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_DAY3_DN, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_DN GSE29164_CD8_TCELL_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP, GSE29164_CD8_TCELL_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP GSE25088_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_UP, GSE25088_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_M 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 GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_UP, GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_UP GSE8868_SPLEEN_VS_INTESTINE_CD11B_POS_CD11C_NEG_DC_UP, GSE8868_SPLEEN_VS_INTESTINE_CD11B_POS_CD11C_NEG_DC_UP GSE411_UNSTIM_VS_100MIN_IL6_STIM_MACROPHAGE_DN, GSE411_UNSTIM_VS_100MIN_IL6_STIM_MACROPHAGE_DN DAZARD_RESPONSE_TO_UV_SCC_DN, DAZARD_RESPONSE_TO_UV_SCC_DN GSE29164_UNTREATED_VS_CD8_TCELL_TREATED_MELANOMA_DAY3_DN, GSE29164_UNTREATED_VS_CD8_TCELL_TREATED_MELANOMA_DAY3_DN GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_45MIN_DN, GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_45MIN_DN GOCC_PHAGOCYTIC_VESICLE, GOCC_PHAGOCYTIC_VESICLE GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_48H_DN, GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_48H_DN GSE17721_CTRL_VS_GARDIQUIMOD_1H_BMDC_DN, GSE17721_CTRL_VS_GARDIQUIMOD_1H_BMDC_DN GSE3982_MAST_CELL_VS_NEUTROPHIL_DN, GSE3982_MAST_CELL_VS_NEUTROPHIL_DN GSE27241_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_TH17_POLARIZING_CONDITIONS_DN, GSE27241_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_TH17_ GSE32034_LY6C_HIGH_VS_LOW_MONOCYTE_UP, GSE32034_LY6C_HIGH_VS_LOW_MONOCYTE_UP MIR6893_5P, MIR6893_5P GSE20366_CD103_POS_VS_CD103_KLRG1_DP_TREG_DN, GSE20366_CD103_POS_VS_CD103_KLRG1_DP_TREG_DN MIR6808_5P, MIR6808_5P MIR4678, MIR4678 MIR6836_5P, MIR6836_5P MIR8066, MIR8066 MIR6131, MIR6131 MIR8082, MIR8082 MIR6132, MIR6132 GSE19941 LPS VS LPS AND IL10 STIM IL10 KO MACROPHAGE DN, GSE19941 LPS VS LPS AND IL10 STIM IL10 KO MACROPHAGE DN MIR6073, MIR6073 BIOCARTA_HIVNEF_PATHWAY, BIOCARTA_HIVNEF_PATHWAY MACAEVA_PBMC_RESPONSE_TO_IR, MACAEVA_PBMC_RESPONSE_TO_IR RAMALHO_STEMNESS_DN, RAMALHO_STEMNESS_DN MIR218_1_3P, MIR218_1_3P GOBP_HOMOTYPIC_CELL_CELL_ADHESION, GOBP_HOMOTYPIC_CELL_CELL_ADHESION BROWNE_HCMV_INFECTION_20HR_DN, BROWNE_HCMV_INFECTION_20HR_DN REACTOME_ONCOGENIC_MAPK_SIGNALING, REACTOME_ONCOGENIC_MAPK_SIGNALING GSE13522_WT_VS_IFNG_KO_SKING_T_CRUZI_Y_STRAIN_INF_DN, GSE13522_WT_VS_IFNG_KO_SKING_T_CRUZI_Y_STRAIN_INF_DN GOBP_LEUKOCYTE_MEDIATED_CYTOTOXICITY, GOBP_LEUKOCYTE_MEDIATED_CYTOTOXICITY MIR4499, MIR4499 MIR10524_5P, MIR10524_5P GOBP_POSITIVE_REGULATION_OF_CELL_KILLING, GOBP_POSITIVE_REGULATION_OF_CELL_KILLING HP_ABNORMALITY_OF_VITAMIN_METABOLISM, HP_ABNORMALITY_OF_VITAMIN_METABOLISM