

OSIGLITAZONE_STIM_MACROPHAGE_6H_UP, GSE16386_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_6H_UP

GSE27786_LSK_VS_ERYTHROBLAST_UP, GSE27786_LSK_VS_ERYTHROBLAST_UP
GSE36392_EOSINOPHIL_VS_MAC_IL25_TREATED_LUNG_DN, GSE36392_EOSINOPHIL_VS_MAC_IL25_TREATED_LUNG_DN
GSE27786_LSK_VS_NEUTROPHIL_UP, GSE27786_LSK_VS_NEUTROPHIL_UP
GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DN, GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DN
GSE27786_NKCELL_VS_NEUTROPHIL_UP, GSE27786_NKCELL_VS_NEUTROPHIL_UP
GSE25123_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_DAY10_UP, GSE25123_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_DAY10_UP
GSE36392_TYPE_2_MYELOID_VS_MAC_IL25_TREATED_LUNG_DN, GSE36392_TYPE_2_MYELOID_VS_MAC_IL25_TREATED_LUNG_DN
GSE10273_HIGH_IL7_VS_HIGH_IL7_AND_IRF4_IN_IRF4_8_NULL_PRE_BCELL_UP, GSE10273_HIGH_IL7_VS_HIGH_IL7_AND_IRF4_IN_IRF4_8_NULL_PRE_BCELL_UP
GSE14308_TH17_VS_NAIVE_CD4_TCELL_UP, GSE14308_TH17_VS_NAIVE_CD4_TCELL_UP
GSE14308_TH2_VS_INDUCED_TREG_DN, GSE14308_TH2_VS_INDUCED_TREG_DN
GSE17721_LPS_VS_GARDIQUIMOD_12H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_12H_BMDC_DN
GSE14308_TH17_VS_INDUCED_TREG_DN, GSE14308_TH17_VS_INDUCED_TREG_DN
FOSTER_KDM1A_TARGETS_DN, FOSTER_KDM1A_TARGETS_DN
GSE21670_TGFB_VS_IL6_TREATED_CD4_TCELL_DN, GSE21670_TGFB_VS_IL6_TREATED_CD4_TCELL_DN
GSE14308_TH1_VS_INDUCED_TREG_DN, GSE14308_TH1_VS_INDUCED_TREG_DN
GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_180MIN_UP, GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_180MIN_UP
GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_MDC_DN, GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_MDC_DN
GSE10147_IL3_AND_HIVP17_VS_IL3_AND_CPG_STIM_PDC_DN, GSE10147_IL3_AND_HIVP17_VS_IL3_AND_CPG_STIM_PDC_DN
HP_BULBOUS_NOSE, HP_BULBOUS_NOSE
WAKASUGI_HAVE_ZNF143_BINDING_SITES, WAKASUGI_HAVE_ZNF143_BINDING_SITES
GSE27786_CD4_TCELL_VS_ERYTHROBLAST_DN, GSE27786_CD4_TCELL_VS_ERYTHROBLAST_DN
ZNF677_TARGET_GENES, ZNF677_TARGET_GENES
GGCNRNWCTTYS_UNKNOWN, GGCNRNWCTTYS_UNKNOWN
GSE27786_NKCELL_VS_ERYTHROBLAST_DN, GSE27786_NKCELL_VS_ERYTHROBLAST_DN
MIR2278, MIR2278
ZHONG_PFC_C4_UNKNOWN_INP, ZHONG_PFC_C4_UNKNOWN_INP
MIR1266_3P, MIR1266_3P
LEE_AGING_NEOCORTEX_UP, LEE_AGING_NEOCORTEX_UP
GOBP_REGULATION_OF_MEMBRANE_LIPID_DISTRIBUTION, GOBP_REGULATION_OF_MEMBRANE_LIPID_DISTRIBUTION
GSE41176_WT_VS_TAK1_KO_ANTLIGM_STIM_BCELL_3H_UP, GSE41176_WT_VS_TAK1_KO_ANTLIGM_STIM_BCELL_3H_UP
GOBP_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION
GOBP_MITOTIC_G2_M_TRANSITION_CHECKPOINT, GOBP_MITOTIC_G2_M_TRANSITION_CHECKPOINT
