

MYELOMA_BY_TACI_DN, MOREAUX_MULTIPLE_MYELOMA_BY_TACI_DN

GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP
GSE22886_NAIVE_CD4_TCELL_VS_NEUTROPHIL_UP, GSE22886_NAIVE_CD4_TCELL_VS_NEUTROPHIL_UP
GSE10239_NAIVE_VS_MEMORY_CD8_TCELL_UP, GSE10239_NAIVE_VS_MEMORY_CD8_TCELL_UP
GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP
GSE2770_UNTREATED_VS_ACT_CD4_TCELL_2H_DN, GSE2770_UNTREATED_VS_ACT_CD4_TCELL_2H_DN
MORF_GSPT1, MORF_GSPT1
GSE5542_UNTREATED_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_24H_UP, GSE5542_UNTREATED_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_24H_UP
GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_UP, GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_UP
ERB2_UP.V1_DN, ERB2_UP.V1_DN
GSE2405_S_AUREUS_VS_UNTREATED_NEUTROPHIL_UP, GSE2405_S_AUREUS_VS_UNTREATED_NEUTROPHIL_UP
GO_REGULATION_OF_TELOMERE_MAINTENANCE, GO_REGULATION_OF_TELOMERE_MAINTENANCE
GSE29164_DAY3_VS_DAY7_CD8_TCELL_AND_IL12_TREATED_MELANOMA_UP, GSE29164_DAY3_VS_DAY7_CD8_TCELL_AND_IL12_TREATED_MELANOMA_UP
GSE17721_POLYIC_VS_CPG_1H_BMDC_UP, GSE17721_POLYIC_VS_CPG_1H_BMDC_UP
GO_RNA_HELICASE_ACTIVITY, GO_RNA_HELICASE_ACTIVITY
GSE22229_RENAL_TRANSPLANT_IMMUNOSUPP_THERAPY_VS_HEALTHY_PBMCDUP, GSE22229_RENAL_TRANSPLANT_IMMUNOSUPP_THERAPY_VS_HEALTHY_PBMCDUP
GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE, GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE
GSE15330_HSC_VS_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_IKAROS_KO_UP, GSE15330_HSC_VS_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_IKAROS_KO_UP
GSE2706_UNSTIM_VS_2H_LPS_DC_UP, GSE2706_UNSTIM_VS_2H_LPS_DC_UP
XU_RESPONSE_TO_TRETINOIN_AND_NSC682994_DN, XU_RESPONSE_TO_TRETINOIN_AND_NSC682994_DN
GSE45365_CD8A_DC_VS_CD11B_DC_UP, GSE45365_CD8A_DC_VS_CD11B_DC_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_2H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_2H_UP
GAL_LEUKEMIC_STEM_CELL_UP, GAL_LEUKEMIC_STEM_CELL_UP
GO_PROTEIN_TRANSMEMBRANE_TRANSPORT, GO_PROTEIN_TRANSMEMBRANE_TRANSPORT
GO_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE, GO_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE
REACTOME_PROTEIN_FOLDING, REACTOME_PROTEIN_FOLDING
GSE7548_NAIVE_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_UP, GSE7548_NAIVE_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_UP
GSE15624_CTRL_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_UP, GSE15624_CTRL_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_UP
SNF5_DN.V1_UP, SNF5_DN.V1_UP
GO_BAF_TYPE_COMPLEX, GO_BAF_TYPE_COMPLEX
JAERVINEN_AMPLIFIED_IN_LARYNGEAL_CANCER, JAERVINEN_AMPLIFIED_IN_LARYNGEAL_CANCER
GSE24210_IL35_TREATED_VS_UNTREATED_TCONV_CD4_TCELL_UP, GSE24210_IL35_TREATED_VS_UNTREATED_TCONV_CD4_TCELL_UP
GO_POSITIVE_REGULATION_OF_TELOMERASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_TELOMERASE_ACTIVITY
