

	GSE27786_LSK_VS_NKCELL_UP, GSE27786_LSK_VS_NKCELL_UP GSE9037_CTRL_VS_LPS_1H_STIM_IRAK1_KO_BMDM_UP, GSE9037_CTRL_VS_LPS_1H_STIM_IRAK1_KO_BMDM_UP GSE44649_NAIVE_VS_ACTIVATED_CD8_TCELL_UP, GSE44649_NAIVE_VS_ACTIVATED_CD8_TCELL_UP GSE4535_BM_DERIVED_DC_VS_FOLLICULAR_DC_UP, GSE4535_BM_DERIVED_DC_VS_FOLLICULAR_DC_UP GSE7460_WT_VS_FOXP3_HET_ACT_TCONV_DN, GSE7460_WT_VS_FOXP3_HET_ACT_TCONV_DN GSE557_WT_VS_CIT1A_KO_DC_UP, GSE557_WT_VS_CIT1A_KO_DC_UP GSE879_PPARG_LIGAND_ROSIGLITAZONE_VS_ROSIGLITAZONE_AND_RARA_AGONIST_AM580_TREATED_DC_UP, GSE879_PPARG_LIGAND_ROSIGLITAZONE_VS_ROSIGLITAZONE_AND_RARA_AGONIST_AM580_TREATED_DC_UP GSE40274_HELLOS_VS_FOXP3_AND_HELLOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_HELLOS_VS_FOXP3_AND_HELLOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN GSE7460_FOXP3_MUT_VS_WT_ACT_TCONV_UP, GSE7460_FOXP3_MUT_VS_WT_ACT_TCONV_UP GSE27786_LSK_VS_BCELL_UP, GSE27786_LSK_VS_BCELL_UP LIU_VMYB_TARGETS_UP, LIU_VMYB_TARGETS_UP GSE16522_ANTL_CD3CD28_STIM_VS_UNSTIM_MEMORY_CD8_TCELL_UP, GSE16522_ANTL_CD3CD28_STIM_VS_UNSTIM_MEMORY_CD8_TCELL_UP GSE21063_3H_VS_16H_ANTL_IIGM_STIM_NFATC1_KOBCELL_UP, GSE21063_3H_VS_16H_ANTL_IIGM_STIM_NFATC1_KOBCELL_UP GSE36078_UNTREATED_VS_AD5_T425A_HEXON_INF_IL1R_KO_MOUSE_LUNG_DC_DN, GSE36078_UNTREATED_VS_AD5_T425A_HEXON_INF_IL1R_KO_MOUSE_LUNG_DC_DN SCHERER_PBMC_APSV_WETVAX_AGE_18_32YO_5_TO_7DY_DN, SCHERER_PBMC_APSV_WETVAX_AGE_18_32YO_5_TO_7DY_DN GSE10239_MEMORY_VS_KLRG1INT_EFF_CD8_TCELL_UP, GSE10239_MEMORY_VS_KLRG1INT_EFF_CD8_TCELL_UP GSE5099_MONOCYTE_VS_ALTERNATIVE_M2_MACROPHAGE_UP, GSE5099_MONOCYTE_VS_ALTERNATIVE_M2_MACROPHAGE_UP MIR6734_3P, MIR6734_3P GSE12845_IJD_POS_BLOOD_VS_DARKZONE_GC_TONSIL_BCELL_UP, GSE12845_IJD_POS_BLOOD_VS_DARKZONE_GC_TONSIL_BCELL_UP GOBP_NUCLEOSIDE_BISPHOSPHATE_METABOLIC_PROCESS, GOBP_NUCLEOSIDE_BISPHOSPHATE_METABOLIC_PROCESS MIR891B, MIR891B GSE2405_0H_VS_6H_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_DN, GSE2405_0H_VS_6H_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_DN HP_RED_EYE_HP_RED_EYE MIR1288_5P, MIR1288_5P MIR1_3P, MIR1_3P GSE19941_UNSTIM_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBp50_KO_MACROPHAGE_UP, GSE19941_UNSTIM_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBp50_KO_MACROPHAGE_UP GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_FOXP3 TRASDUCED_CD4_TCELL_DN, GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_FOXP3 TRASDUCED_CD4_TCELL_DN MIR4769_3P, MIR4769_3P