

	GSE10239_NAIVE_VS_MEMORY_CD8_TCELL_UP, GSE10239_NAIVE_VS_MEMORY_CD8_TCELL_UP GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP GSE411_WT_VS_SOCS3_KO_MACROPHAGE_UP, GSE411_WT_VS_SOCS3_KO_MACROPHAGE_UP GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN GSE7831_UNSTIM_VS_CPG_STIM_PDC_4H_UP, GSE7831_UNSTIM_VS_CPG_STIM_PDC_4H_UP GSE8835_HEALTHY_VS_CLL_CD8_TCELL_UP, GSE8835_HEALTHY_VS_CLL_CD8_TCELL_UP GSE46025_WT_VS_FOXO1_KO_KLRG1_LOW_CD8_EFFECTOR_TCELL_DN, GSE46025_WT_VS_FOXO1_KO_KLRG1_LOW_CD8_EFFECTOR_TCELL_DN GSE21379_WT_VS_SAP_KO_CD4_TCELL_DN, GSE21379_WT_VS_SAP_KO_CD4_TCELL_DN GSE14415_INDUCED_TREG_VS_FAILED_INDUCED_TREG_DN, GSE14415_INDUCED_TREG_VS_FAILED_INDUCED_TREG_DN HALLMARK_HEME_METABOLISM, HALLMARK_HEME_METABOLISM GSE7460_TREG_VS_TCONV_ACT_UP, GSE7460_TREG_VS_TCONV_ACT_UP GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TREG_DN, GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TREG_DN GSE25087_TREG_VS_TCONV_ADULT_UP, GSE25087_TREG_VS_TCONV_ADULT_UP GSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_DN, GSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_DN GSE28737_WT_VS_BCL6_KO_MARGINAL_ZONE_BCELL_DN, GSE28737_WT_VS_BCL6_KO_MARGINAL_ZONE_BCELL_DN MIR3168, MIR3168 GSE7460_WT_VS_FOXP3_HET_ACT_WITH_TGFB_TCONV_DN, GSE7460_WT_VS_FOXP3_HET_ACT_WITH_TGFB_TCONV_DN GSE26669_CD4_VS_CD8_TCELL_IN_MLR_COSTIM_BLOCK_UP, GSE26669_CD4_VS_CD8_TCELL_IN_MLR_COSTIM_BLOCK_UP GSE14386_UNTREATED_VS_IFNA_TREATED_ACT_PBMCS_PATIENT_DN, GSE14386_UNTREATED_VS_IFNA_TREATED_ACT_PBMCS_PATIENT_DN MIR6511A_5P, MIR6511A_5P GOBP_CYTOSKELETON_DEPENDENT_CYTOKINESIS, GOBP_CYTOSKELETON_DEPENDENT_CYTOKINESIS GSE17721_LPS_VS_POLYIC_2H_BMDC_UP, GSE17721_LPS_VS_POLYIC_2H_BMDC_UP GSE7348_UNSTIM_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_DN, GSE7348_UNSTIM_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_DN GSE23114_WT_VS_SLE2C1_MOUSE_PERITONEAL_CAVITY_BIA_BCELL_UP, GSE23114_WT_VS_SLE2C1_MOUSE_PERITONEAL_CAVITY_BIA_BCELL_UP SENESE_HDAC2_TARGETS_UP, SENESE_HDAC2_TARGETS_UP GSE37301_MULTIPOTENT_PROGENITOR_VS_GRAN_MONO_PROGENITOR_UP, GSE37301_MULTIPOTENT_PROGENITOR_VS_GRAN_MONO_PROGENITOR_UP GOCC_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE, GOCC_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE MIR942_3P, MIR942_3P ACTACCT_MIR196A_MIR196B, ACTACCT_MIR196A_MIR196B MIR3674, MIR3674 GOCC_NEURON_PROJECTION_CYTOPLASM, GOCC_NEURON_PROJECTION_CYTOPLASM GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_UP, GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_UP GOCC_H4_HISTONE_ACETYLTRANSFERASE_COMPLEX, GOCC_H4_HISTONE_ACETYLTRANSFERASE_COMPLEX MIR1208, MIR1208 