

MUS_VS_FAT_TREG_UP, GSE7852_THYMUS_VS_FAT_TREG_UP

GSE7852_LN_VS_FAT_TREG_UP, GSE7852_LN_VS_FAT_TREG_UP
GSE8921_3H_VS_24H_TLR1_2_STIM_MONOCYTE_UP, GSE8921_3H_VS_24H_TLR1_2_STIM_MONOCYTE_UP
GSE7852_LN_VS_FAT_TCONV_UP, GSE7852_LN_VS_FAT_TCONV_UP
GSE7852_LN_VS_THYMUS_TREG_DN, GSE7852_LN_VS_THYMUS_TREG_DN
GSE7852_TREG_VS_TCONV_FAT_DN, GSE7852_TREG_VS_TCONV_FAT_DN
GSE7852_THYMUS_VS_FAT_TCONV_UP, GSE7852_THYMUS_VS_FAT_TCONV_UP
MIR4778_5P, MIR4778_5P
GSE19941_UNSTIM_VS_LPS_STIM_IL10_KO_MACROPHAGE_DN, GSE19941_UNSTIM_VS_LPS_STIM_IL10_KO_MACROPHAGE_DN
GSE40666_NAIVE_VS_EFFECTOR_CD8_TCELL_UP, GSE40666_NAIVE_VS_EFFECTOR_CD8_TCELL_UP
GSE37301_CD4_TCELL_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_DN, GSE37301_CD4_TCELL_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_DN
GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_INF_DN, GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_INF_DN
GSE21670_TGFB_VS_TGFB_AND_IL6_TREATED_STAT3_KO_CD4_TCELL_UP, GSE21670_TGFB_VS_TGFB_AND_IL6_TREATED_STAT3_KO_CD4_TCELL_UP
MEK_UP.V1_DN, MEK_UP.V1_DN
GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_DN, GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_DN
GSE22611_UNSTIM_VS_6H_MDP_STIM_NOD2_TRANSDUCED_HEK293T_CELL_DN, GSE22611_UNSTIM_VS_6H_MDP_STIM_NOD2_TRANSDUCED_HEK293T_CELL_DN
GSE20366_CD103_KLRG1_DP_VS_DN_TREG_UP, GSE20366_CD103_KLRG1_DP_VS_DN_TREG_UP
GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN, GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN
GSE14415_FOXP3_KO_NATURAL_TREG_VS_TCONV_DN, GSE14415_FOXP3_KO_NATURAL_TREG_VS_TCONV_DN
GSE40274_FOXP3_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_FOXP3_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE2935_UV_INACTIVATED_VS_LIVE_SENDAI_VIRUS_INF_MACROPHAGE_UP, GSE2935_UV_INACTIVATED_VS_LIVE_SENDAI_VIRUS_INF_MACROPHAGE_UP
GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_UP, GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_UP
GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_2H_UP
MIR6831_5P, MIR6831_5P
GSE14415_ACT_TCONV_VS_ACT_NATURAL_TREG_UP, GSE14415_ACT_TCONV_VS_ACT_NATURAL_TREG_UP
GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_UP, GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_UP
KEGG_PROSTATE_CANCER, KEGG_PROSTATE_CANCER
GOBP_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_POTENTIAL, GOBP_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_POTENTIAL
GSE39820_IL1B_IL6_VS_IL1B_IL6_IL23A_TREATED_CD4_TCELL_UP, GSE39820_IL1B_IL6_VS_IL1B_IL6_IL23A_TREATED_CD4_TCELL_UP
DESCARTES_FETAL_MUSCLE_LYMPHOID_CELLS, DESCARTES_FETAL_MUSCLE_LYMPHOID_CELLS
MIR504_5P, MIR504_5P
MIR4463, MIR4463
MIR4725_5P, MIR4725_5P
GUTIERREZ_CHRONIC_LYMPHOCYTIC_LEUKEMIA_DN, GUTIERREZ_CHRONIC_LYMPHOCYTIC_LEUKEMIA_DN
VANASSE_BCL2_TARGETS_DN, VANASSE_BCL2_TARGETS_DN
YRTCANNRCCG_UNKOWN, YRTCANNRCCG_UNKOWN
BOSCO_TH1_CYTOTOXIC_MODULE, BOSCO_TH1_CYTOTOXIC_MODULE
MIR2116_3P, MIR2116_3P
MIR10524_5P, MIR10524_5P
MEIS1AHOXA9_01, MEIS1AHOXA9_01
GOBP_THYMUS_DEVELOPMENT, GOBP_THYMUS_DEVELOPMENT
GOCC_SWR1_COMPLEX, GOCC_SWR1_COMPLEX
DESCARTES_MAIN_FETAL_THYMOCYTES, DESCARTES_MAIN_FETAL_THYMOCYTES
GOLUB_ALL_VS_AML_UP, GOLUB_ALL_VS_AML_UP
HP_LARGE_BASAL_GANGLIA, HP_LARGE_BASAL_GANGLIA
TENEDINI_MEGAKARYOCYTE_MARKERS, TENEDINI_MEGAKARYOCYTE_MARKERS
GOBP_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_III, GOBP_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_III
MIR549A_5P, MIR549A_5P
GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_UP, GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_UP
