

LITAZONE_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_TREATED_MACROPHAGE_DN

GSE14769_LPS_UNSTIM_VS_80MIN_LPS_BMDM_UP, GSE14769_UNSTIM_VS_80MIN_LPS_BMDM_UP
GSE16385_MONOCYTE_VS_12H_IL4_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_IL4_TREATED_MACROPHAGE_DN
GSE23925_LIGHT_ZONE_VS_NAIVE_BCELL_DN, GSE23925_LIGHT_ZONE_VS_NAIVE_BCELL_DN
GSE6674_ANTI_IGM_VS_CPG_STIM_BCELL_UP, GSE6674_ANTI_IGM_VS_CPG_STIM_BCELL_UP
GSE23925_DARK_ZONE_VS_NAIVE_BCELL_UP, GSE23925_DARK_ZONE_VS_NAIVE_BCELL_UP
GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN
GSE36527_CD62L_HIGH_VS_CD62L_LOW_TREG_CD69_NEG_KLRG1_NEG_UP, GSE36527_CD62L_HIGH_VS_CD62L_LOW_TREG_CD69_NEG_KLRG1_NEG_UP
GSE7460_FOXP3_MUT_VS_WT_ACT_WITH_TGFB_TCONV_UP, GSE7460_FOXP3_MUT_VS_WT_ACT_WITH_TGFB_TCONV_UP
GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP, GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP
GSE26669_CD4_VS_CD8_TCELL_IN_MLR_DN, GSE26669_CD4_VS_CD8_TCELL_IN_MLR_DN
GSE7568_CTRL_VS_24H_TGFB_TREATED_MACROPHAGES_WITH_IL4_AND_DEXAMETHASONE_UP, GSE7568_CTRL_VS_24H_TGFB_TREATED_MACROPHAGES_WITH_IL4_AND_DEXAMETHASONE_UP
GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4_KO_BCELL_DN, GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4_KO_BCELL_DN
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_UP, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_UP
GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_DN, GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_DN
GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP, GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP
GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD4_TCELL_DN, GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD4_TCELL_DN
GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN, GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN
GSE36527_CD62L_HIGH_CD69_NEG_VS_CD62L_LOW_CD69_POS_TREG_KLRG1_NEG_UP, GSE36527_CD62L_HIGH_CD69_NEG_VS_CD62L_LOW_CD69_POS_TREG_KLRG1_NEG_UP
GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP, GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP
GSE17721_PAM3CSK4_VS_CPG_16H_BMDC_UP, GSE17721_PAM3CSK4_VS_CPG_16H_BMDC_UP
JAZAG_TGFB1_SIGNALING_UP, JAZAG_TGFB1_SIGNALING_UP
GO_NUCLEAR_LOCALIZATION_SEQUENCE_BINDING, GO_NUCLEAR_LOCALIZATION_SEQUENCE_BINDING
GSE9946_MATURE_STIMULATORY_VS_LISTERIA_INF_MATURE_DC_UP, GSE9946_MATURE_STIMULATORY_VS_LISTERIA_INF_MATURE_DC_UP
AHRARNT_01, AHRARNT_01
FIGUEROA_AML_METHYLATION_CLUSTER_6_UP, FIGUEROA_AML_METHYLATION_CLUSTER_6_UP
FIGUEROA_AML_METHYLATION_CLUSTER_3_UP, FIGUEROA_AML_METHYLATION_CLUSTER_3_UP
GO_MAINTENANCE_OF_CELL_NUMBER, GO_MAINTENANCE_OF_CELL_NUMBER
GO_SIGNAL_SEQUENCE_BINDING, GO_SIGNAL_SEQUENCE_BINDING
FIGUEROA_AML_METHYLATION_CLUSTER_7_UP, FIGUEROA_AML_METHYLATION_CLUSTER_7_UP
GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_MEMORY_BCELL_DAY40_UP, GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_MEMORY_BCELL_DAY40_UP
GSE2706_UNSTIM_VS_8H_LPS_AND_R848_DC_DN, GSE2706_UNSTIM_VS_8H_LPS_AND_R848_DC_DN
FIGUEROA_AML_METHYLATION_CLUSTER_4_UP, FIGUEROA_AML_METHYLATION_CLUSTER_4_UP
