY_ATRESIA, HP_BILIARY_ATRESIA
- GOBP_ACID_SECRETION, GOBP_ACID_SECRETION
- GOBP_EPITHELIAL_FLUID_TRANSPORT, GOBP_EPITHELIAL_FLUID_TRANSPORT
- REACTOME_CHONDROITIN_SULFATE_BIOSYNTHESIS, REACTOME_CHONDROITIN_SULFATE_BIOSYNTHESIS