	GSE13485_DAY1_VS_DAY3_YF17D_VACCINE_PBMC_DN, GSE13485_DAY1_VS_DAY3_YF17D_VACCINE_PBMC_DN GSE44649_NAIVE_VS_ACTIVATED_CD8_TCELL_MIR155_KO_DN, GSE44649_NAIVE_VS_ACTIVATED_CD8_TCELL_MIR155_KO_DN GSE17974_CTRL_VS_ACT_IL4_AND_ANTL_IL12_2H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTL_IL12_2H_CD4_TCELL_DN GSE10239_NAIVE_VS_MEMORY_CD8_TCELL_UP, GSE10239_NAIVE_VS_MEMORY_CD8_TCELL_UP GSE17974_0H_VS_1H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_1H_IN_VITRO_ACT_CD4_TCELL_DN GSE13485_DAY1_VS_DAY7_YF17D_VACCINE_PBMC_DN, GSE13485_DAY1_VS_DAY7_YF17D_VACCINE_PBMC_DN GSE17721_LPS_VS_GARDIQUIMOD_24H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_24H_BMDC_DN GSE13484_UNSTIM_VS_12H_YF17D_VACCINE_STIM_PBMC_DN, GSE13484_UNSTIM_VS_12H_YF17D_VACCINE_STIM_PBMC_DN
	GSE13484_UNSTIM_VS_12H_YF17D_VACCINE_STIM_PBMC_DN, GSE13484_UNSTIM_VS_12H_YF17D_VACCINE_STIM_PBMC_DN GSE11864_UNTREATED_VS_CSF1_PAM3CYS_IN_MAC_DN, GSE11864_UNTREATED_VS_CSF1_PAM3CYS_IN_MAC_DN MORF_PPP2R5E, MORF_PPP2R5E GCM_HBP1, GCM_HBP1 GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_1H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_1H_CD4_TCELL_DN GSE18804_BRAIN_VS_COLON_TUMORAL_MACROPHAGE_DN, GSE18804_BRAIN_VS_COLON_TUMORAL_MACROPHAGE_DN GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_0.5H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_0.5H_CD4_TCELL_DN GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_12H_ACT_CD4_TCELL_UP, GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_12H_ACT_CD4_TCELL_UP
	GSE2124_CTRL_VS_LYMPHOTOXIN_BETA_TREATED_MLN_DN, GSE2124_CTRL_VS_LYMPHOTOXIN_BETA_TREATED_MLN_DN MIR520A_5P, MIR520A_5P MIR525_5P, MIR525_5P GSE17974_0H_VS_0.5H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_0.5H_IN_VITRO_ACT_CD4_TCELL_DN MIR802, MIR802 GSE17721_CTRL_VS_POLYIC_24H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_24H_BMDC_DN GSE27291_0H_VS_7D_STIM_GAMMADELTA_TCELL_UP, GSE27291_0H_VS_7D_STIM_GAMMADELTA_TCELL_UP GSE15659_RESTING_VS_ACTIVATED_TREG_DN, GSE15659_RESTING_VS_ACTIVATED_TREG_DN
	GSE25123_CTRL_VS_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_DN, GSE25123_CTRL_VS_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_DN MORF_RAB11A, MORF_RAB11A GSE45365_CD8A_DC_VS_CD11B_DC_IFNAR_KO_DN, GSE45365_CD8A_DC_VS_CD11B_DC_IFNAR_KO_DN ELVIDGE_HIF1A_TARGETS_UP, ELVIDGE_HIF1A_TARGETS_UP MIR6849_3P, MIR6849_3P GSE21927_HEALTHY_VS_TUMOROUS_BALBC_MONOCYTE_UP, GSE21927_HEALTHY_VS_TUMOROUS_BALBC_MOUSE_MONOCYTE_UP MIR1285_3P, MIR1285_3P
	MIR196A_3P, MIR196A_3P MIR1910_5P, MIR1910_5P MIR193A_5P, MIR133A_5P GSE13485_PRE_VS_POST_YF17D_VACCINATION_PBMC_UP, GSE13485_PRE_VS_POST_YF17D_VACCINATION_PBMC_UP GSE17721_CTRL_VS_GARDIQUIMOD_4H_BMDC_DN, GSE17721_CTRL_VS_GARDIQUIMOD_4H_BMDC_DN CAFFAREL_RESPONSE_TO_THC_24HR_5_DN, CAFFAREL_RESPONSE_TO_THC_24HR_5_DN GSE15624_3H_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN, GSE15624_3H_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN
	MIR3187_5P, MIR3187_5P MIR5189_5P, MIR5189_5P MIR4793_3P, MIR4793_3P MIR610, MIR610 GCM_BAG5, GCM_BAG5 MIR3166, MIR3166 MIR3166 MIR4699_5P, MIR4699_5P GOBP_HISTONE_H3_K4_METHYLATION, GOBP_HISTONE_H3_K4_METHYLATION
	REICHERT_MITOSIS_LIN9_TARGETS, REICHERT_MITOSIS_LIN9_TARGETS ATF6_01, ATF6_01 MIR556_3P, MIR556_3P GSE18281_SUBCAPSULAR_VS_PERIMEDULLARY_CORTICAL_REGION_OF_THYMUS_DN, GSE18281_SUBCAPSULAR_VS_PERIMEDULLARY_CORTICAL_REGION_OF_THYMUS_DN GOBP_MEIOSIS_I_CELL_CYCLE_PROCESS, GOBP_MEIOSIS_I_CELL_CYCLE_PROCESS MIR4519, MIR4519 MIR216A_5P, MIR216A_5P
	MIR549A_3P GOBP_ERROR_FREE_TRANSLESION_SYNTHESIS, GOBP_ERROR_FREE_TRANSLESION_SYNTHESIS MIR630, MIR630 REACTOME_INOSITOL_PHOSPHATE_METABOLISM, REACTOME_INOSITOL_PHOSPHATE_METABOLISM ELVIDGE_HIF1A_AND_HIF2A_TARGETS_UP, ELVIDGE_HIF1A_AND_HIF2A_TARGETS_UP MIR147A, MIR147A MIR6796_3P, MIR6796_3P SMITH_LIVER_CANCER, SMITH_LIVER_CANCER
	GOBP_ENDOPLASMIC_RETICULUM_TO_CYTOSOL_TRANSPORT, GOBP_ENDOPLASMIC_RETICULUM_TO_CYTOSOL_TRANSPORT GSE13485_CTRL_VS_DAY3_YF17D_VACCINE_PBMC_UP, GSE13485_CTRL_VS_DAY3_YF17D_VACCINE_PBMC_UP MIR7153_3P, MIR7153_3P GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMC_DN, GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMC_DN GAZDA_DIAMOND_BLACKFAN_ANEMIA_PROGENITOR_UP, GAZDA_DIAMOND_BLACKFAN_ANEMIA_PROGENITOR_UP KENNY_CTNNB1_TARGETS_UP, KENNY_CTNNB1_TARGETS_UP MIR1228_3P, MIR1228_3P