BROWNE_HCMV_INFECTION_10HR_UP, BROWNE_HCMV_INFECTION_10HR_UP MIR6817_5P, MIR6817_5P MIR17_3P, MIR17_3P GOBP_THIOESTER_METABOLIC_PROCESS, GOBP_THIOESTER_METABOLIC_PROCESS SHAFFER_IRF4_TARGETS_IN_ACTIVATED_DENDRITIC_CELL, SHAFFER_IRF4_TARGETS_IN_ACTIVATED_DENDRITIC_CELL MIR1304_5P, MIR1304_5P MIR4766_3P, MIR4766_3P MACAEVA_PBMC_RESPONSE_TO_IR, MACAEVA_PBMC_RESPONSE_TO_IR MIR1098_5P, MIR1098_5P MARTINEZ_RESPONSE_TO_TRABECTEDIN, MARTINEZ_RESPONSE_TO_TRABECTEDIN MIR567, MIR567 GOBP_PRIMARY_ALCOHOL_METABOLIC_PROCESS, GOBP_PRIMARY_ALCOHOL_METABOLIC_PROCESS GSE17974_IL4_AND_ANTL_IL12_VS_UNTREATED_2H_ACT_CD4_TCELL_DN, GSE17974_IL4_AND_ANTL_IL12_VS_UNTREATED_2H_ACT_CD4_TCELL_DN HP_BLEPHARITIS, HP_BLEPHARITIS MIR1258, MIR1258 GOCC_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX_PROTON_TRANSPORTING_DOMAIN, GOCC_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX_PROTON_TRANSPORTING_DOMAIN REACTOME_CARBOXYTERMINAL_POST_TRANSLATIONAL_MODIFICATIONS_OF_TUBULIN, REACTOME_CARBOXYTERMINAL_POST_TRANSLATIONAL_MODIFICATIONS_OF_TUBULIN MORE_DMIP, MORE_DMIP GOBP_PPTIDYL_TYROSINE_DEPHOSPHORYLATION, GOBP_PPTIDYL_TYROSINE_DEPHOSPHORYLATION MORE_TNRSF6, MORE_TNRSF6 OUILLETTE_CLL_13Q14_DELETION_DN, OUILLETTE_CLL_13Q14_DELETION_DN GOMF_PROTEIN_TYROSINE_PHOSPHATASE_ACTIVITY, GOMF_PROTEIN_TYROSINE_PHOSPHATASE_ACTIVITY REACTOME_ACTIVATION_OF_NMDA_RECEPTORS_AND_POSTSYNAPTIC_EVENTS, REACTOME_ACTIVATION_OF_NMDA_RECEPTORS_AND_POSTSYNAPTIC_EVENTS MIR108_3P, MIR108_3P MORE_PRKCA, MORE_PRKCA GOBP_FATTY_ACID_BIOSYNTHETIC_PROCESS, GOBP_FATTY_ACID_BIOSYNTHETIC_PROCESS HP_KERATITIS, HP_KERATITIS MORE_THRO, MORE_THRO MORE_MAP2K7, MORE_MAP2K7 MODULE_544, MODULE_544 MODULE_481, MODULE_481 REACTOME_PEROXISOMAL_PROTEIN_IMPORT, REACTOME_PEROXISOMAL_PROTEIN_IMPORT MIR873_3P, MIR873_3P GOCC_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX_COUPLING_FACTOR_F_O, GOCC_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX_COUPLING_FACTOR_F_O YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_4, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_4 MIR12125, MIR12125 GOBP_MIRNA_METABOLIC_PROCESS, GOBP_MIRNA_METABOLIC_PROCESS YANG_BREAST_CANCER_ESR1_DN, YANG_BREAST_CANCER_ESR1_DN GOBP_INTRA_S_DNA_DAMAGE_CHECKPOINT, GOBP_INTRA_S_DNA_DAMAGE_CHECKPOINT REACTOME_DEFECTS_IN_VITAMIN_AND_COFACTOR_METABOLISM, REACTOME_DEFECTS_IN_VITAMIN_AND_COFACTOR_METABOLISM GOMF_C_ACYLTRANSFERASE_ACTIVITY, GOMF_C_ACYLTRANSFERASE_ACTIVITY HP_ABNORMAL_HEAD_MOVEMENTS, HP_ABNORMAL_HEAD_MOVEMENTS