

B_VS_TNFA_IL1B_IL6_PGE_STIM_DC_UP, GSE7509_FCGRIB_VS_TNFA_IL1B_IL6_PGE_STIM_DC_UP

GSE6090_UNSTIM_VS_DC_SIGN_STIM_DC_UP, GSE6090_UNSTIM_VS_DC_SIGN_STIM_DC_UP
GSE7509_UNSTIM_VS_TNFA_IL1B_IL6_PGE_STIM_DC_UP, GSE7509_UNSTIM_VS_TNFA_IL1B_IL6_PGE_STIM_DC_UP
GSE27241_WT_VS_RORGT_KO_TH17_POLARIZED_CD4_TCELL_DN, GSE27241_WT_VS_RORGT_KO_TH17_POLARIZED_CD4_TCELL_DN
GSE6259_DEC205_POS_DC_VS_CD8_TCELL_UP, GSE6259_DEC205_POS_DC_VS_CD8_TCELL_UP
GSE39152_BRAIN_VS_SPLEEN_CD103_NEG_MEMORY_CD8_TCELL_UP, GSE39152_BRAIN_VS_SPLEEN_CD103_NEG_MEMORY_CD8_TCELL_UP
GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN
GSE36891_POLYIC_TLR3_VS_PAM_TLR2_STIM_PERITONEAL_MACROPHAGE_DN, GSE36891_POLYIC_TLR3_VS_PAM_TLR2_STIM_PERITONEAL_MACROPHAGE_DN
GSE27896_HDAC6_KO_VS_WT_TREG_DN, GSE27896_HDAC6_KO_VS_WT_TREG_DN
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP
GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_UP
GOBP_REGULATION_OF_PHOSPHATASE_ACTIVITY, GOBP_REGULATION_OF_PHOSPHATASE_ACTIVITY
KEGG_INSULIN_SIGNALING_PATHWAY, KEGG_INSULIN_SIGNALING_PATHWAY
MIR124_5P, MIR124_5P
PID_MET_PATHWAY, PID_MET_PATHWAY
MIR3135B, MIR3135B
HP_ABNORMALITY_OF_PERIAURICULAR_REGION, HP_ABNORMALITY_OF_PERIAURICULAR_REGION
ACTWSNACTNY_UNKNOWN, ACTWSNACTNY_UNKNOWN
GOCC_AZUROPHIL_GRANULE_MEMBRANE, GOCC_AZUROPHIL_GRANULE_MEMBRANE
CAIRO_LIVER_DEVELOPMENT_UP, CAIRO_LIVER_DEVELOPMENT_UP
GOMF_SNARE_BINDING, GOMF_SNARE_BINDING
GOMF_PROTEIN_SERINE_THREONINE_PHOSPHATASE_ACTIVITY, GOMF_PROTEIN_SERINE_THREONINE_PHOSPHATASE_ACTIVITY
GSE6259_33D1_POS_DC_VS_CD8_TCELL_DN, GSE6259_33D1_POS_DC_VS_CD8_TCELL_DN
GSE6674_UNSTIM_VS_PL2_3_STIM_BCELL_UP, GSE6674_UNSTIM_VS_PL2_3_STIM_BCELL_UP
GOBP_EXOCYTIC_PROCESS, GOBP_EXOCYTIC_PROCESS
HP_ABNORMALITY_OF_THE_ACETABULUM, HP_ABNORMALITY_OF_THE_ACETABULUM
GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MEMBRANE, GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MEMBRANE
MIR4425, MIR4425
ZHAN_MULTIPLE_MYELOMA_CD2_DN, ZHAN_MULTIPLE_MYELOMA_CD2_DN
GOMF_SNAP_RECEPTOR_ACTIVITY, GOMF_SNAP_RECEPTOR_ACTIVITY
MIR4764_5P, MIR4764_5P
GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_PLASMA_MEMBRANE, GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_PLASMA_MEMBRANE
GOCC_SNARE_COMPLEX, GOCC_SNARE_COMPLEX
KEGG_ARGININE_AND_PROLINE_METABOLISM, KEGG_ARGININE_AND_PROLINE_METABOLISM
KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_DEGRADATION, KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_DEGRADATION
MIR3120_5P, MIR3120_5P
WP_WNT_SIGNALING, WP_WNT_SIGNALING
