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| | GSE14350_TREG_VS_TEFF_UP, GSE14350_TREG_VS_TEFF_UP GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_DN, GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_DN GSE5679_CTRL_VS_PPARG_LIGAND_ROSIGLITAZONE_TREATED_DC_DN, GSE5679_CTRL_VS_PPARG_LIGAND_ROSIGLITAZONE_TREATED_DC_DN GSE4590_PRE_BCELL_VS_VPREB_POS_LARGE_PRE_BCELL_DN, GSE4590_PRE_BCELL_VS_VPREB_POS_LARGE_PRE_BCELL_DN GSE7348_UNSTIM_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP, GSE7348_UNSTIM_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP GSE28237_FOLLICULAR_VS_EARLY_GC_BCELL_UP, GSE28237_FOLLICULAR_VS_EARLY_GC_BCELL_UP GSE7764_NKCELL_VS_SPLENOCYTE_DN, GSE7764_NKCELL_VS_SPLENOCYTE_DN GSE7568_IL4_VS_IL4_AND_DEXAMETHASONE_TREATED_MACROPHAGE_DN, GSE7568_IL4_VS_IL4_AND_DEXAMETHASONE_TREATED_MACROPHAGE_DN GOCC_TRANSCRIPTION_REPRESSOR_COMPLEX, GOCC_TRANSCRIPTION_REPRESSOR_COMPLEX MIR9500, MIR9500 WP_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY, WP_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY MIR221_3P, MIR221_3P MIR676_5P, MIR676_5P GSE360_I_DONOVANL_VS_B_MALAYL_HIGH_DOSE_MAC_UP, GSE360_I_DONOVANL_VS_B_MALAYL_HIGH_DOSE_MAC_UP GSE22601_DOUBLE_POSITIVE_VS_CD4_SINGLE_POSITIVE_THYMOCYTE_UP, GSE22601_DOUBLE_POSITIVE_VS_CD4_SINGLE_POSITIVE_THYMOCYTE_UP GSE40274_CTRL_VS_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_CTRL_VS_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN GSE37301_HEMATOPOIETIC_STEM_CELL_VS_GRAN_MONO_PROGENITOR_UP, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_GRAN_MONO_PROGENITOR_UP MIR4642, MIR4642 MIR4700_3P, MIR4700_3P HP_KINETIC_TREMOR, HP_KINETIC_TREMOR MIR6871_3P, MIR6871_3P MIR758_3P, MIR758_3P CSE12963_ENV_NEF_VS_ENV_NEF_AND_VPR_DEFICIENT_HIV1_INF_CD4_TCELL_UP, GSE12963_ENV_NEF_VS_ENV_NEF_AND_VPR_DEFICIENT_HIV1_INF_CD4_TCELL_UP MIR6506_5P, MIR6506_5P CSE37605_NOD_VS_C57BL6_IRES_GFP_TREG_DN, GSE37605_NOD_VS_C57BL6_IRES_GFP_TREG_DN MIR511_3P, MIR511_3P JACKSON_DNMT1_TARGETS_UP, JACKSON_DNMT1_TARGETS_UP REACTOME_RHOC_GTPASE_CYCLE, REACTOME_RHOC_GTPASE_CYCLE SHAFFER_IRF4_TARGETS_IN_ACTIVATED_DENDRITIC_CELL, SHAFFER_IRF4_TARGETS_IN_ACTIVATED_DENDRITIC_CELL MIR1237_3P, MIR1237_3P GCTCTTG, MIR335, GCTCTTG, MIR335 GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_DN, GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_DN REACTOME_INOSITOL_PHOSPHATE_METABOLISM, REACTOME_INOSITOL_PHOSPHATE_METABOLISM HP_AORTIC_ROOT_ANEURYSM, HP_AORTIC_ROOT_ANEURYSM MIR664A_5P, MIR664A_5P GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA_STIMULUS, GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA_STIMULUS GSE45365_NK_CELL_VS_CD8A_DC_DN, GSE45365_NK_CELL_VS_CD8A_DC_DN MIR6754_3P, MIR6754_3P GOBP_HIPPO_SIGNALING, GOBP_HIPPO_SIGNALING MIR1261, MIR1261 MIR4645_3P, MIR4645_3P MIR7156_3P, MIR7156_3P GOCC_APICAL_JUNCTION_COMPLEX, GOCC_APICAL_JUNCTION_COMPLEX MIR5588_5P, MIR5588_5P MIR4720_5P, MIR4720_5P MIR4799_3P, MIR4799_3P MIR4524B_3P, MIR4524B_3P GOBP_ACTIVATION_OF_GTPASE_ACTIVITY, GOBP_ACTIVATION_OF_GTPASE_ACTIVITY MIR655_5P, MIR655_5P GOBP_RESPONSE_TO_ACID_CHEMICAL, GOBP_RESPONSE_TO_ACID_CHEMICAL MED25_TARGET_GENES, MED25_TARGET_GENES REACTOME_DOWNREGULATION_OF_SMAD2_3_SMAD4_TRANSCRIPTIONAL_ACTIVITY, REACTOME_DOWNREGULATION_OF_SMAD2_3_SMAD4_TRANSCRIPTIONAL_ACTIVITY GOBP_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION, GOBP_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_RRNA, GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_RRNA GOBP_SEQUESTERING_OF_CALCIUM_ION, GOBP_SEQUESTERING_OF_CALCIUM_ION GOMF_HISTONE_ACETYLTRANSFERASE_BINDING, GOMF_HISTONE_ACETYLTRANSFERASE_BINDING GOCC_TIGHT_JUNCTION, GOCC_TIGHT_JUNCTION BIOCARTA_PPARG_PATHWAY, BIOCARTA_PPARG_PATHWAY MIR4793_5P, MIR4793_5P MIR4268, MIR4268 GOMF_ACTIVATING_TRANSCRIPTION_FACTOR_BINDING, GOMF_ACTIVATING_TRANSCRIPTION_FACTOR_BINDING SS18_SSX1_FUSION_UNIPROT_Q8IZH1_UNREVIEWED_TARGET_GENES, SS18_SSX1_FUSION_UNIPROT_Q8IZH1_UNREVIEWED_TARGET_GENES GOBP_CALCIUM_ION_TRANSPORT_INTO_CYTOSOL, GOBP_CALCIUM_ION_TRANSPORT_INTO_CYTOSOL PID_NETRIN_PATHWAY, PID_NETRIN_PATHWAY GSE8515_IL1_VS_IL6_4H_STIM_MAC_DN, GSE8515_IL1_VS_IL6_4H_STIM_MAC_DN BROWNE_HCMV_INFECTION_8HR_DN, BROWNE_HCMV_INFECTION_8HR_DN MIR1272, MIR1272 GOMF_UBIQUITIN_CONJUGATING_ENZYME_BINDING, GOMF_UBIQUITIN_CONJUGATING_ENZYME_BINDING GOBP_POSITIVE_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_BY_HOST, GOBP_POSITIVE_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_BY_HOST REACTOME_SIGNALING_BY_HIPPO, REACTOME_SIGNALING_BY_HIPPO HP_ABNORMALITY_OF_THE_PUBIC_BONE, HP_ABNORMALITY_OF_THE_PUBIC_BONE STAMBOLSKY_RESPONSE_TO_VITAMIN_D3_UP, STAMBOLSKY_RESPONSE_TO_VITAMIN_D3_UP MIR3663_5P, MIR3663_5P PID_RETINOIC_ACID_PATHWAY, PID_RETINOIC_ACID_PATHWAY ZIRN_TRETINOIN_RESPONSE_WT1_UP, ZIRN_TRETINOIN_RESPONSE_WT1_UP LI_PROSTATE_CANCER_EPIGENETIC, LI_PROSTATE_CANCER_EPIGENETIC BIOCARTA_PDGF_PATHWAY, BIOCARTA_PDGF_PATHWAY MIR4529_5P, MIR4529_5P REACTOME_CELL_JUNCTION_ORGANIZATION, REACTOME_CELL_JUNCTION_ORGANIZATION ZSCAN5B_TARGET_GENES, ZSCAN5B_TARGET_GENES GOBP_REGULATION_OF_PEPTIDYL_THREONINE_PHOSPHORYLATION, GOBP_REGULATION_OF_PEPTIDYL_THREONINE_PHOSPHORYLATION DELACROIX_RAR_TARGETS_DN, DELACROIX_RAR_TARGETS_DN HP_ENLARGED_KIDNEY, HP_ENLARGED_KIDNEY MIR15A_3P, MIR15A_3P HP_DIPLOPIA, HP_DIPLOPIA REACTOME_CONSTITUTIVE_SIGNALING_BY_LIGAND_RESPONSIVE_EGFR_CANCER_VARIANTS, REACTOME_CONSTITUTIVE_SIGNALING_BY_LIGAND_RESPONSIVE_EGFR_CANCER_VARIANTS GOBP_POSITIVE_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY BIOCARTA_ARENERF2_PATHWAY, BIOCARTA_ARENERF2_PATHWAY MIR6888_5P, MIR6888_5P REACTOME_SIGNALING_BY_NTRK3_TRKC, REACTOME_SIGNALING_BY_NTRK3_TRKC GOBP_RAP_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_RAP_PROTEIN_SIGNAL_TRANSDUCTION GOBP_CELL_JUNCTION_MAINTENANCE, GOBP_CELL_JUNCTION_MAINTENANCE WP_ANGIOGENESIS, WP_ANGIOGENESIS GOBP_REGULATION_OF_NITRIC_OXIDE_METABOLIC_PROCESS, GOBP_REGULATION_OF_NITRIC_OXIDE_METABOLIC_PROCESS HP_MICROPHALLUS, HP_MICROPHALLUS GOBP_CELLULAR_RESPONSE_TO_FLUID_SHEAR_STRESS, GOBP_CELLULAR_RESPONSE_TO_FLUID_SHEAR_STRESS HP_ABNORMALITY_OF_BLADDER_MORPHOLOGY, HP_ABNORMALITY_OF_BLADDER_MORPHOLOGY GOBP_MESENCHYMAL_EPITHELIAL_CELL_SIGNALING, GOBP_MESENCHYMAL_EPITHELIAL_CELL_SIGNALING GOBP_REGULATION_OF_BONE_REMODELING, GOBP_REGULATION_OF_BONE_REMODELING REACTOME_SIGNALING_BY_ERBB2_ECD_MUTANTS, REACTOME_SIGNALING_BY_ERBB2_ECD_MUTANTS BIOCARTA_ETS_PATHWAY, BIOCARTA_ETS_PATHWAY HP_DIFFICULTY_STANDING, HP_DIFFICULTY_STANDING GOCC_BRUSH_BORDER_MEMBRANE, GOCC_BRUSH_BORDER_MEMBRANE GOBP_TRYPTOPHAN_METABOLIC_PROCESS, GOBP_TRYPTOPHAN_METABOLIC_PROCESS GOBP_REGULATION_OF_GLUCOSE_TRANSMEMBRANE_TRANSPORT, GOBP_REGULATION_OF_GLUCOSE_TRANSMEMBRANE_TRANSPORT PID_ECADHERIN_KERATINOCYTE_PATHWAY, PID_ECADHERIN_KERATINOCYTE_PATHWAY GOMF_BHLH_TRANSCRIPTION_FACTOR_BINDING, GOMF_BHLH_TRANSCRIPTION_FACTOR_BINDING GOBP_POSITIVE_REGULATION_OF_CALCIUM_ION_TRANSPORT_INTO_CYTOSOL, GOBP_POSITIVE_REGULATION_OF_CALCIUM_ION_TRANSPORT_INTO_CYTOSOL BRF1_TARGET_GENES, BRF1_TARGET_GENES GOBP_NEGATIVE_REGULATION_OF_MUSCLE_ADAPTATION, GOBP_NEGATIVE_REGULATION_OF_MUSCLE_ADAPTATION GOMF_STRUCTURAL_CONSTITUENT_OF_SYNAPSE, GOMF_STRUCTURAL_CONSTITUENT_OF_SYNAPSE REACTOME_CONSTITUTIVE_SIGNALING_BY_EGFRVIII, REACTOME_CONSTITUTIVE_SIGNALING_BY_EGFRVIII GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_THREONINE_PHOSPHORYLATION, GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_THREONINE_PHOSPHORYLATION REACTOME_CELL_CELL_JUNCTION_ORGANIZATION, REACTOME_CELL_CELL_JUNCTION_ORGANIZATION HP ABDOMINAL_WALL_MUSCLE_WEAKNESS, HP ABDOMINAL_WALL_MUSCLE_WEAKNESS GOBP_REGULATION_OF_RESPONSE_TO_REACTIVE_OXYGEN_SPECIES, GOBP_REGULATION_OF_RESPONSE_TO_REACTIVE_OXYGEN_SPECIES HP_LEFT_BUNDLE_BRANCH_BLOCK, HP_LEFT_BUNDLE_BRANCH_BLOCK GOBP_MESONEPHRIC_DUCT_DEVELOPMENT, GOBP_MESONEPHRIC_DUCT_DEVELOPMENT MIR6784_3P, MIR6784_3P