	GSE22886_NAIVE_CD4_TCELL_VS_12H_ACT_TH2_UP, GSE22886_NAIVE_CD4_TCELL_VS_12H_ACT_TH2_UP GSE29618_MONOCYTE_VS_PDC_DN, GSE29618_MONOCYTE_VS_PDC_DN GSE29618_BCELL_VS_PDC_DAY7_FLU_VACCINE_DN, GSE29618_BCELL_VS_PDC_DAY7_FLU_VACCINE_DN GSE29618_BCELL_VS_PDC_DN, GSE29618_BCELL_VS_PDC_DN GSE29618_PDC_VS_MDC_DAY7_FLU_VACCINE_UP, GSE29618_PDC_VS_MDC_DAY7_FLU_VACCINE_UP REACTOME_DEATH_RECEPTOR_SIGNALLING, REACTOME_DEATH_RECEPTOR_SIGNALLING GSE11961_MARGINAL_ZONE_BCELL_VS_GERMINAL_CENTER_BCELL_DAY7_DN, GSE11961_MARGINAL_ZONE_BCELL_VS_GERMINAL_CENTER_BCELL_DAY7_DN GSE1460_DP_VS_CD4_THYMOCYTE_UP, GSE1460_DP_VS_CD4_THYMOCYTE_UP
	GSE1460_DP_VS_CD4_THYMOCYTE_DP, GSE1460_DP_VS_CD4_THYMOCYTE_DP GSE9006_HEALTHY_VS_TYPE_2_DIABETES_PBMC_AT_DX_DN, GSE9006_HEALTHY_VS_TYPE_2_DIABETES_PBMC_AT_DX_DN GSE43863_DAY6_EFF_VS_DAY150_MEM_TH1_CD4_TCELL_DN, GSE43863_DAY6_EFF_VS_DAY150_MEM_TH1_CD4_TCELL_DN GSE29618_PDC_VS_MDC_UP, GSE29618_PDC_VS_MDC_UP GSE9006_1MONTH_VS_4MONTH_POST_TYPE_1_DIABETES_DX_PBMC_UP, GSE9006_1MONTH_VS_4MONTH_POST_TYPE_1_DIABETES_DX_PBMC_UP GSE30971_WBP7_HET_VS_KO_MACROPHAGE_2H_LPS_STIM_DN, GSE30971_WBP7_HET_VS_KO_MACROPHAGE_2H_LPS_STIM_DN GOBP_ORGANELLE_FUSION, GOBP_ORGANELLE_FUSION GSE2706_2H_VS_8H_R848_STIM_DC_DN, GSE2706_2H_VS_8H_R848_STIM_DC_DN
	GSE3994_WT_VS_PAC1_KO_ACTIVATED_MAST_CELL_DN, GSE3994_WT_VS_PAC1_KO_ACTIVATED_MAST_CELL_DN GSE3982_BCELL_VS_BASOPHIL_DN, GSE3982_BCELL_VS_BASOPHIL_DN GSE17974_0H_VS_4H_IN_VITRO_ACT_CD4_TCELL_UP, GSE17974_0H_VS_4H_IN_VITRO_ACT_CD4_TCELL_UP GSE17974_0H_VS_4H_IN_VITRO_ACT_CD4_TCELL_UP, GSE17974_0H_VS_4H_IN_VITRO_ACT_CD4_TCELL_UP MIR4275, MIR4275 RUIZ_TNC_TARGETS_UP, RUIZ_TNC_TARGETS_UP GSE17721_POLYIC_VS_PAM3CSK4_1H_BMDC_DN, GSE17721_POLYIC_VS_PAM3CSK4_1H_BMDC_DN WP_EBOLA_VIRUS_PATHWAY_ON_HOST, WP_EBOLA_VIRUS_PATHWAY_ON_HOST
	GOBP_NEGATIVE_REGULATION_OF_CELL_PROJECTION_ORGANIZATION, GOBP_NEGATIVE_REGULATION_OF_CELL_PROJECTION_ORGANIZATION HP_FULL_CHEEKS, HP_FULL_CHEEKS REACTOME_SIGNALING_BY_VEGF, REACTOME_SIGNALING_BY_VEGF WP_16P112_PROXIMAL_DELETION_SYNDROME, WP_16P112_PROXIMAL_DELETION_SYNDROME GNF2_PTPRC, GNF2_PTPRC WP_PROLACTIN_SIGNALING_PATHWAY, WP_PROLACTIN_SIGNALING_PATHWAY TRAVAGLINI_LUNG_PLASMACYTOID_DENDRITIC_CELL, TRAVAGLINI_LUNG_PLASMACYTOID_DENDRITIC_CELL HP_AUTOIMMUNITY, HP_AUTOIMMUNITY
	GOMF_LYASE_ACTIVITY, GOMF_LYASE_ACTIVITY GOBP_NEGATIVE_REGULATION_OF_CELL_GROWTH, GOBP_NEGATIVE_REGULATION_OF_CELL_GROWTH MODULE_235, MODULE_235 DACOSTA_UV_RESPONSE_VIA_ERCC3_XPCS_DN, DACOSTA_UV_RESPONSE_VIA_ERCC3_XPCS_DN CSR_LATE_UP.V1_DN, CSR_LATE_UP.V1_DN GCM_IL6ST, GCM_IL6ST PID_VEGFR1_2_PATHWAY, PID_VEGFR1_2_PATHWAY
	HUMMERICH_SKIN_CANCER_PROGRESSION_UP, HUMMERICH_SKIN_CANCER_PROGRESSION_UP GOBP_ALCOHOL_BIOSYNTHETIC_PROCESS, GOBP_ALCOHOL_BIOSYNTHETIC_PROCESS GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS GOBP_NEGATIVE_REGULATION_OF_NEURON_PROJECTION_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_NEURON_PROJECTION_DEVELOPMENT WP_PANCREATIC_ADENOCARCINOMA_PATHWAY, WP_PANCREATIC_ADENOCARCINOMA_PATHWAY SARRIO_EPITHELIAL_MESENCHYMAL_TRANSITION_DN, SARRIO_EPITHELIAL_MESENCHYMAL_TRANSITION_DN HP_LEUKOCYTOSIS, HP_LEUKOCYTOSIS BIOCARTA_HIVNEF_PATHWAY, BIOCARTA_HIVNEF_PATHWAY
	MIR3166, MIR3166 HP_PURPURA, HP_PURPURA GOBP_STRIATED_MUSCLE_CONTRACTION, GOBP_STRIATED_MUSCLE_CONTRACTION GOBP_STRIATED_MUSCLE_CONTRACTION, GOBP_STRIATED_MUSCLE_CONTRACTION FAN_EMBRYONIC_CTX_BIG_GROUPS_GLIAL, FAN_EMBRYONIC_CTX_BIG_GROUPS_GLIAL GOBP_NEGATIVE_REGULATION_OF_CELL_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_CELL_DEVELOPMENT REACTOME_TRANSCRIPTIONAL_ACTIVATION_OF_MITOCHONDRIAL_BIOGENESIS, REACTOME_TRANSCRIPTIONAL_ACTIVATION_OF_MITOCHONDRIAL_BIOGENESIS WP_IL4_SIGNALING_PATHWAY, WP_IL4_SIGNALING_PATHWAY
	MAHAJAN_RESPONSE_TO_IL1A_DN, MAHAJAN_RESPONSE_TO_IL1A_DN MIR6833_5P, MIR6833_5P HP_NEOPLASM_OF_THE_EYE, HP_NEOPLASM_OF_THE_EYE MIR6839_5P, MIR6839_5P REACTOME_ONCOGENIC_MAPK_SIGNALING, REACTOME_ONCOGENIC_MAPK_SIGNALING MIR3977, MIR3977 SIRNA_EIF4GI_DN, SIRNA_EIF4GI_DN GSE3982_DC_VS_NKCELL_DN, GSE3982_DC_VS_NKCELL_DN
