	GSE41867_LCMV_ARMSTRONG_VS_CLONE13_DAY6_EFFECTOR_CD8_TCELL_DN, GSE41867_LCMV_ARMSTRONG_VS_CLONE13_DAY6_EFFECTOR_CD8_TCELL_DN GSE41867_NAIVE_VS_DAY30_LCMV_ARMSTRONG_MEMORY_CD8_TCELL_DN, GSE41867_NAIVE_VS_DAY30_LCMV_ARMSTRONG_MEMORY_CD8_TCELL_DN GSE41867_NAIVE_VS_DAY15_LCMV_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY15_LCMV_EFFECTOR_CD8_TCELL_UP GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_DN, GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_DN GSE1721_CTRL_VS_POLYIC_4H_BMDC_UP, GSE1721_CTRL_VS_POLYIC_4H_BMDC_UP GSE42492_WT_VS_PARG_KO_MACROPHAGE_UP, GSE4492_WT_VS_PARG_KO_MACROPHAGE_UP GSE44649_WT_VS_MIR155_KO_ACTIVATED_CD8_TCELL_UP, GSE44649_WT_VS_MIR155_KO_ACTIVATED_CD8_TCELL_UP GSE2770_UNTREATED_VS_ACT_CD4_TCELL_48H_UP, GSE2770_UNTREATED_VS_ACT_CD4_TCELL_48H_UP GSE3982_NEUTROPHIL_VS_CENT_MEMORY_CD4_TCELL_1N, GSE3982_NEUTROPHIL_VS_CENT_MEMORY_CD4_TCELL_UP
	GSE3039_ALPHAALPHA_VS_ALPHABETA_CD8_TCELL_UP, GSE3039_ALPHAALPHA_VS_ALPHABETA_CD8_TCELL_UP GSE10240_CTRL_VS_ILI7_AND_IL2_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_UP, GSE10240_CTRL_VS_ILI7_AND_IL22_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_UP GSE1052_ANTICD3CD28_STIM_VS_UNSTIM_MEMORY_CD8_TCELL_UP, GSE16522_ANTICD3CD28_STIM_VS_UNSTIM_MEMORY_CD8_TCELL_UP GSE1074_CD56_BRIGHT_VS_DIM_CD621_POSITIVE_NK_CELL_DN, GSE302774_CD56_BRIGHT_VS_DIM_CD621_POSITIVE_NK_CELL_DN GSE8921_3H_VS_24H_TLR1_2_STIM_MONOCYTE_DN, GSE8921_3H_VS_24H_TLR1_2_STIM_MONOCYTE_DN GSE43863_LY6C_INT_CXCR5POS_VS_LY6C_LOW_CXCR5NEG_EFFECTOR_CD4_TCELL_DN, GSE43863_LY6C_INT_CXCR5POS_VS_LY6C_LOW_CXCR5NEG_EFFECTOR_CD4_TCELL_DN GSE40274_CTRL_VS_HEILOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE4374_CTRL_VS_HEILOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN GSE40274_CTRL_VS_VS_DOS_MONOCYTE_UP_GSE4355_CD16_NEC_VS_DSS_MONOCYTE_UP_GSE4375_CD16_NEC_VS_DSS_MONOCYTE_UP_GS_MONOCY
	GSE34515_CD16_NEG_VS_POS_MONOCYTE_UP, GSE34515_CD16_NEG_VS_POS_MONOCYTE_UP GSE30971_ZH_VS_4H_LIPS_STIM_MACROPHAGE_WBP7_HET_UP, GSE30971_ZH_VS_4H_LIPS_STIM_MACROPHAGE_WBP7_HET_UP GSE43863_TFH_VS_LY6C_INT_CXCR5POS_MEMORY_CD4_TCELL_UP, GSE43863_TFH_VS_LY6C_INT_CXCR5POS_MEMORY_CD4_TCELL_UP GSE17721_CTTRL_VS_GARDIQUIMOD_2H_BMDC_DN, GSE17721_CTRL_VS_GRADIQUIMOD_2H_BMDC_DN GSE43863_TFH_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN, GSE43863_TFH_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN GSE43863_TFH_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN, GSE43863_TFH_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN SENESE_HDAC2_TARGFTS_UP, SENESE_HDAC2_TARGFTS_UP GSE11961_FOLLICULAR_BCELL_VS_PLASMA_CELL_DAY7_UP GSE11961_FOLLICULAR_BCELL_VS_PLASMA_CELL_DAY7_UP GOBP_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_ENDONUCLEOLYTIC, GOBP_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_ENDONUCLEOLYTIC NEMETH_INFLAMMATORY_RESPONSE_LPS_UP, NEMETH_INFLAMMATORY_RESPONSE_LPS_UP GSE17721_CTRL_VS_LIPS_2H_BMDC_DN, GSE17721_CTRL_VS_LIPS_2H_BMDC_DN
	GOMF_METAL_CLUSTER_BINDING, GOMF_METAL_CLUSTER_BINDING ICHIBA_CRAFT_VERSUS_HOST_DISEASE_DT_UP, ICHIBA_CRAFT_VERSUS_HOST_DISEASE_DT_UP GSE2585_CTEC_VS_MTEC_THYMUS_DN, GSE2585_CTEC_VS_MTEC_THYMUS_DN GSE3982_BASOPHIL_VS_NKCELL_DN, GSE3982_BASOPHIL_VS_NKCELL_DN MIR122_5P, MIR122_5P GSE40274_CTRL_VS_FOXP3_AND_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_CTRL_VS_FOXP3_AND_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_UP GOBP_MATURATION_OF_SSU_RRNA, GOBP_MATURATION_OF_SSU_RRNA GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_DN RAX2_TARGET_GENES_RAX2_TARGET_GENES