MIR4697_3P, MIR4697_3P SESTO_RESPONSE_TO_UV_C8, SESTO_RESPONSE_TO_UV_C8 MIR218_1_3P, MIR218_1_3P GOBP_CYTOKINETIC_PROCESS, GOBP_CYTOKINETIC_PROCESS GOCC_NUCLEAR_INNER_MEMBRANE, GOCC_NUCLEAR_INNER_MEMBRANE GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_UP, GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_UP GOMF_KINESIN_BINDING, GOMF_KINESIN_BINDING MIR2682_5P, MIR2682_5P GSE25123_CTRL_VS_IL4_STIM_PPARG_KO_MACROPHAGE_DN, GSE25123_CTRL_VS_IL4_STIM_PPARG_KO_MACROPHAGE_DN HP_ABNORMAL_SACRUM_MORPHOLOGY, HP_ABNORMAL_SACRUM_MORPHOLOGY GOBP_MITOTIC_SISTER_CHROMATID_COHESION, GOBP_MITOTIC_SISTER_CHROMATID_COHESION MIR6505_3P, MIR6505_3P MIR4802_3P, MIR4802_3P MIR487A_5P, MIR487A_5P MIR196A_5P_MIR196B_5P, MIR196A_5P_MIR196B_5P MIR487B_5P, MIR487B_5P GOBP_REGULATION_OF_RECEPTOR_MEDIATED_ENDOCYTOSIS, GOBP_REGULATION_OF_RECEPTOR_MEDIATED_ENDOCYTOSIS DESCARTES_FETAL_STOMACH_ERYTHROBLASTS, DESCARTES_FETAL_STOMACH_ERYTHROBLASTS GOCC_TRANS_GOLGI_NETWORK_TRANSPORT_VESICLE, GOCC_TRANS_GOLGI_NETWORK_TRANSPORT_VESICLE GSE21927_SPLENIC_C26GM_TUMOROUS_VS_4T1_TUMOR_MONOCYTES_UP, GSE21927_SPLENIC_C26GM_TUMOROUS_VS_4T1_TUMOR_MONOCYTES_UP MIR875_5P, MIR875_5P MIR4274, MIR4274 MIR33A_5P_MIR33B_5P, MIR33A_5P_MIR33B_5P GSE17721_CTRL_VS_CPG_1H_BMDC_DN, GSE17721_CTRL_VS_CPG_1H_BMDC_DN HP_ABNORMALITY_OF_THE_RADIAL_HEAD, HP_ABNORMALITY_OF_THE_RADIAL_HEAD MIR3935, MIR3935 MIR346, MIR346 GOCC_SWLSNF_COMPLEX, GOCC_SWLSNF_COMPLEX ZNF777_TARGET_GENES, ZNF777_TARGET_GENES GOCC_ADA2_GCN5_ADA3_TRANSCRIPTION_ACTIVATOR_COMPLEX, GOCC_ADA2_GCN5_ADA3_TRANSCRIPTION_ACTIVATOR_COMPLEX GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT, GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATION_MEDIATED_BY_ACTIVATED_HUMAN_TAK1, REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATION_MEDIATED_BY_ACTIVATED_HUMAN_TAK1 MIR488_5P, MIR488_5P MIR4421, MIR4421 HP_PROFOUND_GLOBAL_DEVELOPMENTAL_DELAY, HP_PROFOUND_GLOBAL_DEVELOPMENTAL_DELAY GOBP_NEGATIVE_REGULATION_OF_ERBB_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_ERBB_SIGNALING_PATHWAY MIR106A_3P, MIR106A_3P HP_APLASIA_HYPOPLASIA_AFFECTING_THE_FUNDUS, HP_APLASIA_HYPOPLASIA_AFFECTING_THE_FUNDUS GOBP_PROTEIN_TETRAMERIZATION, GOBP_PROTEIN_TETRAMERIZATION MIR7975, MIR7975 HORTON_SREBF_TARGETS, HORTON_SREBF_TARGETS HP_HYPOCHOLESTEROLEMIA, HP_HYPOCHOLESTEROLEMIA BURTON_ADIPOGENESIS_10, BURTON_ADIPOGENESIS_10 GOMF_CHANNEL_INHIBITOR_ACTIVITY, GOMF_CHANNEL_INHIBITOR_ACTIVITY MIR4522, MIR4522 MIR632, MIR632 GOBP_VENTRICULAR_SYSTEM_DEVELOPMENT, GOBP_VENTRICULAR_SYSTEM_DEVELOPMENT HEIDENBLAD_AMPLICON_12P11_12_UP, HEIDENBLAD_AMPLICON_12P11_12_UP REACTOME_SIGNALING_BY_EGFR, REACTOME_SIGNALING_BY_EGFR HP_OPTIC_NERVE_HYPOPLASIA, HP_OPTIC_NERVE_HYPOPLASIA GOBP_POSITIVE_REGULATION_OF_RECEPTOR_MEDIATED_ENDOCYTOSIS, GOBP_POSITIVE_REGULATION_OF_RECEPTOR_MEDIATED_ENDOCYTOSIS GOBP_GOLGI_TO_ENDOSOME_TRANSPORT, GOBP_GOLGI_TO_ENDOSOME_TRANSPORT GOBP_NUCLEAR_ENVELOPE_REASSEMBLY, GOBP_NUCLEAR_ENVELOPE_REASSEMBLY TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_DUCTAL_NORMAL_UP, TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_DUCTAL_NORMAL_UP GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_24H_CD8_T_CELL_UP, GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_24H_CD8_T_CELL_UP GENTILE_UV_RESPONSE_CLUSTER_D2, GENTILE_UV_RESPONSE_CLUSTER_D2 MIR380_5P, MIR380_5P GOBP_PROTEIN_HOMOTETRAMERIZATION, GOBP_PROTEIN_HOMOTETRAMERIZATION GOBP_REGULATION_OF_RECEPTOR_BINDING, GOBP_REGULATION_OF_RECEPTOR_BINDING HP_SLURRED_SPEECH, HP_SLURRED_SPEECH GCM_PPKAG1, GCM_PPKAG1 GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY MODULE_509, MODULE_509 DESCARTES_FETAL_EYE_SMOOTH_MUSCLE_CELLS, DESCARTES_FETAL_EYE_SMOOTH_MUSCLE_CELLS GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BROWN_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BROWN_UP HP_CHRONIC_LEUKEMIA, HP_CHRONIC_LEUKEMIA HP_ACCESSORY_SPLEEN, HP_ACCESSORY_SPLEEN GOBP_SYNAPTONEMAL_COMPLEX_ORGANIZATION, GOBP_SYNAPTONEMAL_COMPLEX_ORGANIZATION GOBP_GLOMERULAR_VISCERAL_EPITHELIAL_CELL_MIGRATION, GOBP_GLOMERULAR_VISCERAL_EPITHELIAL_CELL_MIGRATION PRDM12_TARGET_GENES, PRDM12_TARGET_GENES GOBP_NEGATIVE_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY BIOCARTA_41BB_PATHWAY, BIOCARTA_41BB_PATHWAY GOMF_PHOSPHATIDYLINOSITOL_4_PHOSPHATE_BINDING, GOMF_PHOSPHATIDYLINOSITOL_4_PHOSPHATE_BINDING REACTOME_ESTABLISHMENT_OF_SISTER_CHROMATID_COHESION, REACTOME_ESTABLISHMENT_OF_SISTER_CHROMATID_COHESION WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_MAPK_SIGNALING, WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_MAPK_SIGNALING PID_EPHRINB_REV_PATHWAY, PID_EPHRINB_REV_PATHWAY HP_METAPHYSEAL_STRIATIONS, HP_METAPHYSEAL_STRIATIONS GOBP_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_APOPTOTIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_APOPTOTIC_PROCESS GOMF_CD4_RECEPTOR_BINDING, GOMF_CD4_RECEPTOR_BINDING MIR6769B_5P, MIR6769B_5P MIR4751, MIR4751 GOBP_POTASSIUM_ION_IMPORT_ACROSS_PLASMA_MEMBRANE, GOBP_POTASSIUM_ION_IMPORT_ACROSS_PLASMA_MEMBRANE LEE_LIVER_CANCER_MYC_TGFA_UP, LEE_LIVER_CANCER_MYC_TGFA_UP GOBP_REGULATION_OF_SPONTANEOUS_SYNAPTIC_TRANSMISSION, GOBP_REGULATION_OF_SPONTANEOUS_SYNAPTIC_TRANSMISSION GOCC_COHESIN_COMPLEX, GOCC_COHESIN_COMPLEX GOBP_TAIL_ANCHORED_MEMBRANE_PROTEIN_INSERTION_INTO_ER_MEMBRANE, GOBP_TAIL_ANCHORED_MEMBRANE_PROTEIN_INSERTION_INTO_ER_MEMBRANE ZNF26_TARGET_GENES, ZNF26_TARGET_GENES GOBP_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE, GOBP_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE GOBP_ADRENERGIC_RECEPTOR_SIGNALING_PATHWAY, GOBP_ADRENERGIC_RECEPTOR_SIGNALING_PATHWAY GOBP_POSITIVE_REGULATION_OF_MUSCLE_CELL_APOPTOTIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_MUSCLE_CELL_APOPTOTIC_PROCESS GOBP_RESPONSE_TO_LEAD_ION, GOBP_RESPONSE_TO_LEAD_ION MIR6769A_5P, MIR6769A_5P REACTOME_INSULIN_PROCESSING, REACTOME_INSULIN_PROCESSING GOBP_POSITIVE_REGULATION_OF_RECEPTOR_BINDING, GOBP_POSITIVE_REGULATION_OF_RECEPTOR_BINDING MIR6800_3P, MIR6800_3P HP_SUBCORTICAL_CEREBRAL_ATROPHY, HP_SUBCORTICAL_CEREBRAL_ATROPHY GOMF_MEMBRANE_INSERTASE_ACTIVITY, GOMF_MEMBRANE_INSERTASE_ACTIVITY GOCC_CALCINEURIN_COMPLEX, GOCC_CALCINEURIN_COMPLEX WANG_BARRETTES_ESOPHAGUS_DN, WANG_BARRETTES_ESOPHAGUS_DN GOBP_POSITIVE_REGULATION_OF_MYOBlast_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_MYOBlast_PROLIFERATION GOBP_RESPONSE_TO_MANGANESE_ION, GOBP_RESPONSE_TO_MANGANESE_ION VISALA_RESPONSE_TO_HEAT_SHOCK_AND_AGING_UP, VISALA_RESPONSE_TO_HEAT_SHOCK_AND_AGING_UP GOBP_TRIGLYCERIDE_METABOLIC_PROCESS, GOBP_TRIGLYCERIDE_METABOLIC_PROCESS HP_DEGENERATION_OF_THE_STRIATUM, HP_DEGENERATION_OF_THE_STRIATUM GOCC_DENDRITE_CYTOPLASM, GOCC_DENDRITE_CYTOPLASM GOBP_BONE_MATURATION, GOBP_BONE_MATURATION chr1p22, chr1p22 GOBP_SODIUM_ION_IMPORT_ACROSS_PLASMA_MEMBRANE, GOBP_SODIUM_ION_IMPORT_ACROSS_PLASMA_MEMBRANE TAKAYAMA_BOUND_BY_AR, TAKAYAMA_BOUND_BY_AR HP_INCREASED_DENSITY_OF_LONG_BONES, HP_INCREASED_DENSITY_OF_LONG_BONES GOBP_REGULATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CONTRACTION, GOBP_REGULATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CONTRACTION GOBP_MEIOTIC_CYTOKINESIS, GOBP_MEIOTIC_CYTOKINESIS GOBP_NEGATIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_RECEPTOR_ACTIVITY GOBP_ESTABLISHMENT_OF_EPITHELIAL_CELL_POLARITY, GOBP_ESTABLISHMENT_OF_EPITHELIAL_CELL_POLARITY chr6q25, chr6q25 LAKE_ADULT_KIDNEY_C1_EPITHELIAL_CELLS_UNASSIGNED, LAKE_ADULT_KIDNEY_C1_EPITHELIAL_CELLS_UNASSIGNED GOBP_NEGATIVE_REGULATION_OF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY WP_OXIDATIVE_STRESS, WP_OXIDATIVE_STRESS IRITANI_MAD1_TARGETS_UP, IRITANI_MAD1_TARGETS_UP WP_METABOLIC_PATHWAY_OF_LDL_HDL_AND_TG_INCLUDING_DISEASES, WP_METABOLIC_PATHWAY_OF_LDL_HDL_AND_TG_INCLUDING_DISEASES GOCC_HOLLIDAY_JUNCTION_RESOLVASE_COMPLEX, GOCC_HOLLIDAY_JUNCTION_RESOLVASE_COMPLEX HP_DIAPHYSEAL_SCLEROSIS, HP_DIAPHYSEAL_SCLEROSIS HP_PREMATURE_ARTERIOSCLEROSIS, HP_PREMATURE_ARTERIOSCLEROSIS