

GSE369_PRE_VS_POST_IL6_INJECTION_S0CS3_KO_LIVER_DN, GSE369_PRE_VS_POST_IL6_INJECTION_S0CS3_KO_LIVER_DN
GSE19401_NAIVE_VS_IMMUNIZED_MOUSE_PLN_FOLLICULAR_DC_DN, GSE19401_NAIVE_VS_IMMUNIZED_MOUSE_PLN_FOLLICULAR_DC_DN
GSE36826_NORMAL_VS_STAPH_AUREUS_INF_SKIN_DN, GSE36826_NORMAL_VS_STAPH_AUREUS_INF_SKIN_DN
GSE36826_NORMAL_VS_STAPH_AUREUS_INF_IL1R_KO_SKIN_UP, GSE36826_NORMAL_VS_STAPH_AUREUS_INF_IL1R_KO_SKIN_UP
GSE2706_2H_VS_8H_R848_AND_LPS_STIM_DC_UP, GSE2706_2H_VS_8H_R848_AND_LPS_STIM_DC_UP
GSE11864_CSF1_VS_CSF1_PAM3CYS_IN_MAC_UP, GSE11864_CSF1_VS_CSF1_PAM3CYS_IN_MAC_UP
GSE21670_UNTREATED_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_UP, GSE21670_UNTREATED_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_UP
GSE41176_UNSTIM_VS_ANT1JGM_STIM_BCELL_6H_UP, GSE41176_UNSTIM_VS_ANT1JGM_STIM_BCELL_6H_UP
GSE45365_NK_CELL_VS_BCELL_MCMV_INFECTION_UP, GSE45365_NK_CELL_VS_BCELL_MCMV_INFECTION_UP
GSE11864_CSF1_VS_CSF1_IFNG_PAM3CYS_IN_MAC_UP, GSE11864_CSF1_VS_CSF1_IFNG_PAM3CYS_IN_MAC_UP
GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_2H_UP, GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_2H_UP
GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG1_AND_FOXP3 TRASDUCED_DN, GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG1_AND_FOXP3 TRASDUCED_DN
GSE360_CTRL_VS_1_DONOVANI_DC_UP, GSE360_CTRL_VS_1_DONOVANI_DC_UP
GSE22886_UNSTIM_VS_IL2_STIM_NKCELL_UP, GSE22886_UNSTIM_VS_IL2_STIM_NKCELL_UP
GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_17H_UP, GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_17H_UP
GSE3982_BCELL_VS_EFF_MEMORY_CD4_TCELL_UP, GSE3982_BCELL_VS_EFF_MEMORY_CD4_TCELL_UP
GSE30971_WBP7_HET_VS_KO_MACROPHAGE_4H_LPS_STIM_DN, GSE30971_WBP7_HET_VS_KO_MACROPHAGE_4H_LPS_STIM_DN
GSE14000_4H_VS_16H_LPS_DC_TRANSLATED_RNA_UP, GSE14000_4H_VS_16H_LPS_DC_TRANSLATED_RNA_UP
GSE19198_1H_VS_24H_IL21_TREATED_TCELL_UP, GSE19198_1H_VS_24H_IL21_TREATED_TCELL_UP
GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN
GSE12845_IGD_POS_BLOOD_VS_NAIVE_TONSIL_BCELL_UP, GSE12845_IGD_POS_BLOOD_VS_NAIVE_TONSIL_BCELL_UP
GSE36826_WT_VS_IL1R_KO_SKIN_UP, GSE36826_WT_VS_IL1R_KO_SKIN_UP
GSE9960_HEALTHY_VS_GRAM_POS_SEPSIS_PBMIC_UP, GSE9960_HEALTHY_VS_GRAM_POS_SEPSIS_PBMIC_UP
GSE19198_6H_VS_24H_IL21_TREATED_TCELL_DN, GSE19198_6H_VS_24H_IL21_TREATED_TCELL_DN
GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_UP, GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_UP