	GOMF_ENDORIBONUCLEASE_ACTIVITY, GOMF_ENDORIBONUCLEASE_ACTIVITY GOBP_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GOBP_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION AR_01, AR_01 GOBP_B_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_B_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE MIR_150_3P, MIR_150_3P GOMF_4_IRON_4_SULFUR_CLUSTER_BINDING, GOMF_4_IRON_4_SULFUR_CLUSTER_BINDING GOMF_1_TRNA_METHYLTRANSFERASE_ACTIVITY, GOMF_TRNA_METHYLTRANSFERASE_ACTIVITY GOCC_ENDORIBONUCLEASE_COMPLEX, GOCC_ENDORIBONUCLEASE_COMPLEX
	GOBP_CELL_REDOX_HOMEOSTASIS, GOBP_CELL_REDOX_HOMEOSTASIS MIR1287_5P FONTAINE_FOLLICULAR_THYROID_ADENOMA_UP, FONTAINE_FOLLICULAR_THYROID_ADENOMA_UP GSE40274_CTRL_VS_FOXP3_AND_GATA1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_CTRL_VS_FOXP3_AND_GATA1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN GOCC_90S_PRERIBOSOME, GOCC_90S_PRERIBOSOME ALTEMETER_RESPONSE_TO_LPS_WITH_MECHANICAL_VENTILATION, ALTEMETER_RESPONSE_TO_LPS_WITH_MECHANICAL_VENTILATION GOBP_MATURATION_OF_5_8S_RRNA, GOBP_MATURATION_OF_5_8S_RRNA REACTOME_CONSTITUTIVE_SIGNALING_BY_AKT1_E17K_IN_CANCER, REACTOME_CONSTITUTIVE_SIGNALING_BY_AKT1_E17K_IN_CANCER, REACTOME_CONSTITUTIVE_SIGNALING_BY_AKT1_E17K_IN_CANCER, GOBP_CELLULAR_CARBOHYDRATE_BIOSYNTHETIC_PROCESS
	MIR7851_3P, MIR7851_3P GOMF_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS, GOMF_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS GOBP_IMMUNOGLOBULIN_PRODUCTION_INVOLVED_IN_IMMUNOGLOBULIN_MEDIATED_IMMUNE_RESPONSE, GOBP_IMMUNOGLOBULIN_PRODUCTION_INVOLVED_IN_IMMUNOGLOBULIN_MEDIATED_IMMUNE_RESPONSE KEGG_MTOR_SIGNALING_PATHWAY, KEGG_MTOR_SIGNALING_PATHWAY GOMF_DISULFIDE_OXIDOREDUCTASE_ACTIVITY, GOMF_DISULFIDE_OXIDOREDUCTASE_ACTIVITY BOYLAN_MULTIPLE_MYELOMA_C_UP, BOYLAN_MULTIPLE_MYELOMA_C_UP GOBP_PROTEIN_DESTABILIZATION, GOBP_PROTEIN_DESTABILIZATION GOBP_PROTEIN_ESTIT_FROM_ENDOPLASMIC_RETICULUM, GOBP_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM NIKOLSKY_BREAST_CANCER_16P13_AMPLICON, NIKOLSKY_BREAST_CANCER_16P13_AMPLICON
	GOBP_MITOCHONDRIAL_RNA_PROCESSING, GOBP_MITOCHONDRIAL_RNA_PROCESSING GOBP_GLUCAN_BIOSYNTHETIC_PROCESS, GOBP_GLUCAN_BIOSYNTHETIC_PROCESS GOBP_PROTEIN_REFOLDING, GOBP_PROTEIN_REFOLDING MODULE_42, MODULE_42 GOBP_PSEUDOURIDINE_SYNTHESIS, GOBP_PSEUDOURIDINE_SYNTHESIS WP_PHOTODYNAMIC_THERAPYINDUCED_HIFI_SURVIVAL_SIGNALING, WP_PHOTODYNAMIC_THERAPYINDUCED_HIFI_SURVIVAL_SIGNALING WAKABAYASHI_ADIPOGENESIS_PPARG_BOUND_36HR, WAKABAYASHI_ADIPOGENESIS_PPARG_BOUND_36HR REACTOME_VEGFR2_MEDIATED_VASCULAR_PERMEABILITY, REACTOME_VEGFR2_MEDIATED_VASCULAR_PERMEABILITY
	GOCC_TRANSCRIPTION_EXPORT_COMPLEX, GOCC_TRANSCRIPTION_EXPORT_COMPLEX GOBP_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_GATION, GOBP_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION GOBP_LYMPHOCYTE_HOMEOSTASIS, GOBP_LYMPHOCYTE_HOMEOSTASIS AMIT_EGF_RESPONSE_120_MCF10A, AMIT_EGF_RESPONSE_120_MCF10A GOBP_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GOBP_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I DESCARTES_MAIN_FETAL_PARIETAL_AND_CHIEF_CELLS, DESCARTES_MAIN_FETAL_PARIETAL_AND_CHIEF_CELLS GOBP_ATTACHMENT_OF_MITOTIC_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GOBP_ATTACHMENT_OF_MITOTIC_SPINDLE_MICROTUBULES_TO_KINETOCHORE HP_IMPAIRED_PROPRIOCEPTION, HP_IMPAIRED_PROPRIOCEPTION GOBP_POSITIVE_REGULATION_OF_ACTIN_FILAMENT_BUNDLE_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_ACTIN_FILAMENT_BUNDLE_ASSEMBLY GOBP_POSITIVE_REGULATION_OF_ACTIN_FILAMENT_BUNDLE_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_ACTIN_FILAMENT_BUNDLE_ASSEMBLY
	GOBP_RNA_5_END_PROCESSING, GOBP_RNA_5_END_PROCESSING HP_POSTERIOR_SUBCAPSULAR_CATARACT, HP_POSTERIOR_SUBCAPSULAR_CATARACT HP_JOINT_CONTRACTURES_INVOLVING_THE_JOINTS_OF_THE_FEET, HP_JOINT_CONTRACTURES_INVOLVING_THE_JOINTS_OF_THE_FEET PPARG_01, PPARG_01 MIR4745_5P, MIR4745_5P GOBP_POSITIVE_REGULATION_OF_STRESS_FIBER_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_STRESS_FIBER_ASSEMBLY GOBP_MRNA_MODIFICATION, GOBP_MRNA_MODIFICATION WP_PURINE_METABOLISM_AND_RELATED_DISORDERS, WP_PURINE_METABOLISM_AND_RELATED_DISORDERS, WP_PURINE_METABOLISM_AND_RELATED_DISORDERS, WP_PURINE_METABOLISM_AND_RELATED_DISORDERS, WP_PURINE_METABOLISM_AND_RELATED_DISORDERS GESERICK_TERT_TARGETS_DN, GESERICK_TERT_TARGETS_DN
	GOMF 2_IRON 2_SULFUR_CLUSTER_BINDING, GOMF 2_IRON 2_SULFUR_CLUSTER_BINDING GOMF 2_IRON 2_SULFUR_CLUSTER_BINDING, GOMF 2_IRON 2_SULFUR_CLUSTER_BINDING GOBP_INTERLEUKIN_12_PRODUCTION, GOBP_INTERLEUKIN_12_PRODUCTION GOBP_IRON_SULFUR_CLUSTER_ASSEMBLY, GOBP_IRON_SULFUR_CLUSTER_ASSEMBLY REACTOME_MITOCHONDRIAL_FATTY_ACID_BETA_OXIDATION, REACTOME_MITOCHONDRIAL_FATTY_ACID_BETA_OXIDATION GOBP_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_IL_PROMOTER, GOBP_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_IL_PROMOTER WP_REGUCALCIN_IN_PROXIMAL_TUBULE_EPITHELIAL_KIDNEY_CELLS, WP_REGUCALCIN_IN_PROXIMAL_TUBULE_EPITHELIAL_KIDNEY_CELLS GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_DN, GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_DN BURTON_ADIPOGENESIS_1, BURTON_ADIPOCENESIS_1
	BENNETT_SYSTEMIC_LUPUS_ERYTHEMATOSUS, BENNETT_SYSTEMIC_LUPUS_ERYTHEMATOSUS MANN_RESPONSE_TO_AMIFOSTINE_UP, MANN_RESPONSE_TO_AMIFOSTINE_UP MIR6855_5P, MIR6855_5P, MIR6855_5P, MIR6855_TO_AMIFOSTINE_UP HP_SUBCAPSULAR_CATARACT, HP_SUBCAPSULAR_CATARACT GOMF_PSEUDOURIDINE_SYNTHASE_ACTIVITY, GOMF_PSEUDOURIDINE_SYNTHASE_ACTIVITY GOCC_TOR_COMPLEX, GOCC_TOR_COMPLEX MCDOWELL_ACUTE_LUNG_INJURY_UP, MCDOWELL_ACUTE_LUNG_INJURY_UP SEIDEN_MET_SICNALING, SEIDEN_MET_SIGNALING MIR647, MIR647
	VANDESLUIS_COMMD1_TARGETS_GROUP_3_UP, VANDESLUIS_COMMD1_TARGETS_GROUP_3_UP GOBP_NEGATIVE_REGULATION_OF_FIBROBLAST_PROLIFERATION, GOBP_NEGATIVE_REGULATION_OF_FIBROBLAST_PROLIFERATION MIR6529_5P, MIR6529_5P MALONEY_RESPONSE_TO_17AAG_UP, MALONEY_RESPONSE_TO_17AAG_UP ZHONG_PEC_C1_ASTROCYTE, ZHONG_PEC_C1_ASTROCYTE HP_NEUROGENIC_BLADDER, HP_NEUROGENIC_BLADDER GOBP_TRNA_5_END_PROCESSING, GOBP_TRNA_5_END_PROCESSING GOBP_MATERNAL_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY, GOBP_MATERNAL_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY
	GOBP_ENDONUCLEOLYTIC_CLEAVAGE_INVOLVED_IN_RRNA_PROCESSING, GOBP_ENDONUCLEOLYTIC_CLEAVAGE_INVOLVED_IN_RRNA_PROCESSING MIR759, MIR759 OZEN_MIR125B1_TARGETS, OZEN_MIR125B1_TARGETS GOBP_MITOCHONDRION_MORPHOGENESIS, GOBP_MITOCHONDRION_MORPHOGENESIS WP_GLYCEROPHOSPHOLIPID_BIOSYNTHETIC_PATHWAY, WP_GLYCEROPHOSPHOLIPID_BIOSYNTHETIC_PATHWAY, WP_GLYCEROPHOSPHOLIPID_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_GLUCAN_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_GLUCAN_BIOSYNTHETIC_PROCESS HP_AQUEDUCTAL_STENOSIS, MP_AQUEDUCTAL_STENOSIS GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_12_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_12_PRODUCTION GOCC_TORC2_COMPLEX, GOCC_TORC2_COMPLEX
