	GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_DN, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_DN GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY7_UP GSE26351_UNSTIM_VS_WNT_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_DN, GSE26351_UNSTIM_VS_WNT_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_DN GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_MEMORY_CD8_TCELL_DN, GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_MEMORY_CD8_TCELL_DN GSE29164_DAY3_VS_DAY7_UNTREATED_MELANOMA_DN, GSE29164_DAY3_VS_DAY7_UNTREATED_MELANOMA_DN GSE369_SOCS3_KO_VS_WT_LIVER_DN, GSE369_SOCS3_KO_VS_WT_LIVER_DN GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_12H_CD4_TCELL_UP, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_12H_CD4_TCELL_UP GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_UP GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_UP, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_UP
	GSE27/0_IL12_AND_TGFI_ACT_VS_ACT_CD4_TCELL_48H_UP, GSE27/0_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_UP GSE4142_GC_BCELL_VS_MEMORY_BCELL_DN, GSE4142_GC_BCELL_VS_MEMORY_BCELL_DN GSE7460_WT_VS_FOXP3_HET_ACT_TCONV_UP, GSE7460_WT_VS_FOXP3_HET_ACT_TCONV_UP GSE21360_NAIVE_VS_PRIMARY_MEMORY_CD8_TCELL_DN, GSE21360_NAIVE_VS_PRIMARY_MEMORY_CD8_TCELL_DN GSE3982_DC_VS_BASOPHIL_UP, GSE3982_DC_VS_BASOPHIL_UP GSE13493_DP_VS_CD8POS_THYMOCYTE_UP, GSE13493_DP_VS_CD8POS_THYMOCYTE_UP MANNE_COVID19_NONICU_VS_HEALTHY_DONOR_PLATELETS_UP, MANNE_COVID19_NONICU_VS_HEALTHY_DONOR_PLATELETS_UP GSE16450_CTRL_VS_IFNA_6H_STIM_IMMATURE_NEURON_CELL_LINE_UP, GSE16450_CTRL_VS_IFNA_6H_STIM_IMMATURE_NEURON_CELL_LINE_UP GSE24210_IL35_TREATED_VS_UNTREATED_TCONV_CD4_TCELL_UP, GSE24210_IL35_TREATED_VS_UNTREATED_TCONV_CD4_TCELL_UP
	MIR12119, MIR12119 GSE8835_CD4_VS_CD8_TCELL_CLL_PATIENT_UP, GSE8835_CD4_VS_CD8_TCELL_CLL_PATIENT_UP GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_4H_DN, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_4H_DN MIR561_5P, MIR561_5P GOBP_ORGANELLE_FUSION, GOBP_ORGANELLE_FUSION GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_UP, GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_UP GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_UP, GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_UP MIR10394_5P, MIR10394_5P
	HP_ABNORMAL_METABOLISM, HP_ABNORMAL_METABOLISM ZNF669_TARGET_GENES, ZNF669_TARGET_GENES MIR6893_5P, MIR6893_5P HP_ABNORMALITY_OF_HAIR_TEXTURE, HP_ABNORMALITY_OF_HAIR_TEXTURE MIR6808_5P, MIR6808_5P GSE21379_WT_VS_SAP_KO_TFH_CD4_TCELL_DN, GSE21379_WT_VS_SAP_KO_TFH_CD4_TCELL_DN GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_UP, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_UP BIOCARTA_MAPK_PATHWAY, BIOCARTA_MAPK_PATHWAY
	GOBP_REGULATION_OF_CELL_PROJECTION_ASSEMBLY, GOBP_REGULATION_OF_CELL_PROJECTION_ASSEMBLY MIR140_3P, MIR140_3P MIR4252, MIR4252 MIR7844_5P, MIR7844_5P MIR4793_3P, MIR4793_3P HP_TAPERED_FINGER, HP_TAPERED_FINGER SUNG_METASTASIS_STROMA_UP, SUNG_METASTASIS_STROMA_UP GOBP_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_ALPHA_BETA_T_CELL_ACTIVATION
	GSE9601_NFKB_INHIBITOR_VS_PI3K_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_DN, GSE9601_NFKB_INHIBITOR_VS_PI3K_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_DN GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_UP, GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_UP CHENG_IMPRINTED_BY_ESTRADIOL, CHENG_IMPRINTED_BY_ESTRADIOL WP_PANCREATIC_ADENOCARCINOMA_PATHWAY, WP_PANCREATIC_ADENOCARCINOMA_PATHWAY GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_DN, GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_DN GOBP_ORGANELLE_MEMBRANE_FUSION, GOBP_ORGANELLE_MEMBRANE_FUSION REACTOME_ONCOGENIC_MAPK_SIGNALING, REACTOME_ONCOGENIC_MAPK_SIGNALING MIR671_5P, MIR671_5P
	MIR4769_3P, MIR4769_3P HP_HALLUCINATIONS, HP_HALLUCINATIONS IRF2_01, IRF2_01 MIR6817_5P, MIR6817_5P ZNF34_TARGET_GENES, ZNF34_TARGET_GENES MIR7703, MIR7703 MIR7703 MIR584_3P, MIR584_3P GOBP_MEMBRANE_FUSION, GOBP_MEMBRANE_FUSION
	GSE37534_GW1929_VS_PIOGLITAZONE_TREATED_CD4_TCELL_PPARG1_FOXP3_TRANSDUCED_UP, GSE37534_GW1929_VS_PIOGLITAZONE_TREATED_CD4_TCELL_PPARG1_FOXP3_TRANSDUCED_UP HP_ASCITES, HP_ASCITES JOHNSTONE_PARVB_TARGETS_1_DN, JOHNSTONE_PARVB_TARGETS_1_DN MIR6500_3P, MIR6500_3P HP_LONG_PALPEBRAL_FISSURE, HP_LONG_PALPEBRAL_FISSURE MIR4509, MIR4509 GOBP_TEMPERATURE_HOMEOSTASIS, GOBP_TEMPERATURE_HOMEOSTASIS HP_2_3_TOE_SYNDACTYLY, HP_2_3_TOE_SYNDACTYLY
	MIR4318, MIR4318 HP_ABNORMAL_VENOUS_MORPHOLOGY, HP_ABNORMAL_VENOUS_MORPHOLOGY MIR4424, MIR4424 PID_FCER1_PATHWAY, PID_FCER1_PATHWAY HP_PRENATAL_MOVEMENT_ABNORMALITY, HP_PRENATAL_MOVEMENT_ABNORMALITY MIR21_3P, MIR21_3P MIR5088_3P, MIR5088_3P GOCC_CUL3_RING_UBIQUITIN_LIGASE_COMPLEX, GOCC_CUL3_RING_UBIQUITIN_LIGASE_COMPLEX
	MIR6755_3P, MIR6755_3P REACTOME_SIGNALING_BY_BRAF_AND_RAF_FUSIONS, REACTOME_SIGNALING_BY_BRAF_AND_RAF_FUSIONS HP_MUSCLE_FIBRILLATION, HP_MUSCLE_FIBRILLATION WP_ONCOSTATIN_M_SIGNALING_PATHWAY, WP_ONCOSTATIN_M_SIGNALING_PATHWAY NAKAYAMA_FRA2_TARGETS, NAKAYAMA_FRA2_TARGETS MIR197_3P, MIR197_3P GOCC_VESICLE_TETHERING_COMPLEX, GOCC_VESICLE_TETHERING_COMPLEX BIOCARTA_FCER1_PATHWAY, BIOCARTA_FCER1_PATHWAY
	GSE13522_CTRL_VS_T_CRUZI_BRAZIL_STRAIN_INF_SKIN_DN, GSE13522_CTRL_VS_T_CRUZI_BRAZIL_STRAIN_INF_SKIN_DN HP_CUTIS_LAXA, HP_CUTIS_LAXA WP_CARDIAC_HYPERTROPHIC_RESPONSE, WP_CARDIAC_HYPERTROPHIC_RESPONSE GOBP_PROTEIN_TARGETING_TO_VACUOLE, GOBP_PROTEIN_TARGETING_TO_VACUOLE GSE29949_CD8_POS_DC_SPLEEN_VS_DC_BRAIN_DN, GSE29949_CD8_POS_DC_SPLEEN_VS_DC_BRAIN_DN MIR141_SP, MIR141_SP GOBP_POSITIVE_REGULATION_OF_DEPHOSPHORYLATION, GOBP_POSITIVE_REGULATION_OF_DEPHOSPHORYLATION HNF3_Q6, HNF3_Q6
	GUO_HEX_TARGETS_UP, GUO_HEX_TARGETS_UP GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE, GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE MIR7113_3P, MIR7113_3P GGGATGC_MIR3245P, GGGATGC_MIR3245P PAX8_B, PAX8_B MIR3912_5P, MIR3912_5P YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_0, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_0 PID_INSULIN_PATHWAY, PID_INSULIN_PATHWAY
	GSE29618_PRE_VS_DAY7_FLU_VACCINE_MONOCYTE_UP, GSE29618_PRE_VS_DAY7_FLU_VACCINE_MONOCYTE_UP HP_HEPATOCELLULAR_CARCINOMA, HP_HEPATOCELLULAR_CARCINOMA HP_ABNORMAL_PULMONARY_VALVE_MORPHOLOGY, HP_ABNORMAL_PULMONARY_VALVE_MORPHOLOGY GOBP_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION GOBP_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_METABOLIC_PROCESS, GOBP_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_METABOLIC_PROCESS GOBP_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_ALPHA_BETA_T_CELL_DIFFERENTIATION HP_ABNORMAL_T_CELL_SUBSET_DISTRIBUTION, HP_ABNORMAL_T_CELL_SUBSET_DISTRIBUTION GOBP_AUTOPHAGOSOME_MATURATION, GOBP_AUTOPHAGOSOME_MATURATION
	GOBP_POSITIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS PID_P38_ALPHA_BETA_DOWNSTREAM_PATHWAY, PID_P38_ALPHA_BETA_DOWNSTREAM_PATHWAY MORF_DMPK, MORF_DMPK MIR4715_3P, MIR4715_3P GSE20366_TREG_VS_NAIVE_CD4_TCELL_DEC205_CONVERSION_DN, GSE20366_TREG_VS_NAIVE_CD4_TCELL_DEC205_CONVERSION_DN GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4HIGH_SORTED_BCELL_UP, GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4HIGH_SORTED_BCELL_UP HP_CONGENITAL_HIP_DISLOCATION, HP_CONGENITAL_HIP_DISLOCATION GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEPHOSPHORYLATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEPHOSPHORYLATION
	GSE2770_IL12_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_UP GOBP_POSITIVE_REGULATION_OF_PLASMA_MEMBRANE_BOUNDED_CELL_PROJECTION_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_PLASMA_MEMBRANE_BOUNDED_CELL_PROJECTION_ASSEMBLY MIR3928_5P, MIR3928_5P HP_URTICARIA, HP_URTICARIA KIM_HYPOXIA, KIM_HYPOXIA GOBP_ACTIVATION_OF_JUN_KINASE_ACTIVITY, GOBP_ACTIVATION_OF_JUN_KINASE_ACTIVITY HP_ABNORMALITY_OF_SUBCUTANEOUS_FAT_TISSUE, HP_ABNORMALITY_OF_SUBCUTANEOUS_FAT_TISSUE MIR4494, MIR4494
	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 TERAO_AOX4_TARGETS_HG_UP, TERAO_AOX4_TARGETS_HG_UP MIR6806_3P, MIR6806_3P REACTOME_FCGAMMA_RECEPTOR_FCGR_DEPENDENT_PHAGOCYTOSIS, REACTOME_FCGAMMA_RECEPTOR_FCGR_DEPENDENT_PHAGOCYTOSIS ZHANG_ANTIVIRAL_RESPONSE_TO_RIBAVIRIN_DN, ZHANG_ANTIVIRAL_RESPONSE_TO_RIBAVIRIN_DN YUAN_ZNF143_PARTNERS, YUAN_ZNF143_PARTNERS GOBP_PHOSPHOLIPID_CATABOLIC_PROCESS, GOBP_PHOSPHOLIPID_CATABOLIC_PROCESS GOBP_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY, GOBP_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY
	HP_CURLY_HAIR, HP_CURLY_HAIR GOMF_LIGASE_ACTIVITY_FORMING_CARBON_SULFUR_BONDS, GOMF_LIGASE_ACTIVITY_FORMING_CARBON_SULFUR_BONDS MORF_BCL2L11, MORF_BCL2L11 GOBP_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION MIR3130_3P, MIR3130_3P MIR5002_3P, MIR5002_3P XU_GH1_EXOGENOUS_TARGETS_DN, XU_GH1_EXOGENOUS_TARGETS_DN GOCC_ORGANELLE_MEMBRANE_CONTACT_SITE, GOCC_ORGANELLE_MEMBRANE_CONTACT_SITE
	GOMF_ADENYLYLTRANSFERASE_ACTIVITY, GOMF_ADENYLYLTRANSFERASE_ACTIVITY BERTUCCI_INVASIVE_CARCINOMA_DUCTAL_VS_LOBULAR_UP, BERTUCCI_INVASIVE_CARCINOMA_DUCTAL_VS_LOBULAR_UP GOBP_POSITIVE_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_METABOLIC_PROCESS MIR1263, MIR1263 GOBP_FILOPODIUM_ASSEMBLY, GOBP_FILOPODIUM_ASSEMBLY REACTOME_NUCLEAR_EVENTS_KINASE_AND_TRANSCRIPTION_FACTOR_ACTIVATION, REACTOME_NUCLEAR_EVENTS_KINASE_AND_TRANSCRIPTION_FACTOR_ACTIVATION HP_DELAYED_CRANIAL_SUTURE_CLOSURE, HP_DELAYED_CRANIAL_SUTURE_CLOSURE VANTVEER_BREAST_CANCER_BRCA1_DN, VANTVEER_BREAST_CANCER_BRCA1_DN
	MIR6816_3P, MIR6816_3P HP_ABNORMAL_MORPHOLOGY_OF_THE_ABDOMINAL_MUSCULATURE, HP_ABNORMAL_MORPHOLOGY_OF_THE_ABDOMINAL_MUSCULATURE GOBP_VIRAL_BUDDING_VIA_HOST_ESCRT_COMPLEX, GOBP_VIRAL_BUDDING_VIA_HOST_ESCRT_COMPLEX HTF_01, HTF_01 POS_HISTAMINE_RESPONSE_NETWORK, POS_HISTAMINE_RESPONSE_NETWORK HP_PERIORBITAL_FULLNESS, HP_PERIORBITAL_FULLNESS REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION, REACTOME_MITOCHONDRIAL_TRNA_AMINOACYLATION GSE43863_NAIVE_VS_MEMORY_THI_CD4_TCELL_D150_LCMV_DN, GSE43863_NAIVE_VS_MEMORY_THI_CD4_TCELL_D150_LCMV_DN
	XU_RESPONSE_TO_TRETINOIN_UP, XU_RESPONSE_TO_TRETINOIN_UP BONOME_OVARIAN_CANCER_POOR_SURVIVAL_UP, BONOME_OVARIAN_CANCER_POOR_SURVIVAL_UP WP_UNFOLDED_PROTEIN_RESPONSE, WP_UNFOLDED_PROTEIN_RESPONSE HP_CHRONIC_PULMONARY_OBSTRUCTION, HP_CHRONIC_PULMONARY_OBSTRUCTION HP_APLASIA_HYPOPLASIA_OF_THE_THYMUS, HP_APLASIA_HYPOPLASIA_OF_THE_THYMUS GOBP_GLYCEROPHOSPHOLIPID_CATABOLIC_PROCESS, GOBP_GLYCEROPHOSPHOLIPID_CATABOLIC_PROCESS GOBP_PEPTIDYL_CYSTEINE_MODIFICATION, GOBP_PEPTIDYL_CYSTEINE_MODIFICATION REACTOME_INACTIVATION_OF_CSF3_G_CSF_SIGNALING, REACTOME_INACTIVATION_OF_CSF3_G_CSF_SIGNALING HP_ABNORMALITY_OF_VITAMIN_B12_METABOLISM, HP_ABNORMALITY_OF_VITAMIN_B12_METABOLISM
	YAN_ESCAPE_FROM_ANOIKIS, YAN_ESCAPE_FROM_ANOIKIS GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_DN, GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_DN GOBP_POSITIVE_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS CHANG_IMMORTALIZED_BY_HPV31_UP, CHANG_IMMORTALIZED_BY_HPV31_UP GOBP_POSITIVE_REGULATION_OF_PHOSPHOPROTEIN_PHOSPHATASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_PHOSPHOPROTEIN_PHOSPHATASE_ACTIVITY HP_IMPAIRED_T_CELL_FUNCTION, HP_IMPAIRED_T_CELL_FUNCTION PID_GMCSF_PATHWAY, PID_GMCSF_PATHWAY GOBP_REGULATION_OF_HISTONE_DEACETYLATION, GOBP_REGULATION_OF_HISTONE_DEACETYLATION
	HP_SHORTENING_OF_ALL_DISTAL_PHALANGES_OF_THE_FINGERS, HP_SHORTENING_OF_ALL_DISTAL_PHALANGES_OF_THE_FINGERS MIDORIKAWA_AMPLIFIED_IN_LIVER_CANCER, MIDORIKAWA_AMPLIFIED_IN_LIVER_CANCER REACTOME_VEGFR2_MEDIATED_VASCULAR_PERMEABILITY, REACTOME_VEGFR2_MEDIATED_VASCULAR_PERMEABILITY GOBP_REGULATION_OF_DNA_DAMAGE_CHECKPOINT, GOBP_REGULATION_OF_DNA_DAMAGE_CHECKPOINT GOBP_PROTEIN_PALMITOYLATION, GOBP_PROTEIN_PALMITOYLATION MIR500B_3P, MIR500B_3P HP_PREMATURE_SKIN_WRINKLING, HP_PREMATURE_SKIN_WRINKLING WP_P38_MAPK_SIGNALING_PATHWAY, WP_P38_MAPK_SIGNALING_PATHWAY
	BIOCARTA_P38MAPK_PATHWAY, BIOCARTA_P38MAPK_PATHWAY GOCC_MICROVILLUS, GOCC_MICROVILLUS GOBP_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS, GOBP_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS GOMF_NON_MEMBRANE_SPANNING_PROTEIN_TYROSINE_KINASE_ACTIVITY, GOMF_NON_MEMBRANE_SPANNING_PROTEIN_TYROSINE_KINASE_ACTIVITY MIR4791, MIR4791 GOBP_T_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_T_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE GOBP_T_CELL_SELECTION, GOBP_T_CELL_SELECTION GOBP_POSITIVE_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA, GOBP_POSITIVE_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA
	GOMF_LYS48_SPECIFIC_DEUBIQUITINASE_ACTIVITY, GOMF_LYS48_SPECIFIC_DEUBIQUITINASE_ACTIVITY HP_JUVENILE_ASEPTIC_NECROSIS, HP_JUVENILE_ASEPTIC_NECROSIS PID_ALPHA_SYNUCLEIN_PATHWAY, PID_ALPHA_SYNUCLEIN_PATHWAY GOBP_PEPTIDYL_L_CYSTEINE_S_PALMITOYLATION, GOBP_PEPTIDYL_L_CYSTEINE_S_PALMITOYLATION REACTOME_CD28_CO_STIMULATION, REACTOME_CD28_CO_STIMULATION BACOLOD_RESISTANCE_TO_ALKYLATING_AGENTS_UP, BACOLOD_RESISTANCE_TO_ALKYLATING_AGENTS_UP GSE33292_DN3_THYMOCYTE_VS_TCF1_KO_TCELL_LYMPHOMA_DN, GSE33292_DN3_THYMOCYTE_VS_TCF1_KO_TCELL_LYMPHOMA_DN WP_KISSPEPTINKISSPEPTIN_RECEPTOR_SYSTEM_IN_THE_OVARY, WP_KISSPEPTINKISSPEPTIN_RECEPTOR_SYSTEM_IN_THE_OVARY
	HP_THICK_HAIR, HP_THICK_HAIR GOBP_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY, GOBP_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY HP_DELAYED_CLOSURE_OF_THE_ANTERIOR_FONTANELLE, HP_DELAYED_CLOSURE_OF_THE_ANTERIOR_FONTANELLE ZNF510_TARGET_GENES, ZNF510_TARGET_GENES GOMF_PALMITOYLTRANSFERASE_ACTIVITY, GOMF_PALMITOYLTRANSFERASE_ACTIVITY MIR1231, MIR1231 HP_MILD_HEARING_IMPAIRMENT, HP_MILD_HEARING_IMPAIRMENT MIR627_5P, MIR627_5P
	GOBP_POSITIVE_T_CELL_SELECTION, GOBP_POSITIVE_T_CELL_SELECTION GOBP_BROWN_FAT_CELL_DIFFERENTIATION, GOBP_BROWN_FAT_CELL_DIFFERENTIATION HP_MICROCYTIC_ANEMIA, HP_MICROCYTIC_ANEMIA MIR4740_5P, MIR4740_5P HP_ABNORMALITY_OF_THE_2ND_TOE, HP_ABNORMALITY_OF_THE_2ND_TOE HP_APLASIA_INVOLVING_BONES_OF_THE_EXTREMITIES, HP_APLASIA_INVOLVING_BONES_OF_THE_EXTREMITIES GOBP_T_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_T_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE HP_CUBITUS_VALGUS, HP_CUBITUS_VALGUS HP_HEMORRHAGE_OF_THE_EYE, HP_HEMORRHAGE_OF_THE_EYE
	GOBP_POSITIVE_REGULATION_OF_PHOSPHATASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_PHOSPHATASE_ACTIVITY HP_HEMATEMESIS, HP_HEMATEMESIS BIOCARTA_IL2_PATHWAY, BIOCARTA_IL2_PATHWAY HP_PROMINENT_SUPERFICIAL_BLOOD_VESSELS, HP_PROMINENT_SUPERFICIAL_BLOOD_VESSELS GOBP_REGULATION_OF_PROTEIN_DEACETYLATION, GOBP_REGULATION_OF_PROTEIN_DEACETYLATION GOBP_POSITIVE_REGULATION_OF_PEPTIDYL_LYSINE_ACETYLATION, GOBP_POSITIVE_REGULATION_OF_PEPTIDYL_LYSINE_ACETYLATION GOBP_GLUCOSE_IMPORT, GOBP_GLUCOSE_IMPORT HP_EEG_WITH_ABNORMALLY_SLOW_FREQUENCIES, HP_EEG_WITH_ABNORMALLY_SLOW_FREQUENCIES