	KOKKINAKIS_METHIONINE_DEPRIVATION_48HR_DN, KOKKINAKIS_METHIONINE_DEPRIVATION_48HR_DN KOKKINAKIS_METHIONINE_DEPRIVATION_48HR_DN, KOKKINAKIS_METHIONINE_DEPRIVATION_48HR_DN KEGG_SNARE_INTERACTIONS_IN_VESICULAR_TRANSPORT, KEGG_SNARE_INTERACTIONS_IN_VESICULAR_TRANSPORT GOBP_INOSITOL_LIPID_MEDIATED_SIGNALING, GOBP_INOSITOL_LIPID_MEDIATED_SIGNALING KEGG_LEUKOCYTE_TRANSENDOTHELIAL_MIGRATION, KEGG_LEUKOCYTE_TRANSENDOTHELIAL_MIGRATION PID_FAK_PATHWAY, PID_FAK_PATHWAY GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_CYTOKINE_STIMULUS, GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_CYTOKINE_STIMULUS MIR2682_5P, MIR2682_5P
	MIR378J, MIR378J GOBP_PROTEIN_DESTABILIZATION, GOBP_PROTEIN_DESTABILIZATION NRL_DN.V1_DN, NRL_DN.V1_DN GOBP_LIPOPROTEIN_BIOSYNTHETIC_PROCESS, GOBP_LIPOPROTEIN_BIOSYNTHETIC_PROCESS HP_HEMOLYTIC_ANEMIA, HP_HEMOLYTIC_ANEMIA REACTOME_TNF_SIGNALING, REACTOME_TNF_SIGNALING PID_P75_NTR_PATHWAY, PID_P75_NTR_PATHWAY WP_PDGF_PATHWAY, WP_PDGF_PATHWAY
	HP_CHRONIC_OTITIS_MEDIA, HP_CHRONIC_OTITIS_MEDIA GOBP_ACTIN_FILAMENT_BASED_MOVEMENT, GOBP_ACTIN_FILAMENT_BASED_MOVEMENT MIR4662A_5P, MIR4662A_5P MIR4662A_5P, MIR4662A_5P HP_FIBROUS_TISSUE_NEOPLASM, HP_FIBROUS_TISSUE_NEOPLASM GOBP_CARDIAC_MUSCLE_CONTRACTION, GOBP_CARDIAC_MUSCLE_CONTRACTION WP_EXERCISEINDUCED_CIRCADIAN_REGULATION, WP_EXERCISEINDUCED_CIRCADIAN_REGULATION GOBP_NEGATIVE_REGULATION_OF_NERVOUS_SYSTEM_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_NERVOUS_SYSTEM_DEVELOPMENT
	GSE3691_IFN_PRODUCING_KILLER_DC_VS_CONVENTIONAL_DC_SPLEEN_DN, GSE3691_IFN_PRODUCING_KILLER_DC_VS_CONVENTIONAL_DC_SPLEEN_DN GOBP_REGULATION_OF_CARBOHYDRATE_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_CARBOHYDRATE_BIOSYNTHETIC_PROCESS GOBP_LAMELLIPODIUM_ORGANIZATION, GOBP_LAMELLIPODIUM_ORGANIZATION GOBP_POSITIVE_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING, GOBP_POSITIVE_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING MIR4529_5P, MIR4529_5P HP_HAMARTOMA_OF_THE_EYE, HP_HAMARTOMA_OF_THE_EYE REACTOME_PLATELET_HOMEOSTASIS, REACTOME_PLATELET_HOMEOSTASIS VANIHARANTA_LITERINE_EIRROUD_DN_VANIHARAN
	VANHARANTA_UTERINE_FIBROID_DN, VANHARANTA_UTERINE_FIBROID_DN GOBP_CYTOPLASMIC_MICROTUBULE_ORGANIZATION, GOBP_CYTOPLASMIC_MICROTUBULE_ORGANIZATION GOBP_POLYSACCHARIDE_METABOLIC_PROCESS, GOBP_POLYSACCHARIDE_METABOLIC_PROCESS WP_HUMAN_THYROID_STIMULATING_HORMONE_TSH_SIGNALING_PATHWAY, WP_HUMAN_THYROID_STIMULATING_HORMONE_TSH_SIGNALING_PATHWAY REACTOME_SIGNALING_BY_ERBB2, REACTOME_SIGNALING_BY_ERBB2 MIR4491, MIR4491 GOBP_NEGATIVE_REGULATION_OF_PROTEIN_POLYMERIZATION, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_POLYMERIZATION REACTOME_RHOU_GTPASE_CYCLE, REACTOME_RHOU_GTPASE_CYCLE
	BIOCARTA_KERATINOCYTE_PATHWAY, BIOCARTA_KERATINOCYTE_PATHWAY GOBP_EPHRIN_RECEPTOR_SIGNALING_PATHWAY, GOBP_EPHRIN_RECEPTOR_SIGNALING_PATHWAY MIR4657, MIR4657 HP_TINNITUS, HP_TINNITUS REACTOME_RAF_ACTIVATION, REACTOME_RAF_ACTIVATION GOCC_GOLGI_ASSOCIATED_VESICLE_MEMBRANE, GOCC_GOLGI_ASSOCIATED_VESICLE_MEMBRANE CCCACAT_MIR2993P, CCCACAT_MIR2993P HP_ABNORMALITY_OF_SUBCUTANEOUS_FAT_TISSUE, HP_ABNORMALITY_OF_SUBCUTANEOUS_FAT_TISSUE
	HP_ABNORMALITY_OF_SUBCUTANEOUS_FAT_TISSUE, HP_ABNORMALITY_OF_SUBCUTANEOUS_FAT_TISSUE GOCC_AP_TYPE_MEMBRANE_COAT_ADAPTOR_COMPLEX, GOCC_AP_TYPE_MEMBRANE_COAT_ADAPTOR_COMPLEX HP_NEOPLASM_OF_THE_ORAL_CAVITY, HP_NEOPLASM_OF_THE_ORAL_CAVITY DELASERNA_MYOD_TARGETS_UP, DELASERNA_MYOD_TARGETS_UP GNF2_JAK1, GNF2_JAK1 GOBP_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY GOBP_NEGATIVE_REGULATION_OF_ACTIN_FILAMENT_POLYMERIZATION, GOBP_NEGATIVE_REGULATION_OF_ACTIN_FILAMENT_POLYMERIZATION PID_AVB3_OPN_PATHWAY, PID_AVB3_OPN_PATHWAY
	HP_RECURRENT_SKIN_INFECTIONS, HP_RECURRENT_SKIN_INFECTIONS GOBP_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING, GOBP_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING GOBP_NON_MOTILE_CILIUM_ASSEMBLY, GOBP_NON_MOTILE_CILIUM_ASSEMBLY REACTOME_RHOV_GTPASE_CYCLE, REACTOME_RHOV_GTPASE_CYCLE GOMF_CARBON_OXYGEN_LYASE_ACTIVITY, GOMF_CARBON_OXYGEN_LYASE_ACTIVITY HP_HEMANGIOBLASTOMA, HP_HEMANGIOBLASTOMA KEGG_TIGHT_JUNCTION, KEGG_TIGHT_JUNCTION
	GAZDA_DIAMOND_BLACKFAN_ANEMIA_MYELOID_UP, GAZDA_DIAMOND_BLACKFAN_ANEMIA_MYELOID_UP WP_NETRINUNC5B_SIGNALING_PATHWAY, WP_NETRINUNC5B_SIGNALING_PATHWAY GOBP_REGULATION_OF_PROTEIN_MATURATION, GOBP_REGULATION_OF_PROTEIN_MATURATION PID_AMB2_NEUTROPHILS_PATHWAY, PID_AMB2_NEUTROPHILS_PATHWAY MIR6778_5P, MIR6778_5P GOBP_REGULATION_OF_ALCOHOL_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_ALCOHOL_BIOSYNTHETIC_PROCESS PID_IL2_1PATHWAY, PID_IL2_1PATHWAY HIR STOMATITIS_HIP_STOMATITIS