GOBP_CEREBELLAR_CORTEX_DEVELOPMENT, GOBP_CEREBELLAR_CORTEX_DEVELOPMENT GOBP_METANEPHRIC_PART_OF_URETERIC_BUD_DEVELOPMENT, GOBP_METANEPHRIC_PART_OF_URETERIC_BUD_DEVELOPMENT GOBP_NUCLEOTIDE_SALVAGE, GOBP_NUCLEOTIDE_SALVAGE chr1q43, chr1q43 GOBP_MULTIORGANISM_CELLULAR_PROCESS, GOBP_MULTIORGANISM_CELLULAR_PROCESS GOBP_BILE_ACID_SIGNALING_PATHWAY, GOBP_BILE_ACID_SIGNALING_PATHWAY WATANABE_COLON_CANCER_MSL_VS_MSS_DN, WATANABE_COLON_CANCER_MSL_VS_MSS_DN GOBP_SPHINGOMYELIN_METABOLIC_PROCESS, GOBP_SPHINGOMYELIN_METABOLIC_PROCESS GOBP_CEREBELLAR_CORTEX_MORPHOGENESIS, GOBP_CEREBELLAR_CORTEX_MORPHOGENESIS HP_FIBULAR_BOWING, HP_FIBULAR_BOWING GOMF_GLUTAMATE_RECEPTOR_BINDING, GOMF_GLUTAMATE_RECEPTOR_BINDING GOBP_REGULATION_OF_BLOOD_BRAIN_BARRIER_PERMEABILITY, GOBP_REGULATION_OF_BLOOD_BRAIN_BARRIER_PERMEABILITY GOBP_DISTAL_TUBULE_DEVELOPMENT, GOBP_DISTAL_TUBULE_DEVELOPMENT OSADA_ASCL1_TARGETS_UP, OSADA_ASCL1_TARGETS_UP HP_OSTEOMALACIA, HP_OSTEOMALACIA GOBP_S_SHAPED_BODY_MORPHOGENESIS, GOBP_S_SHAPED_BODY_MORPHOGENESIS GOBP_NEGATIVE_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATION, GOBP_NEGATIVE_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATION HP_YOUNG_ADULT_ONSET, HP_YOUNG_ADULT_ONSET REACTOME_MET_RECEPTOR_ACTIVATION, REACTOME_MET_RECEPTOR_ACTIVATION GOBP_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_MIGRATION, GOBP_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_MIGRATION GOBP_POSITIVE_REGULATION_OF_LONG_TERM_NEURONAL_SYNAPTIC_PLASTICITY, GOBP_POSITIVE_REGULATION_OF_LONG_TERM_NEURONAL_SYNAPTIC_PLASTICITY GOMF_TRANSMEMBRANE_RECEPTOR_PROTEIN_PHOSPHATASE_ACTIVITY, GOMF_TRANSMEMBRANE_RECEPTOR_PROTEIN_PHOSPHATASE_ACTIVITY GOBP_METANEPHRIC_S_SHAPED_BODY_MORPHOGENESIS, GOBP_METANEPHRIC_S_SHAPED_BODY_MORPHOGENESIS GOBP_POSTSYNAPTIC_ACTIN_CYTOSKELETON_ORGANIZATION, GOBP_POSTSYNAPTIC_ACTIN_CYTOSKELETON_ORGANIZATION HP_PAROXYSMAL_ATRIAL_FIBRILLATION, HP_PAROXYSMAL_ATRIAL_FIBRILLATION MATZUK_CUMULUS_EXPANSION, MATZUK_CUMULUS_EXPANSION HP_EURYBLEPHARON, HP_EURYBLEPHARON FAN_EMBRYONIC_CTX_EX_1_EXCITATORY_NEURON, FAN_EMBRYONIC_CTX_EX_1_EXCITATORY_NEURON GOMF_OUTWARD_RECTIFIER_POTASSIUM_CHANNEL_ACTIVITY, GOMF_OUTWARD_RECTIFIER_POTASSIUM_CHANNEL_ACTIVITY SINGH_KRAS_DEPENDENCY_SIGNATURE, SINGH_KRAS_DEPENDENCY_SIGNATURE GOMF_POTASSIUM_ION_LEAK_CHANNEL_ACTIVITY, GOMF_POTASSIUM_ION_LEAK_CHANNEL_ACTIVITY GOBP_HINDBRAIN_MORPHOGENESIS, GOBP_HINDBRAIN_MORPHOGENESIS GOBP_NEGATIVE_REGULATION_OF_CYTOSOLIC_CALCIUM_ION_CONCENTRATION, GOBP_NEGATIVE_REGULATION_OF_CYTOSOLIC_CALCIUM_ION_CONCENTRATION REACTOME_PHOSPHOLIPASE_C_MEDIATED_CASCADE_FGFR2, REACTOME_PHOSPHOLIPASE_C_MEDIATED_CASCADE_FGFR2 