GSE1432_1H_VS_24H_IFNG_MICROGLIA_UP, GSE1432_1H_VS_24H_IFNG_MICROGLIA_UP
GSE24634_IL4_VS_CTRL_TREATED_NAIVE_CD4_TCELL_DAY5_UP, GSE24634_IL4_VS_CTRL_TREATED_NAIVE_CD4_TCELL_DAY5_UP
GSE3982_MAC_VS_TH2_UP, GSE3982_MAC_VS_TH2_UP
GSE360_T_GONDIL_VS_B_MALAYI_HIGH_DOSE_MAC_DN, GSE360_T_GONDIL_VS_B_MALAYI_HIGH_DOSE_MAC_DN
GSE3982_MEMORY_CD4_TCELL_VS_TH1_UP, GSE3982_MEMORY_CD4_TCELL_VS_TH1_UP
GSE5589_LPS_VS_LPS_AND_IL10_STIM_MACROPHAGE_45MIN_DN, GSE5589_LPS_VS_LPS_AND_IL10_STIM_MACROPHAGE_45MIN_DN
GSE24634_TEFF_VS_TCONV_DAY3_IN_CULTURE_DN, GSE24634_TEFF_VS_TCONV_DAY3_IN_CULTURE_DN
GSE2706_UNSTIM_VS_8H_R848_DC_UP, GSE2706_UNSTIM_VS_8H_R848_DC_UP
GSE11864_CSF1_VS_CSF1_IFNG_IN_MAC_UP, GSE11864_CSF1_VS_CSF1_IFNG_IN_MAC_UP
GSE22886_NAIVE_BCELL_VS_BLOOD_PLASMA_CELL_UP, GSE22886_NAIVE_BCELL_VS_BLOOD_PLASMA_CELL_UP
GOCC_COATED_VESICLE_MEMBRANE, GOCC_COATED_VESICLE_MEMBRANE
DESCARTES_FETAL_EYE_MICROGLIA, DESCARTES_FETAL_EYE_MICROGLIA
GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_60H_UP, GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_60H_UP
DESCARTES_FETAL_MUSCLE_MYELOID_CELLS, DESCARTES_FETAL_MUSCLE_MYELOID_CELLS
GNF2_CASP1, GNF2_CASP1
RODRIGUES_DCC_TARGETS_DN, RODRIGUES_DCC_TARGETS_DN
GSE29618_BCELL_VS_PDC_DAY7_FLU_VACCINE_UP, GSE29618_BCELL_VS_PDC_DAY7_FLU_VACCINE_UP
GSE22886_IGG_IGA_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_UP, GSE22886_IGG_IGA_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_UP
GSE22886_UNSTIM_VS_IL15_STIM_NKCELL_UP, GSE22886_UNSTIM_VS_IL15_STIM_NKCELL_UP
MIR593_5P, MIR593_5P
GSE19888_ADENOSINE_A3R_INH_VS_ACT_IN_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_VS_ACT_IN_MAST_CELL_DN
GOCC_ENDOCYTIC_VESICLE_MEMBRANE, GOCC_ENDOCYTIC_VESICLE_MEMBRANE
ATAAGCT_MIR21, ATAAGCT_MIR21
HP_ABNORMAL_CIRCULATING_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_AMINO_ACID_CONCENTRATION
REACTOME_CLATHRIN_MEDIATED_ENDOCYTOSIS, REACTOME_CLATHRIN_MEDIATED_ENDOCYTOSIS
RUBENSTEIN_SKELETAL_MUSCLE_ENDOTHELIAL_CELLS, RUBENSTEIN_SKELETAL_MUSCLE_ENDOTHELIAL_CELLS
GOCC_CLATHRIN_COATED_VESICLE, GOCC_CLATHRIN_COATED_VESICLE
MIR4679, MIR4679
LEE_NEURAL_CREST_STEM_CELL_DN, LEE_NEURAL_CREST_STEM_CELL_DN
GSE35435_RESTING_VS_IL4_TREATED_MACROPHAGE_DN, GSE35435_RESTING_VS_IL4_TREATED_MACROPHAGE_DN
BYSTROEM_CORRELATED_WITH_IL5_DN, BYSTROEM_CORRELATED_WITH_IL5_DN
GSE3720_LPS_VS_PMA_STIM_VD1_GAMMADELTA_TCELL_UP, GSE3720_LPS_VS_PMA_STIM_VD1_GAMMADELTA_TCELL_UP