	HP_STOMATITIS, HP_STOMATITIS PID_IFNG_PATHWAY, PID_IFNG_PATHWAY BIOCARTA_RACCYCD_PATHWAY, BIOCARTA_RACCYCD_PATHWAY WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_INTERFERON_INDUCTION, WP_HOSTPATHOGEN_INTERACTION_OF_HUMAN_CORONA_VIRUSES_INTERFERON_INDUCTION GOBP_ACTIN_MEDIATED_CELL_CONTRACTION, GOBP_ACTIN_MEDIATED_CELL_CONTRACTION MIR632, MIR632 LEE_AGING_NEOCORTEX_UP, LEE_AGING_NEOCORTEX_UP GOBP_ACTIN_FILAMENT_DEPOLYMERIZATION, GOBP_ACTIN_FILAMENT_DEPOLYMERIZATION
	MIR3613_5P, MIR3613_5P WP_THYMIC_STROMAL_LYMPHOPOIETIN_TSLP_SIGNALING_PATHWAY, WP_THYMIC_STROMAL_LYMPHOPOIETIN_TSLP_SIGNALING_PATHWAY GOBP_CELLULAR_RESPONSE_TO_DRUG, GOBP_CELLULAR_RESPONSE_TO_DRUG PID_ECADHERIN_STABILIZATION_PATHWAY, PID_ECADHERIN_STABILIZATION_PATHWAY HP_LIPOATROPHY, HP_LIPOATROPHY BIOCARTA_EIF4_PATHWAY, BIOCARTA_EIF4_PATHWAY GOBP_REGULATION_OF_DENDRITE_MORPHOGENESIS, GOBP_REGULATION_OF_DENDRITE_MORPHOGENESIS
	WP_PHOSPHOINOSITIDES_METABOLISM, WP_PHOSPHOINOSITIDES_METABOLISM NUTT_GBM_VS_AO_GLIOMA_UP, NUTT_GBM_VS_AO_GLIOMA_UP WP_MICRORNAS_IN_CARDIOMYOCYTE_HYPERTROPHY, WP_MICRORNAS_IN_CARDIOMYOCYTE_HYPERTROPHY GOBP_CARDIAC_MUSCLE_CELL_CONTRACTION, GOBP_CARDIAC_MUSCLE_CELL_CONTRACTION HP_THYROID_CARCINOMA, HP_THYROID_CARCINOMA WOOD_EBV_EBNA1_TARGETS_DN, WOOD_EBV_EBNA1_TARGETS_DN GOBP_REGULATION_OF_ACTIN_FILAMENT_BASED_MOVEMENT, GOBP_REGULATION_OF_ACTIN_FILAMENT_BASED_MOVEMENT GOBP_P38MAPK_CASCADE, GOBP_P38MAPK_CASCADE
	GOBP_LAMELLIPODIUM_ASSEMBLY, GOBP_LAMELLIPODIUM_ASSEMBLY PID_THROMBIN_PAR1_PATHWAY, PID_THROMBIN_PAR1_PATHWAY LUI_THYROID_CANCER_CLUSTER_2, LUI_THYROID_CANCER_CLUSTER_2 GOBP_POLYSACCHARIDE_BIOSYNTHETIC_PROCESS, GOBP_POLYSACCHARIDE_BIOSYNTHETIC_PROCESS HP_NEOPLASM_OF_THE_TONGUE, HP_NEOPLASM_OF_THE_TONGUE PID_IGF1_PATHWAY, PID_IGF1_PATHWAY WP_GLYCOGEN_SYNTHESIS_AND_DEGRADATION, WP_GLYCOGEN_SYNTHESIS_AND_DEGRADATION
	HP_ASYMMETRIC_GROWTH, HP_ASYMMETRIC_GROWTH REACTOME_OTHER_INTERLEUKIN_SIGNALING, REACTOME_OTHER_INTERLEUKIN_SIGNALING MIR3165, MIR3165 HP_ABNORMAL_EOSINOPHIL_MORPHOLOGY, HP_ABNORMAL_EOSINOPHIL_MORPHOLOGY BAELDE_DIABETIC_NEPHROPATHY_UP, BAELDE_DIABETIC_NEPHROPATHY_UP WP_SIGNALING_OF_HEPATOCYTE_GROWTH_FACTOR_RECEPTOR, WP_SIGNALING_OF_HEPATOCYTE_GROWTH_FACTOR_RECEPTOR REACTOME_SIGNALING_BY_BRAF_AND_RAF_FUSIONS, REACTOME_SIGNALING_BY_BRAF_AND_RAF_FUSIONS HB_ABNORMAL_TIALABLE MB_CIGNAL_RETENCTIVE HB_ABNORMAL_TIALABLE MB_CIGNAL_RETENCTIVE HB_ABNORMAL_TIALABLE MB_CIGNAL_TIALABLE MB_CIGNAL_RETENCTIVE HB_ABNORMAL_TIALABLE MB
	HP_ABNORMAL_THALAMIC_MRI_SIGNAL_INTENSITY, HP_ABNORMAL_THALAMIC_MRI_SIGNAL_INTENSITY GOBP_REGULATION_OF_ESTABLISHMENT_OR_MAINTENANCE_OF_CELL_POLARITY, GOBP_REGULATION_OF_ESTABLISHMENT_OR_MAINTENANCE_OF_CELL_POLARITY REACTOME_METABOLISM_OF_COFACTORS, REACTOME_METABOLISM_OF_COFACTORS BIOCARTA_MET_PATHWAY, BIOCARTA_MET_PATHWAY BIOCARTA_CREB_PATHWAY, BIOCARTA_CREB_PATHWAY REACTOME_NEPHRIN_FAMILY_INTERACTIONS, REACTOME_NEPHRIN_FAMILY_INTERACTIONS PID_PTP1B_PATHWAY, PID_PTP1B_PATHWAY PID_ANGIOPOIETIN_RECEPTOR_PATHWAY, PID_ANGIOPOIETIN_RECEPTOR_PATHWAY
	GOBP_NEGATIVE_REGULATION_OF_CARBOHYDRATE_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_CARBOHYDRATE_METABOLIC_PROCESS ABBUD_LIF_SIGNALING_1_UP, ABBUD_LIF_SIGNALING_1_UP SOBOLEV_PBMC_PANDEMRIX_AGE_18_64YO_1DY_UP, SOBOLEV_PBMC_PANDEMRIX_AGE_18_64YO_1DY_UP HP_MYOCLONIC_SPASMS, HP_MYOCLONIC_SPASMS BIOCARTA_GCR_PATHWAY, BIOCARTA_GCR_PATHWAY HP_THYROIDITIS, HP_THYROIDITIS SATO_SILENCED_EPIGENETICALLY_IN_PANCREATIC_CANCER, SATO_SILENCED_EPIGENETICALLY_IN_PANCREATIC_CANCER
	PID_TXA2PATHWAY, PID_TXA2PATHWAY BIOCARTA_HCMV_PATHWAY WP_RESISTIN_AS_A_REGULATOR_OF_INFLAMMATION, WP_RESISTIN_AS_A_REGULATOR_OF_INFLAMMATION PID_ECADHERIN_KERATINOCYTE_PATHWAY, PID_ECADHERIN_KERATINOCYTE_PATHWAY MIR4715_5P, MIR4715_5P HP_ABNORMAL_CIRCULATING_PYRUVATE_FAMILY_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_PYRUVATE_FAMILY_AMINO_ACID_CONCENTRATION GOBP_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING, GOBP_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING, GOBP_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING, GOBP_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING, GOBP_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING, GOBP_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING.