GO_REGULATION_OF_PROTEIN_ACETYLATION, GO_REGULATION_OF_PROTEIN_ACETYLATION
GSE7548_NAIVE_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_UP, GSE7548_NAIVE_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_UP
BROWNE_HCMV_INFECTION_10HR_UP, BROWNE_HCMV_INFECTION_10HR_UP
GSE16385_MONOCYTE_VS_12H_IFNG_TNF_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_IFNG_TNF_TREATED_MACROPHAGE_DN
GROSS_HYPOXIA_VIA_ELK3_ONLY_DN, GROSS_HYPOXIA_VIA_ELK3_ONLY_DN
DELASERNA_MYOD_TARGETS_DN, DELASERNA_MYOD_TARGETS_DN
GO_THYMOCYTE_AGGREGATION, GO_THYMOCYTE_AGGREGATION
GO_MRNA_CLEAVAGE_AND_POLYADENYLATION_SPECIFICITY_FACTOR_COMPLEX, GO_MRNA_CLEAVAGE_AND_POLYADENYLATION_SPECIFICITY_FACTOR_COMPLEX
BRACHAT_RESPONSE_TO_CISPLATIN, BRACHAT_RESPONSE_TO_CISPLATIN
ELLWOOD_MYC_TARGETS_DN, ELLWOOD_MYC_TARGETS_DN
GSE12963_UNINF_VS_ENV_AND_NEF_DEFICIENT_HIV1_INF_CD4_TCELL_UP, GSE12963_UNINF_VS_ENV_AND_NEF_DEFICIENT_HIV1_INF_CD4_TCELL_UP
GO_HISTONE_ACETYLTRANSFERASE_BINDING, GO_HISTONE_ACETYLTRANSFERASE_BINDING
chr15q15, chr15q15
GO_PHOSPHOTRANSFERASE_ACTIVITY_FOR_OTHER_SUBSTITUTED_PHOSPHATE_GROUPS, GO_PHOSPHOTRANSFERASE_ACTIVITY_FOR_OTHER_SUBSTITUTED_PHOSPHATE_GROUPS
BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_UP, BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_UP
GO_HEMATOPOIETIC_STEM_CELL_PROLIFERATION, GO_HEMATOPOIETIC_STEM_CELL_PROLIFERATION
GTGTGAG_MIR342, GTGTGAG_MIR342
SARTIPY_BLUNTED_BY_INSULIN_RESISTANCE_UP, SARTIPY_BLUNTED_BY_INSULIN_RESISTANCE_UP
GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RESPONSE_TO_CHEMICAL_STIMULUS, GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RESPONSE_TO_CHEMICAL_STIMULUS
GO_REGULATION_OF_TRANSFORMING_GROWTH_FACTOR_BETA_PRODUCTION, GO_REGULATION_OF_TRANSFORMING_GROWTH_FACTOR_BETA_PRODUCTION
GO_POSITIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GO_POSITIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
GO_ENDOTHELIUM_DEVELOPMENT, GO_ENDOTHELIUM_DEVELOPMENT
MA_MYELOID_DIFFERENTIATION_DN, MA_MYELOID_DIFFERENTIATION_DN
GO_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR, GO_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR
SNIJDERS_AMPLIFIED_IN_HEAD_AND_NECK_TUMORS, SNIJDERS_AMPLIFIED_IN_HEAD_AND_NECK_TUMORS
GOBERT_CORE_OLIGODENDROCYTE_DIFFERENTIATION, GOBERT_CORE_OLIGODENDROCYTE_DIFFERENTIATION
KUWANO_RNA_STABILIZED_BY_NO, KUWANO_RNA_STABILIZED_BY_NO
GO_FOUR_WAY_JUNCTION_DNA_BINDING, GO_FOUR_WAY_JUNCTION_DNA_BINDING
GNF2_ATM, GNF2_ATM
GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR, GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR
CCCACAT_MIR2993P, CCCACAT_MIR2993P
GO_HEART_VALVE_DEVELOPMENT, GO_HEART_VALVE_DEVELOPMENT
GO_ENDOCARDIAL_CUSHION_FORMATION, GO_ENDOCARDIAL_CUSHION_FORMATION
chr5q21, chr5q21
GO_ISOPRENOID_BIOSYNTHETIC_PROCESS, GO_ISOPRENOID_BIOSYNTHETIC_PROCESS
GO_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_INVOLVED_IN_ENDOCARDIAL_CUSHION_FORMATION, GO_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_INVOLVED_IN_ENDOCARDIAL_CUSHION_FORMATION
PID_S1P_META_PATHWAY, PID_S1P_META_PATHWAY