	PID_CXCR3_PATHWAY, PID_CXCR3_PATHWAY NIKOLSKY_BREAST_CANCER_17Q11_Q21_AMPLICON, NIKOLSKY_BREAST_CANCER_17Q11_Q21_AMPLICON HP_MACULAR_THICKENING, HP_MACULAR_THICKENING REACTOME_MAPPX_AND_MAPK_ACTIVATION, REACTOME_MAPPX_ACTIVATION GOBP_REGULATION_OF_FATTY_ACID_OXIDATION, GOBP_REGULATION_OF_FATTY_ACID_OXIDATION GOMF_TRNA_SPECIFIC_RIBONUCLEASE_ACTIVITY, GOMF_TRNA_SPECIFIC_RIBONUCLEASE_ACTIVITY GOBP_CELLULAR_IRON_ION_HOMEOSTASIS, GOBP_CELLULAR_IRON_ION_HOMEOSTASIS HP_ABNORMAL_SENSORY_NERVE_CONDUCTION_VELOCITY, HP_ABNORMAL_SENSORY_NERVE_CONDUCTION_VELOCITY
	GOBP_AMP_METABOLIC_PROCESS, GOBP_AMP_METABOLIC_PROCESS GOBP_RESPONSE_TO_INTERFERON_BETA, GOBP_RESPONSE_TO_INTERFERON_BETA KANNAN_TP53_TARGETS_DN, KANNAN_TP53_TARGETS_DN GROSS_ELK3_TARGETS_DN, GROSS_ELK3_TARGETS_DN GOSS_ELK3_TARGETS_DN, GROSS_ELK3_TARGETS_DN GOBP_REGULATION_OF_LEUKOCYTE_APOPTOTIC_PROCESS, GOBP_REGULATION_OF_LEUKOCYTE_APOPTOTIC_PROCESS GOMF_PROTEIN_DISULFIDE_OXIDOREDUCTASE_ACTIVITY, GOMF_PROTEIN_DISULFIDE_OXIDOREDUCTASE_ACTIVITY GOCC_RIBONUCLEASE_P_COMPLEX, GOCC_RIBONUCLEASE_P_COMPLEX SCHOEN_NFKB_SIGNALINC, SCHOEN_NFKB_SIGNALING GOBP_TRNA_3_END_PROCESSING, GOBP_TRNA_3_END_PROCESSING
	GOBP_POLYADENT_RNA_CATABOLIC_PROCESS, GOBP_POLYADENYLATION_DEPENDENT_RNA_CATABOLIC_PROCESS GOCC_TORC1_COMPLEX, GOCC_TORC1_COMPLEX HP_STOMACH_CANCER, HP_STOMACH_CANCER GOBP_REGULATION_OF_FATTY_ACID_BETA_OXIDATION, GOBP_REGULATION_OF_FATTY_ACID_BETA_OXIDATION GOBP_POST_EMBRYONIC_ANIMAL_ORGAN_DEVELOPMENT, GOBP_POST_EMBRYONIC_ANIMAL_ORGAN_DEVELOPMENT REACTOME_CD28_DEPENDENT_PI3K_AKT_SIGNALING, REACTOME_CD28_DEPENDENT_PI3K_AKT_SIGNALING GOBP_NEGATIVE_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT, GOBP_NEGATIVE_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT ZHONG_PFC_C2_SOX5_BCL11B_POS_EXCITATORY_NEURON, ZHONG_PFC_C2_SOX5_BCL11B_POS_EXCITATORY_NEURON
	MIR3158_3P, MIR3158_3P GOBP_IMP_METABOLIC_PROCESS, GOBP_IMP_METABOLIC_PROCESS GOBP_CELLULAR_OXIDANT_DETOXIFICATION, GOBP_CELLULAR_OXIDANT_DETOXIFICATION GOBP_MRNA_METHYLATION, GOBP_MRNA_METHYLATION GOCC_LOW_DENSITY_LIPOPROTEIN_PARTICLE, GOCC_LOW_DENSITY_LIPOPROTEIN_PARTICLE GOBP_NUCLEOLAR_LARGE_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GOBP_NUCLEOLAR_LARGE_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_I REACTOME_NUCLEOBASE_CATABOLISM, REACTOME_NUCLEOBASE_CATABOLISM REACTOME_TRNA_PROCESSING_IN_THE_MITOCHONDRION_PROCESSING_IN_THE_MITOCHONDRION CORP_POSITIVE_PECULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I_CORP_POSITIVE
	GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I HP_BONE_SPICULE_PIGMENTATION_OF_THE_RETINA, HP_BONE_SPICULE_PIGMENTATION_OF_THE_RETINA MIR6782_5P, MIR6782_5P GOBP_TRNA_5_LEADER_REMOVAL, GOBP_TRNA_5_LEADER_REMOVAL GOBP_RESPONSE_TO_FOOD, GOBP_RESPONSE_TO_FOOD GOBP_RESPONSE_TO_FOOD, GOBP_RESPONSE_TO_FOOD GOBP_MITOCHONDRIAL_FRAGMENTATION_INVOLVED_IN_APOPTOTIC_PROCESS, GOBP_MITOCHONDRIAL_FRAGMENTATION_INVOLVED_IN_APOPTOTIC_PROCESS GOBP_REGULATION_OF_ISOTYPE_SWITCHING, GOBP_REGULATION_OF_ISOTYPE_SWITCHING INGA_TP53_TARGETS, INGA_TP53_TARGETS GOBP_REGULATION_OF_OLIGODENDROCYTE_DIFFERENTIATION_OF_OLIGODENDROCYTE_DIFFERENTIATION
	COBP_PHOSPHATIDYLCLYCEROL_METABOLIC_PROCESS, GOBP_PHOSPHATIDYLCLYCEROL_METABOLIC_PROCESS GOBP_PURINE_NUCLEOBASE_METABOLIC_PROCESS, GOBP_PURINE_NUCLEOBASE_METABOLIC_PROCESS GOBP_GLOMERULAR_VISCERAL_EPITHELIAL_CELL_MIGRATION, GOBP_GLOMERULAR_VISCERAL_EPITHELIAL_CELL_MIGRATION GOBP_REGULATION_OF_TRANSCRIPTION_OF_NUCLEOLAR_LARGE_RRNA_BY_RNA_POLYMERASE_I, GOBP_REGULATION_OF_TRANSCRIPTION_OF_NUCLEOLAR_LARGE_RRNA_BY_RNA_POLYMERASE_I HP_OLIGURIA, HP_OLIGURIA NOJIMA_SFRPP_TARGETS_DN, NOJIMA_SFRPP2_TARGETS_DN GTGGTGA_MIR197, GTGGTGA_MIR197 GSE43863_NAIVE_VS_LY6C_INT_CXCR5POS_CD4_EFF_TCELL_D6_LCMV_DN, GSE43863_NAIVE_VS_LY6C_INT_CXCR5POS_CD4_EFF_TCELL_D6_LCMV_DN
	GOMF_CD4_RECEPTOR_BINDING, GOMF_CD4_RECEPTOR_BINDING GOBP_PRESYNAPTIC_MODULATION_OF_CHEMICAL_SYNAPTIC_MODULATION_OF_CHEMICAL_SYNAPTIC_TRANSMISSION SCHAVOLT_TARGETS_OF_TP53_AND_TP63, SCHAVOLT_TARGETS_OF_TP53_AND_TP63 PID_LISI_PATHWAY, PID_LISI_PATHWAY GOMF_RIBONUCLEASE_P_ACTIVITY, GOMF_RIBONUCLEASE_P_ACTIVITY MIR4507, MIR4507 GOMF_ANTIOXIDANT_ACTIVITY, GOMF_ANTIOXIDANT_ACTIVITY HP_NONPROGRESSIVE_ENCEPHALOPATHY, HP_NONPROGRESSIVE_ENCEPHALOPATHY CORP_MITOCHONIDPIAL_RNA_2_END_PROCRESSIVE_COR
	GOBP_MITOCHONDRIAL_RNA_3_END_PROCESSING, GOBP_MITOCHONDRIAL_RNA_3_END_PROCESSING GOCC_MITOCHONDRIAL_INTERMEMBRANE_SPACE_PROTEIN_TRANSPORTER_COMPLEX, GOCC_MITOCHONDRIAL_INTERMEMBRANE_SPACE_PROTEIN_TRANSPORTER_COMPLEX GOBP_NEGATIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION, GOBP_NEGATIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION HP_GRANULOCYTOPENIA, HP_GRANULOCYTOPENIA GOBP_INFLAMMATORY_RESPONSE_TO_WOUNDING, GOBP_INFLAMMATORY_RESPONSE_TO_WOUNDING REACTOME_PURINE_CATABOLISM, REACTOME_PURINE_CATABOLISM BIOCARTA_PS3HYPOXIA_PATHWAY, BIOCARTA_PS3HYPOXIA_PATHWAY GOBP_ESTABLISHMENT_OF_MITOCHONDRION_LOCALIZATION, GOBP_ESTABLISHMENT_OF_MITOCHONDRION_LOCALIZATION GOMF_DEATH_DOMAIN_BINDING, GOMF_DEATH_DOMAIN_BINDING
	HP_POSITIVE_ROMBERG_SIGN, HP_POSITIVE_ROMBERG_SIGN GOBP_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_III_ASSEMBLY, GOBP_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_III_ASSEMBLY HP_INTRAALVEOLAR_PHOSPHOLIPID_ACCUMULATION, HP_INTRAALVEOLAR_PHOSPHOLIPID_ACCUMULATION REACTOME_SIGNALING_BY_MAPK_MUTANTS, REACTOME_SIGNALING_BY_MAPK_MUTANTS GOBP_DE_NOVO_IMP_BIOSYNTHETIC_PROCESS, GOBP_DE_NOVO_IMP_BIOSYNTHETIC_PROCESS MIR6802_5P, MIR6802_5P GOBP_POSITIVE_REGULATION_OF_T_HELPER_17_TYPE_IMMUNE_RESPONSE, GOBP_POSITIVE_REGULATION_OF_T_HELPER_17_TYPE_IMMUNE_RESPONSE GOBP_PROLINE_METABOLIC_PROCESS, GOBP_PROLINE_METABOLIC_PROCESS
	GOBP_PHOSPHATIDYLGLYCEROL_BIOSYNTHETIC_PROCESS, GOBP_PHOSPHATIDYLGLYCEROL_BIOSYNTHETIC_PROCESS GOBP_MIRNA_CATABOLIC_PROCESS, GOBP_PHOSPHATIDYLGLYCEROL_BIOSYNTHETIC_PROCESS WP_SPLICING_FACTOR_NOVA_REGULATED_SYNAPTIC_PROTEINS, WP_SPLICING_FACTOR_NOVA_REGULATED_SYNAPTIC_PROTEINS GOBP_PHOSPHATIDYLSERINE_EXPOSURE_ON_APOPTOTIC_CELL_SURFACE, GOBP_PHOSPHATIDYLSERINE_EXPOSURE_ON_APOPTOTIC_CELL_SURFACE CGCAAAA_MIR450, CGCAAAA_MIR450 GOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA, GOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA_GOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_SOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_SOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_SOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_SOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_SOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_SOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_SOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_SOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_SOBP_ENDONUCLEOLYTIC_CLEAVAGE_IN_ITS1_TO_SEPARATE_SSU_RRNA_FROM_5_8S_RRNA_AND_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_SOBP_ENDONUCLEOLYTIC
	GOBP_CARDIOLIPIN_DEVELOPMENTAL PIGMENTATION, GOBP_POSITIVE_REGULATION_OF_DEVELOPMENTAL_PIGMENTATION GOBP_CARDIOLIPIN_BIOSYNTHETIC_PROCESS, GOBP_CARDIOLIPIN_BIOSYNTHETIC_PROCESS GOBP_PRIMITIVE_HEMOPOIESIS, GOBP_PRIMITIVE_HEMOPOIESIS GOMF_PRIMARY_MIRNA_BINDING, GOMF_PRIMARY_MIRNA_BINDING HP_INTERMITTENT_EPISODES_OF_RESPIRATORY_INSUFFICIENCY_DUE_TO_MUSCLE_WEAKNESS, HP_INTERMITTENT_EPISODES_OF_RESPIRATORY_INSUFFICIENCY_DUE_TO_MUSCLE_WEAKNESS GOBP_REGULATION_OF_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM, GOBP_REGULATION_OF_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM HP_NEONATAL_ONSET, HP_NEONATAL_ONSET
	HP_ABNORMALITY_OF_FRONTALIS_MUSCLE_BELLY, HP_ABNORMALITY_OF_FRONTALIS_MUSCLE_BELLY GOCC_MULTIMERIC_RIBONUCLEASE_P_COMPLEX, GOCC_MULTIMERIC_RIBONUCLEASE_P_COMPLEX GOMF_RIBONUCLEASE_P_RNA_BINDING, GOMF_RIBONUCLEASE_P_RNA_BINDING MIR371A_3P, MIR371A_3P GOCC_CILLARY_TRANSITION_FIBER, GOCC_CILLARY_TRANSITION_FIBER GOCC_CILLARY_TRANSITION_OF_IUN_KINASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_IUN_KINASE_ACTIVITY HP_PANCREATIC_ISLET_CELL_HYPERPLASIA, HP_PANCREATIC_ISLET_CELL_HYPERPLASIA HP_APNEIC_EPISODES_PRECIPITATED_BY_ILLNESS_FATIGUE_STRESS, HP_APNEIC_EPISODES_PRECIPITATED_BY_ILLNESS_FATIGUE_STRESS, HP_APNEIC_EPISODES_PRECIPITATED_BY_ILLNESS_FATIGUE_STRESS HP_NARROW_JAW, HP_NARROW_JAW
	GOBP_REGULATION_OF_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY, GOBP_REGULATION_OF_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY GOMF_MAP_KINASE_SCAFFOLD_ACTIVITY, GOMF_MAP_KINASE_SCAFFOLD_ACTIVITY GOBP_L_HISTIDINE_TRANSMEMBRANE_TRANSPORT, GOBP_L_HISTIDINE_TRANSMEMBRANE_TRANSPORT GOBP_POSITIVE_REGULATION_OF_MIRNA_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_MIRNA_METABOLIC_PROCESS GOCC_THO_COMPLEX_GOCC_THO_COMPLEX ZERBINI_RESPONSE_TO_SULINDAC_DN, ZERBINI_RESPONSE_TO_SULINDAC_DN GOMF_AZOLE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_AZOLE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY GOBP_OLIGODENDROCYTE_DIFFERENTIATION, GOBP_OLIGODENDROCYTE_DIFFERENTIATION
	WP_PEROXIREDOXIN_2_INDUCED_OVARIAN_FAILURE, WP_PEROXIREDOXIN_2_INDUCED_OVARIAN_FAILURE HP_PHARYNGITIS, HP_PHARYNGITIS HP_PUNCTATE_CATARACT, HP_PUNCTATE_CATARACT GOCC_MITTOCHONDRIAL_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX_COUPLING_FACTOR_F_O GOCC_MITTOCHONDRIAL_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX_COUPLING_FACTOR_F_O GOCC_MITTOCHONDRIAL_PROTON_TRAN_LIPPORTOTEIN_PARTICLE, GOCC_TRIGLYCERIDE_RICH_PLASMA_LIPPORTOTEIN_PARTICLE GOCC_PROTEIN_LIPPD_COMPLEX, GOCC_PROTEIN_LIPPD_COMPLEX GOCC_PROTEIN_LIPPD_COMPLEX, GOCC_PROTEIN_LIPPD_COMPLEX GOBP_NUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_NUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS REACTOME_MET_RECEPTOR_RECYCLING, REACTOME_MET_RECEPTOR_RECYCLING
6C_INT_CXCR5POS_CD4_TCELL_D150_LCMV_DN, GSE43863_NAIVE_VS_MEMORY_LY6C_INT_CXCR5POS_CD4_TCELL_D150_LCMV_DN	HP_TALIPES_CALCANEOVALGUS, HP_TALIPES_CALCANEOVALGUS GOBP_MARGINAL_ZONE_B_CELL_DIFFERENTIATION, GOBP_MARGINAL_ZONE_B_CELL_DIFFERENTIATION GOBP_ARGININE_TRANSMEMBRANE_TRANSPORT, GOBP_ARGININE_TRANSMEMBRANE_TRANSPORT GOMF_ARGININE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_ARGININE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY GOBP_FATTY_ACID_ELONGATION_SATURATED_FATTY_ACID_GOBP_FATTY_ACID_ELONGATION_SATURATED_FATTY_ACID GOMF_3_OXO_ARACHIDOVL_COA_SYNTHASE_ACTIVITY, GOMF_3_OXO_ARACHIDOYL_COA_SYNTHASE_ACTIVITY GOBP_PROTEIN_PROCESSING_INVOLVED_IN_PROTEIN_TARGETING_TO_MITOCHONDRION, GOBP_PROTEIN_PROCESSING_INVOLVED_IN_PROTEIN_TARGETING_TO_MITOCHONDRION HP_DECREASED_ADENOSYLCOBALAMIN, HP_DECREASED_ADENOSYLCOBALAMIN HP_STIFF_SKIN, HP_STIFF_SKIN