WANG_THOC1_TARGETS_DN, WANG_THOC1_TARGETS_DN HP_DENSE_CALCIFICATIONS_IN_THE_CEREBELLAR_DENTATE_NUCLEUS, HP_DENSE_CALCIFICATIONS_IN_THE_CEREBELLAR_DENTATE_NUCLEUS GOMF_STRUCTURAL_CONSTITUENT_OF_POSTSYNAPTIC_ACTIN_CYTOSKELETON, GOMF_STRUCTURAL_CONSTITUENT_OF_POSTSYNAPTIC_ACTIN_CYTOSKELETON REACTOME_SODIUM_COUPLED_PHOSPHATE_COTRANSPORTERS, REACTOME_SODIUM_COUPLED_PHOSPHATE_COTRANSPORTERS VALK_AML_WITH_T_8_21_TRANSLOCATION, VALK_AML_WITH_T_8_21_TRANSLOCATION HP_SPARSE_BONE TRABECULAE, HP_SPARSE_BONE TRABECULAE GOBP_PROTEIN_LOCALIZATION_TO_BICELLULAR_TIGHT_JUNCTION, GOBP_PROTEIN_LOCALIZATION_TO_BICELLULAR_TIGHT_JUNCTION GOBP_POSITIVE_REGULATION_OF_BLOOD_BRAIN_BARRIER_PERMEABILITY, GOBP_POSITIVE_REGULATION_OF_BLOOD_BRAIN_BARRIER_PERMEABILITY GOBP_REGULATION_OF_LUNG_BLOOD_PRESSURE, GOBP_REGULATION_OF_LUNG_BLOOD_PRESSURE GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_OXIDATIVE_STRESS, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_OXIDATIVE_STRESS GOBP_METANEPHRIC_LOOP_OF_HENLE_DEVELOPMENT, GOBP_METANEPHRIC_LOOP_OF_HENLE_DEVELOPMENT GOBP_POSTSYNAPTIC_CYTOSKELETON_ORGANIZATION, GOBP_POSTSYNAPTIC_CYTOSKELETON_ORGANIZATION GOBP_CELL_DEATH_IN_RESPONSE_TO_HYDROGEN_PEROXIDE, GOBP_CELL_DEATH_IN_RESPONSE_TO_HYDROGEN_PEROXIDE GOBP_POSITIVE_REGULATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_MIGRATION, GOBP_POSITIVE_REGULATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_MIGRATION HP_ENLARGEMENT_OF_THE_ANKLES, HP_ENLARGEMENT_OF_THE_ANKLES HP_ABSENT_BRAINSTEM_AUDITORY_RESPONSES, HP_ABSENT_BRAINSTEM_AUDITORY_RESPONSES REACTOME_TYPE_1_HEMIDESMOSOME_ASSEMBLY, REACTOME_TYPE_1_HEMIDESMOSOME_ASSEMBLY KIM_ALL_DISORDERS_DURATION_CORR_UP, KIM_ALL_DISORDERS_DURATION_CORR_UP GOBP_NEURONAL_SIGNAL_TRANSDUCTION, GOBP_NEURONAL_SIGNAL_TRANSDUCTION GOMF_VITAMIN_D_BINDING, GOMF_VITAMIN_D_BINDING GOBP_METANEPHRIC_DISTAL_TUBULE_DEVELOPMENT, GOBP_METANEPHRIC_DISTAL_TUBULE_DEVELOPMENT GOCC_AXON_INITIAL_SEGMENT, GOCC_AXON_INITIAL_SEGMENT REACTOME_G_ALPHA_S_SIGNALLING_EVENTS, REACTOME_G_ALPHA_S_SIGNALLING_EVENTS HP_RETINOSCHISIS, HP_RETINOSCHISIS GOBP_NEGATIVE_REGULATION_OF_NEUTROPHIL_MIGRATION, GOBP_NEGATIVE_REGULATION_OF_NEUTROPHIL_MIGRATION GOMF_TYPE_1_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_BINDING, GOMF_TYPE_1_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_BINDING REACTOME_MET_INTERACTS_WITH_TNS_PROTEINS, REACTOME_MET_INTERACTS_WITH_TNS_PROTEINS HP_ABNORMALITY_OF_LENS_SHAPE, HP_ABNORMALITY_OF_LENS_SHAPE REACTOME_PHOSPHOLIPASE_C_MEDIATED_CASCADE_FGFR4, REACTOME_PHOSPHOLIPASE_C_MEDIATED_CASCADE_FGFR4 |
| D_AND_LPS_STIM_MACROPHAGE_UP, GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP | |