

	GSE41867_NAIVE_VS_DAY15_LCMV_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY15_LCMV_EFFECTOR_CD8_TCELL_UP GSE41867_NAIVE_VS_DAY30_LCMV_ARMSTRONG_MEMORY_CD8_TCELL_DN, GSE41867_NAIVE_VS_DAY30_LCMV_ARMSTRONG_MEMORY_CD8_TCELL_DN GSE24671_BAKIMULC_VS_SENDAL_VIRUS_INFECTED_MOUSE_SPLENOCYTES_DN, GSE24671_BAKIMULC_VS_SENDAL_VIRUS_INFECTED_MOUSE_SPLENOCYTES_DN GSE41867_NAIVE_VS_DAY30_LCMV_ARMSTRONG_MEMORY_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY30_LCMV_ARMSTRONG_MEMORY_CD8_TCELL_UP GSE17186_CD21LOW_VS_CD21HIGH_TRANSITIONAL_BCELL_CORD_BLOOD_UP, GSE17186_CD21LOW_VS_CD21HIGH_TRANSITIONAL_BCELL_CORD_BLOOD_UP GSE37301_HEMATOPOIETIC_STEM_CELL_VS_PRO_BCELL_DN, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_PRO_BCELL_DN GSE16385_UNTREATED_VS_12H_IFNG_TNF_TREATED_MACROPHAGE_DN, GSE16385_UNTREATED_VS_12H_IFNG_TNF_TREATED_MACROPHAGE_DN GSE10240_IL22_VS_IL17_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN, GSE10240_IL22_VS_IL17_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN GSE21774_CD56_BRIGHT_VS_DIM_CD62L_POSITIVE_NK_CELL_DN, GSE21774_CD56_BRIGHT_VS_DIM_CD62L_POSITIVE_NK_CELL_DN MIR19A_5P, MIR19A_5P GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP, GSE21379_WT_VS_SAP_KO_CD4_TCELL_UP MIR19B_1_5P, MIR19B_2_5P, MIR19B_1_5P, MIR19B_2_5P MIR299_5P, MIR299_5P HP_PROMINENT_NOSE, HP_PROMINENT_NOSE GOCC_MICROBODY, GOCC_MICROBODY GSE17721_LPS_VS_GARDIQUIMOD_2H_BMDC_UP, GSE17721_LPS_VS_GARDIQUIMOD_2H_BMDC_UP GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_UP, GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_UP GOMF_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GOMF_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING BRUINS_UVC_RESPONSE_MIDDLE, BRUINS_UVC_RESPONSE_MIDDLE GSE43955_TH0_VS_TGFB_IL6_THI7_ACT_CD4_TCELL_42H_DN, GSE43955_TH0_VS_TGFB_IL6_THI7_ACT_CD4_TCELL_42H_DN MODULE_192, MODULE_192 PANGAS_TUMOR_SUPPRESSION_BY_SMAD1_AND_SMAD5_DN, PANGAS_TUMOR_SUPPRESSION_BY_SMAD1_AND_SMAD5_DN GOBP_MAINTENANCE_OF_CELL_NUMBER, GOBP_MAINTENANCE_OF_CELL_NUMBER KATSANOUELAVL1_TARGETS_DN, KATSANOUELAVL1_TARGETS_DN GSE19374_UNINF_VS_LISTERIA_INFECTED_MACROPHAGE_DN, GSE19374_UNINF_VS_LISTERIA_INFECTED_MACROPHAGE_DN MIR487A_5P, MIR487A_5P MIR12127, MIR12127 HP_LIMB_GIRDLE_MUSCLE_WEAKNESS, HP_LIMB_GIRDLE_MUSCLE_WEAKNESS MIR487B_5P, MIR487B_5P GSE37301_LYMPHOID_PRIMED_MPP_VS_RAG2_KO_NK_CELL_DN, GSE37301_LYMPHOID_PRIMED_MPP_VS_RAG2_KO_NK_CELL_DN MIR6891_3P, MIR6891_3P