MODULE_188, MODULE_188
HP_LYMPHOPENIA, HP_LYMPHOPENIA
GSE19888_CTRL_VS_TCELL_MEMBRANES_ACT_MAST_CELL_PRETREAT_A3R_INH_UP, GSE19888_CTRL_VS_TCELL_MEMBRANES_ACT_MAST_CELL_PRETREAT_A3R_INH_UP
MODULE_238, MODULE_238
HOEBEKE_LYMPHOID_STEM_CELL_DN, HOEBEKE_LYMPHOID_STEM_CELL_DN
GOBP_RESPONSE_TO_MONOSACCHARIDE, GOBP_RESPONSE_TO_MONOSACCHARIDE
MODULE_301, MODULE_301
GOCC_CLATHRIN_COATED_VESICLE_MEMBRANE, GOCC_CLATHRIN_COATED_VESICLE_MEMBRANE
MODULE_361, MODULE_361
MIR6761_5P, MIR6761_5P
GSE3982_NKCELL_VS_TH1_UP, GSE3982_NKCELL_VS_TH1_UP
MIR874_3P, MIR874_3P
MIR193A_3P, MIR193B_3P, MIR193A_3P, MIR193B_3P
GNF2_MCL1, GNF2_MCL1
MODULE_436, MODULE_436
GOBP_INTEGRIN_MEDIATED_SIGNALING_PATHWAY, GOBP_INTEGRIN_MEDIATED_SIGNALING_PATHWAY
DESCARTES_FETAL_STOMACH_MYELOID_CELLS, DESCARTES_FETAL_STOMACH_MYELOID_CELLS
GOCC_TRANS_GOLGI_NETWORK_MEMBRANE, GOCC_TRANS_GOLGI_NETWORK_MEMBRANE
REACTOME_NEGATIVE_REGULATION_OF_THE_P13K_AKT_NETWORK, REACTOME_NEGATIVE_REGULATION_OF_THE_P13K_AKT_NETWORK
REACTOME_SIGNALING_BY_EGFR, REACTOME_SIGNALING_BY_EGFR
MIR4266, MIR4266
GOBP_CLATHRIN_DEPENDENT_ENDOCYTOSIS, GOBP_CLATHRIN_DEPENDENT_ENDOCYTOSIS
KEGG_INOSITOL_PHOSPHATE_METABOLISM, KEGG_INOSITOL_PHOSPHATE_METABOLISM
GOBP_CELLULAR_GLUCOSE_HOEMEOSTASIS, GOBP_CELLULAR_GLUCOSE_HOEMEOSTASIS
TOMLINS_PROSTATE_CANCER_UP, TOMLINS_PROSTATE_CANCER_UP
GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_PROLIFERATION
GOMF_MANGANESE_ION_BINDING, GOMF_MANGANESE_ION_BINDING
GARCIA_PINERES_PBMIC_HPV_16_L1_VLP_AGE_18_25YO_STIMULATED_VS_UNSTIMULATED_0DY_VACCINATION_INDEPENDENT_DN, GARCIA_PINERES_PBMIC_HPV_16_L1_VLP_AGE_18_25YO_STIMULATED_VS_UNSTIMULATED_0DY_VACCINATION_INDEPENDENT_DN
MODULE_145, MODULE_145
MIR589_5P, MIR589_5P
GOBP_CELLULAR_RESPONSE_TO_CARBOHYDRATE_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_CARBOHYDRATE_STIMULUS
GOMF_PROTEIN_PHOSPHORYLATED_AMINO_ACID_BINDING, GOMF_PROTEIN_PHOSPHORYLATED_AMINO_ACID_BINDING
HP_SKIN_RASH, HP_SKIN_RASH
KIM_GERMINAL_CENTER_T_HELPER_UP, KIM_GERMINAL_CENTER_T_HELPER_UP
TOMLINS_PROSTATE_CANCER_DN, TOMLINS_PROSTATE_CANCER_DN
MIR3616_3P, MIR3616_3P
DAVICIONI_RHABDOMYOSARCOMA_PAX_FOXP01_FUSION_UP, DAVICIONI_RHABDOMYOSARCOMA_PAX_FOXP01_FUSION_UP
GSE13522_WT_VS_IFNG_KO_SKING_T_CRUZI_Y_STRAIN_INF_DN, GSE13522_WT_VS_IFNG_KO_SKING_T_CRUZI_Y_STRAIN_INF_DN
GOBP_CELLULAR_MODIFIED_AMINO_ACID_BIOSYNTHETIC_PROCESS, GOBP_CELLULAR_MODIFIED_AMINO_ACID_BIOSYNTHETIC_PROCESS
