	GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_TREATED_MACROPHAGE_DN
	GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_UP, GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_UP GSE31082_DN_VS_DP_THYMOCYTE_UP, GSE31082_DN_VS_DP_THYMOCYTE_UP GSE24574_NAIVE_VS_TCONV_CD4_TCELL_DN, GSE24574_NAIVE_VS_TCONV_CD4_TCELL_DN MOREAUX_MULTIPLE_MYELOMA_BY_TACI_DN, MOREAUX_MULTIPLE_MYELOMA_BY_TACI_DN GSE24574_BCL6_HIGH_TFH_VS_NAIVE_CD4_TCELL_DN, GSE24574_BCL6_HIGH_TFH_VS_NAIVE_CD4_TCELL_DN GSE23114_WT_VS_SLE2C1_MOUSE_SPLEEN_B1A_BCELL_UP, GSE23114_WT_VS_SLE2C1_MOUSE_SPLEEN_B1A_BCELL_UP
	GSE14000_TRANSLATED_RNA_VS_MRNA_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_DC_UP GSE28726_NAIVE_VS_ACTIVATED_VA24NEG_NKTCELL_UP, GSE28726_NAIVE_VS_ACTIVATED_VA24NEG_NKTCELL_UP GSE411_100MIN_VS_400MIN_IL6_STIM_MACROPHAGE_DN, GSE411_100MIN_VS_400MIN_IL6_STIM_MACROPHAGE_DN GSE46468_LUNG_INNATE_LYMPHOID_CELL_VS_SPLEEN_CD4_TCELL_DN, GSE46468_LUNG_INNATE_LYMPHOID_CELL_VS_SPLEEN_CD4_TCELL_DN GSE28726_NAIVE_VS_ACTIVATED_NKTCELL_DN, GSE28726_NAIVE_VS_ACTIVATED_NKTCELL_DN GOBP_TRNA_METABOLIC_PROCESS, GOBP_TRNA_METABOLIC_PROCESS
	GSE6674_CPG_VS_PL2_3_STIM_BCELL_DN, GSE6674_CPG_VS_PL2_3_STIM_BCELL_DN GSE3982_MAC_VS_NEUTROPHIL_UP, GSE3982_MAC_VS_NEUTROPHIL_UP GSE15330_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_VS_PRO_BCELL_DN, GSE15330_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_VS_PRO_BCELL_DN GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_DN, GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_DN GSE41867_DAY6_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN, GSE41867_DAY6_EFFECTOR_VS_DAY30_EXHAUSTED_CD8_TCELL_LCMV_CLONE13_DN GSE6674_ANTI_IGM_VS_ANTI_IGM_AND_CPG_STIM_BCELL_DN, GSE6674_ANTI_IGM_VS_ANTI_IGM_AND_CPG_STIM_BCELL_DN
	GSE46606_UNSTIM_VS_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_DN, GSE46606_UNSTIM_VS_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_DN GSE37301_LYMPHOID_PRIMED_MPP_VS_GRAN_MONO_PROGENITOR_UP, GSE37301_LYMPHOID_PRIMED_MPP_VS_GRAN_MONO_PROGENITOR_UP GSE17721_LPS_VS_POLYIC_24H_BMDC_UP, GSE17721_LPS_VS_POLYIC_24H_BMDC_UP GSE45739_NRAS_KO_VS_WT_ACD3_ACD28_STIM_CD4_TCELL_UP, GSE45739_NRAS_KO_VS_WT_ACD3_ACD28_STIM_CD4_TCELL_UP GSE14415_TCONV_VS_FOXP3_KO_INDUCED_TREG_UP, GSE14415_TCONV_VS_FOXP3_KO_INDUCED_TREG_UP GSE32986_CURDLAN_LOWDOSE_VS_GMCSF_AND_CURDLAN_LOWDOSE_STIM_DC_DN, GSE32986_CURDLAN_LOWDOSE_VS_GMCSF_AND_CURDLAN_LOWDOSE_STIM_DC_DN
	GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_DN, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_DN GOLDRATH_HOMEOSTATIC_PROLIFERATION, GOLDRATH_HOMEOSTATIC_PROLIFERATION GSE7400_CTRL_VS_CSF3_IN_VIVO_TREATED_PBMC_UP, GSE7400_CTRL_VS_CSF3_IN_VIVO_TREATED_PBMC_UP REACTOME_HOST_INTERACTIONS_OF_HIV_FACTORS, REACTOME_HOST_INTERACTIONS_OF_HIV_FACTORS GSE27786_CD4_TCELL_VS_MONO_MAC_UP, GSE27786_CD4_TCELL_VS_MONO_MAC_UP GSE43863_TH1_VS_TFH_MEMORY_CD4_TCELL_UP, GSE43863_TH1_VS_TFH_MEMORY_CD4_TCELL_UP
	GSE3982_NEUTROPHIL_VS_BCELL_DN, GSE3982_NEUTROPHIL_VS_BCELL_DN GSE22886_NAIVE_CD4_TCELL_VS_NEUTROPHIL_UP, GSE22886_NAIVE_CD4_TCELL_VS_NEUTROPHIL_UP GSE3982_NEUTROPHIL_VS_EFF_MEMORY_CD4_TCELL_DN, GSE3982_NEUTROPHIL_VS_EFF_MEMORY_CD4_TCELL_DN GSE37301_LYMPHOID_PRIMED_MPP_VS_RAG2_KO_NK_CELL_UP, GSE37301_LYMPHOID_PRIMED_MPP_VS_RAG2_KO_NK_CELL_UP GSE17721_LPS_VS_GARDIQUIMOD_8H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_8H_BMDC_DN GSE17721_POLYIC_VS_CPG_12H_BMDC_DN, GSE17721_POLYIC_VS_CPG_12H_BMDC_DN