GOMF_NUCLEAR_IMPORT_SIGNAL_RECEPTOR_ACTIVITY, GOMF_NUCLEAR_IMPORT_SIGNAL_RECEPTOR_ACTIVITY KESHELAVA_MULTIPLE_DRUG_RESISTANCE, KESHELAVA_MULTIPLE_DRUG_RESISTANCE GOBP_INTRACILIARY_TRANSPORT, GOBP_INTRACILIARY_TRANSPORT GSE29615_CTRL_VS_DAY3_LAIV_IFLU_VACCINE_PBMCDN, GSE29615_CTRL_VS_DAY3_LAIV_IFLU_VACCINE_PBMCDN GOCC_MICROBODY_MEMBRANE, GOCC_MICROBODY_MEMBRANE MIR5681B, MIR5681B MIR4763_5P, MIR4763_5P GOMF_AMINOPEPTIDASE_ACTIVITY, GOMF_AMINOPEPTIDASE_ACTIVITY MIR508_3P, MIR508_3P ONGUSAHA_TP53_TARGETS, ONGUSAHA_TP53_TARGETS HP_ABNORMAL_CIRCULATING_BILIRUBIN_CONCENTRATION, HP_ABNORMAL_CIRCULATING_BILIRUBIN_CONCENTRATION GOBP_HISTONE_DEUBIQUITINATION, GOBP_HISTONE_DEUBIQUITINATION GOBP_INTRACILIARY_TRANSPORT_INVOLVED_IN_CILIUM_ASSEMBLY, GOBP_INTRACILIARY_TRANSPORT_INVOLVED_IN_CILIUM_ASSEMBLY ROSS_AML_OF_FAB_M7_TYPE, ROSS_AML_OF_FAB_M7_TYPE REACTOME_HIV_ELONGATION_ARREST_AND_RECOVERY, REACTOME_HIV_ELONGATION_ARREST_AND_RECOVERY GNF2_CDC27, GNF2_CDC27 PID_NETRIN_PATHWAY, PID_NETRIN_PATHWAY GOBP_FIBROBLAST_PROLIFERATION, GOBP_FIBROBLAST_PROLIFERATION KENNY_CTNNB1_TARGETS_DN, KENNY_CTNNB1_TARGETS_DN REACTOME_CELL_CELL_JUNCTION_ORGANIZATION, REACTOME_CELL_CELL_JUNCTION_ORGANIZATION HP_INCREASED_CIRCULATING_IgA_LEVEL, HP_INCREASED_CIRCULATING_IgA_LEVEL GOCC_POST_MRNA_RELEASE_SPLICEOSOMAL_COMPLEX, GOCC_POST_MRNA_RELEASE_SPLICEOSOMAL_COMPLEX BRACHAT_RESPONSE_TO_CAMPTOTHECIN_UP, BRACHAT_RESPONSE_TO_CAMPTOTHECIN_UP HP_DEEP_PALMAR_CREASE, HP_DEEP_PALMAR_CREASE HP_DEPRESSED_NASAL_TIP, HP_DEPRESSED_NASAL_TIP NEUROG3_TARGET_GENES, NEUROG3_TARGET_GENES GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BROWN_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BROWN_UP HP_SMALL_EARLOBE, HP_SMALL_EARLOBE KANG_CISPLATIN_RESISTANCE_UP, KANG_CISPLATIN_RESISTANCE_UP ZNF777_TARGET_GENES, ZNF777_TARGET_GENES GOCC_ENDOPLASMIC_RETICULUM_EXIT_SITE, GOCC_ENDOPLASMIC_RETICULUM_EXIT_SITE HP_MICRONODULAR_CIRRHOSIS, HP_MICRONODULAR_CIRRHOSIS HP_ABNORMALITY_OF_THE_UPPER_ARM, HP_ABNORMALITY_OF_THE_UPPER_ARM chr16q22, chr16q22 KEGG_TIGHT_JUNCTION, KEGG_TIGHT_JUNCTION HP_UNDERDEVELOPED_SUPRAORBITAL_RIDGES, HP_UNDERDEVELOPED_SUPRAORBITAL_RIDGES GOCC_RNA_POLYMERASE_II_CORE_COMPLEX, GOCC_RNA_POLYMERASE_II_CORE_COMPLEX GSE41176_WT_VS_TAK1_KO_UNSTIM_BCELL_DN, GSE41176_WT_VS_TAK1_KO_UNSTIM_BCELL_DN HP_SHOULDER_GIRDLE_MUSCLE_WEAKNESS, HP_SHOULDER_GIRDLE_MUSCLE_WEAKNESS BRACHAT_RESPONSE_TO_CISPLATIN, BRACHAT_RESPONSE_TO_CISPLATIN HP_APLASIA_HYPOPLASIA_OF_THE_FEMUR, HP_APLASIA_HYPOPLASIA_OF_THE_FEMUR