

CELL_VS_NEUTROPHIL_UP, GSE27786_BCELL_VS_NEUTROPHIL_UP

GSE27786_CD8_TCELL_VS_ERYTHROBLAST_UP, GSE27786_CD8_TCELL_VS_ERYTHROBLAST_UP
GSE27786_CD8_TCELL_VS_MONO_MAC_UP, GSE27786_CD8_TCELL_VS_MONO_MAC_UP
GSE27786_NKTCELL_VS_MONO_MAC_UP, GSE27786_NKTCELL_VS_MONO_MAC_UP
GSE27786_NKTCELL_VS_NEUTROPHIL_UP, GSE27786_NKTCELL_VS_NEUTROPHIL_UP
GSE27786_CD4_TCELL_VS_MONO_MAC_UP, GSE27786_CD4_TCELL_VS_MONO_MAC_UP
GSE27786_CD4_TCELL_VS_NEUTROPHIL_UP, GSE27786_CD4_TCELL_VS_NEUTROPHIL_UP
GSE2197_IMMUNOSUPPRESSIVE_DNA_VS_UNTREATEDIN_DC_DN, GSE2197_IMMUNOSUPPRESSIVE_DNA_VS_UNTREATEDIN_DC_DN
GSE27786_BCELL_VS_NKCELL_UP, GSE27786_BCELL_VS_NKCELL_UP
GSE20366_EX_VIVO_VS_HOMEOSTATIC_CONVERSION_NAIVE_CD4_TCELL_UP, GSE20366_EX_VIVO_VS_HOMEOSTATIC_CONVERSION_NAIVE_CD4_TCELL_UP
GSE17721_POLYIC_VS_CPG_24H_BMDC_UP, GSE17721_POLYIC_VS_CPG_24H_BMDC_UP
GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_UP, GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_UP
GSE36392_TYPE_2_MYELOID_VS_EOSINOPHIL_IL25_TREATED_LUNG_UP, GSE36392_TYPE_2_MYELOID_VS_EOSINOPHIL_IL25_TREATED_LUNG_UP
GSE32423_IL7_VS_IL7_IL4_MEMORY_CD8_TCELL_DN, GSE32423_IL7_VS_IL7_IL4_MEMORY_CD8_TCELL_DN
GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_UP
GCM_CASP2, GCM_CASP2
GO_LIGAND_DEPENDENT_NUCLEAR_RECEPTOR_TRANSCRIPTION_COACTIVATOR_ACTIVITY, GO_LIGAND_DEPENDENT_NUCLEAR_RECEPTOR_TRANSCRIPTION_COACTIVATOR_ACTIVITY
GO_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE, GO_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE
GSE22025_UNTREATED_VS_TGFB1_TREATED_CD4_TCELL_DN, GSE22025_UNTREATED_VS_TGFB1_TREATED_CD4_TCELL_DN
GSE13306_RA_VS_UNTREATED_TCONV_DN, GSE13306_RA_VS_UNTREATED_TCONV_DN
GSE42021_CD24INT_TREG_VS_CD24INT_TCONV_THYMUS_DN, GSE42021_CD24INT_TREG_VS_CD24INT_TCONV_THYMUS_DN
GRE_C, GRE_C
GO_PLACENTA_DEVELOPMENT, GO_PLACENTA_DEVELOPMENT
GO_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION, GO_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION
GO_NEGATIVE_REGULATION_OF_VIRAL_PROCESS, GO_NEGATIVE_REGULATION_OF_VIRAL_PROCESS
GO_BRANCHING_INVOLVED_IN_SALIVARY_GLAND_MORPHOGENESIS, GO_BRANCHING_INVOLVED_IN_SALIVARY_GLAND_MORPHOGENESIS
GO_EMBRYONIC_PLACENTA_DEVELOPMENT, GO_EMBRYONIC_PLACENTA_DEVELOPMENT
RYTGCNNRGNAAC_MIF1_01, RYTGCNNRGNAAC_MIF1_01
GO_FERROUS_IRON_BINDING, GO_FERROUS_IRON_BINDING
GO_STEROL_BINDING, GO_STEROL_BINDING
GO_VENTRAL_SPINAL_CORD_DEVELOPMENT, GO_VENTRAL_SPINAL_CORD_DEVELOPMENT
GO_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_I_PROMOTER, GO_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_I_PROMOTER
GO_BIOTIN_METABOLIC_PROCESS, GO_BIOTIN_METABOLIC_PROCESS
YRCCAKNNGNCGC_UNKNOWN, YRCCAKNNGNCGC_UNKNOWN
GO_MISMATCH_REPAIR_COMPLEX, GO_MISMATCH_REPAIR_COMPLEX