HP_ABNORMAL_PLEURA_MORPHOLOGY, HP_ABNORMAL_PLEURA_MORPHOLOGY
HP_DECREASED_ACTIVITY_OF_MITOCHONDRIAL_COMPLEX_IV, HP_DECREASED_ACTIVITY_OF_MITOCHONDRIAL_COMPLEX_IV
KEGG_LEISHMANIA_INFECTION, KEGG_LEISHMANIA_INFECTION
GOBP_PHOSPHOLIPID_CATABOLIC_PROCESS, GOBP_PHOSPHOLIPID_CATABOLIC_PROCESS
HALLMARK_COAGULATION, HALLMARK_COAGULATION
GOBP_PHOSPHATIDYLINOSITOL_PHOSPHORYLATION, GOBP_PHOSPHATIDYLINOSITOL_PHOSPHORYLATION
TRAVAGLINI_LUNG_IGSF21_DENDRITIC_CELL, TRAVAGLINI_LUNG_IGSF21_DENDRITIC_CELL
MIR483_3P, MIR483_3P
KUNINGER_IGF1_VS_PDGF_TARGETS_UP, KUNINGER_IGF1_VS_PDGF_TARGETS_UP
KEGG_VIRAL_MYOCARDITIS, KEGG_VIRAL_MYOCARDITIS
FULLER_PBMIC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_18HR_UP, FULLER_PBMIC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_18HR_UP
KEGG_PYRUVATE_METABOLISM, KEGG_PYRUVATE_METABOLISM
GOCC_PLASMA_MEMBRANE_RAFT, GOCC_PLASMA_MEMBRANE_RAFT
RAMALHO_STEMNESS_DN, RAMALHO_STEMNESS_DN
GCM_AQP4, GCM_AQP4
HP_PULMONARY_FIBROSIS, HP_PULMONARY_FIBROSIS
KIM_LRRC3B_TARGETS, KIM_LRRC3B_TARGETS
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_ALDEHYDE_OR_OXO_GROUP_OF_DONORS_NAD_OR_NADP_AS_ACCEPTOR, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_ALDEHYDE_OR_OXO_GROUP_OF_DONORS_NAD_OR_NADP_AS_ACCEPTOR
NAKAMURA_ADIPOGENESIS_EARLY_UP, NAKAMURA_ADIPOGENESIS_EARLY_UP
SIX4_TARGET_GENES, SIX4_TARGET_GENES
GOBP_ICOSANOID_METABOLIC_PROCESS, GOBP_ICOSANOID_METABOLIC_PROCESS
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_ALDEHYDE_OR_OXO_GROUP_OF_DONORS, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_ALDEHYDE_OR_OXO_GROUP_OF_DONORS
GOBP_POSITIVE_REGULATION_OF_PEPTIDYL_TYROSINE_PHOSPHORYLATION, GOBP_POSITIVE_REGULATION_OF_PEPTIDYL_TYROSINE_PHOSPHORYLATION
AR_02, AR_02
MIR3622B_5P, MIR3622B_5P
HP_ABNORMAL_B_CELL_MORPHOLOGY, HP_ABNORMAL_B_CELL_MORPHOLOGY
HP_ORAL_ULCER, HP_ORAL_ULCER
HP_HEMIPLEGIA, HP_HEMIPLEGIA
GOMF_O_METHYLTRANSFERASE_ACTIVITY, GOMF_O_METHYLTRANSFERASE_ACTIVITY
GYORFFY_MITOXANTHRONE_RESISTANCE, GYORFFY_MITOXANTHRONE_RESISTANCE
GOBP_GLYCEROPHOSPHOLIPID_CATABOLIC_PROCESS, GOBP_GLYCEROPHOSPHOLIPID_CATABOLIC_PROCESS
REACTOME_ANTIGEN_ACTIVATES_B_CELL_RECEPTOR_BCR_LEADING_TO_GENERATION_OF_SECOND_MESSENGERS, REACTOME_ANTIGEN_ACTIVATES_B_CELL_RECEPTOR_BCR_LEADING_TO_GENERATION_OF_SECOND_MESSENGERS
FULLER_PBMIC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_336HR_DN, FULLER_PBMIC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_336HR_DN