PRC2_SUZ12_UP.V1_DN, PRC2_SUZ12_UP.V1_DN
HP_SPASTIC_DYSARTHRIA, HP_SPASTIC_DYSARTHRIA
MIR6813_3P, MIR6813_3P
GOBP_PYRIDINE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_PYRIDINE_CONTAINING_COMPOUND_METABOLIC_PROCESS
MIR3605_5P, MIR3605_5P
ZNF169_TARGET_GENES, ZNF169_TARGET_GENES
HP_ABNORMALITY_OF_LOWER_LIMB_EPIPHYSIS_MORPHOLOGY, HP_ABNORMALITY_OF_LOWER_LIMB_EPIPHYSIS_MORPHOLOGY
GOBP_BENZENE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_BENZENE_CONTAINING_COMPOUND_METABOLIC_PROCESS
MEIS1BHOXA9_01, MEIS1BHOXA9_01
GOBP_PEPTIDYL_THREONINE_DEPHOSPHORYLATION, GOBP_PEPTIDYL_THREONINE_DEPHOSPHORYLATION
GOBP_LYMPHOID_PROGENITOR_CELL_DIFFERENTIATION, GOBP_LYMPHOID_PROGENITOR_CELL_DIFFERENTIATION
GOBP_MEMBRANE_LIPID_CATABOLIC_PROCESS, GOBP_MEMBRANE_LIPID_CATABOLIC_PROCESS
AZARE_STAT3_TARGETS, AZARE_STAT3_TARGETS
SU_LIVER, SU_LIVER
GOBP_PYRIDINE_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS, GOBP_PYRIDINE_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS
DESCARTES_MAIN_FETAL_HEPATOBLASTS, DESCARTES_MAIN_FETAL_HEPATOBLASTS
GNF2_PTX3, GNF2_PTX3
MIR7106_3P, MIR7106_3P
GOBP_AMINE_CATABOLIC_PROCESS, GOBP_AMINE_CATABOLIC_PROCESS
MIR6803_5P, MIR6803_5P
GOMF_GAMMA_TUBULIN_BINDING, GOMF_GAMMA_TUBULIN_BINDING
GOBP_RESPONSE_TO_ARSENIC_CONTAINING_SUBSTANCE, GOBP_RESPONSE_TO_ARSENIC_CONTAINING_SUBSTANCE
MIR6751_5P, MIR6751_5P
GOBP_XENOBIOTIC_CATABOLIC_PROCESS, GOBP_XENOBIOTIC_CATABOLIC_PROCESS
GOBP_GLUTATHIONE_DERIVATIVE_METABOLIC_PROCESS, GOBP_GLUTATHIONE_DERIVATIVE_METABOLIC_PROCESS
SPIRA_SMOKERS_LUNG_CANCER_DN, SPIRA_SMOKERS_LUNG_CANCER_DN
GOCC_PHAGOPHORE_ASSEMBLY_SITE_MEMBRANE, GOCC_PHAGOPHORE_ASSEMBLY_SITE_MEMBRANE
GOCC_CYTOPLASMIC_UBIQUITIN_LIGASE_COMPLEX, GOCC_CYTOPLASMIC_UBIQUITIN_LIGASE_COMPLEX
GOBP_PHOSPHATIDIC_ACID_METABOLIC_PROCESS, GOBP_PHOSPHATIDIC_ACID_METABOLIC_PROCESS
ZNF20_TARGET_GENES, ZNF20_TARGET_GENES
GOMF_GLUTATHIONE_TRANSFERASE_ACTIVITY, GOMF_GLUTATHIONE_TRANSFERASE_ACTIVITY
HALLMARK_APICAL_SURFACE, HALLMARK_APICAL_SURFACE
GOMF_ANNEALING_ACTIVITY, GOMF_ANNEALING_ACTIVITY
GOMF_OLIGOPEPTIDE_BINDING, GOMF_OLIGOPEPTIDE_BINDING
GOBP_CELLULAR_RESPONSE_TO_ARSENIC_CONTAINING_SUBSTANCE, GOBP_CELLULAR_RESPONSE_TO_ARSENIC_CONTAINING_SUBSTANCE
GOBP_POSITIVE_REGULATION_OF_UTERINE_SMOOTH_MUSCLE_CONTRACTION, GOBP_POSITIVE_REGULATION_OF_UTERINE_SMOOTH_MUSCLE_CONTRACTION
GOBP_URATE_METABOLIC_PROCESS, GOBP_URATE_METABOLIC_PROCESS
SESTO_RESPONSE_TO_UV_C3, SESTO_RESPONSE_TO_UV_C3
GOBP_RESPONSE_TO_INTERLEUKIN_6, GOBP_RESPONSE_TO_INTERLEUKIN_6
GOBP_RESPONSE_TO_MANGANESE_ION, GOBP_RESPONSE_TO_MANGANESE_ION