GOMF_PHOSPHOLIPASE_INHIBITOR_ACTIVITY, GOMF_PHOSPHOLIPASE_INHIBITOR_ACTIVITY GOBP_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_APOPTOTIC_PROCESS, GOBP_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_APOPTOTIC_PROCESS HOXB7_TARGET_GENES, HOXB7_TARGET_GENES GOBP_EPITHELIAL_FLUID_TRANSPORT, GOBP_EPITHELIAL_FLUID_TRANSPORT GOBP_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE, GOBP_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE GOBP_RESPONSE_TO_SELENIUM_ION, GOBP_RESPONSE_TO_SELENIUM_ION GOCC_ENDODEOXYRIBONUCLEASE_COMPLEX, GOCC_ENDODEOXYRIBONUCLEASE_COMPLEX GOBP_CELLULAR_RESPONSE_TO_POTASSIUM_ION, GOBP_CELLULAR_RESPONSE_TO_POTASSIUM_ION SEF1_C, SEF1_C GOBP_LEUKOCYTE_ADHESION_TO_VASCULAR_ENDOTHELIAL_CELL, GOBP_LEUKOCYTE_ADHESION_TO_VASCULAR_ENDOTHELIAL_CELL GOCC_SUPER_ELONGATION_COMPLEX, GOCC_SUPER_ELONGATION_COMPLEX GOBP_PHENOTYPIC_SWITCHING, GOBP_PHENOTYPIC_SWITCHING HP_CEREBELLAR_VERMIS_ATROPHY, HP_CEREBELLAR_VERMIS_ATROPHY GOBP_PROTON_TRANSPORTING_V_TYPE_ATPASE_COMPLEX_ASSEMBLY, GOBP_PROTON_TRANSPORTING_V_TYPE_ATPASE_COMPLEX_ASSEMBLY HP_CORONARY_ARTERY_ANEURYSM, HP_CORONARY_ARTERY_ANEURYSM HP_DELAYED_SOCIAL_DEVELOPMENT, HP_DELAYED_SOCIAL_DEVELOPMENT GOBP_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_RECEPTOR_CATABOLIC_PROCESS, GOBP_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_RECEPTOR_CATABOLIC_PROCESS GOBP_NEGATIVE_REGULATION_OF_CELL_PROLIFERATION_INVOLVED_IN_KIDNEY_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_CELL_PROLIFERATION_INVOLVED_IN_KIDNEY_DEVELOPMENT RNTCANNRNNYNATTW_UNKNOWN, RNTCANNRNNYNATTW_UNKNOWN GOMF_NEUROTRANSMITTER_BINDING, GOMF_NEUROTRANSMITTER_BINDING WP_ONE_CARBON_METABOLISM_AND_RELATED_PATHWAYS, WP_ONE_CARBON_METABOLISM_AND_RELATED_PATHWAYS HP_GAZE_EVOKED_HORIZONTAAL_NYSTAGMUS, HP_GAZE_EVOKED_HORIZONTAAL_NYSTAGMUS GCM_PPKCG, GCM_PPKCG HP_PARTIAL_ALBINISM, HP_PARTIAL_ALBINISM HP_VISUAL_FIXATION_INSTABILITY, HP_VISUAL_FIXATION_INSTABILITY GOBP_RESPONSE_TO_L_ASCORBIC_ACID, GOBP_RESPONSE_TO_L_ASCORBIC_ACID GOBP_MODULATION_BY_HOST_OF_SYMBIONT_MOLECULAR_FUNCTION, GOBP_MODULATION_BY_HOST_OF_SYMBIONT_MOLECULAR_FUNCTION GOCC_GAMMA_SECRETASE_COMPLEX, GOCC_GAMMA_SECRETASE_COMPLEX GOBP_ADENYLATE_CYCLASE_INHIBITING_G_PROTEIN_COUPLED_GLUTAMATE_RECEPTOR_SIGNALING_PATHWAY, GOBP_ADENYLATE_CYCLASE_INHIBITING_G_PROTEIN_COUPLED_GLUTAMATE_RECEPTOR_SIGNALING_PATHWAY GOBP_SEBACEOUS_GLAND_DEVELOPMENT, GOBP_SEBACEOUS_GLAND_DEVELOPMENT GRAHAM_CML_QUIESCENT_VS_CML_DIVIDING_UP, GRAHAM_CML_QUIESCENT_VS_CML_DIVIDING_UP NIKOLSKY_BREAST_CANCER_17P11_AMPLICON, NIKOLSKY_BREAST_CANCER_17P11_AMPLICON
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