DURCHDEWALD_SKIN_CARCINOGENESIS_UP, DURCHDEWALD_SKIN_CARCINOGENESIS_UP MIR4774_3P, MIR4774_3P HP_CUTANEOUS_MELANOMA, HP_CUTANEOUS_MELANOMA WP_OVERVIEW_OF_LEUKOCYTEINTRINSIC_HIPPO_PATHWAY_FUNCTIONS, WP_OVERVIEW_OF_LEUKOCYTEINTRINSIC_HIPPO_PATHWAY_FUNCTIONS MIR6826_5P, MIR6826_5P GOMF_CAMP_RESPONSE_ELEMENT_BINDING, GOMF_CAMP_RESPONSE_ELEMENT_BINDING MORE_PPP2R5B, MORE_PPP2R5B NAKAMURA_METASTASIS_MODEL_UP, NAKAMURA_METASTASIS_MODEL_UP MIR4722_3P, MIR4722_3P MORE_BCL2L1, MORE_BCL2L1 WP_SEROTONIN_RECEPTOR_2_AND_ELSKRFGATA4_SIGNALING, WP_SEROTONIN_RECEPTOR_2_AND_ELSKRFGATA4_SIGNALING GOCC_MITOCHONDRIAL_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX_COUPLING_FACTOR_F_O, GOCC_MITOCHONDRIAL_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX_COUPLING_FACTOR_F_O MORE_ITGA2, MORE_ITGA2 MIR7855_5P, MIR7855_5P MIR8078, MIR8078 HP_PTERYGIUM_HP_PTERYGIUM MIR4313, MIR4313 REACTOME_SIGNALING_BY_RETINOIC_ACID, REACTOME_SIGNALING_BY_RETINOIC_ACID BIOCARTA_CXCR4_PATHWAY, BIOCARTA_CXCR4_PATHWAY REACTOME_CONSTITUTIVE_SIGNALING_BY_AKT1_E17K_IN_CANCER, REACTOME_CONSTITUTIVE_SIGNALING_BY_AKT1_E17K_IN_CANCER LEIN_OLIGODENDROCYTE_MARKERS, LEIN_OLIGODENDROCYTE_MARKERS MORE_CD8A, MORE_CD8A GOUYER_TAT1_TARGETS_DN, GOUYER_TAT1_TARGETS_DN MIR520C_5P, MIR520C_5P HP_PAROXYSMAL_DYSTONIA, HP_PAROXYSMAL_DYSTONIA MIR410_5P, MIR410_5P SILIGAN_TARGETS_OF_EWS_FLI1_FUSION_DN, SILIGAN_TARGETS_OF_EWS_FLI1_FUSION_DN GOBP_TERPENOID_METABOLIC_PROCESS, GOBP_TERPENOID_METABOLIC_PROCESS MIR323B_5P, MIR323B_5P MIR494_5P, MIR494_5P GOBP_POSITIVE_REGULATION_OF_WOUND_HEALING, GOBP_POSITIVE_REGULATION_OF_WOUND_HEALING MIR9903, MIR9903 GOMF_STRUCTURAL_CONSTITUENT_OF_CYTOSKELETON, GOMF_STRUCTURAL_CONSTITUENT_OF_CYTOSKELETON CHO_NRA1_TARGETS, CHO_NRA1_TARGETS MORE_CTSB, MORE_CTSB PETRETTO_HEART_MASS_QTL_CIS_UP, PETRETTO_HEART_MASS_QTL_CIS_UP THEODOROU_MAMMARY_TUMORIGENESIS, THEODOROU_MAMMARY_TUMORIGENESIS MORE_TTN, MORE_TTN MIR924, MIR924 REACTOME_PEROXISOMAL_LIPID_METABOLISM, REACTOME_PEROXISOMAL_LIPID_METABOLISM MORE_KDR, MORE_KDR VALK_AML_CLUSTER_I5, VALK_AML_CLUSTER_I5 GOBP_PRENILATION, GOBP_PRENILATION HP_ABNORMAL_GLOBUS_PALLIDUS_MORPHOLOGY, HP_ABNORMAL_GLOBUS_PALLIDUS_MORPHOLOGY HP_INFANTILE_ENCEPHALOPATHY, HP_INFANTILE_ENCEPHALOPATHY REACTOME_TRIGLYCERIDE_METABOLISM, REACTOME_TRIGLYCERIDE_METABOLISM GOBP_POSITIVE_REGULATION_OF_OSTEOCLAST_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_OSTEOCLAST_DIFFERENTIATION GTCTACC_MIR379, GTCTACC_MIR379 