GOBP_CEREBELLAR_PURKINJE_CELL_LAYER_MORPHOGENESIS, GOBP_CEREBELLAR_PURKINJE_CELL_LAYER_MORPHOGENESIS
MCGARVEY_SILENCED_BY_METHYLATION_IN_COLON_CANCER, MCGARVEY_SILENCED_BY_METHYLATION_IN_COLON_CANCER
GOBP_PYRIMIDINE_CONTAINING_COMPOUND_TRANSMEMBRANE_TRANSPORT, GOBP_PYRIMIDINE_CONTAINING_COMPOUND_TRANSMEMBRANE_TRANSPORT
GOBP_CEREBELLAR_PURKINJE_CELL_LAYER_DEVELOPMENT, GOBP_CEREBELLAR_PURKINJE_CELL_LAYER_DEVELOPMENT
HP_T_CELL_LYMPHOMA, HP_T_CELL_LYMPHOMA
GOBP_NEGATIVE_REGULATION_OF_GASTRIC_ACID_SECRETION, GOBP_NEGATIVE_REGULATION_OF_GASTRIC_ACID_SECRETION
HP_INCREASED_URINARY_NON_PROTEINOGENIC_AMINO_ACID_LEVEL, HP_INCREASED_URINARY_NON_PROTEINOGENIC_AMINO_ACID_LEVEL
WP_DEVELOPMENT_OF_PULMONARY_DENDRITIC_CELLS_AND_MACROPHAGE_SUBSETS, WP_DEVELOPMENT_OF_PULMONARY_DENDRITIC_CELLS_AND_MACROPHAGE_SUBSETS
HOEK_PBMCI_NACTIVATED_INFLUENZA_ADULT_IDY_DN, HOEK_PBMCI_NACTIVATED_INFLUENZA_ADULT_IDY_DN
GOBP_PARENTAL_BEHAVIOR, GOBP_PARENTAL_BEHAVIOR
GOBP_PROTEIN_LOCALIZATION_TO_PRESYNAPSE, GOBP_PROTEIN_LOCALIZATION_TO_PRESYNAPSE
GOBP_UTERINE_SMOOTH_MUSCLE_CONTRACTION, GOBP_UTERINE_SMOOTH_MUSCLE_CONTRACTION
GOBP_DETOXIFICATION_OF_NITROGEN_COMPOUND, GOBP_DETOXIFICATION_OF_NITROGEN_COMPOUND
GOBP_HINDBRAIN_MORPHOGENESIS, GOBP_HINDBRAIN_MORPHOGENESIS
GOCC_JUSTAPARANODE_REGION_OF_AXON, GOCC_JUSTAPARANODE_REGION_OF_AXON
HP_MEDIAN_CLEFT_LIP_AND_PALATE, HP_MEDIAN_CLEFT_LIP_AND_PALATE
GOBP_POSITIVE_REGULATION_OF KERATINOCYTE_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF KERATINOCYTE_PROLIFERATION
HP_PAROXYSMAL_DYSKINESIA, HP_PAROXYSMAL_DYSKINESIA
BUCASAS_PBMCI_FLUARIX_FLUVIRIN_CAUCASIAN_MALE_AGE_18_40YO_HIGH_RESPONDERS_1DY_3DY_POSITIVE_PREDICTIVE_OF_TITER, BUCASAS_PBMCI_FLUARIX_FLUVIRIN_CAUCASIAN_MALE_AGE_18_40YO_HIGH_RESPONDERS_1DY_3DY_POSITIVE_PREDICTIVE_OF_TITER
GOBP_NADPH_OXIDATION, GOBP_NADPH_OXIDATION
GOBP_PENILE_ERECTION, GOBP_PENILE_ERECTION
GOBP_POSITIVE_REGULATION_OF EPITHELIAL_CELL_PROLIFERATION_INVOLVED_IN_WOUND_HEALING, GOBP_POSITIVE_REGULATION_OF EPITHELIAL_CELL_PROLIFERATION_INVOLVED_IN_WOUND_HEALING
GOBP_ANTEROGRADE_AXONAL_PROTEIN_TRANSPORT, GOBP_ANTEROGRADE_AXONAL_PROTEIN_TRANSPORT
GOMF_STAT_FAMILY_PROTEIN_BINDING, GOMF_STAT_FAMILY_PROTEIN_BINDING
YANG_BCL3_TARGETS_DN, YANG_BCL3_TARGETS_DN
GOCC_SPERM_FIBROUS_SHEATH, GOCC_SPERM_FIBROUS_SHEATH
GOBP_INSEMINATION, GOBP_INSEMINATION
HP_SOCIAL_AND_OCCUPATIONAL_DETERIORATION, HP_SOCIAL_AND_OCCUPATIONAL_DETERIORATION
HP_LIMITED_ELBOW_FLEXION, HP_LIMITED_ELBOW_FLEXION
GOBP_AGGRESSIVE_BEHAVIOR, GOBP_AGGRESSIVE_BEHAVIOR
GOBP_REGULATION_OF_PENILE_ERECTION, GOBP_REGULATION_OF_PENILE_ERECTION
GOBP_CELL_DIFFERENTIATION_IN_HINDBRAIN, GOBP_CELL_DIFFERENTIATION_IN_HINDBRAIN
GOBP_CATECHOL_CONTAINING_COMPOUND_CATABOLIC_PROCESS, GOBP_CATECHOL_CONTAINING_COMPOUND_CATABOLIC_PROCESS
GOBP_SPERM_EJACULATION, GOBP_SPERM_EJACULATION
GOBP_PYRIMIDINE_NUCLEOBASE_TRANSPORT, GOBP_PYRIMIDINE_NUCLEOBASE_TRANSPORT
GOBP_POSITIVE_REGULATION_OF_PENILE_ERECTION, GOBP_POSITIVE_REGULATION_OF_PENILE_ERECTION
GOMF CARBONYL_REDUCTASE_NADPH_ACTIVITY, GOMF CARBONYL_REDUCTASE_NADPH_ACTIVITY
GOMF_BENZODIAZEPINE_RECEPTOR_BINDING, GOMF_BENZODIAZEPINE_RECEPTOR_BINDING
GOBP_NEGATIVE_REGULATION_OF_SMAD_PROTEIN_COMPLEX_ASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_SMAD_PROTEIN_COMPLEX_ASSEMBLY
GOBP_NEGATIVE_REGULATION_OF_HISTONE_H3_K27_METHYLATION, GOBP_NEGATIVE_REGULATION_OF_HISTONE_H3_K27_METHYLATION
HP_FACIAL_GRIMACING, HP_FACIAL_GRIMACING
TERAMOTO_OPN_TARGETS_CLUSTER_8, TERAMOTO_OPN_TARGETS_CLUSTER_8
REACTOME_DEFECTIVE_B4GALT1_CAUSES_B4GALT1_CDG_CDG_2D, REACTOME_DEFECTIVE_B4GALT1_CAUSES_B4GALT1_CDG_CDG_2D
HP_STARING_GAZE, HP_STARING_GAZE
GOBP_ANATOMICAL_STRUCTURE_REGRESSION, GOBP_ANATOMICAL_STRUCTURE_REGRESSION
WP_METHYLATION_PATHWAYS, WP_METHYLATION_PATHWAYS
GOBP_PLANAR_CELL_POLARITY_PATHWAY_INVOLVED_IN_NEURAL_TUBE_CLOSURE, GOBP_PLANAR_CELL_POLARITY_PATHWAY_INVOLVED_IN_NEURAL_TUBE_CLOSURE
GOBP_CEREBELLAR_CORTEX_MORPHOGENESIS, GOBP_CEREBELLAR_CORTEX_MORPHOGENESIS
GOBP_POSITIVE_REGULATION_OF_DIGESTIVE_SYSTEM_PROCESS, GOBP_POSITIVE_REGULATION_OF_DIGESTIVE_SYSTEM_PROCESS
GOMF_SPHINGOMYELIN_PHOSPHODIESTERASE_ACTIVITY, GOMF_SPHINGOMYELIN_PHOSPHODIESTERASE_ACTIVITY
HP_ONYCHOGRYPOSIS_OF_FINGERNAIL, HP_ONYCHOGRYPOSIS_OF_FINGERNAIL
HP_EURYBLEPHARON, HP_EURYBLEPHARON
GOBP_SPHINGOMYELIN_CATABOLIC_PROCESS, GOBP_SPHINGOMYELIN_CATABOLIC_PROCESS
GOBP_MEMBRANOUS_SEPTUM_MORPHOGENESIS, GOBP_MEMBRANOUS_SEPTUM_MORPHOGENESIS
HP_NEGATIVISM, HP_NEGATIVISM
GOCC_BRUSH_BORDER_MEMBRANE, GOCC_BRUSH_BORDER_MEMBRANE
HOLLEMAN_PREDNISOLONE_RESISTANCE_ALL_DN, HOLLEMAN_PREDNISOLONE_RESISTANCE_ALL_DN
REACTOME_INTERACTION_WITH_CUMULUS_CELLS_AND_THE_ZONA_PELLUCIDA, REACTOME_INTERACTION_WITH_CUMULUS_CELLS_AND_THE_ZONA_PELLUCIDA
KEGG_STEROID_HORMONE_BIOSYNTHESIS, KEGG_STEROID_HORMONE_BIOSYNTHESIS
GOBP_DISACCHARIDE_BIOSYNTHETIC_PROCESS, GOBP_DISACCHARIDE_BIOSYNTHETIC_PROCESS
GOBP_NEUROTRANSMITTER_CATABOLIC_PROCESS, GOBP_NEUROTRANSMITTER_CATABOLIC_PROCESS
GOBP_URATE_TRANSPORT, GOBP_URATE_TRANSPORT
REACTOME_VASOPRESSIN_LIKE_RECEPTORS, REACTOME_VASOPRESSIN_LIKE_RECEPTORS
HP_ONYCHOGRYPOSIS, HP_ONYCHOGRYPOSIS