VANHARANTA_UTERINE_FIBROID_WITH_7Q_DELETION_DN, VANHARANTA_UTERINE_FIBROID_WITH_7Q_DELETION_DN
BIOCARTA_GPCR_PATHWAY, BIOCARTA_GPCR_PATHWAY
GOMF_SYNTAXIN_BINDING, GOMF_SYNTAXIN_BINDING
MIR12115, MIR12115
GOBP_GOLGI_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT, GOBP_GOLGI_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT
GOMF_HYDROLASE_ACTIVITY_ACTING_ON_CARBON_NITROGEN_BUT_NOT_PEPTIDE_BONDS_IN_LINEAR_AMIDES, GOMF_HYDROLASE_ACTIVITY_ACTING_ON_CARBON_NITROGEN_BUT_NOT_PEPTIDE_BONDS_IN_LINEAR_AMIDES
HP_HYPERINTENSITY_OF_CEREBRAL_WHITE_MATTER_ON_MRI, HP_HYPERINTENSITY_OF_CEREBRAL_WHITE_MATTER_ON_MRI
KEGG_FATTY_ACID_METABOLISM, KEGG_FATTY_ACID_METABOLISM
ERWIN_COHEN_BLOOD_VACCINE_TC_83_AGE_23_48YO_VACCINATED_VS_CONTROL_2DY_DN, ERWIN_COHEN_BLOOD_VACCINE_TC_83_AGE_23_48YO_VACCINATED_VS_CONTROL_2DY_DN
GOBP_FACE_DEVELOPMENT, GOBP_FACE_DEVELOPMENT
HP_ATROPHY_DEGENERATION_INVOLVING_THE_CORTICOSPINAL_TRACTS, HP_ATROPHY_DEGENERATION_INVOLVING_THE_CORTICOSPINAL_TRACTS
PID_AR_NONGENOMIC_PATHWAY, PID_AR_NONGENOMIC_PATHWAY
PID_S1P_S1P3_PATHWAY, PID_S1P_S1P3_PATHWAY
SAKAI_TUMOR_INFILTRATING_MONOCYTES_UP, SAKAI_TUMOR_INFILTRATING_MONOCYTES_UP
GOBP TRABECULA MORPHOGENESIS, GOBP TRABECULA MORPHOGENESIS
PID_ER_NONGENOMIC_PATHWAY, PID_ER_NONGENOMIC_PATHWAY
PID_SHP2_PATHWAY, PID_SHP2_PATHWAY
GOBP_REGULATION_OF_GOLGI_ORGANIZATION, GOBP_REGULATION_OF_GOLGI_ORGANIZATION
KAPOSI_LIVER_CANCER_MET_UP, KAPOSI_LIVER_CANCER_MET_UP
NIKOLSKY_BREAST_CANCER_12Q13_Q21_AMPLICON, NIKOLSKY_BREAST_CANCER_12Q13_Q21_AMPLICON
PEPPER_CHRONIC_LYMPHOCYTIC_LEUKEMIA_UP, PEPPER_CHRONIC_LYMPHOCYTIC_LEUKEMIA_UP
BIOCARTA_RAB_PATHWAY, BIOCARTA_RAB_PATHWAY
WP_SIGNAL_TRANSDUCTION_OF_S1P_RECEPTOR, WP_SIGNAL_TRANSDUCTION_OF_S1P_RECEPTOR
GOBP_CALCIIUM_ION_REGULATED_EXOCYTOSIS, GOBP_CALCIIUM_ION_REGULATED_EXOCYTOSIS
GOBP_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_PARTICLE_METABOLIC_PROCESS, GOBP_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_PARTICLE_METABOLIC_PROCESS
WP_CANNABINOID_RECEPTOR_SIGNALING, WP_CANNABINOID_RECEPTOR_SIGNALING
PID_S1P_S1P1_PATHWAY, PID_S1P_S1P1_PATHWAY
BOYAUULT_LIVER_CANCER_SUBCLASS_G3_DN, BOYAUULT_LIVER_CANCER_SUBCLASS_G3_DN
GOBP_REGULATION_OF_VESICLE_SIZE, GOBP_REGULATION_OF_VESICLE_SIZE
HP_SPINAL_CORD_COMPRESSION, HP_SPINAL_CORD_COMPRESSION
STEGEER_ADIPOGENESIS_DN, STEGEER_ADIPOGENESIS_DN
MODULE_440, MODULE_440
REACTOME_PHASE_I_FUNCTIONALIZATION_OF_COMPOUNDS, REACTOME_PHASE_I_FUNCTIONALIZATION_OF_COMPOUNDS
REACTOME_CLEC7A_DECTIN_1_INDUCES_NFAT_ACTIVATION, REACTOME_CLEC7A_DECTIN_1_INDUCES_NFAT_ACTIVATION