	WP_ALPHA_6_BETA_4_SIGNALING_PATHWAY, WP_ALPHA_6_BETA_4_SIGNALING_PATHWAY HP_UVEITIS, HP_UVEITIS BIOCARTA_SPPA_PATHWAY, BIOCARTA_SPPA_PATHWAY REACTOME_FATTY_ACYL_COA_BIOSYNTHESIS, REACTOME_FATTY_ACYL_COA_BIOSYNTHESIS GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ORGANIZATION GOBP_PROTEIN_LOCALIZATION_TO_CELL_CELL_JUNCTION, GOBP_PROTEIN_LOCALIZATION_TO_CELL_CELL_JUNCTION GOMF_MAP_KINASE_KINASE_KINASE_ACTIVITY, GOMF_MAP_KINASE_KINASE_KINASE_ACTIVITY GOCC_IMMUNOLOGICAL_SYNAPSE, GOCC_IMMUNOLOGICAL_SYNAPSE
	GOMF_NAD_DEPENDENT_PROTEIN_DEACETYLASE_ACTIVITY, GOMF_NAD_DEPENDENT_PROTEIN_DEACETYLASE_ACTIVITY HP_MOTOR_POLYNEUROPATHY, HP_MOTOR_POLYNEUROPATHY GOBP_REGULATION_OF_INTEGRIN_ACTIVATION, GOBP_REGULATION_OF_INTEGRIN_ACTIVATION HP_RETICULOCYTOSIS, HP_RETICULOCYTOSIS GOBP_INTEGRIN_ACTIVATION, GOBP_INTEGRIN_ACTIVATION HP_MODERATE_SENSORINEURAL_HEARING_IMPAIRMENT, HP_MODERATE_SENSORINEURAL_HEARING_IMPAIRMENT GOBP_REGULATION_OF_FILOPODIUM_ASSEMBLY, GOBP_REGULATION_OF_FILOPODIUM_ASSEMBLY ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_3DY_DN, ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_3DY_DN GOBP_LAMELLIPODIUM_ASSEMBLY, GOBP_LAMELLIPODIUM_ASSEMBLY
	BIOCARTA_AGR_PATHWAY, BIOCARTA_AGR_PATHWAY HP_ABNORMALITY_OF_THE_NASOLABIAL_REGION, HP_ABNORMALITY_OF_THE_NASOLABIAL_REGION BIOCARTA_SIP_PATHWAY, BIOCARTA_SIP_PATHWAY GOBP_MUSCLE_ATROPHY, GOBP_MUSCLE_ATROPHY HP_LOWER_LIMB_UNDERGROWTH, HP_LOWER_LIMB_UNDERGROWTH REACTOME_PI5P_REGULATES_TP53_ACETYLATION, REACTOME_PI5P_REGULATES_TP53_ACETYLATION MIR6753_3P, MIR6753_3P GOBP_ER_OVERLOAD_RESPONSE, GOBP_ER_OVERLOAD_RESPONSE
	HP_AVASCULAR_NECROSIS_OF_THE_CAPITAL_FEMORAL_EPIPHYSIS, HP_AVASCULAR_NECROSIS_OF_THE_CAPITAL_FEMORAL_EPIPHYSIS MIR4309, MIR4309 HOEGERKORP_CD44_TARGETS_TEMPORAL_DN, HOEGERKORP_CD44_TARGETS_TEMPORAL_DN MIR1225_5P, MIR1225_5P HP_SCHIZOPHRENIA, HP_SCHIZOPHRENIA MORF_BMPR2, MORF_BMPR2 GOBP_CELLULAR_RESPONSE_TO_CADMIUM_ION, GOBP_CELLULAR_RESPONSE_TO_CADMIUM_ION SHIN_B_CELL_LYMPHOMA_CLUSTER_2, SHIN_B_CELL_LYMPHOMA_CLUSTER_2 GOMF_CALCIUM_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GOMF_CALCIUM_DEPENDENT_PROTEIN_KINASE_ACTIVITY
	MIR7107_3P, MIR7107_3P FULLER_PBMC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_18HR_TO_336HR_SUSTAINED_UP, FULLER_PBMC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_18HR_TO_336HR_SUSTAINED_UP GOBP_NEURON_MATURATION, GOBP_NEURON_MATURATION HP_ABNORMAL_PELVIS_BONE_MORPHOLOGY, HP_ABNORMAL_PELVIS_BONE_MORPHOLOGY MIR151A_3P, MIR151A_3P MIR6735_3P, MIR6735_3P GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_PROLIFERATION WP_CALORIC_RESTRICTION_AND_AGING, WP_CALORIC_RESTRICTION_AND_AGING
	GOBP_RESPONSE_TO_COPPER_ION, GOBP_RESPONSE_TO_COPPER_ION GOMF_PROFILIN_BINDING, GOMF_PROFILIN_BINDING KASLER_HDAC7_TARGETS_1_DN, KASLER_HDAC7_TARGETS_1_DN GOBP_RESPONSE_TO_INTERFERON_ALPHA, GOBP_RESPONSE_TO_INTERFERON_ALPHA HP_SEBORRHEIC_DERMATITIS, HP_SEBORRHEIC_DERMATITIS GOBP_PROTOPORPHYRINOGEN_IX_METABOLIC_PROCESS, GOBP_PROTOPORPHYRINOGEN_IX_METABOLIC_PROCESS HP_APLASIA_HYPOPLASIA_OF_THE_ABDOMINAL_WALL_MUSCULATURE, HP_APLASIA_HYPOPLASIA_OF_THE_ABDOMINAL_WALL_MUSCULATURE GOBP_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_SPHINGOLIPID_BIOSYNTHETIC_PROCESS (PRC1_BMI_UP.V1_UP, PRC1_BMI_UP.V1_UP
ELL_AND_IL12_TREATED_MELANOMA_DAY3_UP, GSE29164_CD8_TCELL_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_UP	MIR5196_3P, MIR5196_3P GOBP_LOCOMOTOR_RHYTHM, GOBP_LOCOMOTOR_RHYTHM MIR7109_3P, MIR7109_3P MODULE_236, MODULE_236 HP_ABNORMAL_PROXIMAL_PHALANX_MORPHOLOGY_OF_THE_HAND, HP_ABNORMAL_PROXIMAL_PHALANX_MORPHOLOGY_OF_THE_HAND WP_CELL_DIFFERENTIATION_INDEX, WP_CELL_DIFFERENTIATION_INDEX GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY GOBP_SYNAPTIC_VESICLE_ENDOSOMAL_PROCESSING, GOBP_SYNAPTIC_VESICLE_ENDOSOMAL_PROCESSING
	GOBP_PEPTIDYL_SERINE_AUTOPHOSPHORYLATION, GOBP_PEPTIDYL_SERINE_AUTOPHOSPHORYLATION GOBP_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS, GOBP_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS BIOCARTA_ERK_PATHWAY, BIOCARTA_ERK_PATHWAY MCCABE_HOXC6_TARGETS_CANCER_DN, MCCABE_HOXC6_TARGETS_CANCER_DN HP_TYPE_1_MUSCLE_FIBER_PREDOMINANCE, HP_TYPE_1_MUSCLE_FIBER_PREDOMINANCE GOBP_REGULATION_OF_CELL_ADHESION_MEDIATED_BY_INTEGRIN, GOBP_REGULATION_OF_CELL_ADHESION_MEDIATED_BY_INTEGRIN MORF_CTSB, MORF_CTSB HP_RETINAL_HEMORRHAGE, HP_RETINAL_HEMORRHAGE GOBP_REGULATION_OF_SUPEROXIDE_ANION_GENERATION, GOBP_REGULATION_OF_SUPEROXIDE_ANION_GENERATION
	HP_ELEVATED_CIRCULATING_LONG_CHAIN_FATTY_ACID_CONCENTRATION, HP_ELEVATED_CIRCULATING_LONG_CHAIN_FATTY_ACID_CONCENTRATION GOMF_BETA_1_ADRENERGIC_RECEPTOR_BINDING, GOMF_BETA_1_ADRENERGIC_RECEPTOR_BINDING HP_THICK_CEREBRAL_CORTEX, HP_THICK_CEREBRAL_CORTEX GOBP_T_CELL_LINEAGE_COMMITMENT, GOBP_T_CELL_LINEAGE_COMMITMENT HP_ABNORMAL_MACROSCOPIC_URINE_APPEARANCE, HP_ABNORMAL_MACROSCOPIC_URINE_APPEARANCE GOBP_TRIGLYCERIDE_BIOSYNTHETIC_PROCESS, GOBP_TRIGLYCERIDE_BIOSYNTHETIC_PROCESS HP_SECONDARY_AMENORRHEA, HP_SECONDARY_AMENORRHEA MORF_PRKCA, MORF_PRKCA