	GOBP_PROTEIN_ADP_RIBOSYLATION, GOBP_PROTEIN_ADP_RIBOSYLATION HP_ABNORMALITY_OF_VITAMIN_METABOLISM, HP_ABNORMALITY_OF_VITAMIN_METABOLISM HP_CORNEAL_DYSTROPHY, HP_CORNEAL_DYSTROPHY WP_NONGENOMIC_ACTIONS_OF_125_DIHYDROXYVITAMIN_D3, WP_NONGENOMIC_ACTIONS_OF_125_DIHYDROXYVITAMIN_D3 WP_HEPATITIS_C_AND_HEPATOCELLULAR_CARCINOMA, WP_HEPATITIS_C_AND_HEPATOCELLULAR_CARCINOMA GOBP_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GOBP_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING HP_CELIAC_DISEASE, HP_CELIAC_DISEASE GOBP_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY, GOBP_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY
	GOMF_CLATHRIN_BINDING, GOMF_CLATHRIN_BINDING GOBP_REGULATION_OF_IMMUNOGLOBULIN_PRODUCTION, GOBP_REGULATION_OF_IMMUNOGLOBULIN_PRODUCTION PID_SYNDECAN_4_PATHWAY, PID_SYNDECAN_4_PATHWAY GOBP_ENDOPLASMIC_RETICULUM_CALCIUM_ION_HOMEOSTASIS, GOBP_ENDOPLASMIC_RETICULUM_CALCIUM_ION_HOMEOSTASIS GOMF_PROTEIN_PHOSPHATASE_2A_BINDING, GOMF_PROTEIN_PHOSPHATASE_2A_BINDING GOBP_REGULATION_OF_INSULIN_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_INSULIN_RECEPTOR_SIGNALING_PATHWAY MIR6732_5P, MIR6732_5P
	GOCC_MYELIN_SHEATH, GOCC_MYELIN_SHEATH GOBP_RESPONSE_TO_CADMIUM_ION, GOBP_RESPONSE_TO_CADMIUM_ION REACTOME_CASPASE_ACTIVATION_VIA_EXTRINSIC_APOPTOTIC_SIGNALLING_PATHWAY, REACTOME_CASPASE_ACTIVATION_VIA_EXTRINSIC_APOPTOTIC_SIGNALLING_PATHWAY GOBP_MEMBRANE_LIPID_CATABOLIC_PROCESS, GOBP_MEMBRANE_LIPID_CATABOLIC_PROCESS GOMF_MRNA_3_UTR_AU_RICH_REGION_BINDING, GOMF_MRNA_3_UTR_AU_RICH_REGION_BINDING REACTOME_SIGNALING_BY_PDGF, REACTOME_SIGNALING_BY_PDGF WP_ALPHA_6_BETA_4_SIGNALING_PATHWAY, WP_ALPHA_6_BETA_4_SIGNALING_PATHWAY GOMF_HYDRO_LYASE_ACTIVITY, GOMF_HYDRO_LYASE_ACTIVITY
	HP_ABNORMALITY_OF_TIBIA_MORPHOLOGY, HP_ABNORMALITY_OF_TIBIA_MORPHOLOGY HP_THICK_NASAL_ALAE, HP_THICK_NASAL_ALAE GOBP_REGULATION_OF_CARDIAC_MUSCLE_CONTRACTION, GOBP_REGULATION_OF_CARDIAC_MUSCLE_CONTRACTION HUNSBERGER_EXERCISE_REGULATED_GENES, HUNSBERGER_EXERCISE_REGULATED_GENES BIOCARTA_TALL1_PATHWAY, BIOCARTA_TALL1_PATHWAY GOMF_POTASSIUM_CHANNEL_REGULATOR_ACTIVITY, GOMF_POTASSIUM_CHANNEL_REGULATOR_ACTIVITY VALK_AML_CLUSTER_1, VALK_AML_CLUSTER_1
2_POS_VS_NEG_PDC_DN, GSE15215_CD2_POS_VS_NEG_PDC_DN	HP_BOWEL_DIVERTICULOSIS, HP_BOWEL_DIVERTICULOSIS GOCC_CLATHRIN_ADAPTOR_COMPLEX, GOCC_CLATHRIN_ADAPTOR_COMPLEX TTCCGTT_MIR191, TTCCGTT_MIR191 GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS, GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS GOBP_INTERLEUKIN_12_PRODUCTION, GOBP_INTERLEUKIN_12_PRODUCTION GOBP_MEMBRANE_RAFT_ORGANIZATION, GOBP_MEMBRANE_RAFT_ORGANIZATION PID_VEGFR1_PATHWAY, PID_VEGFR1_PATHWAY VALK_AML_CLUSTER_8, VALK_AML_CLUSTER_8
	HP_ORAL_CAVITY_BLEEDING, HP_ORAL_CAVITY_BLEEDING GOBP_NEGATIVE_REGULATION_OF_ACTIN_FILAMENT_DEPOLYMERIZATION, GOBP_NEGATIVE_REGULATION_OF_ACTIN_FILAMENT_DEPOLYMERIZATION PID_RET_PATHWAY, PID_RET_PATHWAY GOBP_POSITIVE_REGULATION_OF_CILIUM_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_CILIUM_ASSEMBLY HOSHIDA_LIVER_CANCER_LATE_RECURRENCE_DN, HOSHIDA_LIVER_CANCER_LATE_RECURRENCE_DN PID_EPHB_FWD_PATHWAY, PID_EPHB_FWD_PATHWAY PID_NETRIN_PATHWAY, PID_NETRIN_PATHWAY
	HP_CONICAL_TOOTH, HP_CONICAL_TOOTH MODULE_175, MODULE_175 MIR4654, MIR4654 GOBP_ENERGY_HOMEOSTASIS, GOBP_ENERGY_HOMEOSTASIS HP_OVARIAN_CYST, HP_OVARIAN_CYST MIR424_3P, MIR424_3P GOBP_UBIQUINONE_METABOLIC_PROCESS, GOBP_UBIQUINONE_METABOLIC_PROCESS SA_PTEN_PATHWAY, SA_PTEN_PATHWAY
	BIOCARTA_CXCR4_PATHWAY, BIOCARTA_CXCR4_PATHWAY MIR6777_3P, MIR6777_3P LI_PBMC_MENACTRA_AGE_18_45YO_CORRELATED_WITH_ANTI_DT_ANTIBODY_3DY_POSITIVE, LI_PBMC_MENACTRA_AGE_18_45YO_CORRELATED_WITH_ANTI_DT_ANTIBODY_3DY_POSITIVE REACTOME_SIGNALING_BY_WNT_IN_CANCER, REACTOME_SIGNALING_BY_WNT_IN_CANCER WP_APOPTOSIS_MODULATION_BY_HSP70, WP_APOPTOSIS_MODULATION_BY_HSP70 ABE_VEGFA_TARGETS_2HR, ABE_VEGFA_TARGETS_2HR HP_ABNORMALLY_LARGE_GLOBE, HP_ABNORMALLY_LARGE_GLOBE