	GSE43863_NAIVE_VS_MEMORY_TFH_CD4_TCELL_D150_LCMV_DN, GSE43863_NAIVE_VS_MEMORY_TFH_CD4_TCELL_D150_LCMV_DN ZHENG_CORD_BLOOD_C9_GRANULOCYTE_MACROPHAGE_PROGENITOR, ZHENG_CORD_BLOOD_C9_GRANULOCYTE_MACROPHAGE_PROGENITOR GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_TRNA, GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_TRNA GSE17721_POLYIC_VS_GARDIQUIMOD_12H_BMDC_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_12H_BMDC_DN MORF_EIF4E, MORF_EIF4E GSE27786_LSK_VS_NKTCELL_UP, GSE27786_LSK_VS_NKTCELL_UP
	GSE339_CD8POS_VS_CD4CD8DN_DC_IN_CULTURE_DN, GSE339_CD8POS_VS_CD4CD8DN_DC_IN_CULTURE_DN GSE40274_CTRL_VS_FOXP3_AND_SATB1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_CTRL_VS_FOXP3_AND_SATB1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP GSE11973_MIR223_KOVS_WT_BONE_MARROW_NEUTROPHIL_DN, GSE11973_MIR223_KOVS_WT_BONE_MARROW_NEUTROPHIL_DN GOBP_CARBOHYDRATE_CATABOLIC_PROCESS, GOBP_CARBOHYDRATE_CATABOLIC_PROCESS REACTOME_CELLULAR_RESPONSE_TO_HEAT_STRESS, REACTOME_CELLULAR_RESPONSE_TO_HEAT_STRESS
	GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_UP, GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_UP HOSHIDA_LIVER_CANCER_SUBCLASS_S2, HOSHIDA_LIVER_CANCER_SUBCLASS_S2 GSE23308_WT_VS_MINERALCORTICOID_REC_KO_MACROPHAGE_CORTICOSTERONE_TREATED_DN, GSE23308_WT_VS_MINERALCORTICOID_REC_KO_MACROPHAGE_CORTICOSTERONE_TREATED_DN GSE17721_LPS_VS_GARDIQUIMOD_24H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_24H_BMDC_DN GSE339_CD4POS_VS_CD8POS_DC_DN, GSE339_CD4POS_VS_CD8POS_DC_DN GSE360_L_DONOVANI_VS_T_GONDII_MAC_DN, GSE360_L_DONOVANI_VS_T_GONDII_MAC_DN
	REACTOME_TRNA_PROCESSING, REACTOME_TRNA_PROCESSING GSE9316_CD4_TCELL_BALBC_VS_TH17_ENRI_CD4_TCELL_SKG_PMA_IONO_STIM_FR4NEG_DN, GSE9316_CD4_TCELL_BALBC_VS_TH17_ENRI_CD4_TCELL_SKG_PMA_IONO_STIM_FR4NEG_DN GSE4142_PLASMA_CELL_VS_GC_BCELL_DN, GSE4142_PLASMA_CELL_VS_GC_BCELL_DN GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_MAC_DN, GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_MAC_DN GSE21670_STAT3_KO_VS_WT_CD4_TCELL_DN, GSE21670_STAT3_KO_VS_WT_CD4_TCELL_DN GSE22025_UNTREATED_VS_PROGESTERONE_TREATED_CD4_TCELL_DN, GSE22025_UNTREATED_VS_PROGESTERONE_TREATED_CD4_TCELL_DN
	CSR_EARLY_UP.V1_UP, CSR_EARLY_UP.V1_UP KARLSSON_TGFB1_TARGETS_UP, KARLSSON_TGFB1_TARGETS_UP GSE3982_EOSINOPHIL_VS_NKCELL_DN, GSE3982_EOSINOPHIL_VS_NKCELL_DN MODULE_159, MODULE_159 GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_4H_UP, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_4H_UP GSE17721_POLYIC_VS_GARDIQUIMOD_4H_BMDC_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_4H_BMDC_DN
	GOBP_REGULATION_OF_GENERATION_OF_PRECURSOR_METABOLITES_AND_ENERGY, GOBP_REGULATION_OF_GENERATION_OF_PRECURSOR_METABOLITES_AND_ENERGY GOBP_CYTOPLASMIC_TRANSLATION, GOBP_CYTOPLASMIC_TRANSLATION REACTOME_DISORDERS_OF_TRANSMEMBRANE_TRANSPORTERS, REACTOME_DISORDERS_OF_TRANSMEMBRANE_TRANSPORTERS GSE3982_EOSINOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN, GSE3982_EOSINOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN GOBP_PROTEIN_SUMOYLATION, GOBP_PROTEIN_SUMOYLATION