MORF_EPHA7, MORF_EPHA7
BILBAN_B_CLL_LPL_UP, BILBAN_B_CLL_LPL_UP
HP_BONE_PAIN, HP_BONE_PAIN
DESCARTES_FETAL_MUSCLE_LYMPHATIC_ENDOTHELIAL_CELLS, DESCARTES_FETAL_MUSCLE_LYMPHATIC_ENDOTHELIAL_CELLS
KEGG_SULFUR_METABOLISM, KEGG_SULFUR_METABOLISM
HP_MYELOPROLIFERATIVE_DISORDER, HP_MYELOPROLIFERATIVE_DISORDER
chr4q25, chr4q25
BIOCARTA_NUCLEARRS_PATHWAY, BIOCARTA_NUCLEARRS_PATHWAY
HP_WEAKNESS_OF_THE_INTRINSIC_HAND_MUSCLES, HP_WEAKNESS_OF_THE_INTRINSIC_HAND_MUSCLES
HP_MILDLY_ELEVATED_CREATINE_KINASE, HP_MILDLY_ELEVATED_CREATINE_KINASE
GOBP_CYTOPLASMIC_TRANSLATIONAL_ELONGATION, GOBP_CYTOPLASMIC_TRANSLATIONAL_ELONGATION
GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_POTENTIAL, GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_POTENTIAL
GOBP_POSITIVE_REGULATION_OF_MEMBRANE_POTENTIAL, GOBP_POSITIVE_REGULATION_OF_MEMBRANE_POTENTIAL
GOCC_OLIGOSACCHARYLTRANSFERASE_COMPLEX, GOCC_OLIGOSACCHARYLTRANSFERASE_COMPLEX
HP_AGYRIA, HP_AGYRIA
GOBP_POSITIVE_REGULATION_OF_MULTICELLULAR_ORGANISM_GROWTH, GOBP_POSITIVE_REGULATION_OF_MULTICELLULAR_ORGANISM_GROWTH
GREENBAUM_E2A_TARGETS_DN, GREENBAUM_E2A_TARGETS_DN
GOBP_LATERAL_SPROUTING_FROM_AN_EPITHELIUM, GOBP_LATERAL_SPROUTING_FROM_AN_EPITHELIUM
GOBP_SIALYLATION, GOBP_SIALYLATION
WP_NUCLEAR_RECEPTORS_IN_LIPID_METABOLISM_AND_TOXICITY, WP_NUCLEAR_RECEPTORS_IN_LIPID_METABOLISM_AND_TOXICITY
GOMF_SIALYLTRANSFERASE_ACTIVITY, GOMF_SIALYLTRANSFERASE_ACTIVITY
chr6q22, chr6q22
REACTOME_ABC_TRANSPORTERS_IN_LIPID_HOMEOSTASIS, REACTOME_ABC_TRANSPORTERS_IN_LIPID_HOMEOSTASIS
MIR551A_MIR551B_3P, MIR551A_MIR551B_3P
chr12q12, chr12q12
REACTOME_HDL_REMODELING, REACTOME_HDL_REMODELING
HP_HYPOPLASIA_OF_THE_IRIS, HP_HYPOPLASIA_OF_THE_IRIS
WEINBERGER_BLOOD_TWINRIX_AGE_20_40_AND_60_84YO_CORRELATED_WITH_HIGH_ANTI_HBS_CONC_AT_WEEK_4_POST_BOOSTER_VACC_1DY_POSITIVE, WEINBERGER_BLOOD_TWINRIX_AGE_20_40_AND_60_84YO_CORRELATED_WITH_HIGH_ANTI_HBS_CONC_AT_WEEK_4_POST_BOOSTER_VACC_1DY_POSITIVE
GOBP_LACRIMAL_GLAND_DEVELOPMENT, GOBP_LACRIMAL_GLAND_DEVELOPMENT
GOBP_PURINE_DEOXYRIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS, GOBP_PURINE_DEOXYRIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_DENDRITIC_CELL_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_DENDRITIC_CELL_APOPTOTIC_PROCESS
HP_NARROW_FORAMEN_OBTURATORIUM, HP_NARROW_FORAMEN_OBTURATORIUM
GOMF_OLFACTORY_RECEPTOR_BINDING, GOMF_OLFACTORY_RECEPTOR_BINDING
GOBP_SENSORY_PERCEPTION_OF_TASTE, GOBP_SENSORY_PERCEPTION_OF_TASTE
GOMF_LIPASE_INHIBITOR_ACTIVITY, GOMF_LIPASE_INHIBITOR_ACTIVITY
HP_LACRIMAL_DUCT_ATRESIA, HP_LACRIMAL_DUCT_ATRESIA
ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_LOW_IGM_RESPONDERS_6HY_1DY_UP, ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_LOW_IGM_RESPONDERS_6HY_1DY_UP
chr5q14, chr5q14
chr4p12, chr4p12
GOBP_THYMOCYTE_MIGRATION, GOBP_THYMOCYTE_MIGRATION
GOMF_ALPHA_N_ACETYLNEURAMINATE_ALPHA_2_8_SIALYLTRANSFERASE_ACTIVITY, GOMF_ALPHA_N_ACETYLNEURAMINATE_ALPHA_2_8_SIALYLTRANSFERASE_ACTIVITY
GOMF_LIPASE_BINDING, GOMF_LIPASE_BINDING
GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEUBIQUITINATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEUBIQUITINATION
GOBP_NEGATIVE_REGULATION_OF_CHOLESTEROL_STORAGE, GOBP_NEGATIVE_REGULATION_OF_CHOLESTEROL_STORAGE
GOBP_POSITIVE_REGULATION_OF_HYPERSENSITIVITY, GOBP_POSITIVE_REGULATION_OF_HYPERSENSITIVITY
GOBP_REGULATION_OF_HYPERSENSITIVITY, GOBP_REGULATION_OF_HYPERSENSITIVITY
HP_INCREASED_SKULL_OSSIFICATION, HP_INCREASED_SKULL_OSSIFICATION