

DARKZONE_GC_TONSIL_BCELL_DN, GSE12845_NAIVE_VS_DARKZONE_GC_TONSIL_BCELL_DN

- GSE12366_GC_BCELL_VS_PLASMA_CELL_UP, GSE12366_GC_BCELL_VS_PLASMA_CELL_UP
- GSE12845_IGD_NEG_BLOOD_VS_DARKZONE_GC_TONSIL_BCELL_DN, GSE12845_IGD_NEG_BLOOD_VS_DARKZONE_GC_TONSIL_BCELL_DN
- GSE20152_SPHK1_KO_VS_WT_HTNFA_OVERXPRESS_ANKLE_DN, GSE20152_SPHK1_KO_VS_WT_HTNFA_OVERXPRESS_ANKLE_DN
- GSE39152_BRAIN_VS_SPLEEN_CD103_NEG_MEMORY_CD8_TCELL_DN, GSE39152_BRAIN_VS_SPLEEN_CD103_NEG_MEMORY_CD8_TCELL_DN
- GSE12198_NK_VS_NK_ACT_EXPANSION_SYSTEM_DERIVED_NK_CELL_DN, GSE12198_NK_VS_NK_ACT_EXPANSION_SYSTEM_DERIVED_NK_CELL_DN
- GSE1460_INTRATHYMIC_T_PROGENITOR_VS_CD4_THYMOCYTE_UP, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_CD4_THYMOCYTE_UP
- GSE27670_BLIMP1_VS_LMP1_TRANSDUCED_GC_BCELL_DN, GSE27670_BLIMP1_VS_LMP1_TRANSDUCED_GC_BCELL_DN
- GSE3982_EOSINOPHIL_VS_NKCELL_DN, GSE3982_EOSINOPHIL_VS_NKCELL_DN
- GSE29618_PRE_VS_DAY7_POST_LAIV_FLU_VACCINE_MONOCYTE_DN, GSE29618_PRE_VS_DAY7_POST_LAIV_FLU_VACCINE_MONOCYTE_DN
- GSE21033_CTRL_VS_POLYIC_STIM_DC_12H_DN, GSE21033_CTRL_VS_POLYIC_STIM_DC_12H_DN
- GSE45739_UNSTIM_VS_ACD3_ACD28_STIM_NRAS_KO_CD4_TCELL_UP, GSE45739_UNSTIM_VS_ACD3_ACD28_STIM_NRAS_KO_CD4_TCELL_UP
- GSE5679_PPARG_LIGAND_ROSIGLITAZONE_VS_RARA_AGONIST_AM580_TREATED_DC_UP, GSE5679_PPARG_LIGAND_ROSIGLITAZONE_VS_RARA_AGONIST_AM580_TREATED_DC_UP
- GSE360_DC_VS_MAC_B_MALAYI_HIGH_DOSE_DN, GSE360_DC_VS_MAC_B_MALAYI_HIGH_DOSE_DN
- GAVIN_FOXP3_TARGETS_CLUSTER_T7, GAVIN_FOXP3_TARGETS_CLUSTER_T7
- GSE13493_CD4INTCD8POS_VS_CD8POS_THYMOCYTE_UP, GSE13493_CD4INTCD8POS_VS_CD8POS_THYMOCYTE_UP
- GSE12845_IGD_POS_BLOOD_VS_NAIVE_TONSIL_BCELL_UP, GSE12845_IGD_POS_BLOOD_VS_NAIVE_TONSIL_BCELL_UP
- REACTOME_RNA_POL_II_TRANSCRIPTION_PRE_INITIATION_AND_PROMOTER_OPENING, REACTOME_RNA_POL_II_TRANSCRIPTION_PRE_INITIATION_AND_PROMOTER_OPENING
- MYC_Q2, MYC_Q2
- BRCHAT_RESPONSE_TO_CAMPTOTHECIN_DN, BRCHAT_RESPONSE_TO_CAMPTOTHECIN_DN
- GSE2585_THYMIC_MACROPHAGE_VS_MTEC_UP, GSE2585_THYMIC_MACROPHAGE_VS_MTEC_UP
- GSE10240_IL22_VS_IL22_AND_IL17_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN, GSE10240_IL22_VS_IL22_AND_IL17_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN
- GO_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY, GO_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY
- GSE2585_THYMIC_DC_VS_MTEC_UP, GSE2585_THYMIC_DC_VS_MTEC_UP
- GSE360_L_DONOVANI_VS_L_MAJOR_DC_DN, GSE360_L_DONOVANI_VS_L_MAJOR_DC_DN
- GSE21360_PRIMARY_VS_SECONDARY_MEMORY_CD8_TCELL_DN, GSE21360_PRIMARY_VS_SECONDARY_MEMORY_CD8_TCELL_DN
- GSE18281_CORTEX_VS_MEDULLA_THYMUS_UP, GSE18281_CORTEX_VS_MEDULLA_THYMUS_UP
- GO_T_CELL_PROLIFERATION, GO_T_CELL_PROLIFERATION
- GO_NUCLEOSIDE_DIPHOSPHATE_METABOLIC_PROCESS, GO_NUCLEOSIDE_DIPHOSPHATE_METABOLIC_PROCESS
- AKL_HTLV1_INFECTION_DN, AKL_HTLV1_INFECTION_DN
- GENTILE_UV_RESPONSE_CLUSTER_D7, GENTILE_UV_RESPONSE_CLUSTER_D7
- KEGG_FRUCTOSE_AND_MANNOSSE_METABOLISM, KEGG_FRUCTOSE_AND_MANNOSSE_METABOLISM
- FONTAINE_FOLLICULAR_THYROID_ADENOMA_UP, FONTAINE_FOLLICULAR_THYROID_ADENOMA_UP
- GO_MATERNAL_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY, GO_MATERNAL_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY
- GO_IMP_METABOLIC_PROCESS, GO_IMP_METABOLIC_PROCESS
- GO_PEPTIDYL_PROLINE_MODIFICATION, GO_PEPTIDYL_PROLINE_MODIFICATION
- GO_POSITIVE_REGULATION_OF_DEPHOSPHORYLATION, GO_POSITIVE_REGULATION_OF_DEPHOSPHORYLATION
- LA_MEN1_TARGETS, LA_MEN1_TARGETS
- GO_ISOTYPE_SWITCHING, GO_ISOTYPE_SWITCHING
- MOOTHA_GLYCOLYSIS, MOOTHA_GLYCOLYSIS
- GO_IMP_BIOSYNTHETIC_PROCESS, GO_IMP_BIOSYNTHETIC_PROCESS
- GO_IMMUNOGLOBULIN_PRODUCTION_INVOLVED_IN_IMMUNOGLOBULIN_MEDIATED_IMMUNE_RESPONSE, GO_IMMUNOGLOBULIN_PRODUCTION_INVOLVED_IN_IMMUNOGLOBULIN_MEDIATED_IMMUNE_RESPONSE
- NIKOLSKY_BREAST_CANCER_22Q13_AMPLICON, NIKOLSKY_BREAST_CANCER_22Q13_AMPLICON
- GO_REGULATION_OF_ACTIN_FILAMENT_DEPOLYMERIZATION, GO_REGULATION_OF_ACTIN_FILAMENT_DEPOLYMERIZATION
- KIM_HYPOXIA, KIM_HYPOXIA