

L_VS_MONO_MAC_UP, GSE27786_BCELL_VS_MONO_MAC_UP

GSE27786_CD8_TCELL_VS_ERYTHROBLAST_UP, GSE27786_CD8_TCELL_VS_ERYTHROBLAST_UP
GSE27786_CD8_TCELL_VS_MONO_MAC_UP, GSE27786_CD8_TCELL_VS_MONO_MAC_UP
GSE27786_BCELL_VS_NEUTROPHIL_UP, GSE27786_BCELL_VS_NEUTROPHIL_UP
GSE46606_IRF4HIGH_VS_IRF4MID_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_UP, GSE46606_IRF4HIGH_VS_IRF4MID_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_UP
GSE27786_NKTCCELL_VS_MONO_MAC_UP, GSE27786_NKTCCELL_VS_MONO_MAC_UP
GSE27786_ERYTHROBLAST_VS_MONO_MAC_DN, GSE27786_ERYTHROBLAST_VS_MONO_MAC_DN
GSE27786_CD4_TCELL_VS_MONO_MAC_UP, GSE27786_CD4_TCELL_VS_MONO_MAC_UP
GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_DN, GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_DN
GSE27786_NKCELL_VS_MONO_MAC_UP, GSE27786_NKCELL_VS_MONO_MAC_UP
GSE17721_LPS_VS_CPG_1H_BMDC_UP, GSE17721_LPS_VS_CPG_1H_BMDC_UP
GSE17186_NAIVE_VS_CD21HIGH_TRANSITIONAL_BCELL_CORD_BLOOD_UP, GSE17186_NAIVE_VS_CD21HIGH_TRANSITIONAL_BCELL_CORD_BLOOD_UP
GSE17721_LPS_VS_POLYIC_24H_BMDC_UP, GSE17721_LPS_VS_POLYIC_24H_BMDC_UP
GSE6674_UNSTIM_VS_ANTLJGM_STIM_BCELL_UP, GSE6674_UNSTIM_VS_ANTLJGM_STIM_BCELL_UP
GSE27786_LSK_VS_BCELL_DN, GSE27786_LSK_VS_BCELL_DN
GSE27786_BCELL_VS_NKCELL_UP, GSE27786_BCELL_VS_NKCELL_UP
GSE9960_GRAM_POS_VS_GRAM_NEG_AND_POS_SEPSIS_PBMIC_UP, GSE9960_GRAM_POS_VS_GRAM_NEG_AND_POS_SEPSIS_PBMIC_UP
GSE14308_TH1_VS_INDUCED_TREG_UP, GSE14308_TH1_VS_INDUCED_TREG_UP
GSE39022_LN_VS_SPLEEN_DC_DN, GSE39022_LN_VS_SPLEEN_DC_DN
GSE27786_LIN_NEG_VS_CD8_TCELL_DN, GSE27786_LIN_NEG_VS_CD8_TCELL_DN
GSE15767_MED_VS_SCS_MAC_LN_DN, GSE15767_MED_VS_SCS_MAC_LN_DN
GSE14308_TH2_VS_NATURAL_TREG_DN, GSE14308_TH2_VS_NATURAL_TREG_DN
GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_UP, GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_UP
GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_DN, GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_DN
GSE23568_CTRL_VS_ID3_TRANSDUCED_CD8_TCELL_UP, GSE23568_CTRL_VS_ID3_TRANSDUCED_CD8_TCELL_UP
GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_LPS_STIM_UP, GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_LPS_STIM_UP
GSE27786_CD8_TCELL_VS_NKCELL_UP, GSE27786_CD8_TCELL_VS_NKCELL_UP
GSE8621_UNSTIM_VS_LPS_PRIMED_UNSTIM_MACROPHAGE_DN, GSE8621_UNSTIM_VS_LPS_PRIMED_UNSTIM_MACROPHAGE_DN
GSE1448_CTRL_VS_ANTL_VBETA5_DP_THYMOCYTE_UP, GSE1448_CTRL_VS_ANTL_VBETA5_DP_THYMOCYTE_UP
GSE23505_IL6_IL1_VS_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP, GSE23505_IL6_IL1_VS_IL6_IL1_TGFB_TREATED_CD4_TCELL_UP
GO_PROTEIN_TARGETING_TO_MITOCHONDRION, GO_PROTEIN_TARGETING_TO_MITOCHONDRION
GTGTGA_MIR505, GTGTGA_MIR505
GO_PROTEIN_LOCALIZATION_TO_MITOCHONDRION, GO_PROTEIN_LOCALIZATION_TO_MITOCHONDRION
GSE14308_TH17_VS_NAIVE_CD4_TCELL_DN, GSE14308_TH17_VS_NAIVE_CD4_TCELL_DN
GO_DEMETHYLASE_ACTIVITY, GO_DEMETHYLASE_ACTIVITY
GSE10240_CTRL_VS_IL17_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_UP, GSE10240_CTRL_VS_IL17_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_UP
GO_DNA_MODIFICATION, GO_DNA_MODIFICATION
GSE32901_TH1_VS_TH17_ENRICHED_CD4_TCELL_DN, GSE32901_TH1_VS_TH17_ENRICHED_CD4_TCELL_DN
GNF2_RAP1B, GNF2_RAP1B
GO_DEMETHYLATION, GO_DEMETHYLATION
GSE14308_NAIVE_CD4_TCELL_VS_NATURAL_TREG_UP, GSE14308_NAIVE_CD4_TCELL_VS_NATURAL_TREG_UP
GO_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION, GO_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION
GO_DNA_METHYLATION_OR_DEMETHYLATION, GO_DNA_METHYLATION_OR_DEMETHYLATION
NAKAMURA_ADIPOGENESIS_LATE_UP, NAKAMURA_ADIPOGENESIS_LATE_UP
GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN_2_OXOGLUTARATE_AS_ONE_DONOR_AND_INCORPORATION_OF_ONE_ATOM_EACH_OF_OXYGEN_INTO_BOTH_DONORS, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN_2_OXOGLUTARATE_AS_ONE_DONOR_AND_INCORPORATION_O
GSE5542_IFNG_VS_IFNA_TREATED_EPITHELIAL_CELLS_6H_DN, GSE5542_IFNG_VS_IFNA_TREATED_EPITHELIAL_CELLS_6H_DN
GO_POSITIVE_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING, GO_POSITIVE_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING
HUMMERICH_SKIN_CANCER_PROGRESSION_DN, HUMMERICH_SKIN_CANCER_PROGRESSION_DN
GSE25890_CTRL_VS_IL33_IL7_TREATED_NUOCYTES_UP, GSE25890_CTRL_VS_IL33_IL7_TREATED_NUOCYTES_UP
GO_ESTABLISHMENT_OF_CELL_POLARITY, GO_ESTABLISHMENT_OF_CELL_POLARITY
AR_02, AR_02
BYSTRYKH_HEMATOPOIESIS_STEM_CELL_SCP2_QTL_TRANS, BYSTRYKH_HEMATOPOIESIS_STEM_CELL_SCP2_QTL_TRANS
GO_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX_COUPLING_FACTOR_F_O_, GO_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX_COUPLING_FACTOR_F_O_
GENTILE_UV_RESPONSE_CLUSTER_D6, GENTILE_UV_RESPONSE_CLUSTER_D6
GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_DN, GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_DN
GO_CILIARY_TIP, GO_CILIARY_TIP
GO_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX_PROTON_TRANSPORTING_DOMAIN, GO_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX_PROTON_TRANSPORTING_DOMAIN
KORKOLA_SEMINOMA_UP, KORKOLA_SEMINOMA_UP
STAT3_01, STAT3_01
REACTOME_NUCLEAR_EVENTS_KINASE_AND_TRANSCRIPTION_FACTOR_ACTIVATION, REACTOME_NUCLEAR_EVENTS_KINASE_AND_TRANSCRIPTION_FACTOR_ACTIVATION
TCTGATC_MIR383, TCTGATC_MIR383
REACTOME_MAPK_TARGETS_NUCLEAR_EVENTS_MEDIATED_BY_MAP_KINASES, REACTOME_MAPK_TARGETS_NUCLEAR_EVENTS_MEDIATED_BY_MAP_KINASES
STAT1_01, STAT1_01
KEGG_BIOSYNTHESIS_OF_UNSATURATED_FATTY_ACIDS, KEGG_BIOSYNTHESIS_OF_UNSATURATED_FATTY_ACIDS
GSE360_DC_VS_MAC_T_GONDII_UP, GSE360_DC_VS_MAC_T_GONDII_UP
GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE
GO_ARMADILLO_REPEAT_DOMAIN_BINDING, GO_ARMADILLO_REPEAT_DOMAIN_BINDING
GO_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GO_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE
LEIN_PONS_MARKERS, LEIN_PONS_MARKERS
chrXq24, chrXq24
DER_IFN_GAMMA_RESPONSE_DN, DER_IFN_GAMMA_RESPONSE_DN
GO_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION, GO_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION
PID_LYMPH_ANGIOGENESIS_PATHWAY, PID_LYMPH_ANGIOGENESIS_PATHWAY