TCTAGAG_MIR517, TCTAGAG_MIR517 KEGG_BLADDER_CANCER, KEGG_BLADDER_CANCER GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_BCELL_UP, GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_BCELL_UP GOBP_ANOIKIS, GOBP_ANOIKIS HP_BEAKING_OF_VERTEBRAL_BODIES, HP_BEAKING_OF_VERTEBRAL_BODIES HP_ABNORMALITY_OF_PROTHROMBIN, HP_ABNORMALITY_OF_PROTHROMBIN MIR1296_3P, MIR1296_3P MIR6824_5P, MIR6824_5P MIR515_3P, MIR515_3P WP_MITOCHONDRIAL_GENE_EXPRESSION, WP_MITOCHONDRIAL_GENE_EXPRESSION GOMF_ABC_TYPE_TRANSPORTER_ACTIVITY, GOMF_ABC_TYPE_TRANSPORTER_ACTIVITY LIAO_HAVE_SOX4_BINDING_SITES, LIAO_HAVE_SOX4_BINDING_SITES MIR5684, MIR5684 MORF_LCAT, MORF_LCAT GOBP_POSITIVE_REGULATION_OF_DENDRITIC_SPINE_MORPHOGENESIS, GOBP_POSITIVE_REGULATION_OF_DENDRITIC_SPINE_MORPHOGENESIS MIR519E_3P, MIR519E_3P HP_ABNORMALITY_OF_THE_ANTIHELIX, HP_ABNORMALITY_OF_THE_ANTIHELIX GOBP_HISTAMINE_PRODUCTION_INVOLVED_IN_INFLAMMATORY_RESPONSE, GOBP_HISTAMINE_PRODUCTION_INVOLVED_IN_INFLAMMATORY_RESPONSE GAVIN_IL2_RESPONSIVE_FOXP3_TARGETS_UP, GAVIN_IL2_RESPONSIVE_FOXP3_TARGETS_UP MIR33B_3P, MIR33B_3P MIR621, MIR621 GOBP_MODULATION_BY_HOST_OF_VIRAL_PROCESS, GOBP_MODULATION_BY_HOST_OF_VIRAL_PROCESS HP_ABNORMALITY_OF_THE_HUMERUS, HP_ABNORMALITY_OF_THE_HUMERUS GOMF_RACEMASE_AND_EPIMERASE_ACTIVITY, GOMF_RACEMASE_AND_EPIMERASE_ACTIVITY HP_PELVIC_GIRDLE_MUSCLE_WEAKNESS, HP_PELVIC_GIRDLE_MUSCLE_WEAKNESS ICHIBA_GRAFT_VERSUS_HOST_DISEASE_D7_DN, ICHIBA_GRAFT_VERSUS_HOST_DISEASE_D7_DN GOBP_SNRNA_PROCESSING, GOBP_SNRNA_PROCESSING WP_BIOTIN_METABOLISM_INCLUDING_IEMS, WP_BIOTIN_METABOLISM_INCLUDING_IEMS GOBP_BIOTIN_METABOLIC_PROCESS, GOBP_BIOTIN_METABOLIC_PROCESS DESCARTES_MAIN_FETAL_ACINAR_CELLS, DESCARTES_MAIN_FETAL_ACINAR_CELLS PID_REELIN_PATHWAY, PID_REELIN_PATHWAY GOBP_REPLICATIVE_SENESCENCE, GOBP_REPLICATIVE_SENESCENCE MIR664B_5P, MIR664B_5P WP_GLYCEROPHOSPHOLIPID_BIOSYNTHETIC_PATHWAY, WP_GLYCEROPHOSPHOLIPID_BIOSYNTHETIC_PATHWAY HP_GERM_CELL_NEOPLASIA, HP_GERM_CELL_NEOPLASIA MANN_RESPONSE_TO_AMIFOSTINE_UP, MANN_RESPONSE_TO_AMIFOSTINE_UP GOBP_ETHER_METABOLIC_PROCESS, GOBP_ETHER_METABOLIC_PROCESS GOBP_PHOSPHATIDIC_ACID_METABOLIC_PROCESS, GOBP_PHOSPHATIDIC_ACID_METABOLIC_PROCESS WP_PHOTODYNAMIC_THERAPYINDUCED_NFE2L2_NRF2_SURVIVAL_SIGNALING, WP_PHOTODYNAMIC_THERAPYINDUCED_NFE2L2_NRF2_SURVIVAL_SIGNALING REACTOME_HYDROLYSIS_OF_LPC, REACTOME_HYDROLYSIS_OF_LPC GOMF_LIGASE_ACTIVITY_FORMING_CARBON_CARBON_BONDS, GOMF_LIGASE_ACTIVITY_FORMING_CARBON_CARBON_BONDS