	MIR100_3P, MIR100_3P MIR647, MIR647 HP_ABNORMALLY_FOLDED_HELIX, HP_ABNORMALLY_FOLDED_HELIX GSE29618_PRE_VS_DAY7_FLU_VACCINE_BCELL_UP, GSE29618_PRE_VS_DAY7_FLU_VACCINE_BCELL_UP NIKOLSKY_BREAST_CANCER_20Q12_Q13_AMPLICON, NIKOLSKY_BREAST_CANCER_20Q12_Q13_AMPLICON MIR3912_5P, MIR3912_5P MIR3164, MIR3164 MIR3142, MIR3142
	GOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_PHOSPHATASE_ACTIVITY HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_56D_TOP_100_DEG_AFTER_IN_VITRO_RE_STIMULATION_DN, HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_56D_TOP_100_DEG_AFTER_IN_VITRO_Chr15q15 MIR6820_3P, MIR6820_3P PID_AR_TF_PATHWAY, PID_AR_TF_PATHWAY LOPEZ_MESOTELIOMA_SURVIVAL_TIME_UP, LOPEZ_MESOTELIOMA_SURVIVAL_TIME_UP MIR197_3P, MIR197_3P MIR197_3P, MIR197_3P MIR411_5P, MIR411_5P
	GOBP_FIBROBLAST_MIGRATION, GOBP_FIBROBLAST_MIGRATION MIR6807_5P, MIR6807_5P MIR668_3P, MIR668_3P ACATATC_MIR190, ACATATC_MIR190 GOBP_FEMALE_MEIOTIC_NUCLEAR_DIVISION, GOBP_FEMALE_MEIOTIC_NUCLEAR_DIVISION GSE45365_NK_CELL_VS_CD8A_DC_DN, GSE45365_NK_CELL_VS_CD8A_DC_DN MIR4786_5P, MIR4786_5P BIOCARTA_GLEEVEC_PATHWAY, BIOCARTA_GLEEVEC_PATHWAY
	GOMF_UBIQUITIN_LIKE_PROTEIN_SPECIFIC_PROTEASE_ACTIVITY, GOMF_UBIQUITIN_LIKE_PROTEIN_SPECIFIC_PROTEASE_ACTIVITY MIR6776_3P, MIR6776_3P HP_HAPPY_DEMEANOR, HP_HAPPY_DEMEANOR GOBP_NEGATIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT, GOBP_NEGATIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT HP_ABNORMAL_LARGE_INTESTINE_PHYSIOLOGY, HP_ABNORMAL_LARGE_INTESTINE_PHYSIOLOGY HP_ABNORMAL_CIRCULATING_BILIRUBIN_CONCENTRATION, HP_ABNORMAL_CIRCULATING_BILIRUBIN_CONCENTRATION GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_DN, GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_DN GOCC_ENDOPLASMIC_RETICULUM_QUALITY_CONTROL_COMPARTMENT, GOCC_ENDOPLASMIC_RETICULUM_QUALITY_CONTROL_COMPARTMENT
	GOBP_NEGATIVE_REGULATION_OF_INTRACELLULAR_TRANSPORT, GOBP_NEGATIVE_REGULATION_OF_INTRACELLULAR_TRANSPORT MIR921, MIR921 MIR4714_5P, MIR4714_5P MIR6750_5P, MIR6750_5P GOMF_K63_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GOMF_K63_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING MIR3665, MIR3665 GOCC_COPII_VESICLE_COAT, GOCC_COPII_VESICLE_COAT GOBP_REGULATION_OF_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM, GOBP_REGULATION_OF_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM
	BOYAULT_LIVER_CANCER_SUBCLASS_G2, BOYAULT_LIVER_CANCER_SUBCLASS_G2 BIOCARTA_TNFR1_PATHWAY, BIOCARTA_TNFR1_PATHWAY REACTOME_DISEASES_ASSOCIATED_WITH_N_GLYCOSYLATION_OF_PROTEINS, REACTOME_DISEASES_ASSOCIATED_WITH_N_GLYCOSYLATION_OF_PROTEINS MIR6513_5P, MIR6513_5P SEIDEN_MET_SIGNALING, SEIDEN_MET_SIGNALING MIR581, MIR581 REACTOME_SYNTHESIS_OF_IP2_IP_AND_INS_IN_THE_CYTOSOL, REACTOME_SYNTHESIS_OF_IP2_IP_AND_INS_IN_THE_CYTOSOL GOMF_CULLIN_FAMILY_PROTEIN_BINDING, GOMF_CULLIN_FAMILY_PROTEIN_BINDING
	MODULE_226, MODULE_226 MIR3126_3P, MIR3126_3P WP_GLYCOSYLATION_AND_RELATED_CONGENITAL_DEFECTS, WP_GLYCOSYLATION_AND_RELATED_CONGENITAL_DEFECTS MIR4320, MIR4320 HP_CAPILLARY_HEMANGIOMA, HP_CAPILLARY_HEMANGIOMA WP_SPHINGOLIPID_PATHWAY, WP_SPHINGOLIPID_PATHWAY CCAWWNAAGG_SRF_Q4, CCAWWNAAGG_SRF_Q4
	MODULE_372, MODULE_372 ERWIN_COHEN_BLOOD_VACCINE_TC_83_AGE_23_48YO_VACCINATED_VS_CONTROL_2DY_DN, ERWIN_COHEN_BLOOD_VACCINE_TC_83_AGE_23_48YO_VACCINATED_VS_CONTROL_2DY_DN HP_MEGALOCORNEA, HP_MEGALOCORNEA HP_POOR_COORDINATION, HP_POOR_COORDINATION GOBP_ENDOPLASMIC_RETICULUM_TUBULAR_NETWORK_ORGANIZATION, GOBP_ENDOPLASMIC_RETICULUM_TUBULAR_NETWORK_ORGANIZATION NEMETH_INFLAMMATORY_RESPONSE_LPS_DN, NEMETH_INFLAMMATORY_RESPONSE_LPS_DN AMIT_EGF_RESPONSE_40_MCF10A, AMIT_EGF_RESPONSE_40_MCF10A GTAGGCA_MIR189, GTAGGCA_MIR189
	MIR10400_3P_MIR4674, MIR10400_3P_MIR4674 MIR3157_3P, MIR3157_3P MIR6822_5P, MIR6822_5P GOBP_NUCLEAR_PORE_ORGANIZATION, GOBP_NUCLEAR_PORE_ORGANIZATION GOBP_REGULATION_OF_CLATHRIN_DEPENDENT_ENDOCYTOSIS, GOBP_REGULATION_OF_CLATHRIN_DEPENDENT_ENDOCYTOSIS REACTOME_POLB_DEPENDENT_LONG_PATCH_BASE_EXCISION_REPAIR, REACTOME_POLB_DEPENDENT_LONG_PATCH_BASE_EXCISION_REPAIR MIR4683, MIR4683 MIR4289, MIR4289