GOBP_PYRAMIDAL_NEURON_DIFFERENTIATION, GOBP_PYRAMIDAL_NEURON_DIFFERENTIATION GOBP_UDP_N_ACETYLGUCOSAMINE_METABOLIC_PROCESS, GOBP_UDP_N_ACETYLGUCOSAMINE_METABOLIC_PROCESS GOMF_FATTY_ACID_DERIVATIVE_BINDING, GOMF_FATTY_ACID_DERIVATIVE_BINDING REACTOME_PHOSPHATE_BOND_HYDROLYSIS_BY_NUDT_PROTEINS, REACTOME_PHOSPHATE_BOND_HYDROLYSIS_BY_NUDT_PROTEINS GOBP_REGULATION_OF_FATTY_ACID_METABOLIC_PROCESS, GOBP_REGULATION_OF_FATTY_ACID_METABOLIC_PROCESS HP_UPPER_LIMB_HYPERTONIA, HP_UPPER_LIMB_HYPERTONIA GOBP_PONS_DEVELOPMENT, GOBP_PONS_DEVELOPMENT MORE_PTPRR, MORE_PTPRR HP_CENTRALLY_NUCLEATED_SKELETAL_MUSCLE_FIBERS, HP_CENTRALLY_NUCLEATED_SKELETAL_MUSCLE_FIBERS REACTOME_REGULATION_OF_FOXO_TRANSCRIPTIONAL_ACTIVITY_BY_ACETYLATION, REACTOME_REGULATION_OF_FOXO_TRANSCRIPTIONAL_ACTIVITY_BY_ACETYLATION WP_TRIACYLGLYCERIDE_SYNTHESIS, WP_TRIACYLGLYCERIDE_SYNTHESIS HP_ESOPHAGEAL_STRICTURE, HP_ESOPHAGEAL_STRICTURE MODULE_280, MODULE_280 HP_ABNORMAL_URINE_PHOSPHATE_CONCENTRATION, HP_ABNORMAL_URINE_PHOSPHATE_CONCENTRATION GOBP_DESMOSOME_ORGANIZATION, GOBP_DESMOSOME_ORGANIZATION HP_VENOUS_INSUFFICIENCY, HP_VENOUS_INSUFFICIENCY REACTOME_INTERLEUKIN_36_PATHWAY, REACTOME_INTERLEUKIN_36_PATHWAY HP_RETROCOLLIS, HP_RETROCOLLIS MIR181C_3P, MIR181C_3P GOMF_C_ACETYLTRANSFERASE_ACTIVITY, GOMF_C_ACETYLTRANSFERASE_ACTIVITY REACTOME_PURINE_CATABOLISM, REACTOME_PURINE_CATABOLISM NUNODA_RESPONSE_TO_DASATINIB_IMATINIB_DN, NUNODA_RESPONSE_TO_DASATINIB_IMATINIB_DN BIOCARTA_MTA3_PATHWAY, BIOCARTA_MTA3_PATHWAY GOBP_NUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS, GOBP_NUCLEOSIDE_DIPHOSPHATE_CATABOLIC_PROCESS FRASOR_TAMOXIFEN_RESPONSE_DN, FRASOR_TAMOXIFEN_RESPONSE_DN LEE_AGING_MUSCLE_DN, LEE_AGING_MUSCLE_DN GOMF_C_PALMITOYLTRANSFERASE_ACTIVITY, GOMF_C_PALMITOYLTRANSFERASE_ACTIVITY GOMF_3_HYDROXYACYL_COA_DEHYDRATASE_ACTIVITY, GOMF_3_HYDROXYACYL_COA_DEHYDRATASE_ACTIVITY GOBP_ACYLYGLYCEROL_ACYL_CHAIN_REMODELING, GOBP_ACYLYGLYCEROL_ACYL_CHAIN_REMODELING GOBP_MUSCLE_ATROPHY, GOBP_MUSCLE_ATROPHY GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_OXIDATION, GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_OXIDATION MORE_PDCD1, MORE_PDCD1 RUAN_RESPONSE_TO_TROGLITAZONE_UP, RUAN_RESPONSE_TO_TROGLITAZONE_UP GOBP_RESPONSE_TO_INTERLEUKIN_I7, GOBP_RESPONSE_TO_INTERLEUKIN_I7 GOBP_CEREBRAL_CORTIX_TANGENTIAL_MIGRATION, GOBP_CEREBRAL_CORTIX_TANGENTIAL_MIGRATION ZHONG_PFC_C2_ASC1L_POS_ASTROCYTE_ZHONG_PFC_C2_ASC1L_POS_ASTROCYTE