

ERIBOSOME, GOCC_PRERIBOSOME

MYC_UP.V1_UP, MYC_UP.V1_UP
GSE34156_NOD2_LIGAND_VS_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_DN, GSE34156_NOD2_LIGAND_VS_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_DN
GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP, GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP
GSE34156_NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_UP, GSE34156_NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_UP
GOBP_MATURATION_OF_SSU_RRNA, GOBP_MATURATION_OF_SSU_RRNA
GOCC_SMALL_SUBUNIT_PROCESSOME, GOCC_SMALL_SUBUNIT_PROCESSOME
GSE36009_UNSTIM_VS_LPS_STIM_NLRP10_KO_DC_DN, GSE36009_UNSTIM_VS_LPS_STIM_NLRP10_KO_DC_DN
GSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_DN, GSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_DN
GSE14415_ACT_VS_CTRL_NATURAL_TREG_DN, GSE14415_ACT_VS_CTRL_NATURAL_TREG_DN
GSE8678_IL7R_LOW_VS_HIGH_EFF_CD8_TCELL_DN, GSE8678_IL7R_LOW_VS_HIGH_EFF_CD8_TCELL_DN
GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_UP, GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_UP
GSE11961_MARGINAL_ZONE_BCELL_VS_PLASMA_CELL_DAY7_DN, GSE11961_MARGINAL_ZONE_BCELL_VS_PLASMA_CELL_DAY7_DN
GOBP_MATURATION_OF_SSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA, GOBP_MATURATION_OF_SSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA
GOCC_FIBRILLAR_CENTER, GOCC_FIBRILLAR_CENTER
GSE11961_FOLLICULAR_BCELL_VS_PLASMA_CELL_DAY7_UP, GSE11961_FOLLICULAR_BCELL_VS_PLASMA_CELL_DAY7_UP
GSE13547_WT_VS_ZFX_KO_BCELL_ANTI_IGM_STIM_2H_DN, GSE13547_WT_VS_ZFX_KO_BCELL_ANTI_IGM_STIM_2H_DN
GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_FOXP3_TRASDUCED_CD4_TCELL_UP, GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_FOXP3_TRASDUCED_CD4_TCELL_UP
ZNF669_TARGET_GENES, ZNF669_TARGET_GENES
GOBP_RIBOSOMAL_LARGE_SUBUNIT_ASSEMBLY, GOBP_RIBOSOMAL_LARGE_SUBUNIT_ASSEMBLY
CAVIN1_TARGET_GENES, CAVIN1_TARGET_GENES
GOBP_BLASTOCYST_DEVELOPMENT, GOBP_BLASTOCYST_DEVELOPMENT
STEIN_ESRRA_TARGETS_DN, STEIN_ESRRA_TARGETS_DN
GSE9946_IMMATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP, GSE9946_IMMATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP
GOMF_ATPASE_BINDING, GOMF_ATPASE_BINDING
GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNG_KO_DN, GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNG_KO_DN
GOBP_PROTEIN_LOCALIZATION_TO_NUCLEOLUS, GOBP_PROTEIN_LOCALIZATION_TO_NUCLEOLUS
PID_DNA_PK_PATHWAY, PID_DNA_PK_PATHWAY
GOBP_RRNA_BASE_METHYLATION, GOBP_RRNA_BASE_METHYLATION
CHEOK_RESPONSE_TO_MERCAPTOPURINE_UP, CHEOK_RESPONSE_TO_MERCAPTOPURINE_UP
KEGG_NON_HOMOLOGOUS_END_JOINING, KEGG_NON_HOMOLOGOUS_END_JOINING
GOCC_NONHOMOLOGOUS_END_JOINING_COMPLEX, GOCC_NONHOMOLOGOUS_END_JOINING_COMPLEX
GOBP_POSITIVE_REGULATION_BY_HOST_OF_VIRAL_TRANSCRIPTION, GOBP_POSITIVE_REGULATION_BY_HOST_OF_VIRAL_TRANSCRIPTION
GOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_5_ETS_OF_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA, GOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_5_ETS_OF_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA
HP_CONGENITAL_LOCALIZED_ABSENCE_OF_SKIN, HP_CONGENITAL_LOCALIZED_ABSENCE_OF_SKIN
ZNF501_TARGET_GENES, ZNF501_TARGET_GENES
WP_NONHOMOLOGOUS_END_JOINING, WP_NONHOMOLOGOUS_END_JOINING
GOBP_REGULATION_OF_T_CIRCLE_FORMATION, GOBP_REGULATION_OF_T_CIRCLE_FORMATION
GOBP_LYMPHOID_PROGENITOR_CELL_DIFFERENTIATION, GOBP_LYMPHOID_PROGENITOR_CELL_DIFFERENTIATION
PODAR_RESPONSE_TO_ADAPHOSTIN_DN, PODAR_RESPONSE_TO_ADAPHOSTIN_DN
GOBP_NEGATIVE_REGULATION_OF_CELLULAR_SENESCENCE, GOBP_NEGATIVE_REGULATION_OF_CELLULAR_SENESCENCE
GOBP_PRO_B_CELL_DIFFERENTIATION, GOBP_PRO_B_CELL_DIFFERENTIATION
GOCC_TELOMERE_CAP_COMPLEX, GOCC_TELOMERE_CAP_COMPLEX
GOBP_NEGATIVE_REGULATION_OF_HISTONE_ACETYLATION, GOBP_NEGATIVE_REGULATION_OF_HISTONE_ACETYLATION
NIKOLSKY_BREAST_CANCER_12Q24_AMPLICON, NIKOLSKY_BREAST_CANCER_12Q24_AMPLICON
GAJATE_RESPONSE_TO TRABECTEDIN_DN, GAJATE_RESPONSE_TO TRABECTEDIN_DN
GOBP_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_NONHOMOLOGOUS_END_JOINING, GOBP_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_NONHOMOLOGOUS_END_JOINING
chr21p12, chr21p12
GHO_ATF5_TARGETS_UP, GHO_ATF5_TARGETS_UP
MIR548AO_3P, MIR548AO_3P
GOBP_IMMUNOGLOBULIN_V_D_J_RECOMBINATION, GOBP_IMMUNOGLOBULIN_V_D_J_RECOMBINATION
HP_SEVERE_COMBINED_IMMUNODEFICIENCY, HP_SEVERE_COMBINED_IMMUNODEFICIENCY
GOBP_REGULATION_OF_B_CELL_APOPTOTIC_PROCESS, GOBP_REGULATION_OF_B_CELL_APOPTOTIC_PROCESS
HP_LIMB_MYOCOLONUS, HP_LIMB_MYOCOLONUS
MAEKAWA_ATF2_TARGETS, MAEKAWA_ATF2_TARGETS
ZNF529_TARGET_GENES, ZNF529_TARGET_GENES
ZNF292_TARGET_GENES, ZNF292_TARGET_GENES
chr1p12, chr1p12
GOBP_SOMATIC_DIVERSIFICATION_OF_T_CELL_RECEPTOR_GENES, GOBP_SOMATIC_DIVERSIFICATION_OF_T_CELL_RECEPTOR_GENES
GOBP_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR, GOBP_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR
GOCC_GRANULAR_COMPONENT, GOCC_GRANULAR_COMPONENT
MIR6514_5P, MIR6514_5P
chr8p12, chr8p12
chr1q41, chr1q41
ZNF354C_TARGET_GENES, ZNF354C_TARGET_GENES
chr6q15, chr6q15
chr1q43, chr1q43