	HP_ABNORMAL_CIRCULATING_GONADOTROPIN_LEVEL, HP_ABNORMAL_CIRCULATING_GONADOTROPIN_LEVEL GOBP_T_CELL_HOMEOSTASIS, GOBP_T_CELL_HOMEOSTASIS GOMF_ACID_PHOSPHATASE_ACTIVITY, GOMF_ACID_PHOSPHATASE_ACTIVITY MIR10B_3P, MIR10B_3P SIG_REGULATION_OF_THE_ACTIN_CYTOSKELETON_BY_RHO_GTPASES, SIG_REGULATION_OF_THE_ACTIN_CYTOSKELETON_BY_RHO_GTPASES chr9q32, chr9q32 BIOCARTA_VIP_PATHWAY, BIOCARTA_VIP_PATHWAY
	WP_IL1_AND_MEGAKARYOCYTES_IN_OBESITY, WP_IL1_AND_MEGAKARYOCYTES_IN_OBESITY GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H3_K4_SPECIFIC, GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H3_K4_SPECIFIC GOBP_NUCLEAR_MEMBRANE_ORGANIZATION, GOBP_NUCLEAR_MEMBRANE_ORGANIZATION GOMF_PHOSPHATIDYLINOSITOL_MONOPHOSPHATE_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_MONOPHOSPHATE_PHOSPHATASE_ACTIVITY MIR490_5P, MIR490_5P HP_INCREASED_CIRCULATING_GONADOTROPIN_LEVEL, HP_INCREASED_CIRCULATING_GONADOTROPIN_LEVEL BIOCARTA_CDC42RAC_PATHWAY, BIOCARTA_CDC42RAC_PATHWAY GOBP_OOCYTE_MATURATION, GOBP_OOCYTE_MATURATION
	GOBP_PROTEIN_DNA_COVALENT_CROSS_LINKING_REPAIR, GOBP_PROTEIN_DNA_COVALENT_CROSS_LINKING_REPAIR PID_TCR_JNK_PATHWAY, PID_TCR_JNK_PATHWAY HP_STREAK_OVARY, HP_STREAK_OVARY GOBP_OOCYTE_DIFFERENTIATION, GOBP_OOCYTE_DIFFERENTIATION GOCC_NUCLEAR_LAMINA, GOCC_NUCLEAR_LAMINA GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_DN, GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_DN WP_SPHINGOLIPID_METABOLISM_INTEGRATED_PATHWAY, WP_SPHINGOLIPID_METABOLISM_INTEGRATED_PATHWAY FIGUEROA_AML_METHYLATION_CLUSTER_3_DN, FIGUEROA_AML_METHYLATION_CLUSTER_3_DN
	GOBP_DNA_CATABOLIC_PROCESS, GOBP_DNA_CATABOLIC_PROCESS GOBP_FEMALE_MEIOSIS_I, GOBP_FEMALE_MEIOSIS_I GOMF_INOSITOL_TETRAKISPHOSPHATASE_ACTIVITY, GOMF_INOSITOL_TETRAKISPHOSPHATASE_ACTIVITY GOMF_INOSITOL_TETRAKISPHOSPHATASE_ACTIVITY, GOMF_INOSITOL_TETRAKISPHOSPHATASE_ACTIVITY GOBP_TROPHOBLAST_CELL_MIGRATION, GOBP_TROPHOBLAST_CELL_MIGRATION GOMF_STEROID_HORMONE_RECEPTOR_ACTIVITY, GOMF_STEROID_HORMONE_RECEPTOR_ACTIVITY WP_WHITE_FAT_CELL_DIFFERENTIATION, WP_WHITE_FAT_CELL_DIFFERENTIATION GOBP_SKELETAL_MUSCLE_ACETYLCHOLINE_GATED_CHANNEL_CLUSTERING, GOBP_SKELETAL_MUSCLE_ACETYLCHOLINE_GATED_CHANNEL_CLUSTERING
	GOBP_NEGATIVE_REGULATION_OF_MEIOTIC_NUCLEAR_DIVISION, GOBP_NEGATIVE_REGULATION_OF_MEIOTIC_NUCLEAR_DIVISION GOBP_POSITIVE_REGULATION_OF_SISTER_CHROMATID_COHESION, GOBP_POSITIVE_REGULATION_OF_SISTER_CHROMATID_COHESION REACTOME_SMALL_INTERFERING_RNA_SIRNA_BIOGENESIS, REACTOME_SMALL_INTERFERING_RNA_SIRNA_BIOGENESIS GOBP_N_TERMINAL_PEPTIDYL_METHIONINE_ACETYLATION, GOBP_N_TERMINAL_PEPTIDYL_METHIONINE_ACETYLATION GOBP_GOLGI_TO_VACUOLE_TRANSPORT, GOBP_GOLGI_TO_VACUOLE_TRANSPORT GOBP_DNA_CATABOLIC_PROCESS_ENDONUCLEOLYTIC, GOBP_DNA_CATABOLIC_PROCESS_ENDONUCLEOLYTIC WP_SPHINGOLIPID_METABOLISM_GENERAL_OVERVIEW, WP_SPHINGOLIPID_METABOLISM_GENERAL_OVERVIEW WP_INTRACELLULAR_TRAFFICKING_PROTEINS_INVOLVED_IN_CMT_NEUROPATHY, WP_INTRACELLULAR_TRAFFICKING_PROTEINS_INVOLVED_IN_CMT_NEUROPATHY
	GOBP_REGULATION_OF_RETROGRADE_PROTEIN_TRANSPORT_ER_TO_CYTOSOL, GOBP_REGULATION_OF_RETROGRADE_PROTEIN_TRANSPORT_ER_TO_CYTOSOL GOMF_INOSITOL_PHOSPHATE_PHOSPHATASE_ACTIVITY, GOMF_INOSITOL_PHOSPHATE_PHOSPHATASE_ACTIVITY MIR616_3P, MIR616_3P HP_SEGMENTAL_PERIPHERAL_DEMYELINATION_REMYELINATION, HP_SEGMENTAL_PERIPHERAL_DEMYELINATION_REMYELINATION GOMF_PEPTIDE_ALPHA_N_ACETYLTRANSFERASE_ACTIVITY, GOMF_PEPTIDE_ALPHA_N_ACETYLTRANSFERASE_ACTIVITY HP_ABNORMAL_CIRCULATING_ACETYLCARNITINE_CONCENTRATION, HP_ABNORMAL_CIRCULATING_ACETYLCARNITINE_CONCENTRATION GOCC_N_TERMINAL_PROTEIN_ACETYLTRANSFERASE_COMPLEX, GOCC_N_TERMINAL_PROTEIN_ACETYLTRANSFERASE_COMPLEX GOCC_CHROMOSOME_PASSENGER_COMPLEX, GOCC_CHROMOSOME_PASSENGER_COMPLEX
	GOBP_REGULATION_OF_PROTEIN_MONOUBIQUITINATION, GOBP_REGULATION_OF_PROTEIN_MONOUBIQUITINATION HP_ABSENT_AXILLARY_HAIR, HP_ABSENT_AXILLARY_HAIR ACGCACA_MIR210, ACGCACA_MIR210 MIR6503_3P, MIR6503_3P BIOCARTA_SODD_PATHWAY, BIOCARTA_SODD_PATHWAY GOBP_PHOSPHATIDYLGLYCEROL_BIOSYNTHETIC_PROCESS, GOBP_PHOSPHATIDYLGLYCEROL_BIOSYNTHETIC_PROCESS STANELLE_E2F1_TARGETS, STANELLE_E2F1_TARGETS GOCC_PRONUCLEUS, GOCC_PRONUCLEUS
	GOBP_NEGATIVE_REGULATION_OF_CHROMATIN_BINDING, GOBP_NEGATIVE_REGULATION_OF_CHROMATIN_BINDING GOCC_ACTIVIN_RECEPTOR_COMPLEX, GOCC_ACTIVIN_RECEPTOR_COMPLEX GOBP_CELLULAR_SENESCENCE, GOBP_CELLULAR_SENESCENCE GOBP_MEIOTIC_SPINDLE_ORGANIZATION, GOBP_MEIOTIC_SPINDLE_ORGANIZATION GOBP_UV_PROTECTION, GOBP_UV_PROTECTION GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSPORT BIOCARTA_SALMONELLA_PATHWAY, BIOCARTA_SALMONELLA_PATHWAY