MIR4424, MIR4424 BERNARD_PPAPDC1B_TARGETS_UP, BERNARD_PPAPDC1B_TARGETS_UP GOBP NUCLEAR TRANSCRIBED MRNA POLY A TAIL SHORTENING, GOBP NUCLEAR TRANSCRIBED MRNA POLY A TAIL SHORTENING REACTOME_SIGNALING_BY_BRAF_AND_RAF_FUSIONS, REACTOME_SIGNALING_BY_BRAF_AND_RAF_FUSIONS GOBP_PLATELET_AGGREGATION, GOBP_PLATELET_AGGREGATION MIR4718, MIR4718 GOBP_REGULATION_OF_CELL_KILLING, GOBP_REGULATION_OF_CELL_KILLING MIR4266, MIR4266 MIR5591_5P, MIR5591_5P MIR6758_3P, MIR6758_3P MIR203B_3P, MIR203B_3P GOBP_PROTEIN_LOCALIZATION_TO_CELL_JUNCTION, GOBP_PROTEIN_LOCALIZATION_TO_CELL_JUNCTION HALLMARK_IL6_JAK_STAT3_SIGNALING, HALLMARK_IL6_JAK_STAT3_SIGNALING GROSS HYPOXIA VIA ELK3 ONLY UP, GROSS HYPOXIA VIA ELK3 ONLY UP GOBP ACTIVATION OF JUN KINASE ACTIVITY, GOBP ACTIVATION OF JUN KINASE ACTIVITY MIR96_3P, MIR96_3P MIR3188, MIR3188 MIR5591_3P, MIR5591_3P GOMF_MYOSIN_BINDING, GOMF_MYOSIN_BINDING GOBP_SUPEROXIDE_METABOLIC_PROCESS, GOBP_SUPEROXIDE_METABOLIC_PROCESS REACTOME_SIGNALING_BY_MODERATE_KINASE_ACTIVITY_BRAF_MUTANTS, REACTOME_SIGNALING_BY_MODERATE_KINASE_ACTIVITY_BRAF_MUTANTS HP_MANIA, HP_MANIA MIR324_3P, MIR324_3P HP_PREMATURE_SKIN_WRINKLING, HP_PREMATURE_SKIN_WRINKLING MIR4703_3P, MIR4703_3P GOBP_CELL_KILLING, GOBP_CELL_KILLING REACTOME_APOPTOTIC_CLEAVAGE_OF_CELLULAR_PROTEINS, REACTOME_APOPTOTIC_CLEAVAGE_OF_CELLULAR_PROTEINS HP_AUTOIMMUNE_THROMBOCYTOPENIA, HP_AUTOIMMUNE_THROMBOCYTOPENIA MIR2114_5P, MIR2114_5P GOBP_SUPEROXIDE_ANION_GENERATION, GOBP_SUPEROXIDE_ANION_GENERATION HP_DECREASED_MUSCLE_MASS, HP_DECREASED_MUSCLE_MASS GOBP_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING, GOBP_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING HP_ABNORMAL_ENDOCARDIUM_MORPHOLOGY, HP_ABNORMAL_ENDOCARDIUM_MORPHOLOGY REACTOME_REGULATION_OF_IFNA_SIGNALING, REACTOME_REGULATION_OF_IFNA_SIGNALING GOBP_REGULATION_OF_BROWN_FAT_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_BROWN_FAT_CELL_DIFFERENTIATION REACTOME_VXPX_CARGO_TARGETING_TO_CILIUM, REACTOME_VXPX_CARGO_TARGETING_TO_CILIUM HP_BUNDLE_BRANCH_BLOCK, HP_BUNDLE_BRANCH_BLOCK GOBP_CENTRAL_NERVOUS_SYSTEM_NEURON_DEVELOPMENT, GOBP_CENTRAL_NERVOUS_SYSTEM_NEURON_DEVELOPMENT HP_ANTINEUTROPHIL_ANTIBODY_POSITIVITY, HP_ANTINEUTROPHIL_ANTIBODY_POSITIVITY GAUSSMANN_MLL_AF4_FUSION_TARGETS_D_UP, GAUSSMANN_MLL_AF4_FUSION_TARGETS_D_UP GOMF_TOLL_LIKE_RECEPTOR_BINDING, GOMF_TOLL_LIKE_RECEPTOR_BINDING GOBP_REGULATION_OF_VASCULOGENESIS, GOBP_REGULATION_OF_VASCULOGENESIS GOBP POSITIVE REGULATION OF BROWN FAT CELL DIFFERENTIATION, GOBP POSITIVE REGULATION OF BROWN FAT CELL DIFFERENTIATION HP_FIBROADENOMA_OF_THE_BREAST, HP_FIBROADENOMA_OF_THE_BREAST GOBP_NATURAL_KILLER_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_NATURAL_KILLER_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE GOBP_POSITIVE_REGULATION_OF_VASCULOGENESIS, GOBP_POSITIVE_REGULATION_OF_VASCULOGENESIS HP_HYPERSPLENISM, HP_HYPERSPLENISM HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_VESTIBULE_OF_THE_INNER_EAR, HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_VESTIBULE_OF_THE_INNER_EAR HP_ABNORMAL_PROPORTION_OF_CD8_POSITIVE_T_CELLS, HP_ABNORMAL_PROPORTION_OF_CD8_POSITIVE_T_CELLS WP_FGFR3_SIGNALLING_IN_CHONDROCYTE_PROLIFERATION_AND_TERMINAL_DIFFERENTIATION, WP_FGFR3_SIGNALLING_IN_CHONDROCYTE_PROLIFERATION_AND_TERMINAL FOURNIER_ACINAR_DEVELOPMENT_EARLY_UP, FOURNIER_ACINAR_DEVELOPMENT_EARLY_UP GOBP_POSITIVE_REGULATION_OF_KILLING_OF_CELLS_OF_OTHER_ORGANISM, GOBP_POSITIVE_REGULATION_OF_KILLING_OF_CELLS_OF_OTHER_ORGANISM HP_INCREASED_CIRCULATING_IGA_LEVEL, HP_INCREASED_CIRCULATING_IGA_LEVEL HP_THYROID_ADENOMA, HP_THYROID_ADENOMA HP ABNORMAL SERUM INTERLEUKIN LEVEL, HP ABNORMAL SERUM INTERLEUKIN LEVEL HP_FOLLICULAR_HYPERPLASIA, HP_FOLLICULAR_HYPERPLASIA GOBP_REGULATION_OF_KILLING_OF_CELLS_OF_OTHER_ORGANISM, GOBP_REGULATION_OF_KILLING_OF_CELLS_OF_OTHER_ORGANISM HP_APLASIA_OF_THE_SEMICIRCULAR_CANAL, HP_APLASIA_OF_THE_SEMICIRCULAR_CANAL GOBP REGULATION OF DEVELOPMENTAL PIGMENTATION, GOBP REGULATION OF DEVELOPMENTAL PIGMENTATION GOBP_SEROTONIN_TRANSPORT, GOBP_SEROTONIN_TRANSPORT GOBP_RESPONSE_TO_ATP, GOBP_RESPONSE_TO_ATP HP_ABNORMAL_DELAYED_HYPERSENSITIVITY_SKIN_TEST, HP_ABNORMAL_DELAYED_HYPERSENSITIVITY_SKIN_TEST GOBP_ENDOTHELIAL_CELL_MATRIX_ADHESION, GOBP_ENDOTHELIAL_CELL_MATRIX_ADHESION REACTOME_SHC1_EVENTS_IN_ERBB2_SIGNALING, REACTOME_SHC1_EVENTS_IN_ERBB2_SIGNALING GOBP_ESTABLISHMENT_OF_ENDOTHELIAL_INTESTINAL_BARRIER, GOBP_ESTABLISHMENT_OF_ENDOTHELIAL_INTESTINAL_BARRIER REACTOME_REGULATION_BY_C_FLIP, REACTOME_REGULATION_BY_C_FLIP GOBP_MACROPHAGE_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_MACROPHAGE_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE HP_ABNORMALITY_OF_MACULAR_PIGMENTATION, HP_ABNORMALITY_OF_MACULAR_PIGMENTATION GOBP_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_MATRIX_ADHESION_VIA_FIBRONECTIN, GOBP_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_MATRIX_ADHESION_VIA_ HP_PULVERULENT_CATARACT, HP_PULVERULENT_CATARACT HP_LENS_LUXATION, HP_LENS_LUXATION HP ABNORMALITY OF LENS SHAPE, HP ABNORMALITY OF LENS SHAPE GOMF_AMMONIA_LYASE_ACTIVITY, GOMF_AMMONIA_LYASE_ACTIVITY REACTOME FASL CD95L SIGNALING, REACTOME FASL CD95L SIGNALING GOCC_PEPTIDASE_INHIBITOR_COMPLEX, GOCC_PEPTIDASE_INHIBITOR_COMPLEX

GOBP_SUBSTRATE_INDEPENDENT_TELENCEPHALIC_TANGENTIAL_MIGRATION, GOBP_SUBSTRATE_INDEPENDENT_TELENCEPHALIC_TANGENTIAL_MIGRATION

HP_CHRONIC_PAIN, HP_CHRONIC_PAIN

chr12q12, chr12q12

L_AND_IL12_TREATED_MELANOMA_DAY7_UP, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP