

L_AND_IL12_TREATED_MELANOMA_DAY7_UP, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP

GSE29164_CD8_TCELL_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_UP, GSE29164_CD8_TCELL_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP
GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_DN, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_DN
GSE46606_IRF4HIGH_VS_IRF4MID_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_UP, GSE46606_IRF4HIGH_VS_IRF4MID_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_UP
GSE25088_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_UP, GSE25088_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_UP
GSE26351_WNT_VS_BMP_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_DN, GSE26351_WNT_VS_BMP_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_DN
GSE26351_UNSTIM_VS_WNT_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_DN, GSE26351_UNSTIM_VS_WNT_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_DN
GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_2H_UP
GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_UP, GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_UP
GSE557_WT_VS_I_AB_KO_DC_UP, GSE557_WT_VS_I_AB_KO_DC_UP
GSE8868_SPLEEN_VS_INTESTINE_CD11B_POS_CD11C_NEG_DC_UP, GSE8868_SPLEEN_VS_INTESTINE_CD11B_POS_CD11C_NEG_DC_UP
GSE13306_LAMINA_PROPRIA_VS_SPLEEN_TREG_UP, GSE13306_LAMINA_PROPRIA_VS_SPLEEN_TREG_UP
GSE17974_CTRL_VS_ACT_IL4_AND_ANTL_IL12_72H_CD4_TCELL_UP, GSE17974_CTRL_VS_ACT_IL4_AND_ANTL_IL12_72H_CD4_TCELL_UP
GSE411_UNSTIM_VS_100MIN_IL6_STIM_MACROPHAGE_DN, GSE411_UNSTIM_VS_100MIN_IL6_STIM_MACROPHAGE_DN
GSE29164_UNTREATED_VS_CD8_TCELL_TREATED_MELANOMA_DAYS_DN, GSE29164_UNTREATED_VS_CD8_TCELL_TREATED_MELANOMA_DAYS_DN
GSE4142_GC_BCELL_VS_MEMORY_BCELL_UP, GSE4142_GC_BCELL_VS_MEMORY_BCELL_UP
BURTON_ADIPOGENESIS_6, BURTON_ADIPOGENESIS_6
GSE1925_CTRL_VS_24H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_UP, GSE1925_CTRL_VS_24H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_UP
GSE42021_CD24HL_VS_CD24INT_TCONV_THYMUS_UP, GSE42021_CD24HL_VS_CD24INT_TCONV_THYMUS_UP
GRABARCZYK_BCL11B_TARGETS_UP, GRABARCZYK_BCL11B_TARGETS_UP
GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL6_KO_MACROPHAGE_45MIN_DN, GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL6_KO_MACROPHAGE_45MIN_DN
GSE6092_UNSTIM_VS_IFNG_STIM_ENDOTHELIAL_CELL_DN, GSE6092_UNSTIM_VS_IFNG_STIM_ENDOTHELIAL_CELL_DN
HALLMARK_UV_RESPONSE_UP, HALLMARK_UV_RESPONSE_UP
GSE26030_UNSTIM_VS_RESTIM_THI7_DAY15_POST_POLARIZATION_UP, GSE26030_UNSTIM_VS_RESTIM_THI7_DAY15_POST_POLARIZATION_UP
GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_48H_DN, GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_48H_DN
GSE3982_EOSINOPHIL_VS_TH1_UP, GSE3982_EOSINOPHIL_VS_TH1_UP
GO_RHO_GTPASE_BINDING, GO_RHO_GTPASE_BINDING
GSE27786_BCELL_VS_CD4_TCELL_DN, GSE27786_BCELL_VS_CD4_TCELL_DN
GO_REGULATION_OF_JUN_KINASE_ACTIVITY, GO_REGULATION_OF_JUN_KINASE_ACTIVITY
GSE27241_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_THI7_POLARIZING_CONDITIONS_DN, GSE27241_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_THI7_POLARIZING_CONDITIONS_DN
GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_DN, GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_DN
GSE20366_CD103_POS_VS_CD103_KLRG1_DP_TREG_DN, GSE20366_CD103_POS_VS_CD103_KLRG1_DP_TREG_DN
GSE11961_MEMORY_BCELL_DAY7_VS_GERMINAL_CENTER_BCELL_DAY7_DN, GSE11961_MEMORY_BCELL_DAY7_DN
DACOSTA_UV_RESPONSE_VIA_ERCC3_COMMON_UP, DACOSTA_UV_RESPONSE_VIA_ERCC3_COMMON_UP
WARTERS_RESPONSE_TO_IR_SKIN, WARTERS_RESPONSE_TO_IR_SKIN
REACTOME_SIGNALING_BY_ILS, REACTOME_SIGNALING_BY_ILS
GSE7831_1H_VS_4H_INFLUENZA_STIM_PDC_DN, GSE7831_1H_VS_4H_INFLUENZA_STIM_PDC_DN
FIRESTEIN_PROLIFERATION, FIRESTEIN_PROLIFERATION
GO_PLASMA_MEMBRANE_RECEPTOR_COMPLEX, GO_PLASMA_MEMBRANE_RECEPTOR_COMPLEX
GSE1925_CTRL_VS_3H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN, GSE1925_CTRL_VS_3H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN
GO_POSITIVE_REGULATION_OF_JUN_KINASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_JUN_KINASE_ACTIVITY
PID_VEGFR1_2_PATHWAY, PID_VEGFR1_2_PATHWAY
GSE20715_0H_VS_6H_OZONE_LUNG_DN, GSE20715_0H_VS_6H_OZONE_LUNG_DN
BROWNE_HCMV_INFECTION_20HR_DN, BROWNE_HCMV_INFECTION_20HR_DN
GSE22601_DOUBLE_NEGATIVE_VS_DOUBLE_POSITIVE_THYMOCYTE_UP, GSE22601_DOUBLE_NEGATIVE_VS_DOUBLE_POSITIVE_THYMOCYTE_UP
GO_CELL_SUBSTRATE_ADHESION, GO_CELL_SUBSTRATE_ADHESION
GO_RECEPTOR_SIGNALING_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY, GO_RECEPTOR_SIGNALING_PROTEIN_SERINE_THREONINE_KINASE_ACTIVITY
GO_SUBSTRATE_ADHESION_DEPENDENT_CELL_SPREADING, GO_SUBSTRATE_ADHESION_DEPENDENT_CELL_SPREADING
GO_NEGATIVE_REGULATION_OF_MAP_KINASE_ACTIVITY, GO_NEGATIVE_REGULATION_OF_MAP_KINASE_ACTIVITY
GTTATAT_MIR410, GTTATAT_MIR410
WORSCHERCH_TUMOR_EVASION_AND_TOLEROGENICITY_DN, WORSCHERCH_TUMOR_EVASION_AND_TOLEROGENICITY_DN
GSE32034_LY6C_HIGH_VS_LOW_MONOCYTE_UP, GSE32034_LY6C_HIGH_VS_LOW_MONOCYTE_UP
GO_ADAPTIVE_IMMUNE_RESPONSE_BASED_ON_SOMATIC_RECOMBINATION_OF_IMMUNE_RECEPTORS_BUILT_FROM_IMMUNOGLOBULIN_SUPERFAMILY_DOMAINS, GO_ADAPTIVE_IMMUNE_RESPONSE_BASED_ON_SOMATIC_RECOMBINATION_OF_IMMUNE_RECEPTORS_BUILT_F