	HP_BROAD_FEMORAL_NECK, HP_BROAD_FEMORAL_NECK HP_NEONATAL_ASPHYXIA, HP_NEONATAL_ASPHYXIA GOBP_PROTEIN_DESUMOYLATION, GOBP_PROTEIN_DESUMOYLATION GOCC_CDC73_PAF1_COMPLEX, GOCC_CDC73_PAF1_COMPLEX GOCC_VCP_NPL4_UFD1_AAA_ATPASE_COMPLEX, GOCC_VCP_NPL4_UFD1_AAA_ATPASE_COMPLEX GOCC_MRNA_EDITING_COMPLEX, GOCC_MRNA_EDITING_COMPLEX GOMF_SUMO_SPECIFIC_PROTEASE_ACTIVITY, GOMF_SUMO_SPECIFIC_PROTEASE_ACTIVITY GOBP_REGULATION_OF_LYMPHOCYTE_MIGRATION, GOBP_REGULATION_OF_LYMPHOCYTE_MIGRATION
	GOCC_MEIOTIC_SPINDLE, GOCC_MEIOTIC_SPINDLE CHEOK_RESPONSE_TO_MERCAPTOPURINE_AND_LD_MTX_UP, CHEOK_RESPONSE_TO_MERCAPTOPURINE_AND_LD_MTX_UP GOMF_H4_HISTONE_ACETYLTRANSFERASE_ACTIVITY, GOMF_H4_HISTONE_ACETYLTRANSFERASE_ACTIVITY CHASSOT_SKIN_WOUND, CHASSOT_SKIN_WOUND HP_VACUOLATED_LYMPHOCYTES, HP_VACUOLATED_LYMPHOCYTES HOEK_MONOCYTE_2011_2012_TIV_ADULT_7DY_UP, HOEK_MONOCYTE_2011_2012_TIV_ADULT_7DY_UP GOBP_DETECTION_OF_MOLECULE_OF_BACTERIAL_ORIGIN, GOBP_DETECTION_OF_MOLECULE_OF_BACTERIAL_ORIGIN HP_HYPERTROPHIC_NERVE_CHANGES, HP_HYPERTROPHIC_NERVE_CHANGES
	ZNF211_TARGET_GENES, ZNF211_TARGET_GENES GOCC_RAGULATOR_COMPLEX, GOCC_RAGULATOR_COMPLEX GOCC_GATOR1_COMPLEX, GOCC_GATOR1_COMPLEX GOCC_GATOR1_COMPLEX, GOCC_GATOR1_COMPLEX GOMF_SHORT_CHAIN_CARBOXYLESTERASE_ACTIVITY, GOMF_SHORT_CHAIN_CARBOXYLESTERASE_ACTIVITY GOBP_HISTONE_ARGININE_METHYLATION, GOBP_HISTONE_ARGININE_METHYLATION HP_TRUNCUS_ARTERIOSUS, HP_TRUNCUS_ARTERIOSUS MIR711, MIR711 GOBP_POSTSYNAPTIC_SIGNAL_TRANSDUCTION, GOBP_POSTSYNAPTIC_SIGNAL_TRANSDUCTION
	HP_GOWERS_SIGN, HP_GOWERS_SIGN RAMJAUN_APOPTOSIS_BY_TGFB1_VIA_MAPK1_DN, RAMJAUN_APOPTOSIS_BY_TGFB1_VIA_MAPK1_DN GOBP_FOLIC_ACID_TRANSPORT, GOBP_FOLIC_ACID_TRANSPORT GOBP_POSITIVE_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT GOBP_MODIFICATION_OF_POSTSYNAPTIC_STRUCTURE, GOBP_MODIFICATION_OF_POSTSYNAPTIC_STRUCTURE MIR758_5P, MIR758_5P GOBP_BONE_MINERALIZATION_INVOLVED_IN_BONE_MATURATION, GOBP_BONE_MINERALIZATION_INVOLVED_IN_BONE_MATURATION
	HP_MULTIPLE_SUTURE_CRANIOSYNOSTOSIS, HP_MULTIPLE_SUTURE_CRANIOSYNOSTOSIS OSAWA_TNF_TARGETS, OSAWA_TNF_TARGETS GOMF_ALDEHYDE_LYASE_ACTIVITY, GOMF_ALDEHYDE_LYASE_ACTIVITY GOMF_ACTIVIN_ACTIVATED_RECEPTOR_ACTIVITY, GOMF_ACTIVIN_ACTIVATED_RECEPTOR_ACTIVITY HP_RESPIRATORY_INSUFFICIENCY_DUE_TO_DEFECTIVE_CILIARY_CLEARANCE, HP_RESPIRATORY_INSUFFICIENCY_DUE_TO_DEFECTIVE_CILIARY_CLEARANCE REACTOME_MET_RECEPTOR_RECYCLING, REACTOME_MET_RECEPTOR_RECYCLING GOBP_BRANCHED_CHAIN_AMINO_ACID_TRANSPORT, GOBP_BRANCHED_CHAIN_AMINO_ACID_TRANSPORT GOBP_PEPTIDYL_LYSINE_HYDROXYLATION, GOBP_PEPTIDYL_LYSINE_HYDROXYLATION
DAY1_YF17D_VACCINE_PBMC_UP, GSE13485_CTRL_VS_DAY1_YF17D_VACCINE_PBMC_UP	GOMF_AP_2_ADAPTOR_COMPLEX_BINDING, GOMF_AP_2_ADAPTOR_COMPLEX_BINDING MIR4746_5P, MIR4746_5P GOBP_PEPTIDYL_ARGININE_OMEGA_N_METHYLATION, GOBP_PEPTIDYL_ARGININE_OMEGA_N_METHYLATION GOMF_RRNA_ADENINE_METHYLTRANSFERASE_ACTIVITY, GOMF_RRNA_ADENINE_METHYLTRANSFERASE_ACTIVITY GOBP_COBALT_ION_TRANSPORT, GOBP_COBALT_ION_TRANSPORT GOMF_METALLODIPEPTIDASE_ACTIVITY, GOMF_METALLODIPEPTIDASE_ACTIVITY REACTOME_PP2A_MEDIATED_DEPHOSPHORYLATION_OF_KEY_METABOLIC_FACTORS, REACTOME_PP2A_MEDIATED_DEPHOSPHORYLATION_OF_KEY_METABOLIC_FACTORS chr2p16, chr2p16
	HP_DUPLICATION_OF_THE_DISTAL_PHALANX_OF_THE_THUMB, HP_DUPLICATION_OF_THE_DISTAL_PHALANX_OF_THE_THUMB GOMF_SIGNAL_RECOGNITION_PARTICLE_BINDING, GOMF_SIGNAL_RECOGNITION_PARTICLE_BINDING GOMF_BRANCHED_CHAIN_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_BRANCHED_CHAIN_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY GOBP_SKELETAL_MUSCLE_SATELLITE_CELL_ACTIVATION, GOBP_SKELETAL_MUSCLE_SATELLITE_CELL_ACTIVATION GOBP_DOUBLE_STRAND_BREAK_REPAIR_VIA_SYNTHESIS_DEPENDENT_STRAND_ANNEALING, GOBP_DOUBLE_STRAND_BREAK_REPAIR_VIA_SYNTHESIS_DEPENDENT_STRAND_ANNEALING GOBP_REGULATION_BY_VIRUS_OF_VIRAL_PROTEIN_LEVELS_IN_HOST_CELL, GOBP_REGULATION_BY_VIRUS_OF_VIRAL_PROTEIN_LEVELS_IN_HOST_CELL GOCC_NSL_COMPLEX, GOCC_NSL_COMPLEX GOCC_PTW_PP1_PHOSPHATASE_COMPLEX, GOCC_PTW_PP1_PHOSPHATASE_COMPLEX
	GOMF_HISTONE_ARGININE_N_METHYLTRANSFERASE_ACTIVITY, GOMF_HISTONE_ARGININE_N_METHYLTRANSFERASE_ACTIVITY HP_BRANCHIAL_ANOMALY, HP_BRANCHIAL_ANOMALY TOMLINS_METASTASIS_DN, TOMLINS_METASTASIS_DN ABE_VEGFA_TARGETS_30MIN, ABE_VEGFA_TARGETS_30MIN MIR3691_5P, MIR3691_5P PID_NFAT_TFPATHWAY, PID_NFAT_TFPATHWAY TAGHAVI_NEOPLASTIC_TRANSFORMATION, TAGHAVI_NEOPLASTIC_TRANSFORMATION GOBP_INOSITOL_METABOLIC_PROCESS, GOBP_INOSITOL_METABOLIC_PROCESS
	KONG_E2F1_TARGETS, KONG_E2F1_TARGETS MATHEW_FANCONI_ANEMIA_GENES, MATHEW_FANCONI_ANEMIA_GENES GOBP_SECRETORY_GRANULE_ORGANIZATION, GOBP_SECRETORY_GRANULE_ORGANIZATION GOBP_NUCLEAR_PORE_COMPLEX_ASSEMBLY, GOBP_NUCLEAR_PORE_COMPLEX_ASSEMBLY GOBP_VESICLE_TETHERING, GOBP_VESICLE_TETHERING GOMF_CARBOHYDRATE_PROTON_SYMPORTER_ACTIVITY, GOMF_CARBOHYDRATE_PROTON_SYMPORTER_ACTIVITY GOMF_TRIPLET_CODON_AMINO_ACID_ADAPTOR_ACTIVITY, GOMF_TRIPLET_CODON_AMINO_ACID_ADAPTOR_ACTIVITY GOBP_POSITIVE_REGULATION_OF_GROWTH_HORMONE_SECRETION, GOBP_POSITIVE_REGULATION_OF_GROWTH_HORMONE_SECRETION
	GOBP_REGULATION_OF_CELL_AGING, GOBP_REGULATION_OF_CELL_AGING GOMF_ACYLGLYCEROL_LIPASE_ACTIVITY, GOMF_ACYLGLYCEROL_LIPASE_ACTIVITY GOMF_ACTIVIN_RECEPTOR_ACTIVITY_TYPE_I, GOMF_ACTIVIN_RECEPTOR_ACTIVITY_TYPE_I MIR4636, MIR4636 GOBP_PEPTIDYL_ARGININE_METHYLATION_TO_ASYMMETRICAL_DIMETHYL_ARGININE, GOBP_PEPTIDYL_ARGININE_METHYLATION_TO_ASYMMETRICAL_DIMETHYL_ARGININE GOBP_POSITIVE_REGULATION_OF_PROTEIN_MONOUBIQUITINATION, GOBP_POSITIVE_REGULATION_OF_CLATHRIN_DEPENDENT_ENDOCYTOSIS
	GOMF_PROTEIN_ARGININE_OMEGA_N_MONOMETHYLTRANSFERASE_ACTIVITY, GOMF_PROTEIN_ARGININE_OMEGA_N_MONOMETHYLTRANSFERASE_ACTIVITY GOMF_PROTEIN_ARGININE_OMEGA_N_ASYMMETRIC_METHYLTRANSFERASE_ACTIVITY, GOMF_PROTEIN_ARGININE_OMEGA_N_ASYMMETRIC_METHYLTRANSFERASE_ACTIVITY HP_DISTAL_JOINT_LAXITY, HP_DISTAL_JOINT_LAXITY GOBP_WOUND_HEALING_INVOLVED_IN_INFLAMMATORY_RESPONSE, GOBP_WOUND_HEALING_INVOLVED_IN_INFLAMMATORY_RESPONSE GOBP_NEGATIVE_REGULATION_OF_TROPHOBLAST_CELL_MIGRATION, GOBP_NEGATIVE_REGULATION_OF_TROPHOBLAST_CELL_MIGRATION KONDO_HYPOXIA, KONDO_HYPOXIA HP_PERIPHERAL_HYPOMYELINATION, HP_PERIPHERAL_HYPOMYELINATION GOBP_CARDIOLIPIN_BIOSYNTHETIC_PROCESS, GOBP_CARDIOLIPIN_BIOSYNTHETIC_PROCESS
	GOBP_VESICLE_FUSION_WITH_GOLGI_APPARATUS, GOBP_VESICLE_FUSION_WITH_GOLGI_APPARATUS GOCC_MLL3_4_COMPLEX, GOCC_MLL3_4_COMPLEX HP_TICS, HP_TICS HP_ABNORMAL_CIRCULATING_FOLLICLE_STIMULATING_HORMONE_LEVEL, HP_ABNORMAL_CIRCULATING_FOLLICLE_STIMULATING_HORMONE_LEVEL HP_CONJUGATED_HYPERBILIRUBINEMIA, HP_CONJUGATED_HYPERBILIRUBINEMIA REACTOME_REACTIONS_SPECIFIC_TO_THE_COMPLEX_N_GLYCAN_SYNTHESIS_PATHWAY, REACTOME_REACTIONS_SPECIFIC_TO_THE_COMPLEX_N_GLYCAN_SYNTHESIS_PATHWAY GOBP_NEGATIVE_REGULATION_OF_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM GOBP_POSITIVE_REGULATION_OF_MYELINATION, GOBP_POSITIVE_REGULATION_OF_MYELINATION
	HP_TRICUSPID_ATRESIA, HP_TRICUSPID_ATRESIA HP_MINICORE_MYOPATHY, HP_MINICORE_MYOPATHY HP_GENERALIZED_LYMPHADENOPATHY, HP_GENERALIZED_LYMPHADENOPATHY HP_CONSPICUOUSLY_HAPPY_DISPOSITION, HP_CONSPICUOUSLY_HAPPY_DISPOSITION GOBP_POLYPHOSPHATE_METABOLIC_PROCESS, GOBP_POLYPHOSPHATE_METABOLIC_PROCESS GOBP_FOLATE_TRANSMEMBRANE_TRANSPORT, GOBP_FOLATE_TRANSMEMBRANE_TRANSPORT GOBP_GOLGI_TO_LYSOSOME_TRANSPORT, GOBP_GOLGI_TO_LYSOSOME_TRANSPORT MIR933, MIR933
	HP_COLD_INDUCED_MUSCLE_CRAMPS, HP_COLD_INDUCED_MUSCLE_CRAMPS HP_INTRAVENTRICULAR_HEMORRHAGE, HP_INTRAVENTRICULAR_HEMORRHAGE GOBP_HISTONE_H4_R3_METHYLATION, GOBP_HISTONE_H4_R3_METHYLATION GOBP_REGULATION_OF_PHOSPHATIDYLINOSITOL_DEPHOSPHORYLATION, GOBP_REGULATION_OF_PHOSPHATIDYLINOSITOL_DEPHOSPHORYLATION GOBP_HISTONE_H2A_K63_LINKED_UBIQUITINATION, GOBP_HISTONE_H2A_K63_LINKED_UBIQUITINATION GOBP_SPINDLE_ASSEMBLY_INVOLVED_IN_MEIOSIS, GOBP_SPINDLE_ASSEMBLY_INVOLVED_IN_MEIOSIS GOMF_MITOGEN_ACTIVATED_PROTEIN_KINASE_KINASE_KINASE_BINDING, GOMF_MITOGEN_ACTIVATED_PROTEIN_KINASE_KINASE_BINDING
	HP_ABNORMAL_GLOMERULAR_MESANGIUM_MORPHOLOGY, HP_ABNORMAL_GLOMERULAR_MESANGIUM_MORPHOLOGY GOBP_POSITIVE_REGULATION_OF_NERVOUS_SYSTEM_PROCESS, GOBP_POSITIVE_REGULATION_OF_NERVOUS_SYSTEM_PROCESS HP_DIFFUSE_MESANGIAL_SCLEROSIS, HP_DIFFUSE_MESANGIAL_SCLEROSIS MIR4278, MIR4278 GOMF_LYSOPHOSPHATIDIC_ACID_RECEPTOR_ACTIVITY, GOMF_LYSOPHOSPHATIDIC_ACID_RECEPTOR_ACTIVITY HP_REDUCED_FACTOR_XII_ACTIVITY, HP_REDUCED_FACTOR_XII_ACTIVITY TFDP2_TARGET_GENES, TFDP2_TARGET_GENES HP_ABNORMAL_URINE_HORMONE_LEVEL, HP_ABNORMAL_URINE_HORMONE_LEVEL
	GOBP_RESPONSE_TO_CAMPTOTHECIN, GOBP_RESPONSE_TO_CAMPTOTHECIN GOBP_POSITIVE_REGULATION_OF_PROTEIN_NEDDYLATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_NEDDYLATION GOMF_ACETYLESTERASE_ACTIVITY, GOMF_ACETYLESTERASE_ACTIVITY REACTOME_TRAFFICKING_OF_MYRISTOYLATED_PROTEINS_TO_THE_CILIUM, REACTOME_TRAFFICKING_OF_MYRISTOYLATED_PROTEINS_TO_THE_CILIUM HP_BRANCHIAL_FISTULA, HP_BRANCHIAL_FISTULA GOBP_NEGATIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION, GOBP_NEGATIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION REACTOME_SYNTHESIS_OF_PG, REACTOME_SYNTHESIS_OF_PG GRANDVAUX_IRF3_TARGETS_UP, GRANDVAUX_IRF3_TARGETS_UP
	MIR6820_5P, MIR6820_5P GOMF_CARBOXYPEPTIDASE_ACTIVITY, GOMF_CARBOXYPEPTIDASE_ACTIVITY GOBP_LUTEINIZATION, GOBP_LUTEINIZATION GOBP_REGULATION_OF_ANTIMICROBIAL_HUMORAL_RESPONSE, GOBP_REGULATION_OF_ANTIMICROBIAL_HUMORAL_RESPONSE MARIADASON_RESPONSE_TO_BUTYRATE_CURCUMIN_SULINDAC_TSA_8, MARIADASON_RESPONSE_TO_BUTYRATE_CURCUMIN_SULINDAC_TSA_8 HP_ULNAR_CLAW, HP_ULNAR_CLAW GOBP_POSITIVE_REGULATION_OF_T_CELL_DIFFERENTIATION_IN_THYMUS, GOBP_POSITIVE_REGULATION_OF_T_CELL_DIFFERENTIATION_IN_THYMUS HP_TYPE_II_TRANSFERRIN_ISOFORM_PROFILE, HP_TYPE_II_TRANSFERRIN_ISOFORM_PROFILE
	REACTOME_SODIUM_PROTON_EXCHANGERS, REACTOME_SODIUM_PROTON_EXCHANGERS REACTOME_SERINE_BIOSYNTHESIS, REACTOME_SERINE_BIOSYNTHESIS HP_ELEVATED_CIRCULATING_FOLLICLE_STIMULATING_HORMONE_LEVEL, HP_ELEVATED_CIRCULATING_FOLLICLE_STIMULATING_HORMONE_LEVEL GOMF_SELENIUM_BINDING, GOMF_SELENIUM_BINDING MIKKELSEN_DEDIFFERENTIATED_STATE_DN, MIKKELSEN_DEDIFFERENTIATED_STATE_DN MIR6814_3P_MIR6872_5P, MIR6814_3P_MIR6872_5P HP_CYSTIC_RENAL_DYSPLASIA, HP_CYSTIC_RENAL_DYSPLASIA
	GOBP_RHOMBOMERE_DEVELOPMENT, GOBP_RHOMBOMERE_DEVELOPMENT QL_PBMC_ZOSTAVAX_AGE_50_75YO_1DY_UP, QL_PBMC_ZOSTAVAX_AGE_50_75YO_1DY_UP GOBP_PARATHYROID_GLAND_DEVELOPMENT, GOBP_PARATHYROID_GLAND_DEVELOPMENT GOMF_THIOREDOXIN_DISULFIDE_REDUCTASE_ACTIVITY, GOMF_THIOREDOXIN_DISULFIDE_REDUCTASE_ACTIVITY chr8q13, chr8q13 GOBP_TROPHOBLAST_GIANT_CELL_DIFFERENTIATION, GOBP_TROPHOBLAST_GIANT_CELL_DIFFERENTIATION GOMF_FILAMIN_BINDING, GOMF_FILAMIN_BINDING MIR339_3P, MIR339_3P
	MIR1180_3P, MIR1180_3P MIR1204, MIR1204 GOBP_REGULATION_OF_OOCYTE_MATURATION, GOBP_REGULATION_OF_OOCYTE_MATURATION GOMF_FOUR_WAY_JUNCTION_HELICASE_ACTIVITY, GOMF_FOUR_WAY_JUNCTION_HELICASE_ACTIVITY REACTOME_SUMO_IS_PROTEOLYTICALLY_PROCESSED, REACTOME_SUMO_IS_PROTEOLYTICALLY_PROCESSED HP_ABSENT_TOENAIL, HP_ABSENT_TOENAIL AMIT_SERUM_RESPONSE_480_MCF10A, AMIT_SERUM_RESPONSE_480_MCF10A GOMF_SOLUTE_PROTON_ANTIPORTER_ACTIVITY, GOMF_SOLUTE_PROTON_ANTIPORTER_ACTIVITY
	GOBP_ACROSOME_ASSEMBLY, GOBP_ACROSOME_ASSEMBLY HP_SCALING_SKIN, HP_SCALING_SKIN YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_4, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_4 HP_FACIAL_HEMANGIOMA, HP_FACIAL_HEMANGIOMA GOBP_CELLULAR_RESPONSE_TO_UV_A, GOBP_CELLULAR_RESPONSE_TO_UV_A HP_OCCIPITAL_MENINGOCELE, HP_OCCIPITAL_MENINGOCELE GOBP_POSITIVE_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION HP_HYPERTENSIVE_CRISIS, HP_HYPERTENSIVE_CRISIS
	ABE_VEGFA_TARGETS, ABE_VEGFA_TARGETS GOBP_AXIS_SPECIFICATION, GOBP_AXIS_SPECIFICATION REACTOME_DCC_MEDIATED_ATTRACTIVE_SIGNALING, REACTOME_DCC_MEDIATED_ATTRACTIVE_SIGNALING REACTOME_LYSOSPHINGOLIPID_AND_LPA_RECEPTORS, REACTOME_LYSOSPHINGOLIPID_AND_LPA_RECEPTORS GOBP_GROWTH_HORMONE_SECRETION, GOBP_GROWTH_HORMONE_SECRETION GOMF_METAL_ION_PROTON_ANTIPORTER_ACTIVITY, GOMF_METAL_ION_PROTON_ANTIPORTER_ACTIVITY GOBP_DORSAL_VENTRAL_AXIS_SPECIFICATION, GOBP_DORSAL_VENTRAL_AXIS_SPECIFICATION
	MATSUMIYA_PBMC_MODIFIED_VACCINIA_ANKARA_VACCINE_AGE_18_55YO_HIGH_RESPONDERS_VS_LOW_RESPONDERS_0DY_UP MIR4740_3P, MIR4740_3P MIR4497 MIR4497 GOBP_DIAPEDESIS, GOBP_DIAPEDESIS, GOBP_DIAPEDESIS, GOBP_DIAPEDESIS, GOBP_DIAPEDESIS, GOBP_DIAPEDESIS, GOBP_REGULATION, GOBP_REGULATION_OF_MEMBRANE_TUBULATION MIR4497 MIR4497 GOBP_NEGATIVE_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS, GOBP_NEGATIVE_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS
	GOBP_FACIAL_NERVE_MORPHOGENESIS, GOBP_FACIAL_NERVE_MORPHOGENESIS GOBP_MITOTIC_NUCLEAR_MEMBRANE_ORGANIZATION, GOBP_MITOTIC_NUCLEAR_MEMBRANE_ORGANIZATION GOBP_REGULATION_OF_MICROTUBULE_BINDING, GOBP_REGULATION_OF_MICROTUBULE_BINDING GOMF_SERINE_TYPE_EXOPEPTIDASE_ACTIVITY, GOMF_SERINE_TYPE_EXOPEPTIDASE_ACTIVITY GOMF_LIPOPEPTIDE_BINDING, GOMF_LIPOPEPTIDE_BINDING GOBP_NEGATIVE_REGULATION_OF_TYROSINE_PHOSPHORYLATION_OF_STAT_PROTEIN, GOBP_NEGATIVE_REGULATION_OF_TYROSINE_PHOSPHORYLATION_OF_STAT_PROTEIN HARALAMBIEVA_PBMC_TIV_AGE_50_74YO_CORRELATED_WITH_MEMORY_B_CELL_RESPONSE_3DY_POSITIVE, HARALAMBIEVA_PBMC_TIV_AGE_50_74YO_CORRELATED_WITH_MEMORY_B_CELL_RESPONSE_3DY_POSITIVE HP_PAPILLARY_THYROID_CARCINOMA, HP_PAPILLARY_THYROID_CARCINOMA
	GOBP_RESPONSE_TO_BACTERIAL_LIPOPROTEIN, GOBP_RESPONSE_TO_BACTERIAL_LIPOPROTEIN GOCC_ER_UBIQUITIN_LIGASE_COMPLEX, GOCC_ER_UBIQUITIN_LIGASE_COMPLEX HP_METABOLIC_ALKALOSIS, HP_METABOLIC_ALKALOSIS chr5q12, chr5q12 MIR1539, MIR1539 REACTOME_NOSTRIN_MEDIATED_ENOS_TRAFFICKING, REACTOME_NOSTRIN_MEDIATED_ENOS_TRAFFICKING GOBP_SELENIUM_COMPOUND_METABOLIC_PROCESS, GOBP_SELENIUM_COMPOUND_METABOLIC_PROCESS GOBP_RESPONSE_TO_SORBITOL, GOBP_RESPONSE_TO_SORBITOL
	GOBP_REGULATION_OF_SKELETAL_MUSCLE_ACETYLCHOLINE_GATED_CHANNEL_CLUSTERING, GOBP_REGULATION_OF_SKELETAL_MUSCLE_ACETYLCHOLINE_GATED_CHANNEL_CLUSTERING GOCC_FEMALE_GERM_CELL_NUCLEUS, GOCC_FEMALE_GERM_CELL_NUCLEUS GOCC_MITOTIC_COHESIN_COMPLEX, GOCC_MITOTIC_COHESIN_COMPLEX GOMF_SERINE_TYPE_CARBOXYPEPTIDASE_ACTIVITY, GOMF_SERINE_TYPE_CARBOXYPEPTIDASE_ACTIVITY GOMF_INHIBIN_BINDING, GOMF_INHIBIN_BINDING WP_ULTRACONSERVED_REGION_339_MODULATION_OF_TUMOR_SUPPRESSOR_MICRORNAS_IN_CANCER, WP_ULTRACONSERVED_REGION_339_MODULATION_OF_TUMOR_SUPPRESSOR_MICRORNAS_IN_CANCER HP_ELEVATED_CIRCULATING_LONG_CHAIN_FATTY_ACID_CONCENTRATION, HP_ELEVATED_CIRCULATING_LONG_CHAIN_FATTY_ACID_CONCENTRATION HP_RENAL_TUBULAR_EPITHELIAL_NECROSIS, HP_RENAL_TUBULAR_EPITHELIAL_NECROSIS
	HP_SINGLE_FIBER_EMG_ABNORMALITY, HP_SINGLE_FIBER_EMG_ABNORMALITY HP_LINGUAL_DYSTONIA, HP_LINGUAL_DYSTONIA HP_ABNORMAL_FACTOR_IX_ACTIVITY, HP_ABNORMAL_FACTOR_IX_ACTIVITY GOBP_REGULATION_OF_SERINE_C_PALMITOYLTRANSFERASE_ACTIVITY, GOBP_REGULATION_OF_SERINE_C_PALMITOYLTRANSFERASE_ACTIVITY GOCC_CORVET_COMPLEX, GOCC_CORVET_COMPLEX GOCC_MON1_CCZ1_COMPLEX, GOCC_MON1_CCZ1_COMPLEX GOCC_CYTOPLASMIC_SIDE_OF_LYSOSOMAL_MEMBRANE, GOCC_CYTOPLASMIC_SIDE_OF_LYSOSOMAL_MEMBRANE
	GOMF_RNA_POLYMERASE_II_C_TERMINAL_DOMAIN_PHOSPHOSERINE_BINDING, GOMF_RNA_POLYMERASE_II_C_TERMINAL_DOMAIN_PHOSPHOSERINE_BINDING REACTOME_PTK6_EXPRESSION, REACTOME_PTK6_EXPRESSION CHANG_POU5F1_TARGETS_DN, CHANG_POU5F1_TARGETS_DN HP_NON_MEDULLARY_THYROID_CARCINOMA, HP_NON_MEDULLARY_THYROID_CARCINOMA HP_NARROW_ILIAC_WINGS, HP_NARROW_ILIAC_WINGS HP_FEMALE_PSEUDOHERMAPHRODITISM, HP_FEMALE_PSEUDOHERMAPHRODITISM GOBP_REELIN_MEDIATED_SIGNALING_PATHWAY, GOBP_REELIN_MEDIATED_SIGNALING_PATHWAY HP_TORSION_DYSTONIA, HP_TORSION_DYSTONIA
	HP_FATIGABLE_WEAKNESS_OF_DISTAL_LIMB_MUSCLES, HP_FATIGABLE_WEAKNESS_OF_DISTAL_LIMB_MUSCLES GOMF_K48_LINKED_POLYDIRQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GOMF_K48_LINKED_POLYDBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING BIOCARTA_P35ALZHEIMERS_PATHWAY, BIOCARTA_P35ALZHEIMERS_PATHWAY HP_ANKYLOGLOSSIA, HP_ANKYLOGLOSSIA GOBP_POSITIVE_REGULATION_OF_ACTIN_FILAMENT_DEPOLYMERIZATION, GOBP_POSITIVE_REGULATION_OF_ACTIN_FILAMENT_DEPOLYMERIZATION chr5p13, chr5p13 GOBP_DNA_INTEGRATION, GOBP_DNA_INTEGRATION REACTOME_ORGANIC_ANION_TRANSPORTERS, REACTOME_ORGANIC_ANION_TRANSPORTERS
	GOBP_CRANIAL_NERVE_STRUCTURAL_ORGANIZATION, GOBP_CRANIAL_NERVE_STRUCTURAL_ORGANIZATION HP_RIGHT_VENTRICULAR_HYPERTROPHY, HP_RIGHT_VENTRICULAR_HYPERTROPHY chr10p14, chr10p14 PANAPASA_BLOOD_FLUENZ_AGE_03_17YO_3DY_4DY_DN, PANAPASA_BLOOD_FLUENZ_AGE_03_17YO_3DY_4DY_DN GOBP_NEGATIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION GOMF_STEROID_HORMONE_BINDING, GOMF_STEROID_HORMONE_BINDING GOMF_SODIUM_PROTON_ANTIPORTER_ACTIVITY, GOMF_SODIUM_PROTON_ANTIPORTER_ACTIVITY
	HP_PARTIAL_DUPLICATION_OF_THE_DISTAL_PHALANGES_OF_THE_HAND, HP_PARTIAL_DUPLICATION_OF_THE_DISTAL_PHALANGES_OF_THE_HAND HP_ENURESIS_NOCTURNA, HP_ENURESIS_NOCTURNA GOBP_POSITIVE_REGULATION_OF_MICROTUBULE_BINDING, GOBP_POSITIVE_REGULATION_OF_MICROTUBULE_BINDING GOMF_HISTONE_SERINE_KINASE_ACTIVITY, GOMF_HISTONE_SERINE_KINASE_ACTIVITY MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_6, MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_6 HP_REDUCED_FACTOR_XI_ACTIVITY, HP_REDUCED_FACTOR_XI_ACTIVITY GOBP_ACTIVATION_OF_PHOSPHOLIPASE_D_ACTIVITY, GOBP_ACTIVATION_OF_PHOSPHOLIPASE_D_ACTIVITY GOBP_VESICLE_TETHERING_TO_GOLGI, GOBP_VESICLE_TETHERING_TO_GOLGI
	GOBP_REGULATION_OF_ENDOPLASMIC_RETICULUM_TUBULAR_NETWORK_ORGANIZATION, GOBP_REGULATION_OF_ENDOPLASMIC_RETICULUM_TUBULAR_NETWORK_ORGANIZATION GOCC_POSTSYNAPTIC_GOLGI_APPARATUS, GOCC_POSTSYNAPTIC_GOLGI_APPARATUS GOBP_SPLEEN_DEVELOPMENT, GOBP_SPLEEN_DEVELOPMENT HP_MUSCLE_FIBER_INCLUSION_BODIES, HP_MUSCLE_FIBER_INCLUSION_BODIES HP_INCREASED_CIRCULATING_CORTISOL_LEVEL, HP_INCREASED_CIRCULATING_CORTISOL_LEVEL HP_PUNCTATE_CATARACT, HP_PUNCTATE_CATARACT GOBP_HISTONE_SERINE_PHOSPHORYLATION, GOBP_HISTONE_SERINE_PHOSPHORYLATION
	GOBP_CELLULAR_RESPONSE_TO_BACTERIAL_LIPOPROTEIN, GOBP_CELLULAR_RESPONSE_TO_BACTERIAL_LIPOPROTEIN GOCC_HRDIP_UBIQUITIN_LIGASE_ERAD_L_COMPLEX, GOCC_HRDIP_UBIQUITIN_LIGASE_ERAD_L_COMPLEX ERWIN_COHEN_BLOOD_VACCINE_TC_83_AGE_23_48YO_VACCINATED_VS_CONTROL_7DY_DN, ERWIN_COHEN_BLOOD_VACCINE_TC_83_AGE_23_48YO_VACCINATED_VS_CONTROL_7DY_DN HP_FACIAL_CAPILLARY_HEMANGIOMA, HP_FACIAL_CAPILLARY_HEMANGIOMA HP_RESPONSE_TO_DRUGS_ACTING_ON_NEUROMUSCULAR_TRANSMISSION, HP_RESPONSE_TO_DRUGS_ACTING_ON_NEUROMUSCULAR_TRANSMISSION GOBP_POSITIVE_REGULATION_OF_ACTIVIN_RECEPTOR_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_ACTIVIN_RECEPTOR_SIGNALING_PATHWAY REACTOME_ZINC_EFFLUX_AND_COMPARTMENTALIZATION_BY_THE_SLC30_FAMILY, REACTOME_ZINC_EFFLUX_AND_COMPARTMENTALIZATION_BY_THE_SLC30_FAMILY GOBP_ENDOTHELIN_RECEPTOR_SIGNALING_PATHWAY, GOBP_ENDOTHELIN_RECEPTOR_SIGNALING_PATHWAY DHIMAN_PRIC_ATTENLIVAX_AGE_15_25YO_SUBO_7_OR_LADY_DN_DHIMAN_PRIC_ATTENLIVAX_AGE_15_25YO_SUBO_7_OR_LADY_DN_
	DHIMAN_PBMC_ATTENUVAX_AGE_15_25YO_SUBQ_7_OR_14DY_DN, DHIMAN_PBMC_ATTENUVAX_AGE_15_25YO_SUBQ_7_OR_14DY_DN HP_MYOFIBRILLAR_MYOPATHY, HP_MYOFIBRILLAR_MYOPATHY MIR1296_3P, MIR1296_3P GOBP_REGULATION_OF_GROWTH_HORMONE_SECRETION, GOBP_REGULATION_OF_GROWTH_HORMONE_SECRETION GOMF_HMG_BOX_DOMAIN_BINDING, GOMF_HMG_BOX_DOMAIN_BINDING GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEPOLYMERIZATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEPOLYMERIZATION GOBP_DETECTION_OF_BACTERIAL_LIPOPROTEIN, GOBP_DETECTION_OF_BACTERIAL_LIPOPROTEIN GOBP_REGULATION_OF_CHONDROCYTE_DEVELOPMENT, GOBP_REGULATION_OF_CHONDROCYTE_DEVELOPMENT
	GOBP_REGULATION_OF_CHONDROCYTE_DEVELOPMENT, GOBP_REGULATION_OF_CHONDROCYTE_DEVELOPMENT GOBP_REGULATION_OF_BONE_MINERALIZATION_INVOLVED_IN_BONE_MATURATION, GOBP_REGULATION_OF_BONE_MINERALIZATION_INVOLVED_IN_BONE_MATURATION GOBP_SIALIC_ACID_TRANSPORT, GOBP_SIALIC_ACID_TRANSPORT GOBP_THROMBOPOIETIN_MEDIATED_SIGNALING_PATHWAY, GOBP_THROMBOPOIETIN_MEDIATED_SIGNALING_PATHWAY GOMF_SIALIC_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_SIALIC_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY HP_MUSCLE_FIBER_TUBULAR_INCLUSIONS, HP_MUSCLE_FIBER_TUBULAR_INCLUSIONS GOBP_NEGATIVE_REGULATION_OF_HISTONE_DEACETYLATION, GOBP_NEGATIVE_REGULATION_OF_HISTONE_DEACETYLATION GOCC_AXON_HILLOCK, GOCC_AXON_HILLOCK HP_DUPLICATION_OF_THE_DISTAL_PHALANX_OF_HAND, HP_DUPLICATION_OF_THE_DISTAL_PHALANX_OF_HAND
	HP_DUPLICATION_OF_THE_DISTAL_PHALANX_OF_HAND, HP_DUPLICATION_OF_THE_DISTAL_PHALANX_OF_HAND GOBP_ENDODERMAL_CELL_FATE_COMMITMENT, GOBP_ENDODERMAL_CELL_FATE_COMMITMENT GARCIA_PINERES_PBMC_HPV_16_L1_VLP_AGE_18_25YO_7MO_CORRELATED_WITH_ANTIBODY_RESPONSE_NEGATIVE, GARCIA_PINERES_PBMC_HPV_16_L1_VLP_AGE_18_25YO_7MO_CORRELATED_WITH_ANTIBODY_RESPONSE_NEGATIVE CHIN_BREAST_CANCER_COPY_NUMBER_UP, CHIN_BREAST_CANCER_COPY_NUMBER_UP chr6q12, chr6q12 GOBP_REGULATION_OF_RECEPTOR_CLUSTERING, GOBP_REGULATION_OF_RECEPTOR_CLUSTERING GOCC_PERICILIARY_MEMBRANE_COMPARTMENT, GOCC_PERICILIARY_MEMBRANE_COMPARTMENT WP_ZINC_HOMEOSTASIS, WP_ZINC_HOMEOSTASIS
	WP_ZINC_HOMEOSTASIS, WP_ZINC_HOMEOSTASIS GOBP_NEGATIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY chr18q23, chr18q23 GOBP_NEGATIVE_REGULATION_OF_CHONDROCYTE_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_CHONDROCYTE_DIFFERENTIATION GOBP_PARASYMPATHETIC_NERVOUS_SYSTEM_DEVELOPMENT, GOBP_PARASYMPATHETIC_NERVOUS_SYSTEM_DEVELOPMENT GOBP_CELL_FATE_COMMITMENT_INVOLVED_IN_FORMATION_OF_PRIMARY_GERM_LAYER, GOBP_CELL_FATE_COMMITMENT_INVOLVED_IN_FORMATION_OF_PRIMARY_GERM_LAYER GOBP_POSITIVE_REGULATION_OF_CELL_AGING, GOBP_POSITIVE_REGULATION_OF_CELL_AGING GOBP_NEGATIVE_REGULATION_OF_CARTILAGE_DEVELOPMENT, GOBP_NEGATIVE_REGULATION_OF_CARTILAGE_DEVELOPMENT GOBP_NEUROMUSCULAR_SYNAPTIC_TRANSMISSION, GOBP_NEUROMUSCULAR_SYNAPTIC_TRANSMISSION