HP_BROAD_METACARPALS, HP_BROAD_METACARPALS
TERAMOTO_OPN_TARGETS_CLUSTER_6, TERAMOTO_OPN_TARGETS_CLUSTER_6
MOOTHA_GLYCOGEN_METABOLISM, MOOTHA_GLYCOGEN_METABOLISM
WP_IL9_SIGNALING_PATHWAY, WP_IL9_SIGNALING_PATHWAY
GOBP_POSITIVE_REGULATION_OF_CELLULAR_CARBOHYDRATE_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_CELLULAR_CARBOHYDRATE_METABOLIC_PROCESS
GOCC_INTRINSIC_COMPONENT_OF_SYNAPTIC_VESICLE_MEMBRANE, GOCC_INTRINSIC_COMPONENT_OF_SYNAPTIC_VESICLE_MEMBRANE
WP_GABA_RECEPTOR_SIGNALING, WP_GABA_RECEPTOR_SIGNALING
BUSSLINGER_ESOPHAGEAL_LATE_SUPRABASAL_CELLS, BUSSLINGER_ESOPHAGEAL_LATE_SUPRABASAL_CELLS
GSE21033_3H_VS_12H_POLYIC_STIM_DC_DN, GSE21033_3H_VS_12H_POLYIC_STIM_DC_DN
ROZANOV_MMP14_TARGETS_DN, ROZANOV_MMP14_TARGETS_DN
HP_GOITER, HP_GOITER
PID_S1P_S1P4_PATHWAY, PID_S1P_S1P4_PATHWAY
HP_ABNORMAL_HEAD_MOVEMENTS, HP_ABNORMAL_HEAD_MOVEMENTS
GOCC_PSEUDOPODIUM, GOCC_PSEUDOPODIUM
GOBP_TRACHEA_MORPHOGENESIS, GOBP_TRACHEA_MORPHOGENESIS
HP_OROFACIAL_DYSKINESIA, HP_OROFACIAL_DYSKINESIA
DARWICHE_PAPILLOMA_PROGRESSION_RISK, DARWICHE_PAPILLOMA_PROGRESSION_RISK
GOMF_MHC_CLASS_IIB_PROTEIN_BINDING, GOMF_MHC_CLASS_IIB_PROTEIN_BINDING
WP_FGFR3_SIGNALLING_IN_CHONDROCYTE_PROLIFERATION_AND_TERMINAL_DIFFERENTIATION, WP_FGFR3_SIGNALLING_IN_CHONDROCYTE_PROLIFERATION_AND_TERMINAL_DIFFERENTIATION
WP_NO_METABOLISM_IN_CYSTIC_FIBROSIS, WP_NO_METABOLISM_IN_CYSTIC_FIBROSIS
GOBP_THYROID_GLAND_DEVELOPMENT, GOBP_THYROID_GLAND_DEVELOPMENT
GOBP_POSITIVE_REGULATION_OF_GLUCOSE_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_GLUCOSE_METABOLIC_PROCESS
FRASOR_TAMOXIFEN_RESPONSE_DN, FRASOR_TAMOXIFEN_RESPONSE_DN
ZHAN_MULTIPLE_MYELOMA_DN, ZHAN_MULTIPLE_MYELOMA_DN
HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_MIDDLE_EAR, HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_MIDDLE_EAR
HP_BRONCHOMALACIA, HP_BRONCHOMALACIA
HP_LYMPHANGIOMA, HP_LYMPHANGIOMA
TRAVAGLINI_LUNG_MYELOID_DENDRITIC_TYPE_1_CELL, TRAVAGLINI_LUNG_MYELOID_DENDRITIC_TYPE_1_CELL
GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT, GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT
GOBP_NEGATIVE_REGULATION_OF_REGULATED_SECRETORY_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_REGULATED_SECRETORY_PATHWAY
BOYERINAS_ONCOFETAL_TARGETS_OF_LET7A1, BOYERINAS_ONCOFETAL_TARGETS_OF_LET7A1
GOBP_COMMISSURAL_NEURON_AXON_GUIDANCE, GOBP_COMMISSURAL_NEURON_AXON_GUIDANCE
GOMF_SYNTAXIN_1_BINDING, GOMF_SYNTAXIN_1_BINDING
WANG_RESPONSE_TO_BEXAROTENE_UP, WANG_RESPONSE_TO_BEXAROTENE_UP