	GOMF_UNFOLDED_PROTEIN_BINDING, GOMF_UNFOLDED_PROTEIN_BINDING GSE17721_0.5H_VS_4H_PAM3CSK4_BMDC_DN, GSE17721_0.5H_VS_4H_PAM3CSK4_BMDC_DN GSE27786_LIN_NEG_VS_BCELL_UP, GSE27786_LIN_NEG_VS_BCELL_UP GSE11924_TFH_VS_TH2_CD4_TCELL_DN, GSE11924_TFH_VS_TH2_CD4_TCELL_DN GSE3982_DC_VS_BASOPHIL_UP, GSE3982_DC_VS_BASOPHIL_UP GSE37301_COMMON_LYMPHOID_PROGENITOR_VS_GRAN_MONO_PROGENITOR_UP, GSE37301_COMMON_LYMPHOID_PROGENITOR_VS_GRAN_MONO_PROGENITOR_UP
	REACTOME_ANTIVIRAL_MECHANISM_BY_IFN_STIMULATED_GENES, REACTOME_ANTIVIRAL_MECHANISM_BY_IFN_STIMULATED_GENES REACTOME_HCMV_EARLY_EVENTS, REACTOME_HCMV_EARLY_EVENTS GSE33424_CD161_HIGH_VS_INT_CD8_TCELL_UP, GSE33424_CD161_HIGH_VS_INT_CD8_TCELL_UP MORF_SP3, MORF_SP3 GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_UP, GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_UP REACTOME_SUMOYLATION_OF_CHROMATIN_ORGANIZATION_PROTEINS, REACTOME_SUMOYLATION_OF_CHROMATIN_ORGANIZATION_PROTEINS
	GSE13493_DP_VS_CD4INTCD8POS_THYMOCYTE_DN, GSE13493_DP_VS_CD4INTCD8POS_THYMOCYTE_DN REACTOME_HCMV_LATE_EVENTS, REACTOME_HCMV_LATE_EVENTS CCGNMNNTNACG_UNKNOWN, CCGNMNNTNACG_UNKNOWN KAAB_FAILED_HEART_ATRIUM_DN, KAAB_FAILED_HEART_ATRIUM_DN GOBP_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT, GOBP_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT TAKAO_RESPONSE_TO_UVB_RADIATION_DN, TAKAO_RESPONSE_TO_UVB_RADIATION_DN
	GSE360_LOW_DOSE_B_MALAYI_VS_M_TUBERCULOSIS_MAC_DN, GSE360_LOW_DOSE_B_MALAYI_VS_M_TUBERCULOSIS_MAC_DN SEIDEN_ONCOGENESIS_BY_MET, SEIDEN_ONCOGENESIS_BY_MET GSE1460_CORD_VS_ADULT_BLOOD_NAIVE_CD4_TCELL_UP, GSE1460_CORD_VS_ADULT_BLOOD_NAIVE_CD4_TCELL_UP GSE27786_LIN_NEG_VS_NKCELL_UP, GSE27786_LIN_NEG_VS_NKCELL_UP GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEUS, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEUS DER_IFN_BETA_RESPONSE_UP, DER_IFN_BETA_RESPONSE_UP
	GOBP_REGULATION_OF_ATP_METABOLIC_PROCESS, GOBP_REGULATION_OF_ATP_METABOLIC_PROCESS GSE17721_CTRL_VS_PAM3CSK4_12H_BMDC_DN, GSE17721_CTRL_VS_PAM3CSK4_12H_BMDC_DN GOBP_REGULATION_OF_CARBOHYDRATE_CATABOLIC_PROCESS, GOBP_REGULATION_OF_CARBOHYDRATE_CATABOLIC_PROCESS SHAFFER_IRF4_TARGETS_IN_MYELOMA_VS_MATURE_B_LYMPHOCYTE, SHAFFER_IRF4_TARGETS_IN_MYELOMA_VS_MATURE_B_LYMPHOCYTE GSE11864_CSF1_IFNG_VS_CSF1_PAM3CYS_IN_MAC_DN, GSE11864_CSF1_IFNG_VS_CSF1_PAM3CYS_IN_MAC_DN
	HP_DISTAL_SENSORY_IMPAIRMENT, HP_DISTAL_SENSORY_IMPAIRMENT GOMF_TRNA_BINDING, GOMF_TRNA_BINDING JAIN_NFKB_SIGNALING, JAIN_NFKB_SIGNALING GSE22886_NAIVE_CD8_TCELL_VS_NKCELL_UP, GSE22886_NAIVE_CD8_TCELL_VS_NKCELL_UP GSE24142_ADULT_VS_FETAL_DN2_THYMOCYTE_DN, GSE24142_ADULT_VS_FETAL_DN2_THYMOCYTE_DN GSE13306_RA_VS_UNTREATED_MEM_CD4_TCELL_DN, GSE13306_RA_VS_UNTREATED_MEM_CD4_TCELL_DN
	GSE41867_NAIVE_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP GSE360_DC_VS_MAC_L_MAJOR_UP, GSE360_DC_VS_MAC_L_MAJOR_UP REACTOME_SUMOYLATION_OF_SUMOYLATION_PROTEINS, REACTOME_SUMOYLATION_OF_SUMOYLATION_PROTEINS WP_16P112_PROXIMAL_DELETION_SYNDROME, WP_16P112_PROXIMAL_DELETION_SYNDROME ACOSTA_PROLIFERATION_INDEPENDENT_MYC_TARGETS_UP, ACOSTA_PROLIFERATION_INDEPENDENT_MYC_TARGETS_UP GSE43955_1H_VS_20H_ACT_CD4_TCELL_DN, GSE43955_1H_VS_20H_ACT_CD4_TCELL_DN
	GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP BROWN_MYELOID_CELL_DEVELOPMENT_DN, BROWN_MYELOID_CELL_DEVELOPMENT_DN GSE3982_BASOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN, GSE3982_BASOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN REACTOME_SUMOYLATION_OF_RNA_BINDING_PROTEINS, REACTOME_SUMOYLATION_OF_RNA_BINDING_PROTEINS GSE15330_LYMPHOID_MULTIPOTENT_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN, GSE15330_LYMPHOID_MULTIPOTENT_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN GOBP_POSITIVE_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT
	MIR34A_3P, MIR34A_3P MORF_USP5, MORF_USP5 REACTOME_SUMOYLATION_OF_UBIQUITINYLATION_PROTEINS, REACTOME_SUMOYLATION_OF_UBIQUITINYLATION_PROTEINS GSE14350_IL2RB_KO_VS_WT_TEFF_DN, GSE14350_IL2RB_KO_VS_WT_TEFF_DN GOCC_ATPASE_COMPLEX, GOCC_ATPASE_COMPLEX TRAVAGLINI_LUNG_BASAL_CELL, TRAVAGLINI_LUNG_BASAL_CELL
	GSE17721_PAM3CSK4_VS_GADIQUIMOD_0.5H_BMDC_UP, GSE17721_PAM3CSK4_VS_GADIQUIMOD_0.5H_BMDC_UP GOBP_AMINO_ACID_ACTIVATION, GOBP_AMINO_ACID_ACTIVATION GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_UP, GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_UP WP_TRANSLATION_FACTORS, WP_TRANSLATION_FACTORS GOCC_SWI_SNF_SUPERFAMILY_TYPE_COMPLEX, GOCC_SWI_SNF_SUPERFAMILY_TYPE_COMPLEX CCTGTGA_MIR513, CCTGTGA_MIR513
	GSE360_L_DONOVANI_VS_L_MAJOR_DC_UP, GSE360_L_DONOVANI_VS_L_MAJOR_DC_UP MIR4670_3P, MIR4670_3P HP_KINETIC_TREMOR, HP_KINETIC_TREMOR MIR891B, MIR891B YYCATTCAWW_UNKNOWN, YYCATTCAWW_UNKNOWN
	DIRMEIER_LMP1_RESPONSE_LATE_UP, DIRMEIER_LMP1_RESPONSE_LATE_UP REACTOME_SLC_TRANSPORTER_DISORDERS, REACTOME_SLC_TRANSPORTER_DISORDERS GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMC_DN, GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMC_DN KEGG_AMINOACYL_TRNA_BIOSYNTHESIS, KEGG_AMINOACYL_TRNA_BIOSYNTHESIS BRACHAT_RESPONSE_TO_CAMPTOTHECIN_DN, BRACHAT_RESPONSE_TO_CAMPTOTHECIN_DN GSE11864_CSF1_VS_CSF1_PAM3CYS_IN_MAC_DN, GSE11864_CSF1_VS_CSF1_PAM3CYS_IN_MAC_DN
	GSE21360_PRIMARY_VS_SECONDARY_MEMORY_CD8_TCELL_UP, GSE21360_PRIMARY_VS_SECONDARY_MEMORY_CD8_TCELL_UP GOMF_NUCLEOSOME_BINDING, GOMF_NUCLEOSOME_BINDING SESTO_RESPONSE_TO_UV_C2, SESTO_RESPONSE_TO_UV_C2 REACTOME_TRNA_AMINOACYLATION, REACTOME_TRNA_AMINOACYLATION LUI_THYROID_CANCER_PAX8_PPARG_UP, LUI_THYROID_CANCER_PAX8_PPARG_UP GOMF_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS, GOMF_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS
	REACTOME_DNA_DAMAGE_RECOGNITION_IN_GG_NER, REACTOME_DNA_DAMAGE_RECOGNITION_IN_GG_NER GOBP_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS, GOBP_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS GOBP_RIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS, GOBP_RIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_ROSIGLITAZONE_TREATED_UP, GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_ROSIGLITAZONE_TREATED_UP GOBP_REGULATION_OF_PROTEIN_IMPORT, GOBP_REGULATION_OF_PROTEIN_IMPORT CHENG_RESPONSE_TO_NICKEL_ACETATE, CHENG_RESPONSE_TO_NICKEL_ACETATE
	DARWICHE_PAPILLOMA_RISK_LOW_UP, DARWICHE_PAPILLOMA_RISK_LOW_UP MIR1206, MIR1206 ELVIDGE_HYPOXIA_BY_DMOG_DN, ELVIDGE_HYPOXIA_BY_DMOG_DN GOBP_INTERFERON_BETA_PRODUCTION, GOBP_INTERFERON_BETA_PRODUCTION REACTOME_PROTEIN_FOLDING, REACTOME_PROTEIN_FOLDING GOBP_DE_NOVO_PROTEIN_FOLDING, GOBP_DE_NOVO_PROTEIN_FOLDING
	GSE13306_LAMINA_PROPRIA_VS_SPLEEN_TREG_DN, GSE13306_LAMINA_PROPRIA_VS_SPLEEN_TREG_DN HP_ABNORMAL_MOTOR_NERVE_CONDUCTION_VELOCITY, HP_ABNORMAL_MOTOR_NERVE_CONDUCTION_VELOCITY MODULE_299, MODULE_299 BROWNE_HCMV_INFECTION_10HR_UP, BROWNE_HCMV_INFECTION_10HR_UP MARZEC_IL2_SIGNALING_UP, MARZEC_IL2_SIGNALING_UP