	HP_NON_MIDLINE_CLEFT_LIP, HP_NON_MIDLINE_CLEFT_LIP HP_PES_VALGUS, HP_PES_VALGUS GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_LIGHTYELLOW_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_LIGHTYELLOW_UP HP_T_CELL_LYMPHOMA, HP_T_CELL_LYMPHOMA GOBP_RESPONSE_TO_INTERLEUKIN_15, GOBP_RESPONSE_TO_INTERLEUKIN_15 GOBP_ATF6_MEDIATED_UNFOLDED_PROTEIN_RESPONSE, GOBP_ATF6_MEDIATED_UNFOLDED_PROTEIN_RESPONSE GOCC_SCAR_COMPLEX, GOCC_SCAR_COMPLEX GOBP_REGULATION_OF_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS, GOBP_REGULATION_OF_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS HP_FRACTURES_OF_THE_LONG_BONES, HP_FRACTURES_OF_THE_LONG_BONES
	HP_ABNORMAL_HUMERUS_MORPHOLOGY, HP_ABNORMAL_HUMERUS_MORPHOLOGY LIU_CMYB_TARGETS_DN, LIU_CMYB_TARGETS_DN HP_SHORT_LOWER_LIMBS, HP_SHORT_LOWER_LIMBS GOBP_MITOCHONDRIAL_RIBOSOME_ASSEMBLY, GOBP_MITOCHONDRIAL_RIBOSOME_ASSEMBLY GOBP_EPIBOLY, GOBP_EPIBOLY WP_INSULIN_SIGNALLING_IN_HUMAN_ADIPOCYTES_NORMAL_CONDITION, WP_INSULIN_SIGNALLING_IN_HUMAN_ADIPOCYTES_NORMAL_CONDITION WP_INSULIN_SIGNALLING_IN_HUMAN_ADIPOCYTES_DIABETIC_CONDITION, WP_INSULIN_SIGNALLING_IN_HUMAN_ADIPOCYTES_DIABETIC_CONDITION GOBP_PEPTIDYL_LYSINE_DEACETYLATION, GOBP_PEPTIDYL_LYSINE_DEACETYLATION
	GOMF_RDNA_BINDING, GOMF_RDNA_BINDING GOMF_RDNA_BINDING, GOMF_RDNA_BINDING GOMF_K63_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GOMF_K63_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING GOBP_MORPHOGENESIS_OF_AN_EPITHELIAL_SHEET, GOBP_MORPHOGENESIS_OF_AN_EPITHELIAL_SHEET HP_DECREASED_MUSCLE_MASS, HP_DECREASED_MUSCLE_MASS HP_EEG_WITH_MULTIFOCAL_SLOW_ACTIVITY, HP_EEG_WITH_MULTIFOCAL_SLOW_ACTIVITY MORF_MAP2K7, MORF_MAP2K7 HP_ABNORMAL_PROPORTION_OF_CD8_POSITIVE_T_CELLS, HP_ABNORMAL_PROPORTION_OF_CD8_POSITIVE_T_CELLS GOBP_MUSCLE_ADAPTATION, GOBP_MUSCLE_ADAPTATION HP_PROMINENT_NASOLABIAL_FOLD, HP_PROMINENT_NASOLABIAL_FOLD
	REACTOME_SEALING_OF_THE_NUCLEAR_ENVELOPE_NE_BY_ESCRT_III, REACTOME_SEALING_OF_THE_NUCLEAR_ENVELOPE_NE_BY_ESCRT_III GOBP_POSITIVE_REGULATION_OF_PHOSPHOLIPID_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_PHOSPHOLIPID_METABOLIC_PROCESS GOBP_STRIATED_MUSCLE_ATROPHY, GOBP_STRIATED_MUSCLE_ATROPHY GOBP_RESPONSE_TO_CGMP, GOBP_RESPONSE_TO_CGMP GOBP_ENDOSOME_TO_LYSOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_ENDOSOME_TO_LYSOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY GOBP_NEURAL_CREST_CELL_DIFFERENTIATION, GOBP_NEURAL_CREST_CELL_DIFFERENTIATION GOCC_CELL_TRAILING_EDGE, GOCC_CELL_TRAILING_EDGE GOBP_POSITIVE_REGULATION_OF_HISTONE_DEACETYLATION, GOBP_POSITIVE_REGULATION_OF_HISTONE_DEACETYLATION
	HP_FOCAL_CORTICAL_DYSPLASIA_TYPE_II, HP_FOCAL_CORTICAL_DYSPLASIA_TYPE_II HP_ABNORMAL_SARCOMERE_MORPHOLOGY, HP_ABNORMAL_SARCOMERE_MORPHOLOGY HP_PULMONARY_VENOUS_HYPERTENSION, HP_PULMONARY_VENOUS_HYPERTENSION HP_HEMERALOPIA, HP_HEMERALOPIA GOCC_LEWY_BODY, GOCC_LEWY_BODY GOMF_K48_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GOMF_K48_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING GOBP_DEVELOPMENTAL_PIGMENTATION, GOBP_DEVELOPMENTAL_PIGMENTATION GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEACETYLATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEACETYLATION HP_BUNDLE_BRANCH_BLOCK, HP_BUNDLE_BRANCH_BLOCK
	GOBP_REGULATION_OF_PLATELET_ACTIVATION, GOBP_REGULATION_OF_PLATELET_ACTIVATION GOBP_LONG_CHAIN_FATTY_ACYL_COA_METABOLIC_PROCESS, GOBP_LONG_CHAIN_FATTY_ACYL_COA_METABOLIC_PROCESS GOBP_POSITIVE_REGULATION_OF_GLUCOSE_IMPORT, GOBP_POSITIVE_REGULATION_OF_GLUCOSE_IMPORT HOFFMANN_PRE_BI_TO_LARGE_PRE_BII_LYMPHOCYTE_UP, HOFFMANN_PRE_BI_TO_LARGE_PRE_BII_LYMPHOCYTE_UP GOBP_REGULATION_OF_FEEDING_BEHAVIOR, GOBP_REGULATION_OF_FEEDING_BEHAVIOR GOBP_REGULATION_OF_NATURAL_KILLER_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_NATURAL_KILLER_CELL_DIFFERENTIATION REACTOME_CREB3_FACTORS_ACTIVATE_GENES, REACTOME_CREB3_FACTORS_ACTIVATE_GENES GOBP_CELLULAR_RESPONSE_TO_COPPER_ION, GOBP_CELLULAR_RESPONSE_TO_COPPER_ION
	HP_RIGHT_BUNDLE_BRANCH_BLOCK, HP_RIGHT_BUNDLE_BRANCH_BLOCK GOBP_POSITIVE_REGULATION_OF_VASCULOGENESIS, GOBP_POSITIVE_REGULATION_OF_VASCULOGENESIS GOBP_REGULATION_OF_SECONDARY_METABOLIC_PROCESS, GOBP_REGULATION_OF_SECONDARY_METABOLIC_PROCESS IIZUKA_LIVER_CANCER_PROGRESSION_G1_G2_UP, IIZUKA_LIVER_CANCER_PROGRESSION_G1_G2_UP GOBP_CHOLESTEROL_CATABOLIC_PROCESS, GOBP_CHOLESTEROL_CATABOLIC_PROCESS PID_P38_GAMMA_DELTA_PATHWAY, PID_P38_GAMMA_DELTA_PATHWAY HP_CHORIORETINAL_DYSTROPHY, HP_CHORIORETINAL_DYSTROPHY REACTOME_FLT3_SIGNALING_THROUGH_SRC_FAMILY_KINASES, REACTOME_FLT3_SIGNALING_THROUGH_SRC_FAMILY_KINASES
	HP_MACROVESICULAR_HEPATIC_STEATOSIS, HP_MACROVESICULAR_HEPATIC_STEATOSIS GOBP_CD4_POSITIVE_OR_CD8_POSITIVE_ALPHA_BETA_T_CELL_LINEAGE_COMMITMENT, GOBP_CD4_POSITIVE_OR_CD8_POSITIVE_ALPHA_BETA_T_CELL_LINEAGE_COMMITMENT GOBP_LONG_CHAIN_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS, GOBP_LONG_CHAIN_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS MIR6502_5P, MIR6502_5P GOBP_LONG_TERM_MEMORY, GOBP_LONG_TERM_MEMORY ZNF791_TARGET_GENES, ZNF791_TARGET_GENES HP_ACHALASIA, HP_ACHALASIA GOMF_MICROTUBULE_MINUS_END_BINDING, GOMF_MICROTUBULE_MINUS_END_BINDING MIR4750_3P, MIR4750_3P
