

ELL_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_DAY3_UP, GSE29164_CD8_TCELL_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP, GSE26351_UNSTIM_VS_WNT_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_UP, GSE26351_UNSTIM_VS_WNT_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_DOWN, GSE4142_NAIVE_BCELL_VS_PLASMA_CELL_UP, GSE4142_NAIVE_BCELL_VS_PLASMA_CELL_DOWN, GSE45365_NK_CELL_VS_CD11B_DC_MCMV_INFECTION_DN, GSE45365_NK_CELL_VS_CD11B_DC_MCMV_INFECTION_UP, GSE12392_WT_VS_IFNAR_KO_CD8A_NEG_SPLEEN_DC_UP, GSE12392_WT_VS_IFNAR_KO_CD8A_NEG_SPLEEN_DC_DOWN, GSE32423_CTRL_VS_IL4_MEMORY_CD8_TCELL_UP, GSE32423_CTRL_VS_IL4_MEMORY_CD8_TCELL_DOWN, GSE32423_CTRL_VS_IL7_IL4_MEMORY_CD8_TCELL_UP, GSE32423_CTRL_VS_IL7_IL4_MEMORY_CD8_TCELL_DOWN, GSE40666_UNTREATED_VS_IFNA_STIM_STAT1_KO_CD8_TCELL_90MIN_DN, GSE40666_UNTREATED_VS_IFNA_STIM_STAT1_KO_CD8_TCELL_90MIN_UP, GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_BCL6_HET_UP, GSE28737_FOLLICULAR_VS_MARGINAL_ZONE_BCELL_BCL6_HET_DOWN, GSE40274_LEF1_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_LEF1_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_DOWN, GSE17301_CTRL_VS_48H_ACD3_ACD28_STIM_CD8_TCELL_UP, GSE17301_CTRL_VS_48H_ACD3_ACD28_STIM_CD8_TCELL_DOWN, GSE25123_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_UP, GSE25123_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_DOWN, MIR3663_3P, MIR3663_3P, MIR4273, MIR4273, HP_PROMINENT_NOSE, HP_PROMINENT_NOSE, GSE21927_UNTREATED_VS_GMCSF_IL6_TREATED_BONE_MARROW_DN, GSE21927_UNTREATED_VS_GMCSF_IL6_TREATED_BONE_MARROW_UP, REACTOME_RHOG_GTPASE_CYCLE, REACTOME_RHOG_GTPASE_CYCLE, GAGCCTG_MIR484, GAGCCTG_MIR484, MIR4443, MIR4443, BCAT_BILD_ET_AL_DN, BCAT_BILD_ET_AL_UP, GSE6259_FLT3L_INDUCED_DEC205_POS_DC_VS_BCELL_UP, GSE6259_FLT3L_INDUCED_DEC205_POS_DC_VS_BCELL_DOWN, MIR3660, MIR3660, MIR4526, MIR4526, HP_LOWER_LIMB_MUSCLE_WEAKNESS, HP_LOWER_LIMB_MUSCLE_WEAKNESS, HP_ABNORMAL_MITRAL_VALVE_MORPHOLOGY, HP_ABNORMAL_MITRAL_VALVE_MORPHOLOGY, GSE13522_WT_VS_IFNG_KO_SKING_T_CRUZI_Y_STRAIN_INF_DN, GSE13522_WT_VS_IFNG_KO_SKING_T_CRUZI_Y_STRAIN_INF_UP, TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_DUCTAL_NORMAL_DN, TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_DUCTAL_NORMAL_UP, HP_ELBOW_FLEXION_CONTRACTURE, HP_ELBOW_FLEXION_CONTRACTURE, MIR4639_5P, MIR4639_5P, AR_03, AR_03, MIR4791, MIR4791, BOYALT_LIVER_CANCER_SUBCLASS_G2, BOYALT_LIVER_CANCER_SUBCLASS_G2, HP_LIMITED_HIP_MOVEMENT, HP_LIMITED_HIP_MOVEMENT, GOBP_TELOMERIC_LOOP_DISASSEMBLY, GOBP_TELOMERIC_LOOP_DISASSEMBLY, HP_VASCULAR_TORTUOSITY, HP_VASCULAR_TORTUOSITY, MORF_PDPK1, MORF_PDPK1, WP_NEURODEGENERATION_WITH_BRAIN_IRON_ACCUMULATION_NBIA_SUBTYPES_PATHWAY, WP_NEURODEGENERATION_WITH_BRAIN_IRON_ACCUMULATION_NBIA_SUBTYPES_PATHWAY, GOBP_ANTEROGRADE_AXONAL_TRANSPORT, GOBP_ANTEROGRADE_AXONAL_TRANSPORT, GOBP_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS, GOBP_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS, MIR4772_5P, MIR4772_5P, GOBP_EMBRYONIC_PATTERN_SPECIFICATION, GOBP_EMBRYONIC_PATTERN_SPECIFICATION, CUI_TCF21_TARGETS_DN, CUI_TCF21_TARGETS_DN, GOBP_NEUROMUSCULAR_PROCESS, GOBP_NEUROMUSCULAR_PROCESS, GOBP_TELOMERIC_D_LOOP_DISASSEMBLY, GOBP_TELOMERIC_D_LOOP_DISASSEMBLY, HP_FETAL_AKINESIA_SEQUENCE, HP_FETAL_AKINESIA_SEQUENCE, GOBP_REGULATION_OF_NEUTROPHIL_ACTIVATION, GOBP_REGULATION_OF_NEUTROPHIL_ACTIVATION, GOBP_NEGATIVE_REGULATION_OF_STEROID_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_STEROID_METABOLIC_PROCESS, GLI1_UP.V1_DN, GLI1_UP.V1_DN, WP_15Q112_COPY_NUMBER_VARIATION_SYNDROME, WP_15Q112_COPY_NUMBER_VARIATION_SYNDROME, GOBP_NEGATIVE_REGULATION_OF_ALCOHOL_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_ALCOHOL_BIOSYNTHETIC_PROCESS, HP_UNDETECTABLE_ELECTRORETINOGRAM, HP_UNDETECTABLE_ELECTRORETINOGRAM, RUNNE_GENDER_EFFECT_UP, RUNNE_GENDER_EFFECT_UP, GOBP_S_SHAPED_BODY_MORPHOGENESIS, GOBP_S_SHAPED_BODY_MORPHOGENESIS, GOBP_VESTIBULOCOCHLEAR_NERVE_DEVELOPMENT, GOBP_VESTIBULOCOCHLEAR_NERVE_DEVELOPMENT, DESCARTES_MAIN_FETAL_VASCULAR_ENDOTHELIAL_CELLS, DESCARTES_MAIN_FETAL_VASCULAR_ENDOTHELIAL_CELLS, GOBP_MESONEPHRIC_DUCT_DEVELOPMENT, GOBP_MESONEPHRIC_DUCT_DEVELOPMENT, GOBP_METANEPHRIC_S_SHAPED_BODY_MORPHOGENESIS, GOBP_METANEPHRIC_S_SHAPED_BODY_MORPHOGENESIS, GOBP_METANEPHRIC_DISTAL_TUBULE_DEVELOPMENT, GOBP_METANEPHRIC_DISTAL_TUBULE_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_CGMP_MEDIATED_SIGNALING, GOBP_POSITIVE_REGULATION_OF_CGMP_MEDIATED_SIGNALING, HP_REDUCED_FORCED_VITAL_CAPACITY, HP_REDUCED_FORCED_VITAL_CAPACITY, HP_ABNORMAL_UMBILICAL_STUMP_BLEEDING, HP_ABNORMAL_UMBILICAL_STUMP_BLEEDING, GOBP_METANEPHRIC_PART_OF_URETERIC_BUD_DEVELOPMENT, GOBP_METANEPHRIC_PART_OF_URETERIC_BUD_DEVELOPMENT, HP_TYPE_1_FIBERS_RELATIVELY_SMALLER_THAN_TYPE_2_FIBERS, HP_TYPE_1_FIBERS_RELATIVELY_SMALLER_THAN_TYPE_2_FIBERS, HP_LIMITATION_OF_NECK_MOTION, HP_LIMITATION_OF_NECK_MOTION, GOBP_METANEPHRIC_LOOP_OF_HENLE_DEVELOPMENT, GOBP_METANEPHRIC_LOOP_OF_HENLE_DEVELOPMENT, HP_ARTERIAL RUPTURE, HP_ARTERIAL RUPTURE