FOURNIER_ACINAR_DEVELOPMENT_LATE_UP, FOURNIER_ACINAR_DEVELOPMENT_LATE_UP REACTOME_TRIGLYCERIDE_BIOSYNTHESIS, REACTOME_TRIGLYCERIDE_BIOSYNTHESIS HP_LOW_LEVELS_OF_VITAMIN_B1, HP_LOW_LEVELS_OF_VITAMIN_B1 GOBP_POSITIVE_REGULATION_OF_DEACETYLASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_DEACETYLASE_ACTIVITY GOBP_POSITIVE_REGULATION_OF_PEROXISOME_PROLIFERATOR_ACTIVATED_RECEPTOR_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_PEROXISOME_PROLIFERATOR_ACTIVATED_RECEPTOR_SIGNALING_PATHWAY GOMF_ADENYLATE_KINASE_ACTIVITY, GOMF_ADENYLATE_KINASE_ACTIVITY WEBER_METHYLATED_ICT_IN_SPERM_DN, WEBER_METHYLATED_ICT_IN_SPERM_DN HP_ABNORMAL_DLCO, HP_ABNORMAL_DLCO GOBP_STEROID_HOMEOSTASIS, GOBP_STEROID_HOMEOSTASIS GOCC_SYMMETRIC_SYNAPSE, GOCC_SYMMETRIC_SYNAPSE GOMF_CYSINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY, GOMF_CYSINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY GOBP_DIACYLGLYCEROL_BIOSYNTHETIC_PROCESS, GOBP_DIACYLGLYCEROL_BIOSYNTHETIC_PROCESS GOMF_G_PROTEIN_COUPLED_SEROTONIN_RECEPTOR_BINDING, GOMF_G_PROTEIN_COUPLED_SEROTONIN_RECEPTOR_BINDING GOBP_VESICLE_TETHERING, GOBP_VESICLE_TETHERING GOBP_PARAXIAL_MESODERM_FORMATION, GOBP_PARAXIAL_MESODERM_FORMATION GOMF_2_ACYLYGLYCEROL_O_ACYLTRANSFERASE_ACTIVITY, GOMF_2_ACYLYGLYCEROL_O_ACYLTRANSFERASE_ACTIVITY WP_CEREBRAL_ORGANIC_ACIDURIAS_INCLUDING_DISEASES, WP_CEREBRAL_ORGANIC_ACIDURIAS_INCLUDING_DISEASES GOBP_NEGATIVE_REGULATION_OF_MACROPHAGE_CHEMOTAXIS, GOBP_NEGATIVE_REGULATION_OF_MACROPHAGE_CHEMOTAXIS GOBP_CORTICOSPINAL_TRACT_MORPHOGENESIS, GOBP_CORTICOSPINAL_TRACT_MORPHOGENESIS GOBP_PYRUVATE_BIOSYNTHETIC_PROCESS, GOBP_PYRUVATE_BIOSYNTHETIC_PROCESS GOBP_POSITIVE_REGULATION_OF_HISTONE_DEACETYLASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_HISTONE_DEACETYLASE_ACTIVITY GOBP_T_FOLLICULAR_HELPER_CELL_DIFFERENTIATION, GOBP_T_FOLLICULAR_HELPER_CELL_DIFFERENTIATION KAUFFMANN_MELANOMA_RELAPSE_DN, KAUFFMANN_MELANOMA_RELAPSE_DN HP_VIRAL_INFECTION_INDUCED_RHABDOMYOLYSIS, HP_VIRAL_INFECTION_INDUCED_RHABDOMYOLYSIS HP_EXTRAPYRAMIDAL_DYSKINESIA, HP_EXTRAPYRAMIDAL_DYSKINESIA GOBP_CARNITINE_SHUTTLE, GOBP_CARNITINE_SHUTTLE GOMF_OPIOID_RECEPTOR_ACTIVITY, GOMF_OPIOID_RECEPTOR_ACTIVITY GOBP_LUTEOLYSIS, GOBP_LUTEOLYSIS FARDIN_HYPONIA_9, FARDIN_HYPONIA_9 REACTOME_ACYL_CHAIN_REMODELING_OF_DAG_AND_TAG, REACTOME_ACYL_CHAIN_REMODELING_OF_DAG_AND_TAG WP_OXYSTEROLS_DERIVED_FROM_CHOLESTEROL, WP_OXYSTEROLS_DERIVED_FROM_CHOLESTEROL KEGG_RETINOL_METABOLISM, KEGG_RETINOL_METABOLISM