GSE23925_DARK_ZONE_VS_NAIVE_BCELL_UP, GSE23925_DARK_ZONE_VS_NAIVE_BCELL_UP GSE23925 LIGHT ZONE VS NAIVE BCELL DN, GSE23925 LIGHT ZONE VS NAIVE BCELL DN GSE26669_CD4_VS_CD8_TCELL_IN_MLR_DN, GSE26669_CD4_VS_CD8_TCELL_IN_MLR_DN GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP, GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP, GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_DN, GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_DN MIR603, MIR603 GOMF_RRNA_BINDING, GOMF_RRNA_BINDING FIGUEROA_AML_METHYLATION_CLUSTER_3_UP, FIGUEROA_AML_METHYLATION_CLUSTER_3_UP MIR150 3P, MIR150 3P GSE30153_LUPUS_VS_HEALTHY_DONOR_BCELL_UP, GSE30153_LUPUS_VS_HEALTHY_DONOR_BCELL_UP GOCC_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE, GOCC_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE MIR3668, MIR3668 HP_ABNORMALITY_OF_THE_COAGULATION_CASCADE, HP_ABNORMALITY_OF_THE_COAGULATION_CASCADE FIGUEROA_AML_METHYLATION_CLUSTER_4_UP, FIGUEROA_AML_METHYLATION_CLUSTER_4_UP MIR4647_MIR4662B, MIR4647_MIR4662B MIR218_1_3P, MIR218_1_3P BROWNE_HCMV_INFECTION_10HR_UP, BROWNE_HCMV_INFECTION_10HR_UP GOCC_ER_TO_GOLGI_TRANSPORT_VESICLE_MEMBRANE, GOCC_ER_TO_GOLGI_TRANSPORT_VESICLE_MEMBRANE GOBP_T_CELL_DIFFERENTIATION_IN_THYMUS, GOBP_T_CELL_DIFFERENTIATION_IN_THYMUS CCCACAT_MIR2993P, CCCACAT_MIR2993P GOBP_EXTRACELLULAR_VESICLE_BIOGENESIS, GOBP_EXTRACELLULAR_VESICLE_BIOGENESIS MIR200C 5P, MIR200C 5P GOMF_HISTONE_ACETYLTRANSFERASE_BINDING, GOMF_HISTONE_ACETYLTRANSFERASE_BINDING GOBP_PRODUCTION_OF_SIRNA_INVOLVED_IN_RNA_INTERFERENCE, GOBP_PRODUCTION_OF_SIRNA_INVOLVED_IN_RNA_INTERFERENCE GOCC_AGGRESOME, GOCC_AGGRESOME PID_HES_HEY_PATHWAY, PID_HES_HEY_PATHWAY MIR4259, MIR4259 REACTOME_ALPHA_PROTEIN_KINASE_1_SIGNALING_PATHWAY, REACTOME_ALPHA_PROTEIN_KINASE_1_SIGNALING_PATHWAY JIANG_HYPOXIA_VIA_VHL, JIANG_HYPOXIA_VIA_VHL GOBP_ENDOCARDIAL_CUSHION_FORMATION, GOBP_ENDOCARDIAL_CUSHION_FORMATION GOBP_GLOMERULAR_MESANGIUM_DEVELOPMENT, GOBP_GLOMERULAR_MESANGIUM_DEVELOPMENT HP_LACRIMAL_DUCT_STENOSIS, HP_LACRIMAL_DUCT_STENOSIS REACTOME_RECEPTOR_MEDIATED_MITOPHAGY, REACTOME_RECEPTOR_MEDIATED_MITOPHAGY GOBP_AMYLOID_FIBRIL_FORMATION, GOBP_AMYLOID_FIBRIL_FORMATION HAY_BONE_MARROW_CD34_POS_HSC, HAY_BONE_MARROW_CD34_POS_HSC