	GOBP_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION, GOBP_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION TRAVAGLINI_LUNG_LIPOFIBROBLAST_CELL, TRAVAGLINI_LUNG_LIPOFIBROBLAST_CELL GOBP_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GOBP_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING GOBP_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GOBP_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I MIR216B_5P, MIR216B_5P MODULE_355, MODULE_355
ETS_REPRESSED_BY_SERUM, SCHLOSSER_MYC_TARGETS_REPRESSED_BY_SERUM	GSE13887_HEALTHY_VS_LUPUS_RESTING_CD4_TCELL_DN, GSE13887_HEALTHY_VS_LUPUS_RESTING_CD4_TCELL_DN PPARA_02, PPARA_02 CUI_DEVELOPING_HEART_TRABECULAR_VENTRICULAR_CARDIOMYOCYTE, CUI_DEVELOPING_HEART_TRABECULAR_VENTRICULAR_CARDIOMYOCYTE GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_DC_UP, GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_DC_UP GOMF_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_ACTIVITY, GOMF_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_ACTIVITY REACTOME_CYTOSOLIC_TRNA_AMINOACYLATION, REACTOME_CYTOSOLIC_TRNA_AMINOACYLATION
	MODULE_82, MODULE_82 GOMF_CHROMATIN_DNA_BINDING, GOMF_CHROMATIN_DNA_BINDING GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME, GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME HP_EMG_NEUROPATHIC_CHANGES, HP_EMG_NEUROPATHIC_CHANGES GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNG_KO_DN, GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNG_KO_DN CHOW_RASSF1_TARGETS_DN, CHOW_RASSF1_TARGETS_DN
	FAELT_B_CLL_WITH_VH_REARRANGEMENTS_DN, FAELT_B_CLL_WITH_VH_REARRANGEMENTS_DN GOMF_DISULFIDE_OXIDOREDUCTASE_ACTIVITY, GOMF_DISULFIDE_OXIDOREDUCTASE_ACTIVITY TAGAACC_MIR182, TAGAACC_MIR182 GOMF_HYDROLASE_ACTIVITY_ACTING_ON_CARBON_NITROGEN_BUT_NOT_PEPTIDE_BONDS, GOMF_HYDROLASE_ACTIVITY_ACTING_ON_CARBON_NITROGEN_BUT_NOT_PEPTIDE_BONDS MIR3691_3P, MIR3691_3P LUI_THYROID_CANCER_CLUSTER_1, LUI_THYROID_CANCER_CLUSTER_1
	APRELIKOVA_BRCA1_TARGETS, APRELIKOVA_BRCA1_TARGETS ATCTTGC_MIR31, ATCTTGC_MIR31 HP_RELATIVE_MACROCEPHALY, HP_RELATIVE_MACROCEPHALY MIR4764_3P, MIR4764_3P GOBP_INTERFERON_ALPHA_PRODUCTION, GOBP_INTERFERON_ALPHA_PRODUCTION GOBP_RIBONUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_RIBONUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS
	HP_LANGUAGE_IMPAIRMENT, HP_LANGUAGE_IMPAIRMENT REACTOME_CARGO_TRAFFICKING_TO_THE_PERICILIARY_MEMBRANE, REACTOME_CARGO_TRAFFICKING_TO_THE_PERICILIARY_MEMBRANE BIOCARTA_EIF_PATHWAY, BIOCARTA_EIF_PATHWAY GOBP_INTERLEUKIN_6_PRODUCTION, GOBP_INTERLEUKIN_6_PRODUCTION HP_SPINAL_MUSCULAR_ATROPHY, HP_SPINAL_MUSCULAR_ATROPHY GOBP_REGULATION_OF_DNA_BINDING, GOBP_REGULATION_OF_DNA_BINDING
	MIR6781_3P, MIR6781_3P GOMF_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_BINDING, GOMF_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_BINDING GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_6_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_6_PRODUCTION BIOCARTA_IRES_PATHWAY, BIOCARTA_IRES_PATHWAY REACTOME_RESPONSE_OF_EIF2AK1_HRI_TO_HEME_DEFICIENCY, REACTOME_RESPONSE_OF_EIF2AK1_HRI_TO_HEME_DEFICIENCY
	MIR655_5P, MIR655_5P YAMASHITA_LIVER_CANCER_WITH_EPCAM_UP, YAMASHITA_LIVER_CANCER_WITH_EPCAM_UP MIR6832_5P, MIR6832_5P MIR4266, MIR4266 MIR4642, MIR4642 MA_MYELOID_DIFFERENTIATION_UP, MA_MYELOID_DIFFERENTIATION_UP