GO_JNK_CASCADE, GO_JNK_CASCADE
BIOCARTA_SPPA_PATHWAY, BIOCARTA_SPPA_PATHWAY
GO_PHOSPHATIDYLINOSITOL_PHOSPHATE_5_PHOSPHATASE_ACTIVITY, GO_PHOSPHATIDYLINOSITOL_PHOSPHATE_5_PHOSPHATASE_ACTIVITY
GO_LYMPHOCYTE_HOMEOSTASIS, GO_LYMPHOCYTE_HOMEOSTASIS
GO_ACTIVATION_OF_MAPK_ACTIVITY, GO_ACTIVATION_OF_MAPK_ACTIVITY
BOYALT_LIVER_CANCER_SUBCLASS_G123_DN, BOYALT_LIVER_CANCER_SUBCLASS_G123_DN
GO_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS, GO_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS
GO_AMINO_SUGAR_METABOLIC_PROCESS, GO_AMINO_SUGAR_METABOLIC_PROCESS
BIOCARTA_EPO_PATHWAY, BIOCARTA_EPO_PATHWAY
GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_DN, GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_DN
GO_REGULATION_OF_B_CELL_ACTIVATION, GO_REGULATION_OF_B_CELL_ACTIVATION
GO_THIOESTER_METABOLIC_PROCESS, GO_THIOESTER_METABOLIC_PROCESS
PPARG_01, PPARG_01
GO_REGULATION_OF_SUPEROXIDE_ANION_GENERATION, GO_REGULATION_OF_SUPEROXIDE_ANION_GENERATION
GO_LEUKOCYTE_MEDIATED_IMMUNITY, GO_LEUKOCYTE_MEDIATED_IMMUNITY
REACTOME_REGULATION_OF_IFNA_SIGNALING, REACTOME_REGULATION_OF_IFNA_SIGNALING
GNF2_CDC27, GNF2_CDC27
BIOCARTA_NKCELLS_PATHWAY, BIOCARTA_NKCELLS_PATHWAY
GO_RESPONSE_TO_ATP, GO_RESPONSE_TO_ATP
KYNG_DNA_DAMAGE_BY_GAMMA_AND_UV_RADIATION, KYNG_DNA_DAMAGE_BY_GAMMA_AND_UV_RADIATION
GO_MYOSIN_BINDING, GO_MYOSIN_BINDING
GROSS_HYPOXIA_VIA_ELK3_ONLY_UP, GROSS_HYPOXIA_VIA_ELK3_ONLY_UP
GO_FOREBRAIN_CELL_MIGRATION, GO_FOREBRAIN_CELL_MIGRATION
MODULE_419, MODULE_419
GO_MOTILE_CILIUM, GO_MOTILE_CILIUM
GO_N_ACETYLGUCOSAMINE_METABOLIC_PROCESS, GO_N_ACETYLGUCOSAMINE_METABOLIC_PROCESS
chr6p24, chr6p24
GO_MICROVILLUS, GO_MICROVILLUS
GO_REGULATION_OF_NEURON_MIGRATION, GO_REGULATION_OF_NEURON_MIGRATION
GO_POSITIVE_REGULATION_OF_CYTOKINE_SECRETION, GO_POSITIVE_REGULATION_OF_CYTOKINE_SECRETION
REACTOME_PLATELET_SENSITIZATION_BY_LDL, REACTOME_PLATELET_SENSITIZATION_BY_LDL
GO_NEGATIVE_REGULATION_OF_B_CELL_ACTIVATION, GO_NEGATIVE_REGULATION_OF_B_CELL_ACTIVATION
REACTOME_CREB_PHOSPHORYLATION_THROUGH_THE_ACTIVATION_OF_RAS, REACTOME_CREB_PHOSPHORYLATION_THROUGH_THE_ACTIVATION_OF_RAS
BIOCARTA_INTEGRIN_PATHWAY, BIOCARTA_INTEGRIN_PATHWAY
REACTOME_CASPASE_MEDIATED_CLEAVAGE_OF_CYTOSKELETAL_PROTEINS, REACTOME_CASPASE_MEDIATED_CLEAVAGE_OF_CYTOSKELETAL_PROTEINS
GO_CEREBRAL_CORTEX_DEVELOPMENT, GO_CEREBRAL_CORTEX_DEVELOPMENT
GO_B_CELL_HOMEOSTASIS, GO_B_CELL_HOMEOSTASIS
KORKOLA_YOLK_SAC_TUMOR_UP, KORKOLA_YOLK_SAC_TUMOR_UP
REACTOME_POST_NMDA_RECEPTOR_ACTIVATION_EVENTS, REACTOME_POST_NMDA_RECEPTOR_ACTIVATION_EVENTS