	HP_LARGE_FOREHEAD, HP_LARGE_FOREHEAD PID_S1P_S1P2_PATHWAY, PID_S1P_S1P2_PATHWAY BIOCARTA_EDG1_PATHWAY, BIOCARTA_EDG1_PATHWAY HP_HYDROCELE_TESTIS, HP_HYDROCELE_TESTIS REACTOME_SIGNALING_BY_PDGFR_IN_DISEASE, REACTOME_SIGNALING_BY_PDGFR_IN_DISEASE GOBP_POSITIVE_REGULATION_OF_EXOCYTOSIS, GOBP_POSITIVE_REGULATION_OF_EXOCYTOSIS WP_TOLLLIKE_RECEPTOR_SIGNALING_RELATED_TO_MYD88, WP_TOLLLIKE_RECEPTOR_SIGNALING_RELATED_TO_MYD88, WP_TOLLLIKE_RECEPTOR_SIGNALING_RELATED_TO_MYD88 HP_BONE_MARROW_HYPERCELLULARITY, HP_BONE_MARROW_HYPERCELLULARITY
	GOMF_PHOSPHATASE_ACTIVATOR_ACTIVITY, GOMF_PHOSPHATASE_ACTIVATOR_ACTIVITY HP_HAMARTOMATOUS_POLYPOSIS, HP_HAMARTOMATOUS_POLYPOSIS HP_MODERATE_GLOBAL_DEVELOPMENTAL_DELAY, HP_MODERATE_GLOBAL_DEVELOPMENTAL_DELAY SIMBULAN_UV_RESPONSE_NORMAL_DN, SIMBULAN_UV_RESPONSE_NORMAL_DN PID_NECTIN_PATHWAY, PID_NECTIN_PATHWAY ZHONG_PFC_MAJOR_TYPES_OPC, ZHONG_PFC_MAJOR_TYPES_OPC BIOCARTA_TOLL_PATHWAY, BIOCARTA_TOLL_PATHWAY
	HP_DISSEMINATED_INTRAVASCULAR_COAGULATION, HP_DISSEMINATED_INTRAVASCULAR_COAGULATION REACTOME_DOWNSTREAM_SIGNAL_TRANSDUCTION, REACTOME_DOWNSTREAM_SIGNAL_TRANSDUCTION YAN_ESCAPE_FROM_ANOIKIS, YAN_ESCAPE_FROM_ANOIKIS GOBP_QUINONE_METABOLIC_PROCESS, GOBP_QUINONE_METABOLIC_PROCESS BIOCARTA_PAR1_PATHWAY, BIOCARTA_PAR1_PATHWAY GCGCTTT_MIR518B_MIR518C_MIR518D, GCGCTTT_MIR518B_MIR518C_MIR518D HP_APHTHOUS_ULCER, HP_APHTHOUS_ULCER
	WP_ALZHEIMERS_DISEASE, WP_ALZHEIMERS_DISEASE HP_BONE_CYST, HP_BONE_CYST GOBP_NEGATIVE_REGULATION_OF_ERK1_AND_ERK2_CASCADE, GOBP_NEGATIVE_REGULATION_OF_ERK1_AND_ERK2_CASCADE HP_ABNORMALITY_OF_VITAMIN_B_METABOLISM, HP_ABNORMALITY_OF_VITAMIN_B_METABOLISM GOBP_POSITIVE_REGULATION_OF_PROTEIN_MATURATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_MATURATION GOBP_REGULATION_OF_MITOCHONDRIAL_FUSION, GOBP_REGULATION_OF_MITOCHONDRIAL_FUSION HP_COLONIC_DIVERTICULA, HP_COLONIC_DIVERTICULA HP_COLONIC_AL_INCISOR_HP_CONICAL_INCISOR
	HP_CONICAL_INCISOR, HP_CONICAL_INCISOR WP_PHYSIOLOGICAL_AND_PATHOLOGICAL_HYPERTROPHY_OF_THE_HEART, WP_PHYSIOLOGICAL_AND_PATHOLOGICAL_HYPERTROPHY_OF_THE_HEART AMIT_DELAYED_EARLY_GENES, AMIT_DELAYED_EARLY_GENES PID_SYNDECAN_2_PATHWAY, PID_SYNDECAN_2_PATHWAY REACTOME_SYNAPTIC_ADHESION_LIKE_MOLECULES, REACTOME_SYNAPTIC_ADHESION_LIKE_MOLECULES GOBP_RESPONSE_TO_ZINC_ION, GOBP_RESPONSE_TO_ZINC_ION WP_NEURAL_CREST_CELL_MIGRATION_IN_CANCER, WP_NEURAL_CREST_CELL_MIGRATION_IN_CANCER HP_ABNORMAL_VITAMIN_B12_LEVEL, HP_ABNORMAL_VITAMIN_B12_LEVEL
	REACTOME_SIGNALING_BY_KIT_IN_DISEASE, REACTOME_SIGNALING_BY_KIT_IN_DISEASE GOBP_MEMBRANE_PROTEIN_INTRACELLULAR_DOMAIN_PROTEOLYSIS, GOBP_MEMBRANE_PROTEIN_INTRACELLULAR_DOMAIN_PROTEOLYSIS GOBP_NEGATIVE_REGULATION_OF_SMALL_MOLECULE_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_SMALL_MOLECULE_METABOLIC_PROCESS GOBP_POSITIVE_REGULATION_OF_TELOMERE_CAPPING, GOBP_POSITIVE_REGULATION_OF_TELOMERE_CAPPING GOBP_PUTRESCINE_METABOLIC_PROCESS, GOBP_PUTRESCINE_METABOLIC_PROCESS REACTOME_SEMA4D_IN_SEMAPHORIN_SIGNALING, REACTOME_SEMA4D_IN_SEMAPHORIN_SIGNALING HP_FOLLICULAR_THYROID_CARCINOMA, HP_FOLLICULAR_THYROID_CARCINOMA
	WP_ANGIOGENESIS, WP_ANGIOGENESIS HP_REDUNDANT_NECK_SKIN, HP_REDUNDANT_NECK_SKIN GOBP_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS PODAR_RESPONSE_TO_ADAPHOSTIN_DN, PODAR_RESPONSE_TO_ADAPHOSTIN_DN HP_DIFFUSE_ALVEOLAR_HEMORRHAGE, HP_DIFFUSE_ALVEOLAR_HEMORRHAGE BIOCARTA_TFF_PATHWAY, BIOCARTA_TFF_PATHWAY GOBP_REGULATION_OF_HORMONE_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_HORMONE_BIOSYNTHETIC_PROCESS HR_HEMANCIOMATORIS_HP_HEMANCIOMATORIS_
	HP_HEMANGIOMATOSIS, HP_HEMANGIOMATOSIS REACTOME_RHO_GTPASES_ACTIVATE_RHOTEKIN_AND_RHOPHILINS, REACTOME_RHO_GTPASES_ACTIVATE_RHOTEKIN_AND_RHOPHILINS GOBP_MEMBRANE_BIOGENESIS, GOBP_MEMBRANE_BIOGENESIS HP_ABNORMAL_NUMBER_OF_INCISORS, HP_ABNORMAL_NUMBER_OF_INCISORS GOMF_CALCIUM_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GOMF_CALCIUM_DEPENDENT_PROTEIN_KINASE_ACTIVITY GOBP_LOCOMOTOR_RHYTHM, GOBP_LOCOMOTOR_RHYTHM OVSYANNIKOVA_PBMC_FLUARIX_AGE_50_74YO_COMMON_WITH_BOTH_HAI_AND_VNA_28DY_VS_3DY_USED_IN_HAI_AND_VNA_RESPONSE_MODELS_UP, OVSYANNIKOVA_PBMC_FLUARIX_AGE_50_74YO_COMMON_WITH_BOTH_HAI_AND_VNA_28DY_VS_3DY_USED_IN_HAI_A MIR6841_5P, MIR6841_5P
	GOBP_KETONE_BIOSYNTHETIC_PROCESS, GOBP_KETONE_BIOSYNTHETIC_PROCESS WP_NEURAL_CREST_CELL_MIGRATION_DURING_DEVELOPMENT, WP_NEURAL_CREST_CELL_MIGRATION_DURING_DEVELOPMENT GOBP_NEGATIVE_REGULATION_OF_HORMONE_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_HORMONE_METABOLIC_PROCESS GOBP_FAT_SOLUBLE_VITAMIN_BIOSYNTHETIC_PROCESS, GOBP_FAT_SOLUBLE_VITAMIN_BIOSYNTHETIC_PROCESS GOCC_PROTEIN_LIPID_COMPLEX, GOCC_PROTEIN_LIPID_COMPLEX HP_MEGALENCEPHALY, HP_MEGALENCEPHALY HP_CHRONIC_PULMONARY_OBSTRUCTION, HP_CHRONIC_PULMONARY_OBSTRUCTION
	HP_NEUTROPENIA_IN_PRESENCE_OF_ANTI_NEUTROPIL_ANTIBODIES, HP_NEUTROPENIA_IN_PRESENCE_OF_ANTI_NEUTROPIL_ANTIBODIES HP_CUTANEOUS_ABSCESS, HP_CUTANEOUS_ABSCESS GOBP_INTERFERON_ALPHA_PRODUCTION, GOBP_INTERFERON_ALPHA_PRODUCTION DAVICIONI_TARGETS_OF_PAX_FOXO1_FUSIONS_DN, DAVICIONI_TARGETS_OF_PAX_FOXO1_FUSIONS_DN HP_SOMATIC_MOSAICISM, HP_SOMATIC_MOSAICISM GOBP_PROTEIN_AUTO_ADP_RIBOSYLATION, GOBP_PROTEIN_AUTO_ADP_RIBOSYLATION GOBP_GOLGI_TO_LYSOSOME_TRANSPORT, GOBP_GOLGI_TO_LYSOSOME_TRANSPORT HP_HYPERTHYROIDISM, HP_HYPERTHYROIDISM
	BIOCARTA_AKAPCENTROSOME_PATHWAY, BIOCARTA_AKAPCENTROSOME_PATHWAY GOCC_ENDOLYSOSOME, GOCC_ENDOLYSOSOME REACTOME_SEMA4D_INDUCED_CELL_MIGRATION_AND_GROWTH_CONE_COLLAPSE, REACTOME_SEMA4D_INDUCED_CELL_MIGRATION_AND_GROWTH_CONE_COLLAPSE GOBP_ANIMAL_ORGAN_MATURATION, GOBP_ANIMAL_ORGAN_MATURATION GOBP_POSITIVE_REGULATION_OF_INTERFERON_ALPHA_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERFERON_ALPHA_PRODUCTION HP_METATARSUS_VALGUS, HP_METATARSUS_VALGUS HP_RIB_FUSION, HP_RIB_FUSION
	GOBP_INTERLEUKIN_6_MEDIATED_SIGNALING_PATHWAY, GOBP_INTERLEUKIN_6_MEDIATED_SIGNALING_PATHWAY GOBP_VITAMIN_D_BIOSYNTHETIC_PROCESS, GOBP_VITAMIN_D_BIOSYNTHETIC_PROCESS REACTOME_INTERLEUKIN_9_SIGNALING, REACTOME_INTERLEUKIN_9_SIGNALING HP_ABNORMALITY_OF_FACIAL_ADIPOSE_TISSUE, HP_ABNORMALITY_OF_FACIAL_ADIPOSE_TISSUE WP_RAS_AND_BRADYKININ_PATHWAYS_IN_COVID19, WP_RAS_AND_BRADYKININ_PATHWAYS_IN_COVID19 PID_EPHA2_FWD_PATHWAY, PID_EPHA2_FWD_PATHWAY GOBP_POSITIVE_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS
	GOBP_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS, GOBP_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_INSULIN_STIMULUS REACTOME_PTK6_REGULATES_RHO_CTPASES_RAS_CTPASE_AND_MAP_KINASES, REACTOME_PTK6_REGULATES_RHO_CTPASES_RAS_CTPASE_AND_MAP_KINASES GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_12_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_12_PRODUCTION GOBP_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOBP_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY WP_SARSCOV2_MITOCHONDRIAL_INTERACTIONS, WP_SARSCOV2_MITOCHONDRIAL_INTERACTIONS GOBP_BONE_MATURATION, GOBP_BONE_MATURATION HP_SEVERE_INFECTION, HP_SEVERE_INFECTION GOMF_PROTEIN_PHOSPHATASE_ACTIVATOR_ACTIVITY, GOMF_PROTEIN_PHOSPHATASE_ACTIVATOR_ACTIVITY
	GOMF_UBIQUITINATION_LIKE_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GOMF_UBIQUITINATION_LIKE_MODIFICATION_DEPENDENT_PROTEIN_BINDING GOBP_PODOSOME_ASSEMBLY, GOBP_PODOSOME_ASSEMBLY GOBP_NEGATIVE_REGULATION_OF_ALCOHOL_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_ALCOHOL_BIOSYNTHETIC_PROCESS REACTOME_PLATELET_SENSITIZATION_BY_LDL, REACTOME_PLATELET_SENSITIZATION_BY_LDL GOBP_REGULATION_OF_PEPTIDYL_SERINE_PHOSPHORYLATION, GOBP_REGULATION_OF_PEPTIDYL_SERINE_PHOSPHORYLATION GOBP_POSTSYNAPTIC_SPECIALIZATION_ORGANIZATION, GOBP_POSTSYNAPTIC_SPECIALIZATION_ORGANIZATION GOBP_CELLULAR_RESPONSE_TO_VIRUS, GOBP_CELLULAR_RESPONSE_TO_VIRUS
	LEE_LIVER_CANCER_MYC_DN, LEE_LIVER_CANCER_MYC_DN STARK_HYPPOCAMPUS_22Q11_DELETION_DN, STARK_HYPPOCAMPUS_22Q11_DELETION_DN HP_ABNORMAL_NATURAL_KILLER_CELL_PHYSIOLOGY, HP_ABNORMAL_NATURAL_KILLER_CELL_PHYSIOLOGY GOBP_PHAGOSOME_LYSOSOME_FUSION, GOBP_PHAGOSOME_LYSOSOME_FUSION GOBP_SKELETAL_MUSCLE_CONTRACTION, GOBP_SKELETAL_MUSCLE_CONTRACTION WP_RIBOFLAVIN_AND_COQ_DISORDERS, WP_RIBOFLAVIN_AND_COQ_DISORDERS MIR4433B_3P, MIR4433B_3P HP_HODGKIN_LYMPHOMA, HP_HODGKIN_LYMPHOMA
	REACTOME_INTERLEUKIN_21_SIGNALING, REACTOME_INTERLEUKIN_21_SIGNALING BIOCARTA_DSP_PATHWAY, BIOCARTA_DSP_PATHWAY, GOBP_POSITIVE_REGULATION_OF_IMMUNOGLOBULIN_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_IMMUNOGLOBULIN_PRODUCTION GOBP_REGULATION_OF_SYNAPTIC_VESICLE_CLUSTERING, GOBP_REGULATION_OF_SYNAPTIC_VESICLE_CLUSTERING REACTOME_NRCAM_INTERACTIONS, REACTOME_NRCAM_INTERACTIONS REACTOME_SIGNALING_BY_PDGFRA_TRANSMEMBRANE_JUXTAMEMBRANE_AND_KINASE_DOMAIN_MUTANTS, REACTOME_SIGNALING_BY_PDGFRA_TRANSMEMBRANE_JUXTAMEMBRANE_AND_KINASE_DOMAIN_MUTANTS WP_CELL_MIGRATION_AND_INVASION_THROUGH_P75NTR, WP_CELL_MIGRATION_AND_INVASION_THROUGH_P75NTR
	HP_NARROW_PELVIS_BONE, HP_NARROW_PELVIS_BONE BIOCARTA_HER2_PATHWAY, BIOCARTA_HER2_PATHWAY HP_TRACHEAL_STENOSIS, HP_TRACHEAL_STENOSIS GOBP_CELLULAR_RESPONSE_TO_DSRNA, GOBP_CELLULAR_RESPONSE_TO_DSRNA WP_MBDNF_AND_PROBDNF_REGULATION_OF_GABA_NEUROTRANSMISSION, WP_MBDNF_AND_PROBDNF_REGULATION_OF_GABA_NEUROTRANSMISSION SA_TRKA_RECEPTOR, SA_TRKA_RECEPTOR GOBP_RESPONSE_TO_INTERLEUKIN_9, GOBP_RESPONSE_TO_INTERLEUKIN_9 WP_COMPOSITION_OF_LIPID_PARTICLES, WP_COMPOSITION_OF_LIPID_PARTICLES
	WF_COMPOSITION_OF_LIFID_FARTICLES, WF_COMPOSITION_OF_LIFID_FARTICLES HP_RECURRENT_BRONCHITIS, HP_RECURRENT_BRONCHITIS SCHWAB_TARGETS_OF_BMYB_POLYMORPHIC_VARIANTS_DN, SCHWAB_TARGETS_OF_BMYB_POLYMORPHIC_VARIANTS_DN HP_ABNORMAL_AUDITORY_EVOKED_POTENTIALS, HP_ABNORMAL_AUDITORY_EVOKED_POTENTIALS REACTOME_SIGNALING_BY_CTNNB1_PHOSPHO_SITE_MUTANTS, REACTOME_SIGNALING_BY_CTNNB1_PHOSPHO_SITE_MUTANTS WP_FIBRIN_COMPLEMENT_RECEPTOR_3_SIGNALING_PATHWAY, WP_FIBRIN_COMPLEMENT_RECEPTOR_3_SIGNALING_PATHWAY GOBP_GLIAL_CELL_APOPTOTIC_PROCESS, GOBP_GLIAL_CELL_APOPTOTIC_PROCESS BIOCARTA_IL22BP_PATHWAY, BIOCARTA_IL22BP_PATHWAY
	REACTOME_DEFECTIVE_RIPK1_MEDIATED_REGULATED_NECROSIS, REACTOME_DEFECTIVE_RIPK1_MEDIATED_REGULATED_NECROSIS HP_LOSS_OF_SUBCUTANEOUS_ADIPOSE_TISSUE_IN_LIMBS, HP_LOSS_OF_SUBCUTANEOUS_ADIPOSE_TISSUE_IN_LIMBS HP_MONOCYTOSIS, HP_MONOCYTOSIS GOBP_RESPONSE_TO_ANGIOTENSIN, GOBP_RESPONSE_TO_ANGIOTENSIN HP_ABNORMALITY_OF_COMPLEMENT_SYSTEM, HP_ABNORMALITY_OF_COMPLEMENT_SYSTEM LI_PBMC_MENOMUNE_A_C_Y_W_135_AGE_18_45YO_CORRELATED_WITH_ANTIBODY_RESPONSE_3DY_NEGATIVE, LI_PBMC_MENOMUNE_A_C_Y_W_135_AGE_18_45YO_CORRELATED_WITH_ANTIBODY_RESPONSE_3DY_NEGATIVE HP_THROMBOPHLEBITIS, HP_THROMBOPHLEBITIS
	GOBP_SPHINGOMYELIN_METABOLIC_PROCESS, GOBP_SPHINGOMYELIN_METABOLIC_PROCESS MODULE_202, MODULE_202 IKEDA_MIR30_TARGETS_DN, IKEDA_MIR30_TARGETS_DN REACTOME_VITAMINS, REACTOME_VITAMINS GOCC_GOLGI_TRANS_CISTERNA, GOCC_GOLGI_TRANS_CISTERNA HP_CENTRAL_HETEROCHROMIA, HP_CENTRAL_HETEROCHROMIA GOMF_UBIQUITIN_DEPENDENT_PROTEIN_BINDING, GOMF_UBIQUITIN_DEPENDENT_PROTEIN_BINDING HP_PLANTAR_HYPERKERATOSIS, HP_PLANTAR_HYPERKERATOSIS
	HP_PLANTAR_HYPERKERATOSIS, HP_PLANTAR_HYPERKERATOSIS GOBP_PHOSPHATIDYLINOSITOL_3_PHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_PHOSPHATIDYLINOSITOL_3_PHOSPHATE_BIOSYNTHETIC_PROCESS GOBP_FAT_SOLUBLE_VITAMIN_METABOLIC_PROCESS, GOBP_FAT_SOLUBLE_VITAMIN_METABOLIC_PROCESS HP_ENLARGED_OVARIES, HP_ENLARGED_OVARIES GOMF_PROTEIN_KINASE_C_ACTIVITY, GOMF_PROTEIN_KINASE_C_ACTIVITY YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_3, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_3 GOBP_REGULATION_OF_MYD88_INDEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_MYD88_INDEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY GOMF_DEAD_H_BOX_RNA_HELICASE_BINDING, GOMF_DEAD_H_BOX_RNA_HELICASE_BINDING
	HP_ENTEROCOLITIS, HP_ENTEROCOLITIS GOBP_POSITIVE_REGULATION_OF_POTASSIUM_ION_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_POTASSIUM_ION_TRANSPORT GOBP_REGULATION_OF_POTASSIUM_ION_TRANSPORT, GOBP_REGULATION_OF_POTASSIUM_ION_TRANSPORT GOBP_FOAM_CELL_DIFFERENTIATION, GOBP_FOAM_CELL_DIFFERENTIATION GOBP_RESPONSE_TO_IMMOBILIZATION_STRESS, GOBP_RESPONSE_TO_IMMOBILIZATION_STRESS BIOCARTA_CARDIACEGF_PATHWAY, BIOCARTA_CARDIACEGF_PATHWAY GOBP_EPIBOLY, GOBP_EPIBOLY
	GOBP_REGULATION_OF_VITAMIN_METABOLIC_PROCESS, GOBP_REGULATION_OF_VITAMIN_METABOLIC_PROCESS GOBP_NEGATIVE_REGULATION_OF_MEMBRANE_DEPOLARIZATION, GOBP_NEGATIVE_REGULATION_OF_MEMBRANE_DEPOLARIZATION BIOCARTA_EEA1_PATHWAY, BIOCARTA_EEA1_PATHWAY BIOCARTA_TSP1_PATHWAY, BIOCARTA_TSP1_PATHWAY HP_MEMBRANOPROLIFERATIVE_GLOMERULONEPHRITIS, HP_MEMBRANOPROLIFERATIVE_GLOMERULONEPHRITIS HP_OPTIC_NERVE_COMPRESSION, HP_OPTIC_NERVE_COMPRESSION GOBP_RESPONSE_TO_GONADOTROPIN_RELEASING_HORMONE, GOBP_RESPONSE_TO_GONADOTROPIN_RELEASING_HORMONE HP_FIBULAR_BOWING, HP_FIBULAR_BOWING
	HP_PYODERMA_GANGRENOSUM, HP_PYODERMA_GANGRENOSUM GOBP_NOTCH_RECEPTOR_PROCESSING_LIGAND_DEPENDENT, GOBP_NOTCH_RECEPTOR_PROCESSING_LIGAND_DEPENDENT GOCC_GAMMA_SECRETASE_COMPLEX, GOCC_GAMMA_SECRETASE_COMPLEX GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_DERIVED_FOAM_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_DERIVED_FOAM_CELL_DIFFERENTIATION GOCC_PROTEIN_PHOSPHATASE_TYPE_2A_COMPLEX, GOCC_PROTEIN_PHOSPHATASE_TYPE_2A_COMPLEX GOBP_REGULATION_OF_3_UTR_MEDIATED_MRNA_STABILIZATION, GOBP_REGULATION_OF_3_UTR_MEDIATED_MRNA_STABILIZATION HP_TETANY, HP_TETANY
	MIR4769_5P, MIR4769_5P HP_ANGIOID_STREAKS_OF_THE_FUNDUS, HP_ANGIOID_STREAKS_OF_THE_FUNDUS GOBP_RESPONSE_TO_LIPOTEICHOIC_ACID, GOBP_RESPONSE_TO_LIPOTEICHOIC_ACID HP_VISCERAL_ANGIOMATOSIS, HP_VISCERAL_ANGIOMATOSIS GOBP_REGULATION_OF_SUBSTRATE_ADHESION_DEPENDENT_CELL_SPREADING, GOBP_REGULATION_OF_SUBSTRATE_ADHESION_DEPENDENT_CELL_SPREADING LUND_SILENCED_BY_METHYLATION, LUND_SILENCED_BY_METHYLATION GOBP_REGULATION_OF_LIPOPROTEIN_METABOLIC_PROCESS, GOBP_REGULATION_OF_LIPOPROTEIN_METABOLIC_PROCESS
	LANDEMAINE_LUNG_METASTASIS, LANDEMAINE_LUNG_METASTASIS GOBP_POSITIVE_REGULATION_OF_INTEGRIN_ACTIVATION, GOBP_POSITIVE_REGULATION_OF_INTEGRIN_ACTIVATION REACTOME_SIGNALING_BY_LEPTIN, REACTOME_SIGNALING_BY_LEPTIN REACTOME_REGULATED_PROTEOLYSIS_OF_P75NTR, REACTOME_REGULATED_PROTEOLYSIS_OF_P75NTR HP_AGENESIS_OF_LATERAL_INCISOR, HP_AGENESIS_OF_LATERAL_INCISOR HP_THORACIC_DYSPLASIA, HP_THORACIC_DYSPLASIA GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_DEGRANULATION, GOBP_POSITIVE_REGULATION_OF_LEUKOCYTE_DEGRANULATION WP_SIMPLIFIED_DEPICTION_OF_MYD88_DISTINCT_INPUTOUTPUT_PATHWAY, WP_SIMPLIFIED_DEPICTION_OF_MYD88_DISTINCT_INPUTOUTPUT_PATHWAY
	WP_SIMPLIFIED_DEPICTION_OF_MYD88_DISTINCT_INPUTOUTPUT_PATHWAY, WP_SIMPLIFIED_DEPICTION_OF_MYD88_DISTINCT_INPUTOUTPUT_PATHWAY GOBP_POSITIVE_REGULATION_OF_BONE_RESORPTION, GOBP_POSITIVE_REGULATION_OF_BONE_RESORPTION REACTOME_RHO_GTPASES_ACTIVATE_NADPH_OXIDASES, REACTOME_RHO_GTPASES_ACTIVATE_NADPH_OXIDASES