	GOBP_SKELETAL_MUSCLE_ADAPTATION, GOBP_SKELETAL_MUSCLE_ADAPTATION HP_PRELINGUAL_SENSORINEURAL_HEARING_IMPAIRMENT, HP_PRELINGUAL_SENSORINEURAL_HEARING_IMPAIRMENT HP_SPONDYLOEPIPHYSEAL_DYSPLASIA, HP_SPONDYLOEPIPHYSEAL_DYSPLASIA VERRECCHIA_RESPONSE_TO_TGFB1_C3, VERRECCHIA_RESPONSE_TO_TGFB1_C3 GOBP_PEROXISOME_FISSION, GOBP_PEROXISOME_FISSION GOMF_INSULIN_RECEPTOR_SUBSTRATE_BINDING, GOMF_INSULIN_RECEPTOR_SUBSTRATE_BINDING BIOCARTA_IL22BP_PATHWAY, BIOCARTA_IL22BP_PATHWAY HP_MACRODONTIA_OF_PERMANENT_MAXILLARY_CENTRAL_INCISOR, HP_MACRODONTIA_OF_PERMANENT_MAXILLARY_CENTRAL_INCISOR
	BIOCARTA_IFNA_PATHWAY, BIOCARTA_IFNA_PATHWAY REACTOME_SYNTHESIS_OF_VERY_LONG_CHAIN_FATTY_ACYL_COAS, REACTOME_SYNTHESIS_OF_VERY_LONG_CHAIN_FATTY_ACYL_COAS GOBP_POSITIVE_REGULATION_OF_GLUCOSE_TRANSMEMBRANE_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_GLUCOSE_TRANSMEMBRANE_TRANSPORT MIR1301_5P, MIR1301_5P TURASHVILI_BREAST_NORMAL_DUCTAL_VS_LOBULAR_DN, TURASHVILI_BREAST_NORMAL_DUCTAL_VS_LOBULAR_DN GOBP_LATE_ENDOSOME_TO_LYSOSOME_TRANSPORT, GOBP_LATE_ENDOSOME_TO_LYSOSOME_TRANSPORT GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_WOUNDING, GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_WOUNDING CARM1_TARGET_GENES, CARM1_TARGET_GENES GOBP_POSITIVE_REGULATION_OF_CELL_GROWTH_INVOLVED_IN_CARDIAC_MUSCLE_CELL_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_CELL_GROWTH_INVOLVED_IN_CARDIAC_MUSCLE_CELL_DEVELOPMENT
	GOBP_ICOSANOID_BIOSYNTHETIC_PROCESS, GOBP_ICOSANOID_BIOSYNTHETIC_PROCESS GOBP_PROSTANOID_METABOLIC_PROCESS, GOBP_PROSTANOID_METABOLIC_PROCESS ZHU_SKIL_TARGETS_UP, ZHU_SKIL_TARGETS_UP GOBP_CELLULAR_RESPONSE_TO_THYROID_HORMONE_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_THYROID_HORMONE_STIMULUS GOBP_ICOSANOID_SECRETION, GOBP_ICOSANOID_SECRETION chr6p24, chr6p24 HOLLEMAN_PREDNISOLONE_RESISTANCE_ALL_DN, HOLLEMAN_PREDNISOLONE_RESISTANCE_ALL_DN MIR3690, MIR3690
	HP_OCCIPITAL_MENINGOCELE, HP_OCCIPITAL_MENINGOCELE HP_POLAR_CATARACT, HP_POLAR_CATARACT HOXA3_01, HOXA3_01 GOBP_EMBRYONIC_DIGIT_MORPHOGENESIS, GOBP_EMBRYONIC_DIGIT_MORPHOGENESIS WP_TYPE_II_DIABETES_MELLITUS, WP_TYPE_II_DIABETES_MELLITUS HP_CONE_CONE_ROD_DYSTROPHY, HP_CONE_CONE_ROD_DYSTROPHY HP_BILATERAL_CLEFT_LIP_AND_PALATE, HP_BILATERAL_CLEFT_LIP_AND_PALATE GOCC_EARLY_PHAGOSOME, GOCC_EARLY_PHAGOSOME
	REACTOME_INTERLEUKIN_2_SIGNALING, REACTOME_INTERLEUKIN_2_SIGNALING HP_EPISCLERITIS, HP_EPISCLERITIS GOBP_IODIDE_TRANSPORT, GOBP_IODIDE_TRANSPORT GOCC_NATA_COMPLEX, GOCC_NATA_COMPLEX GOCC_CORVET_COMPLEX, GOCC_CORVET_COMPLEX GOMF_CYSTEINE_TYPE_EXOPEPTIDASE_ACTIVITY, GOMF_CYSTEINE_TYPE_EXOPEPTIDASE_ACTIVITY GOMF_RNA_POLYMERASE_II_C_TERMINAL_DOMAIN_PHOSPHOSERINE_BINDING, GOMF_RNA_POLYMERASE_II_C_TERMINAL_DOMAIN_PHOSPHOSERINE_BINDING HP_RECURRENT_INTRAPULMONARY_HEMORRHAGE, HP_RECURRENT_INTRAPULMONARY_HEMORRHAGE ZNF514_TARGET_GENES, ZNF514_TARGET_GENES
	GOBP_POSITIVE_REGULATION_OF_CHOLESTEROL_STORAGE, GOBP_POSITIVE_REGULATION_OF_CHOLESTEROL_STORAGE GOBP_POSITIVE_REGULATION_OF_DEACETYLASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_DEACETYLASE_ACTIVITY HP_SCHIZENCEPHALY, HP_SCHIZENCEPHALY GOBP_TRNA_AMINOACYLATION_FOR_MITOCHONDRIAL_PROTEIN_TRANSLATION, GOBP_TRNA_AMINOACYLATION_FOR_MITOCHONDRIAL_PROTEIN_TRANSLATION GOCC_GROWTH_CONE_MEMBRANE, GOCC_GROWTH_CONE_MEMBRANE HP_BILATERAL_CLEFT_LIP, HP_BILATERAL_CLEFT_LIP GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION, GOBP_GOLGI_RIBBON_FORMATION, GOBP_GOLGI_RIBBON_FORMATION
	HP_ABNORMAL_LYMPHATIC_VESSEL_MORPHOLOGY, HP_ABNORMAL_LYMPHATIC_VESSEL_MORPHOLOGY GOBP_FATTY_ACID_ELONGATION, GOBP_FATTY_ACID_ELONGATION GOBP_REGULATION_OF_BROWN_FAT_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_BROWN_FAT_CELL_DIFFERENTIATION GOBP_RHYTHMIC_BEHAVIOR, GOBP_RHYTHMIC_BEHAVIOR GOBP_GLYCEROLIPID_CATABOLIC_PROCESS, GOBP_GLYCEROLIPID_CATABOLIC_PROCESS GOBP_FOREBRAIN_NEURON_DEVELOPMENT, GOBP_FOREBRAIN_NEURON_DEVELOPMENT GOBP_POSITIVE_REGULATION_OF_WOUND_HEALING, GOBP_POSITIVE_REGULATION_OF_WOUND_HEALING GOBP_ADULT_FEEDING_BEHAVIOR, GOBP_ADULT_FEEDING_BEHAVIOR GOBP_NEGATIVE_REGULATION_OF_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM
	ADCYAP1_TARGET_GENES, ADCYAP1_TARGET_GENES GOBP_PEPTIDYL_LYSINE_MONOMETHYLATION, GOBP_PEPTIDYL_LYSINE_MONOMETHYLATION GOBP_POSITIVE_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_GOBP_VITAMIN_TRANSPORT, GOBP_VITAMIN_TRANSPORT NIKOLSKY_BREAST_CANCER_21Q22_AMPLICON, NIKOLSKY_BREAST_CANCER_21Q22_AMPLICON HP_RECURRENT_CORNEAL_EROSIONS, HP_RECURRENT_CORNEAL_EROSIONS HP_NARROW_PALM, HP_NARROW_PALM GOBP_PHOSPHATIDYLCHOLINE_CATABOLIC_PROCESS, GOBP_PHOSPHATIDYLCHOLINE_CATABOLIC_PROCESS