GOBP_LYMPHOID_PROGENITOR_CELL_DIFFERENTIATION, GOBP_LYMPHOID_PROGENITOR_CELL_DIFFERENTIATION BIOCARTA_P53HYPOXIA_PATHWAY, BIOCARTA_P53HYPOXIA_PATHWAY GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_DN, GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_DN HP_ALVEOLAR_RIDGE_OVERGROWTH, HP_ALVEOLAR_RIDGE_OVERGROWTH GOBP_POSITIVE_REGULATION_OF_TRIGLYCERIDE_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_TRIGLYCERIDE_CATABOLIC_PROCESS GOBP_SHORT_CHAIN_FATTY_ACID_METABOLIC_PROCESS, GOBP_SHORT_CHAIN_FATTY_ACID_METABOLIC_PROCESS GOBP_REGULATION_OF_RAB_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_REGULATION_OF_RAB_PROTEIN_SIGNAL_TRANSDUCTION GOBP_TRIGLYCERIDE_METABOLIC_PROCESS, GOBP_TRIGLYCERIDE_METABOLIC_PROCESS WANG_RECURRENT_LIVER_CANCER_UP, WANG_RECURRENT_LIVER_CANCER_UP WP_NANOPARTICLEMEDIATED_ACTIVATION_OF_RECEPTOR_SIGNALING, WP_NANOPARTICLEMEDIATED_ACTIVATION_OF_RECEPTOR_SIGNALING GOBP_NEGATIVE_REGULATION_OF_SEQUESTERING_OF_TRIGLYCERIDE, GOBP_NEGATIVE_REGULATION_OF_SEQUESTERING_OF_TRIGLYCERIDE GOBP_CELLULAR_RESPONSE_TO_UV_B, GOBP_CELLULAR_RESPONSE_TO_UV_B GOCC_DERLIN_1_RETROTRANSLOCATION_COMPLEX, GOCC_DERLIN_1_RETROTRANSLOCATION_COMPLEX GOMF_O_ACETYLTRANSFERASE_ACTIVITY, GOMF_O_ACETYLTRANSFERASE_ACTIVITY HP_SHORT_COLUMELLA, HP_SHORT_COLUMELLA ANDERSEN_LIVER_CANCER_KRT19_DN, ANDERSEN_LIVER_CANCER_KRT19_DN GOMF_RACEMASE_AND_EPIMERASE_ACTIVITY_ACTING_ON_CARBOHYDRATES_AND_DERIVATIVES, GOMF_RACEMASE_AND_EPIMERASE_ACTIVITY_ACTING_ON_CARBOHYDRATES_AND_DERIVATIVES WP_STRIATED_MUSCLE_CONTRACTION_PATHWAY, WP_STRIATED_MUSCLE_CONTRACTION_PATHWAY GOMF_RETROMER_COMPLEX_BINDING, GOMF_RETROMER_COMPLEX_BINDING GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_OXIDATIVE_STRESS, GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_OXIDATIVE_STRESS REACTOME_DEFECTS_IN_BIOTIN_BTN_METABOLISM, REACTOME_DEFECTS_IN_BIOTIN_BTN_METABOLISM GOBP_HISTONE_H3_K14_ACETYLTATION, GOBP_HISTONE_H3_K14_ACETYLTATION PID_EPHA2_FWD_PATHWAY, PID_EPHA2_FWD_PATHWAY SA_TRKA_RECEPTOR, SA_TRKA_RECEPTOR GOMF_SPHINGOLIPID_TRANSFER_ACTIVITY, GOMF_SPHINGOLIPID_TRANSFER_ACTIVITY HP_APLASIA_HYPOPLASIA_INVOLVING_THE_CARPAL_BONES, HP_APLASIA_HYPOPLASIA_INVOLVING_THE_CARPAL_BONES HP_PHOCOMELIA, HP_PHOCOMELIA GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_II, GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_II HP_APLASIA_HYPOPLASIA_OF_THE_HUMERUS, HP_APLASIA_HYPOPLASIA_OF_THE_HUMERUS GOBP_CERAMIDE_1_PHOSPHATE_TRANSPORT, GOBP_CERAMIDE_1_PHOSPHATE_TRANSPORT