GOBP_DENSE_CORE_GRANULE_TRANSPORT, GOBP_DENSE_CORE_GRANULE_TRANSPORT GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_5_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_5_PRODUCTION HP_TRANSIENT_NEONATAL_DIABETES_MELLITUS, HP_TRANSIENT_NEONATAL_DIABETES_MELLITUS GOMF_COA_HYDROLASE_ACTIVITY, GOMF_COA_HYDROLASE_ACTIVITY GOBP_ACTIN_MODIFICATION, GOBP_ACTIN_MODIFICATION GOBP_PARANODAL_JUNCTION_ASSEMBLY, GOBP_PARANODAL_JUNCTION_ASSEMBLY GOBP_FAT_PAD_DEVELOPMENT, GOBP_FAT_PAD_DEVELOPMENT GOBP_POSITIVE_REGULATION_OF_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS, GOBP_POSITIVE_REGULATION_OF_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS GOBP_REGULATION_OF_SKELETAL_MUSCLE_ACETYLCHOLINE_GATED_CHANNEL_CLUSTERING, GOBP_REGULATION_OF_SKELETAL_MUSCLE_ACETYLCHOLINE_GATED_CHANNEL_CLUSTERING GOMF_DIACYLGLYCEROL_O_ACYLTRANSFERASE_ACTIVITY, GOMF_DIACYLGLYCEROL_O_ACYLTRANSFERASE_ACTIVITY HP_RENAL_SODIUM_WASTING, HP_RENAL_SODIUM_WASTING HP_WIDENED_CEREBELLAR_SUBARACHNOID_SPACE, HP_WIDENED_CEREBELLAR_SUBARACHNOID_SPACE GOBP_THREONINE_CATABOLIC_PROCESS, GOBP_THREONINE_CATABOLIC_PROCESS GOBP_FATTY_ACID_OMEGA_OXIDATION, GOBP_FATTY_ACID_OMEGA_OXIDATION GOMF_TYPE_2A_SEROTONIN_RECEPTOR_BINDING, GOMF_TYPE_2A_SEROTONIN_RECEPTOR_BINDING FUJIWARA_PARK2_IN_LIVER_CANCER_UP, FUJIWARA_PARK2_IN_LIVER_CANCER_UP GOBP_REGULATION_OF_EXCITATORY_SYNAPSE_ASSEMBLY, GOBP_REGULATION_OF_EXCITATORY_SYNAPSE_ASSEMBLY NR5A1_TARGET_GENES, NR5A1_TARGET_GENES GOMF_CYSINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_EXECUTION_PHASE_OF_APOPTOSIS, GOMF_CYSINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_EXECUTION_PHASE_OF_APOPTOSIS HP_INCREASED_URINARY_POTASSIUM, HP_INCREASED_URINARY_POTASSIUM PHESSIE_TARGETS_OF_APC_AND_MBD2_UP, PHESSIE_TARGETS_OF_APC_AND_MBD2_UP GOBP_REGULATION_OF_RECEPTOR_CLUSTERING, GOBP_REGULATION_OF_RECEPTOR_CLUSTERING GOBP_GERMINAL_CENTER_B_CELL_DIFFERENTIATION, GOBP_GERMINAL_CENTER_B_CELL_DIFFERENTIATION HP_UVEAL_MELANOMA, HP_UVEAL_MELANOMA GOBP_PURINE_NUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_PURINE_NUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS GOBP_RIBONUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_RIBONUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS GOBP_ANTERIOR_COMMISURE_MORPHOGENESIS, GOBP_ANTERIOR_COMMISURE_MORPHOGENESIS GOMF_NUCLEOSIDE_DIPHOSPHATASE_ACTIVITY, GOMF_NUCLEOSIDE_DIPHOSPHATASE_ACTIVITY GOMF_RETINOL_DEHYDROGENASE_ACTIVITY, GOMF_RETINOL_DEHYDROGENASE_ACTIVITY