	GOBP_GAMMA_DELTA_T_CELL_DIFFERENTIATION, GOBP_GAMMA_DELTA_T_CELL_DIFFERENTIATION MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_8, MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_8 GOCC_TORC2_COMPLEX, GOCC_TORC2_COMPLEX KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_4NQO_IN_OLD, KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_4NQO_IN_OLD MIR615_3P, MIR615_3P HP_COLD_INDUCED_MUSCLE_CRAMPS, HP_COLD_INDUCED_MUSCLE_CRAMPS HP_HEMOGLOBINURIA, HP_HEMOGLOBINURIA HP_BONE_FRACTURE, HP_BONE_FRACTURE HP_MINICORE_MYOPATHY, HP_MINICORE_MYOPATHY
	GOBP_REGULATION_OF_LIPID_TRANSPORTER_ACTIVITY, GOBP_REGULATION_OF_LIPID_TRANSPORTER_ACTIVITY GOBP_DENDRITIC_CELL_APOPTOTIC_PROCESS, GOBP_DENDRITIC_CELL_APOPTOTIC_PROCESS MORF_TTN, MORF_TTN chr8p21 WP_CELL_DIFFERENTIATION_INDEX_EXPANDED, WP_CELL_DIFFERENTIATION_INDEX_EXPANDED GOBP_NEUTRAL_LIPID_BIOSYNTHETIC_PROCESS, GOBP_NEUTRAL_LIPID_BIOSYNTHETIC_PROCESS GOMF_PROTEIN_KINASE_C_ACTIVITY, GOMF_PROTEIN_KINASE_C_ACTIVITY GOBP_POSITIVE_REGULATION_OF_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS, GOBP_POSITIVE_REGULATION_OF_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS
	MORF_PTPRR, MORF_PTPRR HP_ABNORMAL_PROPORTION_OF_CLASS_SWITCHED_MEMORY_B_CELLS, HP_ABNORMAL_PROPORTION_OF_CLASS_SWITCHED_MEMORY_B_CELLS GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_HYDROGEN_PEROXIDE, GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_HYDROGEN_PEROXIDE GOBP_NEGATIVE_REGULATION_OF_SECONDARY_METABOLITE_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_SECONDARY_METABOLITE_BIOSYNTHETIC_PROCESS GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATIONAL_INITIATION, GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATIONAL_INITIATION TURJANSKI_MAPK11_TARGETS, TURJANSKI_MAPK11_TARGETS REACTOME_HDL_CLEARANCE, REACTOME_HDL_CLEARANCE HP_CONGENITAL_HEMOLYTIC_ANEMIA, HP_CONGENITAL_HEMOLYTIC_ANEMIA GOBP_GLYCERALDEHYDE_3_PHOSPHATE_METABOLIC_PROCESS, GOBP_GLYCERALDEHYDE_3_PHOSPHATE_METABOLIC_PROCESS
	HP_POSTERIOR_POLAR_CATARACT, HP_POSTERIOR_POLAR_CATARACT REACTOME_REGULATION_OF_CYTOSKELETAL_REMODELING_AND_CELL_SPREADING_BY_IPP_COMPLEX_COMPONENTS, REACTOME_REGULATION_OF_CYTOSKELETAL_REMODELING_BY_IPP_COMPLEX_COMPONENTS, REACTOME_REGULATION_OF_CYTOSKELETAL_REMODELING_BY_IPP_COMPLEX_COMPONENTS, REACTOME_REGULATION_OF_CYTOSKELETAL_REMODELING_BY_IPP_COMPLEX_COMPONENTS, REACTOME_REGULATION_OF_CYTOSKELETAL_REMODELING_BY_IPP_COMPLEX_COMPONENTS, REACTOME_REGULATION_OF_CYTOSKELETAL_REMODELING_BY_IPP_COMPONENTS, REACTOME_REGULATION_OF_CYTOSKELETAL_REMODELING_BY_IPP_COMPONENTS, REACTOME_REGULATION_OF_CYTOSKELETAL_REMODELING_BY_IPP_COMPONENTS, REACTOME_REGULATION_OF_CYTOSKELETAL_REMODELING_BY_IPP_COMPONENTS, REACTOME_REGULATION_OF_LANG_IPP_C
	PIEPOLI_LGI1_TARGETS_UP, PIEPOLI_LGI1_TARGETS_UP GOBP_REGULATION_OF_HYDROGEN_PEROXIDE_METABOLIC_PROCESS, GOBP_REGULATION_OF_HYDROGEN_PEROXIDE_METABOLIC_PROCESS HP_INCREASED_MEAN_CORPUSCULAR_HEMOGLOBIN_CONCENTRATION, HP_INCREASED_MEAN_CORPUSCULAR_HEMOGLOBIN_CONCENTRATION REACTOME_SYNTHESIS_SECRETION_AND_DEACYLATION_OF_GHRELIN, REACTOME_SYNTHESIS_SECRETION_AND_DEACYLATION_OF_GHRELIN GOBP_REGULATION_OF_GLIAL_CELL_APOPTOTIC_PROCESS, GOBP_REGULATION_OF_GLIAL_CELL_APOPTOTIC_PROCESS GOBP_ELASTIC_FIBER_ASSEMBLY, GOBP_ELASTIC_FIBER_ASSEMBLY HP_APLASIA_INVOLVING_BONES_OF_THE_LOWER_LIMBS, HP_APLASIA_INVOLVING_BONES_OF_THE_LOWER_LIMBS HP_INTERMITTENT_JAUNDICE, HP_INTERMITTENT_JAUNDICE
	ZEILSTRA_CD44_TARGETS_DN, ZEILSTRA_CD44_TARGETS_DN MAP2K1_TARGET_GENES, MAP2K1_TARGET_GENES HP_RESPIRATORY_PARALYSIS, HP_RESPIRATORY_PARALYSIS GOMF_GROWTH_HORMONE_RECEPTOR_BINDING, GOMF_GROWTH_HORMONE_RECEPTOR_BINDING GOBP_REGULATION_OF_KILLING_OF_CELLS_OF_OTHER_ORGANISM, GOBP_REGULATION_OF_KILLING_OF_CELLS_OF_OTHER_ORGANISM GOBP_HIGH_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE, GOBP_HIGH_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE MATTIOLI_MULTIPLE_MYELOMA_SUBGROUPS, MATTIOLI_MULTIPLE_MYELOMA_SUBGROUPS KOBAYASHI_EGFR_SIGNALING_6HR_UP, KOBAYASHI_EGFR_SIGNALING_6HR_UP HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICULITIS_HP_FOLLICU
	HP_CYSTIC_RENAL_DYSPLASIA, HP_CYSTIC_RENAL_DYSPLASIA HP_FIRST_DORSAL_INTEROSSEI_MUSCLE_WEAKNESS, HP_FIRST_DORSAL_INTEROSSEI_MUSCLE_WEAKNESS HP_THENAR_MUSCLE_ATROPHY, HP_THENAR_MUSCLE_ATROPHY GOBP_FOREBRAIN_GENERATION_OF_NEURONS, GOBP_FOREBRAIN_GENERATION_OF_NEURONS GOBP_ACYLGLYCEROL_HOMEOSTASIS, GOBP_ACYLGLYCEROL_HOMEOSTASIS GOBP_TRYPTOPHAN_METABOLIC_PROCESS, GOBP_TRYPTOPHAN_METABOLIC_PROCESS MIR933, MIR933 HP_PILI_CANALICULI, HP_PILI_CANALICULI
	HP_PERSISTENT_EBV_VIREMIA, HP_PERSISTENT_EBV_VIREMIA GOBP_VOLUNTARY_MUSCULOSKELETAL_MOVEMENT, GOBP_VOLUNTARY_MUSCULOSKELETAL_MOVEMENT GOBP_POSITIVE_REGULATION_OF_CAMP_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_CAMP_DEPENDENT_PROTEIN_KINASE_ACTIVITY HP_PSYCHOTIC_EPISODES, HP_PSYCHOTIC_EPISODES GOBP_COPPER_ION_IMPORT, GOBP_COPPER_ION_IMPORT GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_DIFFERENTIATION REACTOME_METHIONINE_SALVAGE_PATHWAY, REACTOME_METHIONINE_SALVAGE_PATHWAY GOBP_POSITIVE_REGULATION_OF_BEHAVIOR, GOBP_POSITIVE_REGULATION_OF_BEHAVIOR HP_CENTRAL_ADRENAL_INSUFFICIENCY, HP_CENTRAL_ADRENAL_INSUFFICIENCY
	GOBP_POSITIVE_REGULATION_OF_CELL_ADHESION_MEDIATED_BY_INTEGRIN, GOBP_POSITIVE_REGULATION_OF_CELL_ADHESION_MEDIATED_BY_INTEGRIN HP_PERIPHERAL_VISUAL_FIELD_LOSS, HP_PERIPHERAL_VISUAL_FIELD_LOSS GOBP_STRIATED_MUSCLE_ADAPTATION, GOBP_STRIATED_MUSCLE_ADAPTATION GOBP_STRIATED_MUSCLE_ADAPTATION, GOBP_STRIATED_MUSCLE_ADAPTATION GOBP_KILLING_OF_CELLS_OF_OTHER_ORGANISM, GOBP_KILLING_OF_CELLS_OF_OTHER_ORGANISM REACTOME_CD209_DC_SIGN_SIGNALING, REACTOME_CD209_DC_SIGN_SIGNALING GOCC_ASTROCYTE_PROJECTION, GOCC_ASTROCYTE_PROJECTION KEGG_SULFUR_METABOLISM, KEGG_SULFUR_METABOLISM GOMF_DIACYLGLYCEROL_BINDING, GOMF_DIACYLGLYCEROL_BINDING
	GOBP_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION, GOBP_GRANULOCYTE_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION GOBP_T_HELPER_CELL_LINEAGE_COMMITMENT, GOBP_T_HELPER_CELL_LINEAGE_COMMITMENT GSE13411_IGM_VS_SWITCHED_MEMORY_BCELL_UP, GSE13411_IGM_VS_SWITCHED_MEMORY_BCELL_UP GOBP_ACTIVATION_OF_PROTEIN_KINASE_B_ACTIVITY, GOBP_ACTIVATION_OF_PROTEIN_KINASE_B_ACTIVITY HP_ELLIPTOCYTOSIS, HP_ELLIPTOCYTOSIS GOBP_REGULATION_OF_LAMELLIPODIUM_MORPHOGENESIS, GOBP_REGULATION_OF_LAMELLIPODIUM_MORPHOGENESIS HP_ABNORMAL_MUSCLE_GLYCOGEN_CONTENT, HP_ABNORMAL_MUSCLE_GLYCOGEN_CONTENT HP_ABNORMALITY_OF_THE_GLABELLA, HP_ABNORMALITY_OF_THE_GLABELLA
	FERRANDO_LYL1_NEIGHBORS, FERRANDO_LYL1_NEIGHBORS GOBP_T_HELPER_1_CELL_DIFFERENTIATION, GOBP_T_HELPER_1_CELL_DIFFERENTIATION HP_SPONTANEOUS_HEMATOMAS, HP_SPONTANEOUS_HEMATOMAS DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_PLASMA_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_PLASMA_CELLS GOBP_REGULATION_OF_BEHAVIOR, GOBP_REGULATION_OF_BEHAVIOR GOBP_REGULATION_OF_GAMMA_DELTA_T_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_GAMMA_DELTA_T_CELL_DIFFERENTIATION GOBP_POSITIVE_REGULATION_OF_GAMMA_DELTA_T_CELL_ACTIVATION, GOBP_POSITIVE_REGULATION_OF_GAMMA_DELTA_T_CELL_ACTIVATION MIR4707_5P, MIR4707_5P GOBP_NATURAL_KILLER_CELL_DEGRANULATION, GOBP_NATURAL_KILLER_CELL_DEGRANULATION
	GOBP_NATURAL_KILLER_CELL_DEGRANULATION, GOBP_NATURAL_KILLER_CELL_DEGRANULATION GOBP_NATURAL_KILLER_CELL_DEGRANULATION, GOBP_NATURAL_KILLER_CELL_DEGRANULATION GOBP_PHINGOMYELIN_CATABOLIC_PROCESS, GOBP_SPHINGOMYELIN_CATABOLIC_PROCESS GOBP_REGULATION_OF_SYNAPTIC_TRANSMISSION_DOPAMINERGIC GOBP_POSITIVE_REGULATION_OF_GAMMA_DELTA_T_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_GAMMA_DELTA_T_CELL_DIFFERENTIATION GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_MORPHOGENESIS, GOBP_REGULATION_OF_INORGANIC_ANION_TRANSMEMBRANE_TRANSPORT GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_MORPHOGENESIS, GOBP_POSITIVE_REGULATION_OF_LAMELLIPODIUM_MORPHOGENESIS HP_BULL_S_EYE_MACULOPATHY, HP_BULL_S_EYE_MACULOPATHY GOCC_DENDRITIC_SPINE_NECK, GOCC_DENDRITIC_SPINE_NECK GOMF_RNA_POLYMERASE_II_INTRONIC_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING, GOMF_RNA_POLYMERASE_II_INTRONIC_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING, GOMF_RNA_POLYMERASE_II_INTRONIC_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING.
	REACTOME_RUNX3_REGULATES_CDKN1A_TRANSCRIPTION, REACTOME_RUNX3_REGULATES_CDKN1A_TRANSCRIPTION HP_SEVERE_DEMYELINATION_OF_THE_WHITE_MATTER, HP_SEVERE_DEMYELINATION_OF_THE_WHITE_MATTER HP_SHORT_PROXIMAL_PHALANX_OF_FINGER, HP_SHORT_PROXIMAL_PHALANX_OF_FINGER HP_KELOIDS, HP_KELOIDS PRC1_BMI_UP.V1_DN, PRC1_BMI_UP.V1_DN HP_VARICOSE_VEINS, HP_VARICOSE_VEINS SHIN_B_CELL_LYMPHOMA_CLUSTER_6, SHIN_B_CELL_LYMPHOMA_CLUSTER_6 HP_DECREASED_PROPORTION_OF_CD8_POSITIVE_T_CELLS, HP_DECREASED_PROPORTION_OF_CD8_POSITIVE_T_CELLS INAMURA_LUNG_CANCER_SCC_DN, INAMURA_LUNG_CANCER_SCC_DN
	GOBP_FEEDING_BEHAVIOR, GOBP_FEEDING_BEHAVIOR GOBP_CELLULAR_RESPONSE_TO_VITAMIN_D, GOBP_CELLULAR_RESPONSE_TO_VITAMIN_D GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_OXIDATIVE_STRESS, GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_OXIDATIVE_STRESS MORF_ITGA2, MORF_ITGA2 HP_ABNORMALITY_OF_THE_SUBUNGUAL_REGION, HP_ABNORMALITY_OF_THE_SUBUNGUAL_REGION GOMF_INTRONIC_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING, GOMF_INTRONIC_TRANSCRIPTION_REGULATORY_REGION_SEQUENCE_SPECIFIC_DNA_BINDING GOBP_SEROTONIN_SECRETION, GOBP_SEROTONIN_SECRETION GOBP_REGULATION_OF_GASTRIC_ACID_SECRETION, GOBP_REGULATION_OF_GASTRIC_ACID_SECRETION
	GOBP_REGULATION_OF_GASTRIC_ACID_SECRETION, GOBP_REGULATION_OF_GASTRIC_ACID_SECRETION GOBP_REGULATION_OF_SUPEROXIDE_METABOLIC_PROCESS, GOBP_REGULATION_OF_SUPEROXIDE_METABOLIC_PROCESS GOBP_VITAMIN_TRANSMEMBRANE_TRANSPORT, GOBP_VITAMIN_TRANSMEMBRANE_TRANSPORT GOMF_GABA_RECEPTOR_BINDING, GOMF_GABA_RECEPTOR_BINDING GOBP_POSITIVE_REGULATION_OF_MYOTUBE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_MYOTUBE_DIFFERENTIATION WP_GALANIN_RECEPTOR_PATHWAY, WP_GALANIN_RECEPTOR_PATHWAY