HOQUE_METHYLATED_IN_CANCER, HOQUE_METHYLATED_IN_CANCER
GOBP_REGULATION_OF_CELL_ADHESION_MEDIATED_BY_INTEGRIN, GOBP_REGULATION_OF_CELL_ADHESION_MEDIATED_BY_INTEGRIN
MIR6748_5P, MIR6748_5P
GOMF_NADP_BINDING, GOMF_NADP_BINDING
YAO_HOXA10_TARGETS_VIA_PROGESTERONE_UP, YAO_HOXA10_TARGETS_VIA_PROGESTERONE_UP
FULLER_PBMIC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_192HR_DN, FULLER_PBMIC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_192HR_DN
DURANTE_ADULT_OLFATORY_NEUROEPITHELIUM_MONOCYTES, DURANTE_ADULT_OLFATORY_NEUROEPITHELIUM_MONOCYTES
HP_ABNORMAL_CIRCULATING_NON_PROTEINOGENIC_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_NON_PROTEINOGENIC_AMINO_ACID_CONCENTRATION
MIR454_5P, MIR454_5P
GOBP_B_CELL_RECEPTOR_SIGNALING_PATHWAY, GOBP_B_CELL_RECEPTOR_SIGNALING_PATHWAY
ATM_TARGET_GENES, ATM_TARGET_GENES
MODULE_448, MODULE_448
AGARWAL_AKT_PATHWAY_TARGETS, AGARWAL_AKT_PATHWAY_TARGETS
HP_INCREASED_URINARY_NON_PROTEINOGENIC_AMINO_ACID_LEVEL, HP_INCREASED_URINARY_NON_PROTEINOGENIC_AMINO_ACID_LEVEL
REACTOME_AFLATOXIN_ACTIVATION_AND_DETOXIFICATION, REACTOME_AFLATOXIN_ACTIVATION_AND_DETOXIFICATION
REACTOME_HS_GAG_DEGRADATION, REACTOME_HS_GAG_DEGRADATION
GOMF_ALDEHYDE_DEHYDROGENASE_NAD_P_PLUS_ACTIVITY, GOMF_ALDEHYDE_DEHYDROGENASE_NAD_P_PLUS_ACTIVITY
GOCC_CLATHRIN_COATED_ENDOCYTIC_VESICLE, GOCC_CLATHRIN_COATED_ENDOCYTIC_VESICLE
GOBP_INOSITOL_METABOLIC_PROCESS, GOBP_INOSITOL_METABOLIC_PROCESS
HP_CEREBELLAR_VERMIS_ATROPHY, HP_CEREBELLAR_VERMIS_ATROPHY
GUILLAUMOND_KLF10_TARGETS_DN, GUILLAUMOND_KLF10_TARGETS_DN
GOBP_REGULATION_OF_CELL_MIGRATION_INVOLVED_IN_SPROUTING_ANGIOGENESIS, GOBP_REGULATION_OF_CELL_MIGRATION_INVOLVED_IN_SPROUTING_ANGIOGENESIS
HP_URINARY_GLYCOSAMINOGLYCAN_EXCRETION, HP_URINARY_GLYCOSAMINOGLYCAN_EXCRETION
KEGG_TYPE_II_DIABETES_MELLITUS, KEGG_TYPE_II_DIABETES_MELLITUS
SMID_BREAST_CANCER_ERBB2_UP, SMID_BREAST_CANCER_ERBB2_UP
GOMF_PROLINE_RICH_REGION_BINDING, GOMF_PROLINE_RICH_REGION_BINDING
HP_RECURRENT_ASPIRATION_PNEUMONIA, HP_RECURRENT_ASPIRATION_PNEUMONIA
GOBP_REGULATION_OF_ADIPOSE_TISSUE_DEVELOPMENT, GOBP_REGULATION_OF_ADIPOSE_TISSUE_DEVELOPMENT
GOCC_BETA_CATENIN_DESTRUCTION_COMPLEX, GOCC_BETA_CATENIN_DESTRUCTION_COMPLEX
GOBP_ENDOTHELIAL_CELL_PROLIFERATION, GOBP_ENDOTHELIAL_CELL_PROLIFERATION
HP_INCREASED_CIRCULATING_IGG_LEVEL, HP_INCREASED_CIRCULATING_IGG_LEVEL
GOMF_LIPOPROTEIN_PARTICLE_RECEPTOR_BINDING, GOMF_LIPOPROTEIN_PARTICLE_RECEPTOR_BINDING