	MODULE_4, MODULE_4 GOBP_SKIN_EPIDERMIS_DEVELOPMENT, GOBP_SKIN_EPIDERMIS_DEVELOPMENT BIOCARTA_MTOR_PATHWAY, BIOCARTA_MTOR_PATHWAY GOBP_POSITIVE_REGULATION_OF_INTERFERON_ALPHA_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERFERON_ALPHA_PRODUCTION MIR4786_5P, MIR4786_5P GOMF_PROTEIN_FOLDING_CHAPERONE, GOMF_PROTEIN_FOLDING_CHAPERONE
	MIR4274, MIR4274 GENTILE_UV_RESPONSE_CLUSTER_D2, GENTILE_UV_RESPONSE_CLUSTER_D2 NAKAYA_PBMC_FLUMIST_AGE_18_50YO_7DY_IFN_SUBSET_UP, NAKAYA_PBMC_FLUMIST_AGE_18_50YO_7DY_IFN_SUBSET_UP FU_INTERACT_WITH_ALKBH8, FU_INTERACT_WITH_ALKBH8 WP_NSP1_FROM_SARSCOV2_INHIBITS_TRANSLATION_INITIATION_IN_THE_HOST_CELL, WP_NSP1_FROM_SARSCOV2_INHIBITS_TRANSLATION_INITIATION_IN_THE_HOST_CELL GOMF_NUCLEAR_IMPORT_SIGNAL_RECEPTOR_ACTIVITY, GOMF_NUCLEAR_IMPORT_SIGNAL_RECEPTOR_ACTIVITY
	REACTOME_COOPERATION_OF_PREFOLDIN_AND_TRIC_CCT_IN_ACTIN_AND_TUBULIN_FOLDING, REACTOME_COOPERATION_OF_PREFOLDIN_AND_TRIC_CCT_IN_ACTIN_AND_TUBULIN_FOLDING HP_PROGEROID_FACIAL_APPEARANCE, HP_PROGEROID_FACIAL_APPEARANCE CHEOK_RESPONSE_TO_MERCAPTOPURINE_AND_HD_MTX_DN, CHEOK_RESPONSE_TO_MERCAPTOPURINE_AND_HD_MTX_DN GOBP_REGULATION_OF_TRANSLATIONAL_FIDELITY, GOBP_REGULATION_OF_TRANSLATIONAL_FIDELITY GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_TELOMERE, GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_TELOMERE GOBP_TOXIN_TRANSPORT, GOBP_TOXIN_TRANSPORT
	GOCC_POLYSOMAL_RIBOSOME, GOCC_POLYSOMAL_RIBOSOME MIR6826_5P, MIR6826_5P REACTOME_THE_NLRP3_INFLAMMASOME, REACTOME_THE_NLRP3_INFLAMMASOME GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATION, GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATION REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION, REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION
	PID_P38_ALPHA_BETA_DOWNSTREAM_PATHWAY, PID_P38_ALPHA_BETA_DOWNSTREAM_PATHWAY MODULE_447, MODULE_447 MIR4715_5P, MIR4715_5P GOMF_LIGASE_ACTIVITY_FORMING_CARBON_SULFUR_BONDS, GOMF_LIGASE_ACTIVITY_FORMING_CARBON_SULFUR_BONDS HP_FACIAL_HYPOTONIA, HP_FACIAL_HYPOTONIA GOBP_POSITIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION
	GOMF_RNA_POLYMERASE_II_REPRESSING_TRANSCRIPTION_FACTOR_BINDING, GOMF_RNA_POLYMERASE_II_REPRESSING_TRANSCRIPTION_FACTOR_BINDING CHANGOLKAR_H2AFY_TARGETS_DN, CHANGOLKAR_H2AFY_TARGETS_DN GOCC_TRANSLATION_PREINITIATION_COMPLEX, GOCC_TRANSLATION_PREINITIATION_COMPLEX HP_ABNORMAL_HAIR_LABORATORY_EXAMINATION, HP_ABNORMAL_HAIR_LABORATORY_EXAMINATION GOBP_STRESS_GRANULE_ASSEMBLY, GOBP_STRESS_GRANULE_ASSEMBLY GOBP_PROTEIN_NEDDYLATION, GOBP_PROTEIN_NEDDYLATION
	HP_ADIPOSE_TISSUE_LOSS, HP_ADIPOSE_TISSUE_LOSS GOBP_MOLTING_CYCLE, GOBP_MOLTING_CYCLE WP_NEPHROTIC_SYNDROME, WP_NEPHROTIC_SYNDROME GOBP_GLUCOSE_6_PHOSPHATE_METABOLIC_PROCESS, GOBP_GLUCOSE_6_PHOSPHATE_METABOLIC_PROCESS HP_THORACIC_SCOLIOSIS, HP_THORACIC_SCOLIOSIS MIR4312, MIR4312
	CHEN_LUNG_CANCER_SURVIVAL, CHEN_LUNG_CANCER_SURVIVAL NAM_FXYD5_TARGETS_DN, NAM_FXYD5_TARGETS_DN GOBP_FERTILIZATION, GOBP_FERTILIZATION CHOI_ATL_CHRONIC_VS_ACUTE_DN, CHOI_ATL_CHRONIC_VS_ACUTE_DN GOMF_METALLOENDOPEPTIDASE_ACTIVITY, GOMF_METALLOENDOPEPTIDASE_ACTIVITY GOMF_PEPTIDE_DISULFIDE_OXIDOREDUCTASE_ACTIVITY, GOMF_PEPTIDE_DISULFIDE_OXIDOREDUCTASE_ACTIVITY
	REACTOME_INFLAMMASOMES, REACTOME_INFLAMMASOMES RIZKI_TUMOR_INVASIVENESS_2D_DN, RIZKI_TUMOR_INVASIVENESS_2D_DN GOBP_POLY_A_PLUS_MRNA_EXPORT_FROM_NUCLEUS, GOBP_POLY_A_PLUS_MRNA_EXPORT_FROM_NUCLEUS FONTAINE_THYROID_TUMOR_UNCERTAIN_MALIGNANCY_UP, FONTAINE_THYROID_TUMOR_UNCERTAIN_MALIGNANCY_UP NEBEN_AML_WITH_FLT3_OR_NRAS_DN, NEBEN_AML_WITH_FLT3_OR_NRAS_DN
	DHIMAN_PBMC_ATTENUVAX_AGE_15_25YO_SUBQ_7_OR_14DY_UP, DHIMAN_PBMC_ATTENUVAX_AGE_15_25YO_SUBQ_7_OR_14DY_UP GOBP_PROTEIN_K63_LINKED_DEUBIQUITINATION, GOBP_PROTEIN_K63_LINKED_DEUBIQUITINATION GOBP_BINDING_OF_SPERM_TO_ZONA_PELLUCIDA, GOBP_BINDING_OF_SPERM_TO_ZONA_PELLUCIDA GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION MIR8071, MIR8071 GOBP_POSITIVE_REGULATION_OF_TELOMERASE_RNA_LOCALIZATION_TO_CAJAL_BODY, GOBP_POSITIVE_REGULATION_OF_TELOMERASE_RNA_LOCALIZATION_TO_CAJAL_BODY
	REACTOME_FORMATION_OF_TUBULIN_FOLDING_INTERMEDIATES_BY_CCT_TRIC, REACTOME_FORMATION_OF_TUBULIN_FOLDING_INTERMEDIATES_BY_CCT_TRIC CHEN_NEUROBLASTOMA_COPY_NUMBER_GAINS, CHEN_NEUROBLASTOMA_COPY_NUMBER_GAINS GOMF_G_QUADRUPLEX_DNA_BINDING, GOMF_G_QUADRUPLEX_DNA_BINDING GOBP_U4_SNRNA_3_END_PROCESSING, GOBP_U4_SNRNA_3_END_PROCESSING GOBP_CELLULAR_RESPONSE_TO_MANGANESE_ION, GOBP_CELLULAR_RESPONSE_TO_MANGANESE_ION GOBP_NLS_BEARING_PROTEIN_IMPORT_INTO_NUCLEUS, GOBP_NLS_BEARING_PROTEIN_IMPORT_INTO_NUCLEUS
	GOCC_DNA_HELICASE_COMPLEX, GOCC_DNA_HELICASE_COMPLEX GOBP_RRNA_3_END_PROCESSING, GOBP_RRNA_3_END_PROCESSING GOCC_NPBAF_COMPLEX, GOCC_NPBAF_COMPLEX GOBP_PROLINE_BIOSYNTHETIC_PROCESS, GOBP_PROLINE_BIOSYNTHETIC_PROCESS GOBP_RESPONSE_TO_MANGANESE_ION, GOBP_RESPONSE_TO_MANGANESE_ION GOBP_CHAPERONE_MEDIATED_PROTEIN_COMPLEX_ASSEMBLY, GOBP_CHAPERONE_MEDIATED_PROTEIN_COMPLEX_ASSEMBLY
	HP_BIRD_LIKE_FACIES, HP_BIRD_LIKE_FACIES GOMF_NUCLEOSOMAL_DNA_BINDING, GOMF_NUCLEOSOMAL_DNA_BINDING HP_ABNORMAL_HAIR_WHORL, HP_ABNORMAL_HAIR_WHORL BIOCARTA_IGF1MTOR_PATHWAY, BIOCARTA_IGF1MTOR_PATHWAY MIR657, MIR657
	GOMF_P53_BINDING, GOMF_P53_BINDING WP_MAMMARY_GLAND_DEVELOPMENT_PATHWAY_PREGNANCY_AND_LACTATION_STAGE_3_OF_4, WP_MAMMARY_GLAND_DEVELOPMENT_PATHWAY_PREGNANCY_AND_LACTATION_STAGE_3_OF_4 MIR7705, MIR7705 FXR_Q3, FXR_Q3 HP_ABSENCE_OF_SUBCUTANEOUS_FAT, HP_ABSENCE_OF_SUBCUTANEOUS_FAT GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_RED_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_RED_UP
	MIR214_5P, MIR214_5P HP_DERMAL_TRANSLUCENCY, HP_DERMAL_TRANSLUCENCY GOCC_EUKARYOTIC_48S_PREINITIATION_COMPLEX, GOCC_EUKARYOTIC_48S_PREINITIATION_COMPLEX ZNF586_TARGET_GENES, ZNF586_TARGET_GENES GOBP_NEGATIVE_REGULATION_OF_UBIQUITIN_PROTEIN_LIGASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_UBIQUITIN_PROTEIN_LIGASE_ACTIVITY GOCC_SWI_SNF_COMPLEX, GOCC_SWI_SNF_COMPLEX
	GOMF_UBIQUITIN_LIKE_MODIFIER_ACTIVATING_ENZYME_ACTIVITY, GOMF_UBIQUITIN_LIKE_MODIFIER_ACTIVATING_ENZYME_ACTIVITY WP_THE_INFLUENCE_OF_LAMINOPATHIES_ON_WNT_SIGNALING, WP_THE_INFLUENCE_OF_LAMINOPATHIES_ON_WNT_SIGNALING GOBP_REGULATION_OF_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GOBP_REGULATION_OF_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR GOBP_POSITIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY HP_ABNORMAL_TRICUSPID_VALVE_MORPHOLOGY, HP_ABNORMAL_TRICUSPID_VALVE_MORPHOLOGY MIR5192, MIR5192
	MIR1468_5P, MIR1468_5P GOBP_PROTEIN_LOCALIZATION_TO_NUCLEOPLASM, GOBP_PROTEIN_LOCALIZATION_TO_NUCLEOPLASM GOBP_NEGATIVE_REGULATION_OF_INTRACELLULAR_TRANSPORT, GOBP_NEGATIVE_REGULATION_OF_INTRACELLULAR_TRANSPORT POS_HISTAMINE_RESPONSE_NETWORK, POS_HISTAMINE_RESPONSE_NETWORK GOMF_HYDROLASE_ACTIVITY_ACTING_ON_CARBON_NITROGEN_BUT_NOT_PEPTIDE_BONDS_IN_CYCLIC_AMIDINES, GOMF_HYDROLASE_ACTIVITY_ACTING_ON_CARBON_NITROGEN_BUT_NOT_PEPTIDE_BONDS_IN_CYCLIC_AMIDINES.
	GOBP_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME, GOBP_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME GOBP_POSITIVE_REGULATION_OF_INFLAMMATORY_RESPONSE, GOBP_POSITIVE_REGULATION_OF_INFLAMMATORY_RESPONSE REACTOME_CAMK_IV_MEDIATED_PHOSPHORYLATION_OF_CREB, REACTOME_CAMK_IV_MEDIATED_PHOSPHORYLATION_OF_CREB HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_7DY_TOP_100_DEG_EX_VIVO_DN, HOFT_CD4_POSITIVE_ALPHA_BETA_HORD_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_7DY_TOP_100_DEG_EX_VIVO_DN, HOFT_CD4_POSITIVE_ALPHA_BETA_HORD_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_7DY_TOP_100_DEG_EX_VIVO_DN, HOFT_CD4_POSITIVE_AGA_T_CD
	GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSLATION, GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSLATION GOBP_NUCLEOLAR_LARGE_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GOBP_NUCLEOLAR_LARGE_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_I MIR7151_5P, MIR7151_5P GOBP_HISTONE_MRNA_CATABOLIC_PROCESS, GOBP_HISTONE_MRNA_CATABOLIC_PROCESS HP_LONG_TRACT_SIGNS, HP_LONG_TRACT_SIGNS HP_ANGINA_PECTORIS, HP_ANGINA_PECTORIS
	GOCC_ENDOPLASMIC_RETICULUM_CHAPERONE_COMPLEX, GOCC_ENDOPLASMIC_RETICULUM_CHAPERONE_COMPLEX HP_THENAR_MUSCLE_ATROPHY, HP_THENAR_MUSCLE_ATROPHY AIYAR_COBRA1_TARGETS_UP, AIYAR_COBRA1_TARGETS_UP GOMF_PROTEASOME_BINDING, GOMF_PROTEASOME_BINDING TOMIDA_METASTASIS_DN, TOMIDA_METASTASIS_DN HP_ABNORMAL_CIRCULATING_NON_PROTEINOGENIC_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_NON_PROTEINOGENIC_AMINO_ACID_CONCENTRATION
	GOCC_SMN_COMPLEX, GOCC_SMN_COMPLEX MIR1245A, MIR1245A REACTOME_FOLDING_OF_ACTIN_BY_CCT_TRIC, REACTOME_FOLDING_OF_ACTIN_BY_CCT_TRIC HP_ABNORMALITY_OF_FACIAL_ADIPOSE_TISSUE, HP_ABNORMALITY_OF_FACIAL_ADIPOSE_TISSUE GOMF_PHOSPHATIDIC_ACID_TRANSFER_ACTIVITY, GOMF_PHOSPHATIDIC_ACID_TRANSFER_ACTIVITY STARK_HYPPOCAMPUS_22Q11_DELETION_DN, STARK_HYPPOCAMPUS_22Q11_DELETION_DN
	YANG_BREAST_CANCER_ESR1_DN, YANG_BREAST_CANCER_ESR1_DN MIR1322, MIR1322 MIR8079, MIR8079 HP_URINARY_URGENCY, HP_URINARY_URGENCY HP_ABNORMALITY_OF_EPIDERMAL_MORPHOLOGY, HP_ABNORMALITY_OF_EPIDERMAL_MORPHOLOGY MIR5704, MIR5704
	MIR5704, MIR5704 HP_SLENDER_BUILD, HP_SLENDER_BUILD WP_HIJACK_OF_UBIQUITINATION_BY_SARSCOV2, WP_HIJACK_OF_UBIQUITINATION_BY_SARSCOV2 HP_SCANNING_SPEECH, HP_SCANNING_SPEECH GOBP_AMINOACYL_TRNA_METABOLISM_INVOLVED_IN_TRANSLATIONAL_FIDELITY, GOBP_AMINOACYL_TRNA_METABOLISM_INVOLVED_IN_TRANSLATIONAL_FIDELITY GOMF_AMINOACYL_TRNA_EDITING_ACTIVITY, GOMF_AMINOACYL_TRNA_EDITING_ACTIVITY