

_AND_IL12_TREATED_MELANOMA_DAY3_DN, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_DN

- GSE29164_CD8_TCELL_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_UP, GSE29164_CD8_TCELL_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_UP
- GSE29164_UNTREATED_VS_CD8_TCELL_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_DAY7_UP, GSE29164_CD8_TCELL_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP
- GSE26030_UNSTIM_VS_RESTIM_TH1_DAY5_POST_POLARIZATION_DN, GSE26030_UNSTIM_VS_RESTIM_TH1_DAY5_POST_POLARIZATION_DN
- GSE26351_UNSTIM_VS_BMP_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_UP, GSE26351_UNSTIM_VS_BMP_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_UP
- GSE17721_POLYIC_VS_GARDIQUIMOD_0.5H_BMDC_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_0.5H_BMDC_DN
- GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_LPS_STIM_DN, GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_LPS_STIM_DN
- GSE8868_SPLEEN_VS_INTESTINE_CD11B_POS_CD11C_NEG_DC_UP, GSE8868_SPLEEN_VS_INTESTINE_CD11B_POS_CD11C_NEG_DC_UP
- GSE20366_EX_VIVO_VS_HOMEOSTATIC_CONVERSION_NAIVE_CD4_TCELL_DN, GSE20366_EX_VIVO_VS_HOMEOSTATIC_CONVERSION_NAIVE_CD4_TCELL_DN
- GSE18893_CTRL_VS_TNF_TREATED_TCONV_24H_DN, GSE18893_CTRL_VS_TNF_TREATED_TCONV_24H_DN
- MIR6165, MIR6165
- CAGCACT_MIR5123P, CAGCACT_MIR5123P
- HP_PRENATAL_MOVEMENT_ABNORMALITY, HP_PRENATAL_MOVEMENT_ABNORMALITY
- MIR548AT_3P, MIR548AT_3P
- GSE13887_RESTING_VS_NO_TREATED_CD4_TCELL_DN, GSE13887_RESTING_VS_NO_TREATED_CD4_TCELL_DN
- GOBP_NEGATIVE_REGULATION_OF_SUPRAMOLECULAR_FIBER_ORGANIZATION, GOBP_NEGATIVE_REGULATION_OF_SUPRAMOLECULAR_FIBER_ORGANIZATION
- MIR548AY_3P, MIR548AY_3P
- MIR6509_3P, MIR6509_3P
- MIR7155_5P, MIR7155_5P
- GOBP_PEPTIDYL_LYSINE_TRIMETHYLATION, GOBP_PEPTIDYL_LYSINE_TRIMETHYLATION
- GOBP_REGULATION_OF_GLUCOSE_METABOLIC_PROCESS, GOBP_REGULATION_OF_GLUCOSE_METABOLIC_PROCESS
- GOBP_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION, GOBP_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION
- MIR599, MIR599
- GSE3720_UNSTIM_VS_LPS_STIM_VD2_GAMMADELTA_TCELL_DN, GSE3720_UNSTIM_VS_LPS_STIM_VD2_GAMMADELTA_TCELL_DN
- MIR637, MIR637
- MIR3150A_3P, MIR3150A_3P
- MIR6763_5P, MIR6763_5P
- GOBP_POSITIVE_REGULATION_OF_CARBOHYDRATE_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_CARBOHYDRATE_METABOLIC_PROCESS
- HP_BUNDLE_BRANCH_BLOCK, HP_BUNDLE_BRANCH_BLOCK
- GOBP_MONOCARBOXYLIC_ACID_TRANSPORT, GOBP_MONOCARBOXYLIC_ACID_TRANSPORT
- GOBP_NEGATIVE_REGULATION_OF_DNA_REPLICATION, GOBP_NEGATIVE_REGULATION_OF_DNA_REPLICATION
- HP_APLASIA_HYPOPLASIA_OF_THE_TONGUE, HP_APLASIA_HYPOPLASIA_OF_THE_TONGUE
- GSE11367_CTRL_VS_IL17_TREATED_SMOOTH_MUSCLE_CELL_UP, GSE11367_CTRL_VS_IL17_TREATED_SMOOTH_MUSCLE_CELL_UP
- MIR6503_5P, MIR6503_5P
- GSE13522_WT_VS_IFNG_KO_SKIN_DN, GSE13522_WT_VS_IFNG_KO_SKIN_DN
- GOBP_CELLULAR_RESPONSE_TO_ELECTRICAL_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_ELECTRICAL_STIMULUS
- WP_GALANIN_RECEPTOR_PATHWAY, WP_GALANIN_RECEPTOR_PATHWAY
- GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_LIGHTYELLOW_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_LIGHTYELLOW_UP
- MODULE_462, MODULE_462
- GOBP_NEGATIVE_REGULATION_OF_LYASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_LYASE_ACTIVITY
- HP_ADRENAL_HYPOPLASIA, HP_ADRENAL_HYPOPLASIA
- KEGG_PANTOTHENATE_AND_COA_BIOSYNTHESIS, KEGG_PANTOTHENATE_AND_COA_BIOSYNTHESIS
- GOMF_AXON_GUIDANCE_RECEPTOR_ACTIVITY, GOMF_AXON_GUIDANCE_RECEPTOR_ACTIVITY
- MIR4462, MIR4462
- GOBP_NEGATIVE_REGULATION_OF_ADENYLATE_CYCLASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_ADENYLATE_CYCLASE_ACTIVITY