GOMF_SERINE_HYDROLASE_ACTIVITY, GOMF_SERINE_HYDROLASE_ACTIVITY
HP_PERIVENTRICULAR_NODULAR_HETEROTOPIA, HP_PERIVENTRICULAR_NODULAR_HETEROTOPIA
GOBP_INSULIN_LIKE_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_INSULIN_LIKE_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
REACTOME_CELL_SURFACE_INTERACTIONS_AT_THE_VASCULAR_WALL, REACTOME_CELL_SURFACE_INTERACTIONS_AT_THE_VASCULAR_WALL
REACTOME_P13K_EVENTS_IN_ERBB4_SIGNALING, REACTOME_P13K_EVENTS_IN_ERBB4_SIGNALING
GEISS_RESPONSE_TO_DSRNA_DN, GEISS_RESPONSE_TO_DSRNA_DN
BIOCARTA_PGIC1A_PATHWAY, BIOCARTA_PGIC1A_PATHWAY
GOBP_LEARNED_VOCALIZATION_BEHAVIOR_OR_VOCAL_LEARNING, GOBP_LEARNED_VOCALIZATION_BEHAVIOR_OR_VOCAL_LEARNING
WP_METAPATHWAY_BIOTRANSFORMATION_PHASE_I_AND_II, WP_METAPATHWAY_BIOTRANSFORMATION_PHASE_I_AND_II
GOBP_CALCITONIN_FAMILY_RECEPTOR_SIGNALING_PATHWAY, GOBP_CALCITONIN_FAMILY_RECEPTOR_SIGNALING_PATHWAY
DESCARTES_MAIN_FETAL_AFP_ALB_POSITIVE_CELLS, DESCARTES_MAIN_FETAL_AFP_ALB_POSITIVE_CELLS
GOBP_MUSCLE_ATROPHY, GOBP_MUSCLE_ATROPHY
GOMF_CYCLASE_ACTIVITY, GOMF_CYCLASE_ACTIVITY
MODULE_194, MODULE_194
GOMF_NADPH_BINDING, GOMF_NADPH_BINDING
GOBP_VOICAL_LEARNING, GOBP_VOICAL_LEARNING
BUCASAS_PBMIC_FLUARIX_FLUVIRIN_CAUCASIAN_MALE_AGE_18_40YO_HIGH_RESPONDERS_1DY_3DY_POSITIVE_PREDICTIVE_OF_TITER, BUCASAS_PBMIC_FLUARIX_FLUVIRIN_CAUCASIAN_MALE_AGE_18_40YO_HIGH_RESPONDERS_1DY_3DY_POSITIVE_PREDICTIVE_OF_TITER
GOBP_MACROMOLECULE_DEPALMITOYLATION, GOBP_MACROMOLECULE_DEPALMITOYLATION
GOMF_PALMITOYL_PROTEIN_HYDROLASE_ACTIVITY, GOMF_PALMITOYL_PROTEIN_HYDROLASE_ACTIVITY
HP_ABNORMAL_ANTERIOR_CHAMBER_MORPHOLOGY, HP_ABNORMAL_ANTERIOR_CHAMBER_MORPHOLOGY
HP_RECURRENT_INTRA-PULMONARY_HEMORRHAGE, HP_RECURRENT_INTRA-PULMONARY_HEMORRHAGE
GOBP_POSITIVE_REGULATION_OF_MORPHOGENESIS_OF_AN_EPITHELIUM, GOBP_POSITIVE_REGULATION_OF_MORPHOGENESIS_OF_AN_EPITHELIUM
GOBP_URONIC_ACID_METABOLIC_PROCESS, GOBP_URONIC_ACID_METABOLIC_PROCESS
GOMF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_RECEPTOR_BINDING, GOMF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_RECEPTOR_BINDING
HP_HYDROXYPROLINURIA, HP_HYDROXYPROLINURIA
GOBP_POSITIVE_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATION, GOBP_POSITIVE_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATION
GOBP_MONOACYLGLYCEROL_CATABOLIC_PROCESS, GOBP_MONOACYLGLYCEROL_CATABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION