	GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_10H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_10H_UP GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_UP GSE14000_UNSTIM_VS_4H_LPS_DC_UP, GSE14000_UNSTIM_VS_4H_LPS_DC_UP GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_14H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_14H_UP GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_6H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_6H_UP GSE14000_UNSTIM_VS_4H_LPS_DC_TRANSLATED_RNA_UP, GSE14000_UNSTIM_VS_4H_LPS_DC_TRANSLATED_RNA_UP
	GSE14000_UNSTIM_VS_16H_LPS_DC_TRANSLATED_RNA_UP, GSE14000_UNSTIM_VS_16H_LPS_DC_TRANSLATED_RNA_UP GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_18H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_18H_UP GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_6H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_6H_UP GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_4H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_4H_UP GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_10H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_10H_UP GOBP_TRNA_PROCESSING, GOBP_TRNA_PROCESSING GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_18H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_18H_UP GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_18H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_18H_UP GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_18H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_18H_UP
	GSE2706_UNSTIM_VS_2H_LPS_AND_R848_DC_UP, GSE2706_UNSTIM_VS_2H_LPS_AND_R848_DC_UP GSE2706_UNSTIM_VS_2H_LPS_DC_UP, GSE2706_UNSTIM_VS_2H_LPS_DC_UP MYC_UP.V1_UP, MYC_UP.V1_UP GSE29618_PRE_VS_DAY7_POST_LAIV_FLU_VACCINE_MDC_DN, GSE29618_PRE_VS_DAY7_POST_LAIV_FLU_VACCINE_MDC_DN GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_2H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_2H_UP GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_BCELL_DN, GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_BCELL_DN GSE27859_MACROPHAGE_VS_CD11C_INT_F480_INT_DC_DN, GSE27859_MACROPHAGE_VS_CD11C_INT_F480_INT_DC_DN
	GSE3982_BCELL_VS_BASOPHIL_UP, GSE3982_BCELL_VS_BASOPHIL_UP GSE16385_IFNG_TNF_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP, GSE16385_IFNG_TNF_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP MIR330_5P, MIR330_5P GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_UP, GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_UP GSE8515_CTRL_VS_IL1_4H_STIM_MAC_UP, GSE8515_CTRL_VS_IL1_4H_STIM_MAC_UP GOCC_SMALL_SUBUNIT_PROCESSOME, GOCC_SMALL_SUBUNIT_PROCESSOME MIR4802_3P, MIR4802_3P GSE301_NIR4802_3P GSE301_NIR4802_3P GSE301_NIR4802_3P GSE301_NIR4802_3P
	GSE8921_UNSTIM_OH_VS_TLR1_2_STIM_MONOCYTE_6H_UP, GSE8921_UNSTIM_OH_VS_TLR1_2_STIM_MONOCYTE_6H_UP  MARZEC_IL2_SIGNALING_UP, MARZEC_IL2_SIGNALING_UP  GOBP_MITOCHONDRIAL_RNA_METABOLIC_PROCESS, GOBP_MITOCHONDRIAL_RNA_METABOLIC_PROCESS  GOBP_TRNA_METHYLATION, GOBP_TRNA_METHYLATION  ZHAN_MULTIPLE_MYELOMA_CD1_AND_CD2_DN, ZHAN_MULTIPLE_MYELOMA_CD1_AND_CD2_DN  REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION, REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION  MIR326, MIR326  GOSTEMED A RESETS OF VOLVE AND A REACTOME A RESETS OF VOLVE AND A REACTOME.
	GOTTWEIN_TARGETS_OF_KSHV_MIR_K12_11, GOTTWEIN_TARGETS_OF_KSHV_MIR_K12_11  GOMF_TRNA_METHYLTRANSFERASE_ACTIVITY, GOMF_TRNA_METHYLTRANSFERASE_ACTIVITY  GOBP_NUCLEOTIDE_EXCISION_REPAIR_PREINCISION_COMPLEX_ASSEMBLY, GOBP_NUCLEOTIDE_EXCISION_REPAIR_PREINCISION_COMPLEX_ASSEMBLY  WCTCNATGGY_UNKNOWN, WCTCNATGGY_UNKNOWN  GSE43956_WT_VS_SGK1_KO_TH17_DIFFERENTIATED_CD4_TCELL_UP, GSE43956_WT_VS_SGK1_KO_TH17_DIFFERENTIATED_CD4_TCELL_UP  GOBP_RRNA_MODIFICATION, GOBP_RRNA_MODIFICATION  WP_GENOTOXICITY_PATHWAY, WP_GENOTOXICITY_PATHWAY
	GOMF_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS, GOMF_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS, GOMF_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS, GOMF_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS, GOMF_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_SERVERSE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_SERVERSE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_SERVERSE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_SERVERSE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_SERVERSE_ACTIVITY_ACTIVE_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_SERVERSE_ACTIVITY_ACTIVE_ACTI
	ZNF708_TARGET_GENES, ZNF708_TARGET_GENES MIR6799_3P, MIR6799_3P HP_PRURITUS, HP_PRURITUS HP_ABNORMALITY_OF_THE_SCROTUM, HP_ABNORMALITY_OF_THE_SCROTUM HP_ABNORMALITY_OF_T_CELL_PHYSIOLOGY, HP_ABNORMALITY_OF_T_CELL_PHYSIOLOGY MIR3118, MIR3118 MIR134_5P, MIR134_5P
	GOCC_ENDORIBONUCLEASE_COMPLEX, GOCC_ENDORIBONUCLEASE_COMPLEX GSE34156_UNTREATED_VS_6H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP, GSE34156_UNTREATED_VS_6H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP MIR1272, MIR1272 MIR1265, MIR1265 CTCNANGTGNY_UNKNOWN, CTCNANGTGNY_UNKNOWN GOMF_PEPTIDE_LYSINE_N_ACETYLTRANSFERASE_ACTIVITY, GOMF_PEPTIDE_LYSINE_N_ACETYLTRANSFERASE_ACTIVITY MIR764, MIR764
	GOBP_NUCLEOTIDE_EXCISION_REPAIR_PREINCISION_COMPLEX_STABILIZATION, GOBP_NUCLEOTIDE_EXCISION_REPAIR_PREINCISION_COMPLEX_STABILIZATION REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION_INITIATION_FROM_TYPE_3_PROMOTER, REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION_INITIATION_FROM_TYPE_3_PROMOTER HP_PERIVENTRICULAR_LEUKOMALACIA, HP_PERIVENTRICULAR_LEUKOMALACIA GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE GOBP_PSEUDOURIDINE_SYNTHESIS, GOBP_PSEUDOURIDINE_SYNTHESIS MIR4646_3P, MIR4646_3P HEIDENBLAD_AMPLICON_12P11_12_DN, HEIDENBLAD_AMPLICON_12P11_12_DN
	WP_MITOCHONDRIAL_GENE_EXPRESSION, WP_MITOCHONDRIAL_GENE_EXPRESSION MIR125A_3P, MIR125A_3P MIR3173_5P, MIR3173_5P GOBP_HEPARAN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_HEPARAN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS MIR552_3P, MIR552_3P HP_FRONTAL_CORTICAL_ATROPHY, HP_FRONTAL_CORTICAL_ATROPHY GOCC_RNA_POLYMERASE_III_COMPLEX, GOCC_RNA_POLYMERASE_III_COMPLEX
	REACTOME_RNA_POLYMERASE_III_CHAIN_ELONGATION, REACTOME_RNA_POLYMERASE_III_CHAIN_ELONGATION HP_ABNORMAL_SPERM_MOTILITY, HP_ABNORMAL_SPERM_MOTILITY HP_IMPAIRED_SMOOTH_PURSUIT, HP_IMPAIRED_SMOOTH_PURSUIT GSE2706_R848_VS_LPS_8H_STIM_DC_UP, GSE2706_R848_VS_LPS_8H_STIM_DC_UP HP_CONGENITAL_NYSTAGMUS, HP_CONGENITAL_NYSTAGMUS FULLER_PBMC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_336HR_UP, FULLER_PBMC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_336HR_UP GOBP_POSITIVE_REGULATION_OF_MYELOID_LEUKOCYTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MYELOID_LEUKOCYTE_DIFFERENTIATION
	PLASARI_TGFB1_SIGNALING_VIA_NFIC_10HR_DN, PLASARI_TGFB1_SIGNALING_VIA_NFIC_10HR_DN MIR657, MIR657 WEINMANN_ADAPTATION_TO_HYPOXIA_UP, WEINMANN_ADAPTATION_TO_HYPOXIA_UP GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_CORTEX_THYMUS_DN, GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_CORTEX_THYMUS_DN GOBP_STEM_CELL_PROLIFERATION, GOBP_STEM_CELL_PROLIFERATION SHETH_LIVER_CANCER_VS_TXNIP_LOSS_PAM5, SHETH_LIVER_CANCER_VS_TXNIP_LOSS_PAM5 MIR3934_5P, MIR3934_5P
	GOBP_POSITIVE_REGULATION_OF_INTERFERON_BETA_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERFERON_BETA_PRODUCTION  GOBP_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GOBP_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE  WP_HOMOLOGOUS_RECOMBINATION, WP_HOMOLOGOUS_RECOMBINATION  HOLLERN_MICROACINAR_BREAST_TUMOR_DN, HOLLERN_MICROACINAR_BREAST_TUMOR_DN  FULLER_PBMC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_192HR_UP, FULLER_PBMC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_192HR_UP  GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_RRNA, GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_RRNA  GOBP_TRNA_THREONYLCARBAMOYLADENOSINE_METABOLIC_PROCESS, GOBP_TRNA_THREONYLCARBAMOYLADENOSINE_METABOLIC_PROCESS
	GOBP_RESPONSE_TO_RETINOIC_ACID, GOBP_RESPONSE_TO_RETINOIC_ACID HP_ABNORMAL_PLATELET_AGGREGATION, HP_ABNORMAL_PLATELET_AGGREGATION MIR592, MIR592 HP_DOWN_SLOPING_SHOULDERS, HP_DOWN_SLOPING_SHOULDERS HP_RECEPTIVE_LANGUAGE_DELAY, HP_RECEPTIVE_LANGUAGE_DELAY REACTOME_HS_GAG_BIOSYNTHESIS, REACTOME_HS_GAG_BIOSYNTHESIS HP_CHRONIC_LEUKEMIA, HP_CHRONIC_LEUKEMIA
	KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_HEPARAN_SULFATE, KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_HEPARAN_SULFATE MIR769_5P, MIR769_5P HP_ABNORMAL_PULMONARY_THORACIC_IMAGING_FINDING, HP_ABNORMAL_PULMONARY_THORACIC_IMAGING_FINDING MIR1915_5P, MIR1915_5P HP_PULMONARY_INFILTRATES, HP_PULMONARY_INFILTRATES HP_INCREASED_CIRCULATING_IGM_LEVEL, HP_INCREASED_CIRCULATING_IGM_LEVEL DACOSTA_UV_RESPONSE_VIA_ERCC3_XPCS_UP, DACOSTA_UV_RESPONSE_VIA_ERCC3_XPCS_UP
	HP_HYPOTENSION, HP_HYPOTENSION MIR4260, MIR4260 GOCC_NUCLEOTIDE_EXCISION_REPAIR_COMPLEX, GOCC_NUCLEOTIDE_EXCISION_REPAIR_COMPLEX HP_IMPAIRED_T_CELL_FUNCTION, HP_IMPAIRED_T_CELL_FUNCTION INGRAM_SHH_TARGETS_DN, INGRAM_SHH_TARGETS_DN HP_IMPAIRED_TANDEM_GAIT, HP_IMPAIRED_TANDEM_GAIT KIM_MYCN_AMPLIFICATION_TARGETS_UP, KIM_MYCN_AMPLIFICATION_TARGETS_UP
	GOBP_PYRIMIDINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS, GOBP_PYRIMIDINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS  MIR1238_3P, MIR1238_3P  WP_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION, WP_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION  GOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_5_ETS_OF_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA, GOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_5_ETS_OF_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA  GOBP_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_III_PROMOTER, GOBP_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_III_PROMOTER  HEIDENBLAD_AMPLICON_12P11_12_UP, HEIDENBLAD_AMPLICON_12P11_12_UP  GOBP_PIGMENT_GRANULE_ORGANIZATION, GOBP_PIGMENT_GRANULE_ORGANIZATION
	FARMER_BREAST_CANCER_CLUSTER_5, FARMER_BREAST_CANCER_CLUSTER_5  VETTER_TARGETS_OF_PRKCA_AND_ETS1_UP, VETTER_TARGETS_OF_PRKCA_AND_ETS1_UP  PETRETTO_HEART_MASS_QTL_CIS_UP, PETRETTO_HEART_MASS_QTL_CIS_UP  GOBP_DEOXYRIBOSE_PHOSPHATE_CATABOLIC_PROCESS, GOBP_DEOXYRIBOSE_PHOSPHATE_CATABOLIC_PROCESS  GOMF_CARBOXY_LYASE_ACTIVITY, GOMF_CARBOXY_LYASE_ACTIVITY  MIR4322, MIR4322  WP_ATR_SIGNALING, WP_ATR_SIGNALING
	GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_12_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_12_PRODUCTION GOBP_RRNA_PSEUDOURIDINE_SYNTHESIS, GOBP_RRNA_PSEUDOURIDINE_SYNTHESIS MIR4265, MIR4265 PAX5_02, PAX5_02 MIR1226_5P, MIR1226_5P HP_AUDITORY_HALLUCINATIONS, HP_AUDITORY_HALLUCINATIONS
	HP_ABNORMAL_T_CELL_PROLIFERATION, HP_ABNORMAL_T_CELL_PROLIFERATION  HP_INCOMITANT_STRABISMUS, HP_INCOMITANT_STRABISMUS  HP_MEGALOBLASTIC_ANEMIA, HP_MEGALOBLASTIC_ANEMIA  HP_INAPPROPRIATE_LAUGHTER, HP_INAPPROPRIATE_LAUGHTER  HP_BLADDER_DIVERTICULUM, HP_BLADDER_DIVERTICULUM  HP_APLASIA_HYPOPLASIA_OF_THE_EYELID, HP_APLASIA_HYPOPLASIA_OF_THE_EYELID  GOBP_ESTABLISHMENT_OR_MAINTENANCE_OF_BIPOLAR_CELL_POLARITY, GOBP_ESTABLISHMENT_OR_MAINTENANCE_OF_BIPOLAR_CELL_POLARITY  GOOG_DIVA_PERLICATION_EACTOR_A_COMPLEY_COCG_DIVA_PERLICATION_EACTOR_A_COMPLEY
	GOCC_DNA_REPLICATION_FACTOR_A_COMPLEX, GOCC_DNA_REPLICATION_FACTOR_A_COMPLEX HP_MYELOPROLIFERATIVE_DISORDER, HP_MYELOPROLIFERATIVE_DISORDER GOBP_PEPTIDYL_GLUTAMIC_ACID_MODIFICATION, GOBP_PEPTIDYL_GLUTAMIC_ACID_MODIFICATION REACTOME_MATURATION_OF_PROTEIN_3A, REACTOME_MATURATION_OF_PROTEIN_3A GOBP_AXONEMAL_DYNEIN_COMPLEX_ASSEMBLY, GOBP_AXONEMAL_DYNEIN_COMPLEX_ASSEMBLY GOMF_TRNA_GUANINE_METHYLTRANSFERASE_ACTIVITY, GOMF_TRNA_GUANINE_METHYLTRANSFERASE_ACTIVITY GOBP_EXTRACELLULAR_TRANSPORT, GOBP_EXTRACELLULAR_TRANSPORT
	YAP1_DN, YAP1_DN  GOMF_NADH_PYROPHOSPHATASE_ACTIVITY, GOMF_NADH_PYROPHOSPHATASE_ACTIVITY  HP_INCISOR_MACRODONTIA, HP_INCISOR_MACRODONTIA  HP_VISUAL_AGNOSIA, HP_VISUAL_AGNOSIA  GOBP_RESPONSE_TO_MURAMYL_DIPEPTIDE, GOBP_RESPONSE_TO_MURAMYL_DIPEPTIDE  GOMF_TROPONIN_I_BINDING, GOMF_TROPONIN_I_BINDING  REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_DIFFERENTIATION_OF_KERATINOCYTES, REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_DIFFERENTIATION_OF_KERATINOCYTES
	HP_BRAIN_NEOPLASM, HP_BRAIN_NEOPLASM  GOBP_GLYCOSYLCERAMIDE_CATABOLIC_PROCESS, GOBP_GLYCOSYLCERAMIDE_CATABOLIC_PROCESS  GOMF_AMIDINE_LYASE_ACTIVITY, GOMF_AMIDINE_LYASE_ACTIVITY  MIR1180_5P, MIR1180_5P  GOBP_REGULATION_OF_CENTROMERE_COMPLEX_ASSEMBLY, GOBP_REGULATION_OF_CENTROMERE_COMPLEX_ASSEMBLY  HP_SMALL_SCROTUM, HP_SMALL_SCROTUM  GOCC_EXTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_MEMBRANE, GOCC_EXTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_MEMBRANE
	WANG_RESPONSE_TO_BEXAROTENE_UP, WANG_RESPONSE_TO_BEXAROTENE_UP GOBP_PYRIMIDINE_NUCLEOTIDE_CATABOLIC_PROCESS, GOBP_PYRIMIDINE_NUCLEOTIDE_CATABOLIC_PROCESS GOBP_RNA_DECAPPING, GOBP_RNA_DECAPPING REACTOME_UREA_CYCLE, REACTOME_UREA_CYCLE GOBP_CEREBELLAR_PURKINJE_CELL_DIFFERENTIATION, GOBP_CEREBELLAR_PURKINJE_CELL_DIFFERENTIATION SRC_UP.V1_UP, SRC_UP.V1_UP MIR7114_3P, MIR7114_3P
	EHLERS_ANEUPLOIDY_DN, EHLERS_ANEUPLOIDY_DN  HP_APPENDICULAR_HYPOTONIA, HP_APPENDICULAR_HYPOTONIA  GOBP_N_ACYLETHANOLAMINE_METABOLIC_PROCESS, GOBP_N_ACYLETHANOLAMINE_METABOLIC_PROCESS  GAO_STOMACH_24W_C4_PARIETAL_PROGENITOR, GAO_STOMACH_24W_C4_PARIETAL_PROGENITOR  GOBP_TRNA_TRANSCRIPTION, GOBP_TRNA_TRANSCRIPTION  GOBP_DEPYRIMIDINATION, GOBP_DEPYRIMIDINATION  MIR3657, MIR3657
	GOMF_CARBON_NITROGEN_LYASE_ACTIVITY, GOMF_CARBON_NITROGEN_LYASE_ACTIVITY  BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_48HR_UP, BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_48HR_UP  HP_ABNORMAL_RESPIRATORY_MOTILE_CILIUM_PHYSIOLOGY, HP_ABNORMAL_RESPIRATORY_MOTILE_CILIUM_PHYSIOLOGY  GOBP_OUTER_DYNEIN_ARM_ASSEMBLY, GOBP_OUTER_DYNEIN_ARM_ASSEMBLY  AMIT_EGF_RESPONSE_480_MCF10A, AMIT_EGF_RESPONSE_480_MCF10A  HP_PULMONARY_ARTERY_DILATATION, HP_PULMONARY_ARTERY_DILATATION  REACTOME_ORGANIC_CATION_TRANSPORT, REACTOME_ORGANIC_CATION_TRANSPORT
	GOBP_SMALL_NUCLEOLAR_RIBONUCLEOPROTEIN_COMPLEX_ASSEMBLY, GOBP_SMALL_NUCLEOLAR_RIBONUCLEOPROTEIN_COMPLEX_ASSEMBLY WP_BIOMARKERS_FOR_UREA_CYCLE_DISORDERS, WP_BIOMARKERS_FOR_UREA_CYCLE_DISORDERS GOBP_CELLULAR_RESPONSE_TO_OXIDISED_LOW_DENSITY_LIPOPROTEIN_PARTICLE_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_OXIDISED_LOW_DENSITY_LIPOPROTEIN_PARTICLE_STIMULUS GOBP_REGULATION_OF_INOSITOL_TRISPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_INOSITOL_TRISPHOSPHATE_BIOSYNTHETIC_PROCESS HP_ATLANTOAXIAL_ABNORMALITY, HP_ATLANTOAXIAL_ABNORMALITY MIR516A_5P, MIR516A_5P GOBP_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION_INITIATION, GOBP_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION_INITIATION
	GOBP_DNA_PROTECTION, GOBP_DNA_PROTECTION GOCC_PWP2P_CONTAINING_SUBCOMPLEX_OF_90S_PRERIBOSOME, GOCC_PWP2P_CONTAINING_SUBCOMPLEX_OF_90S_PRERIBOSOME GOMF_BILE_ACID_BINDING, GOMF_BILE_ACID_BINDING MODULE_448, MODULE_448 SABATES_COLORECTAL_ADENOMA_SIZE_DN, SABATES_COLORECTAL_ADENOMA_SIZE_DN HP_RECURRENT_UPPER_AND_LOWER_RESPIRATORY_TRACT_INFECTIONS, HP_RECURRENT_UPPER_AND_LOWER_RESPIRATORY_TRACT_INFECTIONS MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_22, MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_22
	HP_FLAT_CORNEA, HP_FLAT_CORNEA HP_OVERFRIENDLINESS, HP_OVERFRIENDLINESS GOBP_SEQUESTERING_OF_ACTIN_MONOMERS, GOBP_SEQUESTERING_OF_ACTIN_MONOMERS MIR6736_5P, MIR6736_5P PODAR_RESPONSE_TO_ADAPHOSTIN_DN, PODAR_RESPONSE_TO_ADAPHOSTIN_DN HP_SUBCORTICAL_HETEROTOPIA, HP_SUBCORTICAL_HETEROTOPIA GOBP_PROTEIN_LOCALIZATION_TO_EARLY_ENDOSOME, GOBP_PROTEIN_LOCALIZATION_TO_EARLY_ENDOSOME
	HP_MILD_POSTNATAL_GROWTH_RETARDATION, HP_MILD_POSTNATAL_GROWTH_RETARDATION HP_TALIPES_EQUINOVALGUS, HP_TALIPES_EQUINOVALGUS HP_NYSTAGMUS_INDUCED_HEAD_NODDING, HP_NYSTAGMUS_INDUCED_HEAD_NODDING HP_ABNORMAL_PLATELET_GRANULES, HP_ABNORMAL_PLATELET_GRANULES GOBP_SMALL_RNA_LOADING_ONTO_RISC, GOBP_SMALL_RNA_LOADING_ONTO_RISC GOBP_PROTEIN_K6_LINKED_UBIQUITINATION, GOBP_PROTEIN_K6_LINKED_UBIQUITINATION GOCC_CHROMATIN_SILENCING_COMPLEX, GOCC_CHROMATIN_SILENCING_COMPLEX
	GOCC_RISC_LOADING_COMPLEX, GOCC_RISC_LOADING_COMPLEX chr14q24 chr14q24, chr14q24 LOPEZ_MESOTELIOMA_SURVIVAL_TIME_DN, LOPEZ_MESOTELIOMA_SURVIVAL_TIME_DN GOBP_NEGATIVE_REGULATION_OF_MOLECULAR_FUNCTION_IN_OTHER_ORGANISM, GOBP_NEGATIVE_REGULATION_OF_MOLECULAR_FUNCTION_IN_OTHER_ORGANISM GOBP_AMYLOID_BETA_CLEARANCE_BY_CELLULAR_CATABOLIC_PROCESS, GOBP_AMYLOID_BETA_CLEARANCE_BY_CELLULAR_CATABOLIC_PROCESS GOBP_GLUCOSYLCERAMIDE_METABOLIC_PROCESS, GOBP_GLUCOSYLCERAMIDE_METABOLIC_PROCESS HP_IMPAIRED_HORIZONTAL_SMOOTH_PURSUIT, HP_IMPAIRED_HORIZONTAL_SMOOTH_PURSUIT
	GOBP_MIRNA_LOADING_ONTO_RISC_INVOLVED_IN_GENE_SILENCING_BY_MIRNA, GOBP_MIRNA_LOADING_ONTO_RISC_INVOLVED_IN_GENE_SILENCING_BY_MIRNA GOBP_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION HP_ONYCHOMYCOSIS, HP_ONYCHOMYCOSIS GOBP_HEPARIN_METABOLIC_PROCESS, GOBP_HEPARIN_METABOLIC_PROCESS GOBP_AZOLE_TRANSMEMBRANE_TRANSPORT, GOBP_AZOLE_TRANSMEMBRANE_TRANSPORT HP_OPEN_ANGLE_GLAUCOMA, HP_OPEN_ANGLE_GLAUCOMA GOCC_HOPS_COMPLEX, GOCC_HOPS_COMPLEX
NEWCASTLE_VIRUS_DC_8H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_	GOMF_NEDD8_TRANSFERASE_ACTIVITY, GOMF_NEDD8_TRANSFERASE_ACTIVITY  HP_ABNORMAL_RESPIRATORY_MOTILE_CILIUM_MORPHOLOGY, HP_ABNORMAL_RESPIRATORY_MOTILE_CILIUM_MORPHOLOGY  MIR1262, MIR1262  HP_SUPRAVENTRICULAR_TACHYCARDIA, HP_SUPRAVENTRICULAR_TACHYCARDIA  GOBP_SEQUESTERING_OF_EXTRACELLULAR_LIGAND_FROM_RECEPTOR, GOBP_SEQUESTERING_OF_EXTRACELLULAR_LIGAND_FROM_RECEPTOR  BIOCARTA_PCAF_PATHWAY, BIOCARTA_PCAF_PATHWAY  GOBP_PROTEIN_POLYGLUTAMYLATION, GOBP_PROTEIN_POLYGLUTAMYLATION
	REACTOME_TRANSLATION_OF_REPLICASE_AND_ASSEMBLY_OF_THE_REPLICATION_TRANSCRIPTION_COMPLEX GOMF_LEUCINE_ZIPPER_DOMAIN_BINDING, GOMF_LEUCINE_ZIPPER_DOMAIN_BINDING chr3q29, chr3q29 GOBP_HISTONE_H3_K14_ACETYLATION, GOBP_HISTONE_H3_K14_ACETYLATION GOBP_CONVERGENT_EXTENSION_INVOLVED_IN_GASTRULATION, GOBP_CONVERGENT_EXTENSION_INVOLVED_IN_GASTRULATION MAP2K1_TARGET_GENES, MAP2K1_TARGET_GENES GOBP_N_ACYLPHOSPHATIDYLETHANOLAMINE_METABOLIC_PROCESS, GOBP_N_ACYLPHOSPHATIDYLETHANOLAMINE_METABOLIC_PROCESS GOBP_N_ACYLPHOSPHATIDYLETHANOLAMINE_METABOLIC_PROCESS, GOBP_N_ACYLPHOSPHATIDYLETHANOLAMINE_METABOLIC_PROCESS GOBP_N_ACYLPHOSPHATIDYLETHANOLAMINE_METABOLIC_PROCESS, GOBP_N_ACYLPHOSPHATIDYLETHANOLAMINE_METABOLIC_PROCESS
	GOBP_POSITIVE_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_POSITIVE_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION  GOMF_LIM_DOMAIN_BINDING, GOMF_LIM_DOMAIN_BINDING  GOBP_REGULATION_OF_MICROTUBULE_NUCLEATION, GOBP_REGULATION_OF_MICROTUBULE_NUCLEATION  GOBP_NEGATIVE_REGULATION_OF_CENTRIOLE_REPLICATION, GOBP_NEGATIVE_REGULATION_OF_CENTRIOLE_REPLICATION  GOCC_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX_CLASS_III, GOCC_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX_CLASS_III  GOMF_ACTIN_MONOMER_BINDING, GOMF_ACTIN_MONOMER_BINDING  GOBP_POSITIVE_REGULATION_OF_INOSITOL_TRISPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_INOSITOL_TRISPHOSPHATE_BIOSYNTHETIC_PROCESS  GOBP_POSITIVE_REGULATION_OF_MESENCHYMAL_STEM_CELL_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_MESENCHYMAL_STEM_CELL_PROLIFERATION
	GOBP_MESENCHYMAL_STEM_CELL_PROLIFERATION, GOBP_MESENCHYMAL_STEM_CELL_PROLIFERATION  HP_ABNORMAL_CIRCULATING_GLUTAMINE_CONCENTRATION, HP_ABNORMAL_CIRCULATING_GLUTAMINE_CONCENTRATION  HP_ARTERIAL_RUPTURE, HP_ARTERIAL_RUPTURE  GOBP_HEPARAN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS_POLYSACCHARIDE_CHAIN_BIOSYNTHETIC_PROCESS, GOBP_HEPARAN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS_POLYSACCHARIDE_CHAIN_BIOSYNTHETIC_PROCESS  GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_DIFFERENTIATION  GOBP_SNRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_III, GOBP_SNRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_III  GOMF_5_3_EXODEOXYRIBONUCLEASE_ACTIVITY, GOMF_5_3_EXODEOXYRIBONUCLEASE_ACTIVITY
	GOBP_TRANSFORMING_GROWTH_FACTOR_BETA1_PRODUCTION, GOBP_TRANSFORMING_GROWTH_FACTOR_BETA1_PRODUCTION GOMF_PROTEIN_GLUTAMIC_ACID_LIGASE_ACTIVITY, GOMF_PROTEIN_GLUTAMIC_ACID_LIGASE_ACTIVITY REACTOME_RUNX3_REGULATES_P14_ARF, REACTOME_RUNX3_REGULATES_P14_ARF HP_SPINAL_CORD_TUMOR, HP_SPINAL_CORD_TUMOR HOEK_NEUTROPHIL_2011_2012_TIV_ADULT_3DY_UP, HOEK_NEUTROPHIL_2011_2012_TIV_ADULT_3DY_UP GOBP_PIGMENT_GRANULE_MATURATION, GOBP_PIGMENT_GRANULE_MATURATION GOBP_BASE_EXCISION_REPAIR_AP_SITE_FORMATION, GOBP_BASE_EXCISION_REPAIR_AP_SITE_FORMATION
	GOCC_H3_HISTONE_ACETYLTRANSFERASE_COMPLEX, GOCC_H3_HISTONE_ACETYLTRANSFERASE_COMPLEX GOMF_HEPARAN_SULFATE_SULFOTRANSFERASE_ACTIVITY, GOMF_HEPARAN_SULFATE_SULFOTRANSFERASE_ACTIVITY ZNF626_TARGET_GENES, ZNF626_TARGET_GENES GOBP_CEREBELLAR_PURKINJE_CELL_LAYER_FORMATION, GOBP_CEREBELLAR_PURKINJE_CELL_LAYER_FORMATION HP_ELFIN_FACIES, HP_ELFIN_FACIES HP_OROTICACIDURIA, HP_OROTICACIDURIA HP_NARROW_FORAMEN_OBTURATORIUM, HP_NARROW_FORAMEN_OBTURATORIUM
	GOBP_CORTICOSPINAL_TRACT_MORPHOGENESIS, GOBP_CORTICOSPINAL_TRACT_MORPHOGENESIS GOCC_CUL5_RING_UBIQUITIN_LIGASE_COMPLEX, GOCC_CUL5_RING_UBIQUITIN_LIGASE_COMPLEX HP_ULNAR_DEVIATION_OF_THE_2ND_FINGER, HP_ULNAR_DEVIATION_OF_THE_2ND_FINGER GOBP_SPLICEOSOMAL_CONFORMATIONAL_CHANGES_TO_GENERATE_CATALYTIC_CONFORMATION, GOBP_SPLICEOSOMAL_CONFORMATIONAL_CHANGES_TO_GENERATE_CATALYTIC_CONFORMATION GOBP_ANTERIOR_COMMISSURE_MORPHOGENESIS, GOBP_ANTERIOR_COMMISSURE_MORPHOGENESIS GOBP_HISTONE_H3_K9_DIMETHYLATION, GOBP_HISTONE_H3_K9_DIMETHYLATION
	REACTOME_SYNTHESIS_OF_WYBUTOSINE_AT_G37_OF_TRNA_PHE, REACTOME_SYNTHESIS_OF_WYBUTOSINE_AT_G37_OF_TRNA_PHE MIKKELSEN_PLURIPOTENT_STATE_UP, MIKKELSEN_PLURIPOTENT_STATE_UP HP_ABNORMAL_TEMPER_TANTRUMS, HP_ABNORMAL_TEMPER_TANTRUMS GOMF_G_PROTEIN_BETA_GAMMA_SUBUNIT_COMPLEX_BINDING, GOMF_G_PROTEIN_BETA_GAMMA_SUBUNIT_COMPLEX_BINDING EBAUER_TARGETS_OF_PAX3_FOXO1_FUSION_DN, EBAUER_TARGETS_OF_PAX3_FOXO1_FUSION_DN HP_HYPEREOSINOPHILIA, HP_HYPEREOSINOPHILIA HP_UPPER_EYELID_EDEMA, HP_UPPER_EYELID_EDEMA
	GOBP_ACTIN_MODIFICATION, GOBP_ACTIN_MODIFICATION  GOCC_TRANSCRIPTION_EXPORT_COMPLEX_2, GOCC_TRANSCRIPTION_EXPORT_COMPLEX_2  HP_INCREASED_C_PEPTIDE_LEVEL, HP_INCREASED_C_PEPTIDE_LEVEL  HP_HYDRANENCEPHALY, HP_HYDRANENCEPHALY  HP_TRICHORRHEXIS_NODOSA, HP_TRICHORRHEXIS_NODOSA  HP_FLAT_NASAL_ALAE, HP_FLAT_NASAL_ALAE  MIR6886_5P, MIR6886_5P
	GOMF_LAMIN_BINDING, GOMF_LAMIN_BINDING  HOEK_T_CELL_2011_2012_TIV_ADULT_1DY_UP, HOEK_T_CELL_2011_2012_TIV_ADULT_1DY_UP  GOMF_AZOLE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_AZOLE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY  MATZUK_SPERMATID_DIFFERENTIATION, MATZUK_SPERMATID_DIFFERENTIATION  HAEGERSTRAND_RESPONSE_TO_IMATINIB, HAEGERSTRAND_RESPONSE_TO_IMATINIB  GOBP_MODULATION_OF_MOLECULAR_FUNCTION_IN_OTHER_ORGANISM, GOBP_MODULATION_OF_MOLECULAR_FUNCTION_IN_OTHER_ORGANISM  VANLOO_SP3_TARGETS_UP, VANLOO_SP3_TARGETS_UP
	WP_UREA_CYCLE_AND_RELATED_DISEASES, WP_UREA_CYCLE_AND_RELATED_DISEASES  GOBP_ETHANOLAMINE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_ETHANOLAMINE_CONTAINING_COMPOUND_METABOLIC_PROCESS  GOBP_NEGATIVE_REGULATION_OF_FEEDING_BEHAVIOR, GOBP_NEGATIVE_REGULATION_OF_FEEDING_BEHAVIOR  HAY_BONE_MARROW_CD34_POS_ERP, HAY_BONE_MARROW_CD34_POS_ERP  GOMF_RACEMASE_AND_EPIMERASE_ACTIVITY, GOMF_RACEMASE_AND_EPIMERASE_ACTIVITY  GOBP_MAINTENANCE_OF_PROTEIN_LOCATION_IN_EXTRACELLULAR_REGION, GOBP_MAINTENANCE_OF_PROTEIN_LOCATION_IN_EXTRACELLULAR_REGION  GOMF_RNA_2_O_METHYLTRANSFERASE_ACTIVITY, GOMF_RNA_2_O_METHYLTRANSFERASE_ACTIVITY
	MIKI_COEXPRESSED_WITH_CYP19A1, MIKI_COEXPRESSED_WITH_CYP19A1 GOBP_REGULATION_OF_GRANULOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_GRANULOCYTE_DIFFERENTIATION HP_ABNORMALITY_OF_THE_EPIDIDYMIS, HP_ABNORMALITY_OF_THE_EPIDIDYMIS GOBP_BONE_TRABECULA_MORPHOGENESIS, GOBP_BONE_TRABECULA_MORPHOGENESIS HP_LONG_HALLUX, HP_LONG_HALLUX HP_KNEE_CLONUS, HP_KNEE_CLONUS HP_DECREASED_SERUM_IRON, HP_DECREASED_SERUM_IRON
	GOBP_MOTOR_LEARNING, GOBP_MOTOR_LEARNING  HP_ABNORMALITY_OF_THE_BRAINSTEM_WHITE_MATTER, HP_ABNORMALITY_OF_THE_BRAINSTEM_WHITE_MATTER  GOBP_PROTEIN_LOCALIZATION_TO_ADHERENS_JUNCTION, GOBP_PROTEIN_LOCALIZATION_TO_ADHERENS_JUNCTION  GOBP_POSITIVE_REGULATION_OF_SENSORY_PERCEPTION_OF_PAIN, GOBP_POSITIVE_REGULATION_OF_SENSORY_PERCEPTION_OF_PAIN  GOBP_NEGATIVE_REGULATION_OF_NON_CANONICAL_WNT_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_NON_CANONICAL_WNT_SIGNALING_PATHWAY  HP_RESPIRATORY_ALKALOSIS, HP_RESPIRATORY_ALKALOSIS  HP_UNUSUAL_SKIN_INFECTION, HP_UNUSUAL_SKIN_INFECTION
	GOBP_ENZYME_DIRECTED_RRNA_PSEUDOURIDINE_SYNTHESIS, GOBP_ENZYME_DIRECTED_RRNA_PSEUDOURIDINE_SYNTHESIS GOBP_CEREBRAL_CORTEX_TANGENTIAL_MIGRATION, GOBP_CEREBRAL_CORTEX_TANGENTIAL_MIGRATION GOBP_5S_CLASS_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_III, GOBP_5S_CLASS_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_III GOCC_TRANSCRIPTION_FACTOR_TFIIIC_COMPLEX, GOCC_TRANSCRIPTION_FACTOR_TFIIIC_COMPLEX GOMF_RNA_POLYMERASE_III_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY, GOMF_RNA_POLYMERASE_III_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_PURPLE_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_PURPLE_DN HP_HYPERACUSIS, HP_HYPERACUSIS
	IVANOV_MUTATED_IN_COLON_CANCER, IVANOV_MUTATED_IN_COLON_CANCER  GOBP_LABYRINTHINE_LAYER_BLOOD_VESSEL_DEVELOPMENT, GOBP_LABYRINTHINE_LAYER_BLOOD_VESSEL_DEVELOPMENT  GOBP_HORMONE_CATABOLIC_PROCESS, GOBP_HORMONE_CATABOLIC_PROCESS  GOCC_OUTER_DYNEIN_ARM, GOCC_OUTER_DYNEIN_ARM  HP_SHOULDER_GIRDLE_MUSCLE_ATROPHY, HP_SHOULDER_GIRDLE_MUSCLE_ATROPHY  GOBP_INTERLEUKIN_1_ALPHA_PRODUCTION, GOBP_INTERLEUKIN_1_ALPHA_PRODUCTION  GOMF_SIRNA_BINDING, GOMF_SIRNA_BINDING
	GOMF_UDP_XYLOSYLTRANSFERASE_ACTIVITY, GOMF_UDP_XYLOSYLTRANSFERASE_ACTIVITY  HP_PHYSICAL_URTICARIA, HP_PHYSICAL_URTICARIA  GOERING_BLOOD_HDL_CHOLESTEROL_QTL_TRANS, GOERING_BLOOD_HDL_CHOLESTEROL_QTL_TRANS  HP_APLASIA_HYPOPLASIA_OF_THE_THYROID_GLAND, HP_APLASIA_HYPOPLASIA_OF_THE_THYROID_GLAND  GOBP_BONE_TRABECULA_FORMATION, GOBP_BONE_TRABECULA_FORMATION  GOMF_CD4_RECEPTOR_BINDING, GOMF_CD4_RECEPTOR_BINDING  GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT_ENDOSOME_TO_PLASMA_MEMBRANE, GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT_ENDOSOME_TO_PLASMA_MEMBRANE
	HP_PRECOCIOUS_PUBERTY_IN_FEMALES, HP_PRECOCIOUS_PUBERTY_IN_FEMALES  GOBP_CELLULAR_RESPONSE_TO_HISTAMINE, GOBP_CELLULAR_RESPONSE_TO_HISTAMINE  GOBP_REGULATION_OF_MICROVILLUS_LENGTH, GOBP_REGULATION_OF_MICROVILLUS_LENGTH  GOBP_REGULATION_OF_CILIUM_BEAT_FREQUENCY_INVOLVED_IN_CILIARY_MOTILITY, GOBP_REGULATION_OF_CILIUM_BEAT_FREQUENCY_INVOLVED_IN_CILIARY_MOTILITY  GOMF_PHOSPHATIDIC_ACID_TRANSFER_ACTIVITY, GOMF_PHOSPHATIDIC_ACID_TRANSFER_ACTIVITY  GOBP_DOUBLE_STRAND_BREAK_REPAIR_VIA_SINGLE_STRAND_ANNEALING, GOBP_DOUBLE_STRAND_BREAK_REPAIR_VIA_SINGLE_STRAND_ANNEALING  GOBP_PERIPHERAL_NERVOUS_SYSTEM_NEURON_DIFFERENTIATION, GOBP_PERIPHERAL_NERVOUS_SYSTEM_NEURON_DIFFERENTIATION
	GOBP_PERIPHERAL_NERVOUS_SYSTEM_NEURON_DIFFERENTIATION, GOBP_PERIPHERAL_NERVOUS_SYSTEM_NEURON_DIFFERENTIATION  MARIADASON_RESPONSE_TO_BUTYRATE_CURCUMIN_SULINDAC_TSA_1, MARIADASON_RESPONSE_TO_BUTYRATE_CURCUMIN_SULINDAC_TSA_1  GOBP_REGULATION_OF_AEROBIC_RESPIRATION, GOBP_REGULATION_OF_AEROBIC_RESPIRATION  GOMF_H3_HISTONE_ACETYLTRANSFERASE_ACTIVITY, GOMF_H3_HISTONE_ACETYLTRANSFERASE_ACTIVITY  NUMATA_CSF3_SIGNALING_VIA_STAT3, NUMATA_CSF3_SIGNALING_VIA_STAT3  chr16q21, chr16q21  GOMF_RACEMASE_AND_EPIMERASE_ACTIVITY_ACTING_ON_CARBOHYDRATES_AND_DERIVATIVES, GOMF_RACEMASE_AND_EPIMERASE_ACTIVITY_ACTING_ON_CARBOHYDRATES_AND_DERIVATIVES  REACTOME_RHOBTB3_ATPASE_CYCLE, REACTOME_RHOBTB3_ATPASE_CYCLE
	chr10p13, chr10p13 GOBP_REGULATION_OF_NON_CANONICAL_WNT_SIGNALING_PATHWAY, GOBP_REGULATION_OF_NON_CANONICAL_WNT_SIGNALING_PATHWAY WP_MIRNAS_INVOLVED_IN_DNA_DAMAGE_RESPONSE, WP_MIRNAS_INVOLVED_IN_DNA_DAMAGE_RESPONSE GOBP_N_ACETYLNEURAMINATE_METABOLIC_PROCESS, GOBP_N_ACETYLNEURAMINATE_METABOLIC_PROCESS HP_LIMB_DYSMETRIA, HP_LIMB_DYSMETRIA GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_POSTSYNAPTIC_MEMBRANE, GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_POSTSYNAPTIC_MEMBRANE GOBP_RESPONSE_TO_DIACYL_BACTERIAL_LIPOPEPTIDE, GOBP_RESPONSE_TO_DIACYL_BACTERIAL_LIPOPEPTIDE
	HP_HIGH_FREQUENCY_SENSORINEURAL_HEARING_IMPAIRMENT, HP_HIGH_FREQUENCY_SENSORINEURAL_HEARING_IMPAIRMENT HP_HYPERGLUTAMINEMIA, HP_HYPERGLUTAMINEMIA HP_IERKY_OCULAR_PURSUIT_MOVEMENTS, HP_IERKY_OCULAR_PURSUIT_MOVEMENTS GOBP_BRADYKININ_CATABOLIC_PROCESS, GOBP_BRADYKININ_CATABOLIC_PROCESS GOBP_ANTEROGRADE_AXONAL_TRANSPORT_OF_MITOCHONDRION, GOBP_ANTEROGRADE_AXONAL_TRANSPORT_OF_MITOCHONDRION GOBP_REGULATION_OF_MEMBRANE_DEPOLARIZATION_DURING_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL, GOBP_REGULATION_OF_MEMBRANE_DEPOLARIZATION_DURING_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL
	GOBP_POSITIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_NEURON_DEATH, GOBP_POSITIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_NEURON_DEATH GOMF_AP_1_ADAPTOR_COMPLEX_BINDING, GOMF_AP_1_ADAPTOR_COMPLEX_BINDING GOMF_INSULIN_BINDING, GOMF_INSULIN_BINDING GOMF_INSULIN_BINDING, GOMF_INSULIN_BINDING GOMF_ADP_RIBOSE_DIPHOSPHATASE_ACTIVITY, GOMF_ADP_RIBOSE_DIPHOSPHATASE_ACTIVITY TURJANSKI_MAPK11_TARGETS, TURJANSKI_MAPK11_TARGETS REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_INTERLEUKIN_SIGNALING, REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_INTERLEUKIN_SIGNALING HP_PROTEIN_AVOIDANCE, HP_PROTEIN_AVOIDANCE
	HP_TRUNCAL_TITUBATION, HP_TRUNCAL_TITUBATION  GOBP_POSITIVE_REGULATION_OF_ALKALINE_PHOSPHATASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_ALKALINE_PHOSPHATASE_ACTIVITY  GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_1_ALPHA_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_1_ALPHA_PRODUCTION  GOBP_POSITIVE_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_CD8_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION  GOBP_ANTEROGRADE_DENDRITIC_TRANSPORT, GOBP_ANTEROGRADE_DENDRITIC_TRANSPORT  GOCC_GRANULAR_COMPONENT, GOCC_GRANULAR_COMPONENT  GOCC_FHF_COMPLEX, GOCC_FHF_COMPLEX
	LOPEZ_MESOTHELIOMA_SURVIVAL_OVERALL_UP, LOPEZ_MESOTHELIOMA_SURVIVAL_OVERALL_UP  GOMF_HYDROXYMETHYL_FORMYL_AND_RELATED_TRANSFERASE_ACTIVITY, GOMF_HYDROXYMETHYL_FORMYL_AND_RELATED_TRANSFERASE_ACTIVITY  HP_VENTRICULAR_SEPTAL_HYPERTROPHY, HP_VENTRICULAR_SEPTAL_HYPERTROPHY  REACTOME_RETINOID_CYCLE_DISEASE_EVENTS, REACTOME_RETINOID_CYCLE_DISEASE_EVENTS  HP_STATUS_EPILEPTICUS_WITH_PROMINENT_MOTOR_SYMPTOMS, HP_STATUS_EPILEPTICUS_WITH_PROMINENT_MOTOR_SYMPTOMS  GOBP_ENDOSOME_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT, GOBP_ENDOSOME_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT  MEISSNER_NPC_HCP_WITH_H3K27ME3, MEISSNER_NPC_HCP_WITH_H3K27ME3
	GOCC_APICAL_CORTEX, GOCC_APICAL_CORTEX GOBP_REGULATION_OF_CELL_PROJECTION_SIZE, GOBP_REGULATION_OF_CELL_PROJECTION_SIZE HP_ABNORMALITY_OF_THE_SUPERIOR_CEREBELLAR_PEDUNCLE, HP_ABNORMALITY_OF_THE_SUPERIOR_CEREBELLAR_PEDUNCLE GOCC_BASEMENT_MEMBRANE_COLLAGEN_TRIMER, GOCC_BASEMENT_MEMBRANE_COLLAGEN_TRIMER AMIT_EGF_RESPONSE_20_HELA, AMIT_EGF_RESPONSE_20_HELA GOBP_LIPID_GLYCOSYLATION, GOBP_LIPID_GLYCOSYLATION HP_SHORT_FACE, HP_SHORT_FACE
	GOMF_HISTAMINE_RECEPTOR_ACTIVITY, GOMF_HISTAMINE_RECEPTOR_ACTIVITY  HP_ABNORMAL_CIRCULATING_ARGININE_CONCENTRATION, HP_ABNORMAL_CIRCULATING_ARGININE_CONCENTRATION  GOBP_MICROGLIA_DIFFERENTIATION, GOBP_MICROGLIA_DIFFERENTIATION  GOBP_REGULATION_OF_MEMBRANE_DEPOLARIZATION_DURING_ACTION_POTENTIAL, GOBP_REGULATION_OF_MEMBRANE_DEPOLARIZATION_DURING_ACTION_POTENTIAL  GOMF_BETA_GLUCOSIDASE_ACTIVITY, GOMF_BETA_GLUCOSIDASE_ACTIVITY  REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_BCR_SIGNALING, REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_BCR_SIGNALING  REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_WNT_SIGNALING, REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_WNT_SIGNALING
	GOBP_HEPARAN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS_ENZYMATIC_MODIFICATION, GOBP_HEPARAN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS_ENZYMATIC_MODIFICATION GOBP_REGULATION_OF_EATING_BEHAVIOR, GOBP_REGULATION_OF_EATING_BEHAVIOR HP_NARROW_INTERNAL_AUDITORY_CANAL, HP_NARROW_INTERNAL_AUDITORY_CANAL HP_CONCAVE_NASAL_RIDGE, HP_CONCAVE_NASAL_RIDGE chr6q16 GOBP_APOPTOTIC_PROCESS_INVOLVED_IN_MORPHOGENESIS, GOBP_APOPTOTIC_PROCESS_INVOLVED_IN_MORPHOGENESIS GOBP_CONNECTIVE_TISSUE_REPLACEMENT, GOBP_CONNECTIVE_TISSUE_REPLACEMENT
	WP_SEROTONIN_AND_ANXIETYRELATED_EVENTS, WP_SEROTONIN_AND_ANXIETYRELATED_EVENTS  ZAIDI_OSTEOBLAST_TRANSCRIPTION_FACTORS, ZAIDI_OSTEOBLAST_TRANSCRIPTION_FACTORS  GOBP_POSITIVE_REGULATION_OF_FEEDING_BEHAVIOR, GOBP_POSITIVE_REGULATION_OF_FEEDING_BEHAVIOR  GOMF_SINGLE_STRANDED_DNA_EXODEOXYRIBONUCLEASE_ACTIVITY, GOMF_SINGLE_STRANDED_DNA_EXODEOXYRIBONUCLEASE_ACTIVITY  GOMF_DYNEIN_HEAVY_CHAIN_BINDING, GOMF_DYNEIN_HEAVY_CHAIN_BINDING  LOPEZ_MESOTHELIOMA_SURVIVAL_WORST_VS_BEST_DN, LOPEZ_MESOTHELIOMA_SURVIVAL_WORST_VS_BEST_DN  GOBP_THIAMINE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_THIAMINE_CONTAINING_COMPOUND_METABOLIC_PROCESS
	GOBP_THIAMINE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_THIAMINE_CONTAINING_COMPOUND_METABOLIC_PROCESS  JOHANSSON_BRAIN_CANCER_EARLY_VS_LATE_UP, JOHANSSON_BRAIN_CANCER_EARLY_VS_LATE_UP  REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_DIFFERENTIATION_OF_MYELOID_CELLS, REACTOME_RUNX1_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_DIFFERENTIATION_OF_MYELOID_CELLS  MIR181A_3P, MIR181A_3P  HP_ABNORMAL_LIVER_PARENCHYMA_MORPHOLOGY, HP_ABNORMAL_LIVER_PARENCHYMA_MORPHOLOGY  GOBP_POSITIVE_REGULATION_OF_ANOIKIS, GOBP_POSITIVE_REGULATION_OF_ANOIKIS  MIR1282, MIR1282  HP_ABNORMAL_SERUM_IRON_CONCENTRATION, HP_ABNORMAL_SERUM_IRON_CONCENTRATION
	HP_ABNORMAL_SERUM_IRON_CONCENTRATION, HP_ABNORMAL_SERUM_IRON_CONCENTRATION  GOBP_NEGATIVE_REGULATION_OF_BEHAVIOR, GOBP_NEGATIVE_REGULATION_OF_BEHAVIOR  GOBP_INNER_DYNEIN_ARM_ASSEMBLY, GOBP_INNER_DYNEIN_ARM_ASSEMBLY  GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT_TO_PLASMA_MEMBRANE, GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT_TO_PLASMA_MEMBRANE  GOMF_SIALYLTRANSFERASE_ACTIVITY, GOMF_SIALYLTRANSFERASE_ACTIVITY  GOBP_RESPONSE_TO_INTERLEUKIN_17, GOBP_RESPONSE_TO_INTERLEUKIN_17  HP_EPILEPSIA_PARTIALIS_CONTINUA, HP_EPILEPSIA_PARTIALIS_CONTINUA  LY_AGING_MIDDLE_UP, LY_AGING_MIDDLE_UP
	LY_AGING_MIDDLE_UP, LY_AGING_MIDDLE_UP  GOBP_CEREBELLAR_CORTEX_FORMATION, GOBP_CEREBELLAR_CORTEX_FORMATION  HP_PAROXYSMAL_SUPRAVENTRICULAR_TACHYCARDIA, HP_PAROXYSMAL_SUPRAVENTRICULAR_TACHYCARDIA  GOBP_ARGININE_BIOSYNTHETIC_PROCESS, GOBP_ARGININE_BIOSYNTHETIC_PROCESS  GOBP_HEAT_ACCLIMATION, GOBP_HEAT_ACCLIMATION  GOBP_SOFT_PALATE_DEVELOPMENT, GOBP_SOFT_PALATE_DEVELOPMENT  GOBP_MUCOCILIARY_CLEARANCE, GOBP_MUCOCILIARY_CLEARANCE  GOBP_INSULIN_METABOLIC_PROCESS, GOBP_INSULIN_METABOLIC_PROCESS
	GOBP_INSULIN_METABOLIC_PROCESS, GOBP_INSULIN_METABOLIC_PROCESS  GOBP_REGULATION_OF_LYMPHANGIOGENESIS, GOBP_REGULATION_OF_LYMPHANGIOGENESIS  GOBP_MELANOSOME_ASSEMBLY, GOBP_MELANOSOME_ASSEMBLY  GOMF_SINGLE_STRANDED_DNA_3_5_EXODEOXYRIBONUCLEASE_ACTIVITY, GOMF_SINGLE_STRANDED_DNA_3_5_EXODEOXYRIBONUCLEASE_ACTIVITY  GOMF_SUMO_POLYMER_BINDING, GOMF_SUMO_POLYMER_BINDING  REACTOME_RUNX1_REGULATES_EXPRESSION_OF_COMPONENTS_OF_TIGHT_JUNCTIONS, REACTOME_RUNX1_REGULATES_EXPRESSION_OF_COMPONENTS_OF_TIGHT_JUNCTIONS  MIR10396A_3P_MIR10396B_3P, MIR10396A_3P_MIR10396B_3P  GOBP_HISTONE_H3_K23_ACETYLATION, GOBP_HISTONE_H3_K23_ACETYLATION
	GOBP_HISTONE_H3_K23_ACETYLATION, GOBP_HISTONE_H3_K23_ACETYLATION  GOMF_CALCIUM_DEPENDENT_PROTEIN_SERINE_THREONINE_PHOSPHATASE_REGULATOR_ACTIVITY, GOMF_CALCIUM_DEPENDENT_PROTEIN_SERINE_THREONINE_PHOSPHATASE_REGULATOR_ACTIVITY  GOMF_HISTONE_ACETYLTRANSFERASE_ACTIVITY_H3_K23_SPECIFIC, GOMF_HISTONE_ACETYLTRANSFERASE_ACTIVITY_H3_K23_SPECIFIC  REACTOME_VITAMIN_B1_THIAMIN_METABOLISM, REACTOME_VITAMIN_B1_THIAMIN_METABOLISM  GOBP_CARDIAC_MUSCLE_TISSUE_REGENERATION, GOBP_CARDIAC_MUSCLE_TISSUE_REGENERATION  GOBP_REGULATION_OF_ALKALINE_PHOSPHATASE_ACTIVITY, GOBP_REGULATION_OF_ALKALINE_PHOSPHATASE_ACTIVITY  GOBP_REGULATION_OF_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL_INVOLVED_IN_REGULATION_OF_CONTRACTION  GOBP_REGULATION_OF_FEEDING_BEHAVIOR, GOBP_REGULATION_OF_FEEDING_BEHAVIOR
	GOBP_REGULATION_OF_FEEDING_BEHAVIOR, GOBP_REGULATION_OF_FEEDING_BEHAVIOR GOMF_METALLOCARBOXYPEPTIDASE_ACTIVITY, GOMF_METALLOCARBOXYPEPTIDASE_ACTIVITY chr17q23, chr17q23 REACTOME_RUNX1_REGULATES_ESTROGEN_RECEPTOR_MEDIATED_TRANSCRIPTION, REACTOME_RUNX1_REGULATES_ESTROGEN_RECEPTOR_MEDIATED_TRANSCRIPTION GOBP_AUTOPHAGY_OF_PEROXISOME, GOBP_AUTOPHAGY_OF_PEROXISOME GOCC_DESMOSOME, GOCC_DESMOSOME VANDESLUIS_COMMD1_TARGETS_GROUP_4_UP, VANDESLUIS_COMMD1_TARGETS_GROUP_4_UP GOMF_ATP_GATED_ION_CHANNEL_ACTIVITY, GOMF_ATP_GATED_ION_CHANNEL_ACTIVITY
	GOBP_RESPONSE_TO_ISOLATION_STRESS, GOBP_RESPONSE_TO_ISOLATION_STRESS GOBP_BUNDLE_OF_HIS_CELL_TO_PURKINJE_MYOCYTE_SIGNALING, GOBP_BUNDLE_OF_HIS_CELL_TO_PURKINJE_MYOCYTE_SIGNALING HP_ABNORMAL_PANCREAS_SIZE, HP_ABNORMAL_PANCREAS_SIZE GOBP_RESPONSE_TO_PEPTIDOGLYCAN, GOBP_RESPONSE_TO_PEPTIDOGLYCAN GOBP_POSITIVE_REGULATION_OF_CYTOKINE_PRODUCTION_INVOLVED_IN_INFLAMMATORY_RESPONSE, GOBP_POSITIVE_REGULATION_OF_CYTOKINE_PRODUCTION_UP, FUJIWARA_PARK2_HEPATOCYTE_PROLIFERATION_UP
	GOBP_BUNDLE_OF_HIS_CELL_TO_PURKINJE_MYOCYTE_COMMUNICATION, GOBP_BUNDLE_OF_HIS_CELL_TO_PURKINJE_MYOCYTE_COMMUNICATION HP_FACIAL_CLEFT, HP_FACIAL_CLEFT MIR6090, MIR6090 chr3p13, chr3p13 REACTOME_RUNX1_AND_FOXP3_CONTROL_THE_DEVELOPMENT_OF_REGULATORY_T_LYMPHOCYTES_TREGS, REACTOME_RUNX1_AND_FOXP3_CONTROL_THE_DEVELOPMENT_OF_REGULATORY_T_LYMPHOCYTES_TREGS