GOMF_CERAMIDE_1_PHOSPHATE_TRANSFER_ACTIVITY, GOMF_CERAMIDE_1_PHOSPHATE_TRANSFER_ACTIVITY GOBP_SIGNAL_COMPLEX_ASSEMBLY, GOBP_SIGNAL_COMPLEX_ASSEMBLY GOBP_STRESS_INDUCED_PREMATURE_SENESCENCE, GOBP_STRESS_INDUCED_PREMATURE_SENESCENCE MIR1193, MIR1193 BLANCO_MELO_MERS_COV_INFECTION_MCR5_CELLS_DN, BLANCO_MELO_MERS_COV_INFECTION_MCR5_CELLS_DN GOBP_ETHER_BIOSYNTHETIC_PROCESS, GOBP_ETHER_BIOSYNTHETIC_PROCESS MIR184, MIR184 HP_NARROW_NASAL_RIDGE, HP_NARROW_NASAL_RIDGE WEBER_METHYLATED_HCP_IN_FIBROBLAST_UP, WEBER_METHYLATED_HCP_IN_FIBROBLAST_UP KUROKAWA_LIVER_CANCER_EARLY_RECURRENCE_DN, KUROKAWA_LIVER_CANCER_EARLY_RECURRENCE_DN HP_MUSCLE_FIBER_INCLUSION_BODIES, HP_MUSCLE_FIBER_INCLUSION_BODIES GOBP_GLYCEROL_ETHER_METABOLIC_PROCESS, GOBP_GLYCEROL_ETHER_METABOLIC_PROCESS MIR549A_5P, MIR549A_5P GOBP_EYE_PHOTORECEPTOR_CELL_DIFFERENTIATION, GOBP_EYE_PHOTORECEPTOR_CELL_DIFFERENTIATION WP_TRIACYLGLYCERIDE_SYNTHESIS, WP_TRIACYLGLYCERIDE_SYNTHESIS XPO1_TARGET_GENES, XPO1_TARGET_GENES GOBP_CELLULAR_DETOXIFICATION_OF_ALDEHYDE, GOBP_CELLULAR_DETOXIFICATION_OF_ALDEHYDE REACTOME_P75NTR_RECRUITS_SIGNALLING_COMPLEXES, REACTOME_P75NTR_RECRUITS_SIGNALLING_COMPLEXES HP_DUPLICATION_OF_THE_DISTAL_PHALANX_OF_HAND, HP_DUPLICATION_OF_THE_DISTAL_PHALANX_OF_HAND GOBP_HISTAMINE_SECRETION, GOBP_HISTAMINE_SECRETION MIR6820_5P, MIR6820_5P GOCC_COMPACT_MYELIN, GOCC_COMPACT_MYELIN GOBP_REGULATION_OF_RESPONSE_TO_REACTIVE_OXYGEN_SPECIES, GOBP_REGULATION_OF_RESPONSE_TO_REACTIVE_OXYGEN_SPECIES CONRAD_STEM_CELL, CONRAD_STEM_CELL GOBP_DNA_DEAMINATION, GOBP_DNA_DEAMINATION CREBL2_TARGET_GENES, CREBL2_TARGET_GENES GOMF_JUN_KINASE_BINDING, GOMF_JUN_KINASE_BINDING BIOCARTA_REELIN_PATHWAY, BIOCARTA_REELIN_PATHWAY GOCC_PI_BODY, GOCC_PI_BODY GOMF_INOSITOL_TETRAKISPHOSPHATE_PHOSPHATASE_ACTIVITY, GOMF_INOSITOL_TETRAKISPHOSPHATE_PHOSPHATASE_ACTIVITY COLLER_MYC_TARGETS_DN, COLLER_MYC_TARGETS_DN REACTOME_WAX_AND_PLASMALOGEN_BIOSYNTHESIS, REACTOME_WAX_AND_PLASMALOGEN_BIOSYNTHESIS GOBP_REGULATION_OF_GONAD_DEVELOPMENT, GOBP_REGULATION_OF_GONAD_DEVELOPMENT GOBP_POSITIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_CELL_DEATH, GOBP_POSITIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_CELL_DEATH WHITESIDE_CISPLATIN_RESISTANCE_UP, WHITESIDE_CISPLATIN_RESISTANCE_UP GOBP_INOSITOL_METABOLIC_PROCESS, GOBP_INOSITOL_METABOLIC_PROCESS GOBP_LAYER_FORMATION_IN_CEREBRAL_CORTEX, GOBP_LAYER_FORMATION_IN_CEREBRAL_CORTEX GOBP_REGULATION_OF_PROTEIN_COMPLEX_STABILITY, GOBP_REGULATION_OF_PROTEIN_COMPLEX_STABILITY