WP_NEURAL_CREST_DIFFERENTIATION, WP_NEURAL_CREST_DIFFERENTIATION GOMF_VOLUME_SENSITIVE_ANION_CHANNEL_ACTIVITY, GOMF_VOLUME_SENSITIVE_ANION_CHANNEL_ACTIVITY ✓ GOBP_EAR_MORPHOGENESIS, GOBP_EAR_MORPHOGENESIS GOBP_DORSAL_AORTA_MORPHOGENESIS, GOBP_DORSAL_AORTA_MORPHOGENESIS GOBP_RENAL_SYSTEM_VASCULATURE_DEVELOPMENT, GOBP_RENAL_SYSTEM_VASCULATURE_DEVELOPMENT GOBP_ENDOCARDIUM_DEVELOPMENT, GOBP_ENDOCARDIUM_DEVELOPMENT HP_DESQUAMATION_OF_SKIN_SOON_AFTER_BIRTH, HP_DESQUAMATION_OF_SKIN_SOON_AFTER_BIRTH GOBP_DORSAL_AORTA_DEVELOPMENT, GOBP_DORSAL_AORTA_DEVELOPMENT GOBP_PRE_MIRNA_PROCESSING, GOBP_PRE_MIRNA_PROCESSING GOMF_LACTATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_LACTATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY GOBP_B_CELL_DIFFERENTIATION, GOBP_B_CELL_DIFFERENTIATION HP_PSORIASIFORM_DERMATITIS, HP_PSORIASIFORM_DERMATITIS GOMF HIGH VOLTAGE GATED CALCIUM CHANNEL ACTIVITY, GOMF HIGH VOLTAGE GATED CALCIUM CHANNEL ACTIVITY WP_PTF1A_RELATED_REGULATORY_PATHWAY, WP_PTF1A_RELATED_REGULATORY_PATHWAY GOBP_ENDOCARDIAL_CUSHION_MORPHOGENESIS, GOBP_ENDOCARDIAL_CUSHION_MORPHOGENESIS GUTIERREZ_WALDENSTROEMS_MACROGLOBULINEMIA_1_DN, GUTIERREZ_WALDENSTROEMS_MACROGLOBULINEMIA_1_DN GOBP_LACTATE_TRANSMEMBRANE_TRANSPORT, GOBP_LACTATE_TRANSMEMBRANE_TRANSPORT GOBP_CARDIAC_MUSCLE_TISSUE_MORPHOGENESIS, GOBP_CARDIAC_MUSCLE_TISSUE_MORPHOGENESIS HP_SUBGLOTTIC_STENOSIS, HP_SUBGLOTTIC_STENOSIS GOBP_3_PHOSPHOADENOSINE_5_PHOSPHOSULFATE_BIOSYNTHETIC_PROCESS, GOBP_3_PHOSPHOADENOSINE_5_PHOSPHOSULFATE_BIOSYNTHETIC_PROCESS REACTOME_TRANSPORT_AND_SYNTHESIS_OF_PAPS, REACTOME_TRANSPORT_AND_SYNTHESIS_OF_PAPS GOBP_CARDIAC_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_CARDIAC_EPITHELIAL_TO_MESENCHYMAL_TRANSITION GOBP_VENTRICULAR_CARDIAC_MUSCLE_TISSUE_DEVELOPMENT, GOBP_VENTRICULAR_CARDIAC_MUSCLE_TISSUE_DEVELOPMENT GOBP_RESPONSE_TO_NICOTINE, GOBP_RESPONSE_TO_NICOTINE HP_CHRONIC_AXONAL_NEUROPATHY, HP_CHRONIC_AXONAL_NEUROPATHY HP_HYPOPROTEINEMIA, HP_HYPOPROTEINEMIA HP_FOLLICULAR_HYPERKERATOSIS, HP_FOLLICULAR_HYPERKERATOSIS HP_ABNORMAL_CELL_PROLIFERATION, HP_ABNORMAL_CELL_PROLIFERATION HP_INTERMITTENT_DIARRHEA, HP_INTERMITTENT_DIARRHEA HP_ABNORMAL_NATURAL_KILLER_CELL_MORPHOLOGY, HP_ABNORMAL_NATURAL_KILLER_CELL_MORPHOLOGY