GSE369_PRE_VS_POST_IL6_INJECTION_IFNG_KO_LIVER_DN, GSE369_PRE_VS_POST_IL6_INJECTION_IFNG_KO_LIVER_DN
GO_NPBAF_COMPLEX, GO_NPBAF_COMPLEX
ZHAN_MULTIPLE_MYELOMA_MS_DN, ZHAN_MULTIPLE_MYELOMA_MS_DN
GO_H4_HISTONE_ACETYLTRANSFERASE_COMPLEX, GO_H4_HISTONE_ACETYLTRANSFERASE_COMPLEX
GO_SWI_SNF_COMPLEX, GO_SWI_SNF_COMPLEX
GO_TRANSCRIPTION_FACTOR_TFIID_COMPLEX, GO_TRANSCRIPTION_FACTOR_TFIID_COMPLEX
PID_MYC_PATHWAY, PID_MYC_PATHWAY
CADWELL_ATG16L1_TARGETS_DN, CADWELL_ATG16L1_TARGETS_DN
ATAAGCT_MIR21, ATAAGCT_MIR21
GO_ADENYLYLTRANSFERASE_ACTIVITY, GO_ADENYLYLTRANSFERASE_ACTIVITY
GSE13547_CTRL_VS_ANTL_IGM_STIM_ZFX_KO_BCELL_12H_UP, GSE13547_CTRL_VS_ANTI_IGM_STIM_ZFX_KO_BCELL_12H_UP
GO_HISTONE_H2A_ACETYLTATION, GO_HISTONE_H2A_ACETYLTATION
GSE45365_CD8A_DC_VS_CD11B_DC_IFNAR_KO_UP, GSE45365_CD8A_DC_VS_CD11B_DC_IFNAR_KO_UP
REACTOME_CONVERSION_FROM_APC_C_CDC20_TO_APC_C_CDH1_IN_LATE_ANAPHASE, REACTOME_CONVERSION_FROM_APC_C_CDC20_TO_APC_C_CDH1_IN_LATE_ANAPHASE
GO_RNA_SPLICING_VIA_ENDONUCLEOLYTIC_CLEAVAGE_AND_LIGATION, GO_RNA_SPLICING_VIA_ENDONUCLEOLYTIC_CLEAVAGE_AND_LIGATION
GSE21360_PRIMARY_VS_SECONDARY_MEMORY_CD8_TCELL_UP, GSE21360_PRIMARY_VS_SECONDARY_MEMORY_CD8_TCELL_UP
GO_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER
GSE15659_NAIVE_CD4_TCELL_VS_ACTIVATED_TREG_DN, GSE15659_NAIVE_CD4_TCELL_VS_ACTIVATED_TREG_DN
chr18q11, chr18q11
GO_INOSITOL_PHOSPHATE_METABOLIC_PROCESS, GO_INOSITOL_PHOSPHATE_METABOLIC_PROCESS
MARIADASON_RESPONSE_TO_BUTYRATE_SULINDAC_4, MARIADASON_RESPONSE_TO_BUTYRATE_SULINDAC_4
LOPEZ_TRANSLATION_VIA_FN1_SIGNALING, LOPEZ_TRANSLATION_VIA_FN1_SIGNALING
GO_CELLULAR_RESPONSE_TO_HYDROGEN_PEROXIDE, GO_CELLULAR_RESPONSE_TO_HYDROGEN_PEROXIDE
GO_POSITIVE_REGULATION_OF_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION, GO_POSITIVE_REGULATION_OF_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION
GO_INTRACILIARY_TRANSPORT_PARTICLE_B, GO_INTRACILIARY_TRANSPORT_PARTICLE_B
NEWMAN_ERCC6_TARGETS_UP, NEWMAN_ERCC6_TARGETS_UP
FRASOR_TAMOXIFEN_RESPONSE_DN, FRASOR_TAMOXIFEN_RESPONSE_DN
GO_PROTEIN_COMPLEX_LOCALIZATION, GO_PROTEIN_COMPLEX_LOCALIZATION
chr15q14, chr15q14
GO_PROTEIN_TRANSPORT_ALONG_MICROTUBULE, GO_PROTEIN_TRANSPORT_ALONG_MICROTUBULE
GO_MANGANESE_ION_BINDING, GO_MANGANESE_ION_BINDING
GSE37301_HEMATOPOIETIC_STEM_CELL_VS_COMMON_LYMPHOID_PROGENITOR_UP, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_COMMON_LYMPHOID_PROGENITOR_UP
GO_DNA_POLYMERASE_BINDING, GO_DNA_POLYMERASE_BINDING
chr5q34, chr5q34
GO_POLYOL_METABOLIC_PROCESS, GO_POLYOL_METABOLIC_PROCESS
MODULE_256, MODULE_256
GO_DNA_HELICASE_COMPLEX, GO_DNA_HELICASE_COMPLEX
REACTOME_MRNA_DECAY_BY_5_TO_3_EXORIBONUCLEASE, REACTOME_MRNA_DECAY_BY_5_TO_3_EXORIBONUCLEASE
NIKOLSKY_BREAST_CANCER_7Q21_Q22_AMPLICON, NIKOLSKY_BREAST_CANCER_7Q21_Q22_AMPLICON
GO_PEROXISOME_ORGANIZATION, GO_PEROXISOME_ORGANIZATION
GO_TELOMERE_CAP_COMPLEX, GO_TELOMERE_CAP_COMPLEX