GOBP_REGULATION_OF_UNSATURATED_FATTY_ACID_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_UNSATURATED_FATTY_ACID_BIOSYNTHETIC_PROCESS MIR4787_5P, MIR4787_5P GOBP_OPIOID_RECEPTOR_SIGNALING_PATHWAY, GOBP_OPIOID_RECEPTOR_SIGNALING_PATHWAY GOBP_RESPONSE_TO_GRAVITY, GOBP_RESPONSE_TO_GRAVITY GOBP_SEGMENT_SPECIFICATION, GOBP_SEGMENT_SPECIFICATION GOMF_RECEPTOR_SERINE_THREONINE_KINASE_BINDING, GOMF_RECEPTOR_SERINE_THREONINE_KINASE_BINDING HP_ABNORMAL_ORAL_GLUCOSE_TOLERANCE, HP_ABNORMAL_ORAL_GLUCOSE_TOLERANCE HP_AXIAL_DYSTONIA, HP_AXIAL_DYSTONIA GOBP_REGULATION_OF_POSTSYNAPTIC_SPECIALIZATION_ASSEMBLY, GOBP_REGULATION_OF_POSTSYNAPTIC_SPECIALIZATION_ASSEMBLY HP_HYPOPHOSPHATEMIA, HP_HYPOPHOSPHATEMIA HP_DESQUAMATION_OF_SKIN_SOON_AFTER_BIRTH, HP_DESQUAMATION_OF_SKIN_SOON_AFTER_BIRTH HP_INFERIOR_LENS_SUBLUXATION, HP_INFERIOR_LENS_SUBLUXATION GOMF_INTERLEUKIN_2_RECEPTOR_BINDING, GOMF_INTERLEUKIN_2_RECEPTOR_BINDING HP_BICARBONATE_WASTING_RENAL_TUBULAR_ACIDOSIS, HP_BICARBONATE_WASTING_RENAL_TUBULAR_ACIDOSIS HP_BIVENTRICULAR_HYPERTROPHY, HP_BIVENTRICULAR_HYPERTROPHY GOBP_ADP_BIOSYNTHETIC_PROCESS, GOBP_ADP_BIOSYNTHETIC_PROCESS GOBP_ENUCLEATE_ERYTHROCYTE_DIFFERENTIATION, GOBP_ENUCLEATE_ERYTHROCYTE_DIFFERENTIATION GOCC_MSL_COMPLEX, GOCC_MSL_COMPLEX GOMF_BIOTIN_BINDING, GOMF_BIOTIN_BINDING HP_LOW_MOLECULAR_WEIGHT_PROTEINURIA, HP_LOW_MOLECULAR_WEIGHT_PROTEINURIA GOBP_TRANS_SYNAPTIC_SIGNALING_BY_TRANS_SYNAPTIC_COMPLEX, GOBP_TRANS_SYNAPTIC_SIGNALING_BY_TRANS_SYNAPTIC_COMPLEX GOBP_SLEEP, GOBP_SLEEP SETLUR_PROSTATE_CANCER_TMPRS2_ERG_FUSION_DN, SETLUR_PROSTATE_CANCER_TMPRS2_ERG_FUSION_DN HP_BICARBONATURIA, HP_BICARBONATURIA HP_ABNORMAL_CILIARY_BODY_MORPHOLOGY, HP_ABNORMAL_CILIARY_BODY_MORPHOLOGY GOBP_POSITIVE_REGULATION_OF_RECEPTOR_CLUSTERING, GOBP_POSITIVE_REGULATION_OF_RECEPTOR_CLUSTERING GOMF_AMINOACYLASE_ACTIVITY, GOMF_AMINOACYLASE_ACTIVITY GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_I3_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_I3_PRODUCTION GOMF_ALCOHOL_DEHYDROGENASE_ACTIVITY_ZINC_DEPENDENT, GOMF_ALCOHOL_DEHYDROGENASE_ACTIVITY_ZINC_DEPENDENT chr14q13, chr14q13 GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_METABOLIC_PROCESS TUOMISTO_TUMOR_SUPPRESSION_BY_COL13A1_DN, TUOMISTO_TUMOR_SUPPRESSION_BY_COL13A1_DN GOBP_STEROID_ESTERIFICATION, GOBP_STEROID_ESTERIFICATION HP_RENAL_PHOSPHATE_WASTING, HP_RENAL_PHOSPHATE_WASTING WP_SLEEP_REGULATION, WP_SLEEP_REGULATION
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