GOCC_L_TYPE_VOLTAGE_GATED_CALCIUM_CHANNEL_COMPLEX, GOCC_L_TYPE_VOLTAGE_GATED_CALCIUM_CHANNEL_COMPLEX HP_EXERCISE_INDUCED_MUSCLE_STIFFNESS, HP_EXERCISE_INDUCED_MUSCLE_STIFFNESS GOBP_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION, GOBP_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION GOBP_ATRIOVENTRICULAR_VALVE_FORMATION, GOBP_ATRIOVENTRICULAR_VALVE_FORMATION MARSHALL_VIRAL_INFECTION_RESPONSE_UP, MARSHALL_VIRAL_INFECTION_RESPONSE_UP GOMF_STRUCTURAL_CONSTITUENT_OF_MYELIN_SHEATH, GOMF_STRUCTURAL_CONSTITUENT_OF_MYELIN_SHEATH GOBP_ACTOMYOSIN_CONTRACTILE_RING_ORGANIZATION, GOBP_ACTOMYOSIN_CONTRACTILE_RING_ORGANIZATION GOBP_ENDOCARDIAL_CELL_DIFFERENTIATION, GOBP_ENDOCARDIAL_CELL_DIFFERENTIATION GOMF_EPOXIDE_HYDROLASE_ACTIVITY, GOMF_EPOXIDE_HYDROLASE_ACTIVITY HP_ABNORMAL_CARDIOMYOCYTE_MORPHOLOGY, HP_ABNORMAL_CARDIOMYOCYTE_MORPHOLOGY HP_ABNORMALITY_OF_THE_CRUS_OF_THE_HELIX, HP_ABNORMALITY_OF_THE_CRUS_OF_THE_HELIX GOBP_ENDOTHELIAL_CELL_FATE_COMMITMENT, GOBP_ENDOTHELIAL_CELL_FATE_COMMITMENT REACTOME_NONCANONICAL_ACTIVATION_OF_NOTCH3, REACTOME_NONCANONICAL_ACTIVATION_OF_NOTCH3 GOBP_SKELETAL_MUSCLE_SATELLITE_CELL_DIFFERENTIATION, GOBP_SKELETAL_MUSCLE_SATELLITE_CELL_DIFFERENTIATION

GSE14769_UNSTIM_VS_80MIN_LPS_BMDM_UP, GSE14769_UNSTIM_VS_80MIN_LPS_BMDM_UP GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN, GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RESPONSE_TO_CHEMICAL_STIMULUS, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_ REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_CYTOCHROME_C_RELEASE, REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_CYTOCHROME_C_RELEASE MATSUMIYA_PBMC_MODIFIED_VACCINIA_ANKARA_VACCINE_AGE_4_6MO_BCG_PRIMED_28DY_UP, MATSUMIYA_PBMC_MODIFIED_VACCINIA_ANKARA_VACCINE_AGE_4_6MO_BCG_PRIMED_28DY_UP GOBP_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_INVOLVED_IN_ENDOCARDIAL_CUSHION_FORMATION, GOBP_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_INVOLVED_IN_ENDOCARDIAL_CUSHION_FORMATION GOBP_POSITIVE_REGULATION_OF_HIGH_VOLTAGE_GATED_CALCIUM_CHANNEL_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_HIGH_VOLTAGE_GATED_CALCIUM_CHANNEL_ACTIVITY GOBP CARDIAC CELL FATE COMMITMENT, GOBP CARDIAC CELL FATE COMMITMENT GOBP_MYOBLAST_MIGRATION, GOBP_MYOBLAST_MIGRATION