GOBP_HISTAMINE_TRANSPORT, GOBP_HISTAMINE_TRANSPORT MIR1277_3P, MIR1277_3P GOCC_ENDOPLASMIC_RETICULUM_CHAPERONE_COMPLEX, GOCC_ENDOPLASMIC_RETICULUM_CHAPERONE_COMPLEX GOCC_STEREOCILUM_BASE, GOCC_STEREOCILUM_BASE GOBP_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_RECEPTOR_2_SIGNALING_PATHWAY, GOBP_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_RECEPTOR_2_SIGNALING_PATHWAY REACTOME_PROPIONYL_COA_CATABOLISM, REACTOME_PROPIONYL_COA_CATABOLISM GOMF_CERAMIDE_1_PHOSPHATE_BINDING, GOMF_CERAMIDE_1_PHOSPHATE_BINDING HP_PROMINENT_CALCANEUS, HP_PROMINENT_CALCANEUS WP_MFAP5MEDIATED_OVARIAN_CANCER_CELL_MOTILITY_AND_INVASIVENESS, WP_MFAP5MEDIATED_OVARIAN_CANCER_CELL_MOTILITY_AND_INVASIVENESS GOBP_DESMOSOME_ORGANIZATION, GOBP_DESMOSOME_ORGANIZATION GOBP_POSITIVE_REGULATION_OF_GONAD_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_GONAD_DEVELOPMENT GOBP_REGULATION_OF_MALE_GONAD_DEVELOPMENT, GOBP_REGULATION_OF_MALE_GONAD_DEVELOPMENT SHIRAIISHI_PLZF_TARGETS_UP, SHIRAIISHI_PLZF_TARGETS_UP MIKKELSEN_PARTIALLY_REPROGRAMMED_TO_PLURIPOTENCY, MIKKELSEN_PARTIALLY_REPROGRAMMED_TO_PLURIPOTENCY REACTOME_ACYL_CHAIN_REMODELLING_OF_PI, REACTOME_ACYL_CHAIN_REMODELLING_OF_PI RUAN_RESPONSE_TO_TNF_UP, RUAN_RESPONSE_TO_TNF_UP JL_CARCINOGENESIS_BY_KRAS_AND_STK11_UP, JL_CARCINOGENESIS_BY_KRAS_AND_STK11_UP HP_EMG_AXONAL_ABNORMALITY, HP_EMG_AXONAL_ABNORMALITY HP_ANISOCORIA, HP_ANISOCORIA HP_CEREBELLAR_HEMORRHAGE, HP_CEREBELLAR_HEMORRHAGE GOBP_SPLICEOSOMAL_CONFORMATIONAL_CHANGES_TO_GENERATE_CATALYTIC_CONFORMATION, GOBP_SPLICEOSOMAL_CONFORMATIONAL_CHANGES_TO_GENERATE_CATALYTIC_CONFORMATION GOBP_POSITIVE_REGULATION_OF_MEIOTIC_NUCLEAR_DIVISION, GOBP_POSITIVE_REGULATION_OF_MEIOTIC_NUCLEAR_DIVISION GOBP_REGULATION_OF_TRIGLYCERIDE_CATABOLIC_PROCESS, GOBP_REGULATION_OF_TRIGLYCERIDE_CATABOLIC_PROCESS GOBP_ASPARAGINE_METABOLIC_PROCESS, GOBP_ASPARAGINE_METABOLIC_PROCESS GOBP_DEPHOSPHORYLATION_OF_RNA_POLYMERASE_II_C_TERMINAL_DOMAIN, GOBP_DEPHOSPHORYLATION_OF_RNA_POLYMERASE_II_C_TERMINAL_DOMAIN GOBP_PROTEIN_LOCALIZATION_TO_BICELLULAR_TIGHT_JUNCTION, GOBP_PROTEIN_LOCALIZATION_TO_BICELLULAR_TIGHT_JUNCTION GOMF_TAP1_BINDING, GOMF_TAP1_BINDING GOBP_REELIN_MEDIATED_SIGNALING_PATHWAY, GOBP_REELIN_MEDIATED_SIGNALING_PATHWAY GOMF_INSULIN_LIKE_GROWTH_FACTOR_II_BINDING, GOMF_INSULIN_LIKE_GROWTH_FACTOR_II_BINDING GOBP_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_APOPTOTIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_APOPTOTIC_PROCESS
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