	HP_VEGETATIVE_STATE, HP_VEGETATIVE_STATE 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_5_8S_RRNA_LSU_RRNA GOCC_CELL_POLE, GOCC_CELL_POLE FARDIN_HYPOXIA_9, FARDIN_HYPOXIA_9 REACTOME_RSK_ACTIVATION, REACTOME_RSK_ACTIVATION REACTOME_DEFECTIVE_FACTOR_VIII_CAUSES_HEMOPHILIA_A, REACTOME_DEFECTIVE_FACTOR_VIII_CAUSES_HEMOPHILIA_A GOBP_RESPONSE_TO_OLEIC_ACID, GOBP_RESPONSE_TO_OLEIC_ACID GOMF_LACTATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_LACTATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
	WP_TP53_NETWORK, WP_TP53_NETWORK GOBP_VAGINA_DEVELOPMENT, GOBP_VAGINA_DEVELOPMENT HP_LIMITED_WRIST_MOVEMENT, HP_LIMITED_WRIST_MOVEMENT GOBP_WRIST_MOVEMENT, HP_LIMITED_WRIST_MOVEMENT GOBP_WRIST_MOVEMENT, HP_LIMITED_WRIST_MOVEMENT GOBP_COBP_WRIST_MOVEMENT, HP_LIMITED_WRIST_MOVEMENT GOBP_NEURON, GOBP_DEVELON, GOBP_WRITE_FAT_CELL_DIFFERENTIATION GOBP_NEURONAL_ION_CHANNEL_CLUSTERING, GOBP_NEURONAL_ION_CHANNEL_CLUSTERING GOMF_ACETYLCHOLINE_RECEPTOR_BINDING, GOMF_ACETYLCHOLINE_RECEPTOR_BINDING HP_SENSORY_ATAXIA, HP_SENSORY_ATAXIA MIR151A_5P_MIR151B, MIR151A_5P_MIR151B
	GOBP_CELLULAR_RESPONSE_TO_INTERFERON_BETA, GOBP_CELLULAR_RESPONSE_TO_INTERFERON_BETA GOBP_CELLULAR_COMPONENT_DISASSEMBLY_INVOLVED_IN_EXECUTION_PHASE_OF_APOPTOSIS, GOBP_CELLULAR_COMPONENT_DISASSEMBLY_INVOLVED_IN_EXECUTION_PHASE_OF_APOPTOSIS GOBP_LONA_TOPOLOGICAL_CHANGE, GOBP_DNA_TOPOLOGICAL_CHANGE GOBP_LACTATE_TRANSMEMBRANE_TRANSPORT, GOBP_LACTATE_TRANSMEMBRANE_TRANSPORT GOBP_REGULATION_OF_MITOCHONDRIAL_MRN_STABILITY, GOBP_REGULATION_OF_MITOCHONDRIAL_MRNA_STABILITY GOMF_SODIUM_ION_BINDING, GOMF_SODIUM_ION_BINDING HP_PERIPORTAL_FIBROSIS, HP_PERIPORTAL_FIBROSIS HP_DECREASED_METHYLMALONYL_COA_MUTASE_ACTIVITY, HP_DECREASED_METHYLMALONYL_COA_MUTASE_ACTIVITY GOBP_DNALA_TERD_ROSEROES GOB_NOVE_SODIUM_GOA_MUTASE_ACTIVITY
	GOBP_RRNA_5_END_PROCESSING, GOBP_RRNA_5_END_PROCESSING GOMF_L_HISTIDINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_L_HISTIDINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY GOBP_POSITIVE_REGULATION_OF_T_HELPER_17_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_T_HELPER_17_CELL_DIFFERENTIATION REACTOME_LINOLEIC_ACID_LA_METABOLISM, REACTOME_LINOLEIC_ACID_LA_METABOLISM GOBP_NEGATIVE_REGULATION_OF_P38MAPK_CASCADE, GOBP_NEGATIVE_REGULATION_OF_P38MAPK_CASCADE GOCC_DERLIN_1_RETROTRANSLOCATION_COMPLEX, GOCC_DERLIN_1_RETROTRANSLOCATION_COMPLEX GOCC_CORE_MEDIATOR_COMPLEX, GOCC_CORE_MEDIATOR_COMPLEX GOMF_ADENYLATE_KINASE_ACTIVITY, GOMF_ADENYLATE_KINASE_ACTIVITY chr15q22, chr15q22
	GYORFFY_DOXORUBICIN_RESISTANCE, GYORFFY_DOXORUBICIN_RESISTANCE GOMF_DIPEPTIDYL_PEPTIDASE_ACTIVITY, GOMF_DIPEPTIDYL_PEPTIDASE_ACTIVITY HP_HYPOPLASIA_OF_THE_EAR_CARTILAGE, HP_HYPOPLASIA_OF_THE_EAR_CARTILAGE REACTOME_ABACAVIR_TRANSPORT_AND_METABOLISM, REACTOME_ABACAVIR_TRANSPORT_AND_METABOLISM_RESULTING_IN_PRODUCTION_OF_ROS_BY_CYP2E1 WP_ETHANOL_METABOLISM_RESULTING_IN_PRODUCTION_FOS_BY_CYP2E1, WP_ETHANOL_METABOLISM_RESULTING_IN_PRODUCTION_OF_ROS_BY_CYP2E1 GOBP_MRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II, GOBP_MRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II HP_TRITANOMALY, HP_TRITANOMALY GOBP_FOLATE_TRANSMEMBRANE_TRANSPORT, GOBP_FOLATE_TRANSMEMBRANE_TRANSPORT
	HP_SCLERODERMA, HP_SCLERODERMA ISHIDA_TARGETS_OF_SYT_SSX_FUSIONS, ISHIDA_TARGETS_OF_SYT_SSX_FUSIONS GOBP_POST_EMBRYONIC_CAMERA_TYPE_EYE_DEVELOPMENT, GOBP_POST_EMBRYONIC_CAMERA_TYPE_EYE_DEVELOPMENT MIR4520_5P, MIR4520_5P HP_CORTICAL_CATARACT, HP_CORTICAL_CATARACT GOBP_PROTEIN_IMPORT_INTO_PEROXISOME_MEMBRANE, GOBP_PROTEIN_IMPORT_INTO_PEROXISOME_MEMBRANE GOBP_CELLULAR_RESPONSE_TO_NITROSATIVE_STRESS, GOBP_CELLULAR_RESPONSE_TO_NITROSATIVE_STRESS GOMF_NEDDB_TRANSFERASE_ACTIVITY, GOMF_NEDDB_TRANSFERASE_ACTIVITY HP_DYSESTHESIA, HP_DYSESTHESIA
	PID_IL5_PATHWAY, PID_IL5_PATHWAY GOBP_NEGATIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE_TO_ANTIGENIC_STIMULUS, GOBP_NEGATIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE_TO_ANTIGENIC_STIMULUS GOBP_ADP_BIOSYNTHETIC_PROCESS, GOBP_ADP_BIOSYNTHETIC_PROCESS GOBP_ACTIN_MODIFICATION, GOBP_ACTIN_MODIFICATION GOBP_TRNA_PSEUDOURIDINE_SYNTHESIS, GOBP_TRNA_PSEUDOURIDINE_SYNTHESIS HP_UPPER_LIMB_POSTURAL_TREMOR, HP_UPPER_LIMB_POSTURAL_TREMOR GOCC_VARICOSITY, GOCC_VARICOSITY HP_PORTAL_FIBROSIS, HP_PORTAL_FIBROSIS
	GOME_PHOSPHATIDYLINOSITOL_4_5_BISPHOSPHATE_PHOSPHATASE_ACTIVITY, GOME_PHOSPHATIDYLINOSITOL_4_5_BISPHOSPHATASE_ACTIVITY GOBP_REGULATION_OF_APOPTOTIC_DNA_FRAGMENTATION, GOBP_REGULATION_OF_APOPTOTIC_DNA_FRAGMENTATION HP_ULNAR_CLAW, HP_ULNAR_CLAW GOBP_DIACYLGLYCEROL_BIOSYNTHETIC_PROCESS, GOBP_DIACYLGLYCEROL_BIOSYNTHETIC_PROCESS GOBP_DIACYLGLYCEROL_BIOSYNTHETIC_PROCESS, GOBP_DEOXYRIBONUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS GOBP_DEOXYRIBONUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS, GOBP_DEOXYRIBONUCLEOSIDE_MONOPHOSPHATE_CATABOLIC_PROCESS GOBP_CELLULAR_RESPONSE_TO_TESTOSTERONE_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_TESTOSTERONE_STIMULUS GOMF_JUN_KINASE_BINDING, GOMF_JUN_KINASE_BINDING GOMF_JUN_KINASE_BINDING, GOMF_JUN_KINASE_BINDING HP_LEFT_TO_RIGHT_SHUNT, HP_LEFT_TO_RIGHT_SHUNT
	GOBP_NEGATIVE_REGULATION_OF_MRNA_3_END_PROCESSING, GOBP_NEGATIVE_REGULATION_OF_MRNA_3_END_PROCESSING COBP_RESPONSE_TO_NITROSATIVE_STRESS, GOBP_RESPONSE_TO_NITROSATIVE_STRESS COBP_L_ARGININE_TRANSPORT, GOBP_L_ARGININE_TRANSPORT HP_RED_GREEN_DYSCHROMATOPSIA, HP_RED_GREEN_DYSCHROMATOPSIA CAO_BLOOD_FLUZONE_AGE_05_14YO_30DY_UP, CAO_BLOOD_FLUZONE_AGE_05_14YO_30DY_UP HP_CHRONIC_ATROPHIC_GASTRITIS, HP_CHRONIC_ATROPHIC_GASTRITIS HP_MALAR_PROMINENCE, HP_MALAR_PROMINENCE GOBP_ACTIVATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS_BY_CYTOCHROME_C, GOBP_ACTIVATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS_BY_CYTOCHROME_C
	GOCC_PINOSOME, GOCC_PINOSOME GOMF_THIOREDOXIN_DISULIFIDE_REDUCTASE_ACTIVITY, GOMF_THIOREDOXIN_DISULFIDE_REDUCTASE_ACTIVITY GOBP_CELLULAR_RESPONSE_TO_CHOLESTEROL, GOBP_CELLULAR_RESPONSE_TO_CHOLESTEROL REACTOME_TRANSPORT_OF_NUCLEOSIDES_AND_FREE_PURINE_AND_PYRIMIDINE_BASES_ACROSS_THE_PLASMA_MEMBRANE, REACTOME_TRANSPORT_OF_NUCLEOSIDES_AND_FREE_PURINE_AND_PYRIMIDINE_BASES_ACROSS_THE_PLASMA_MEMBRANE HP_HYPERTROPHIC_NERVE_CHANGES, HP_HYPERTROPHIC_NERVE_CHANGES GOBP_ECTOPIC_GERM_CELL_PROGRAMMED_CELL_DEATH, GOBP_ECTOPIC_GERM_CELL_PROGRAMMED_CELL_DEATH GOBP_REGULATION_OF_INFLAMMATORY_RESPONSE_TO_WOUNDING, GOBP_REGULATION_OF_INFLAMMATORY_RESPONSE_TO_WOUNDING GOBP_REGULATION_OF_DNA_CATABOLIC_PROCESS_GOBP_REGULATION_OF_DNA_CATABOLIC_PROCESS HERNANDEZ_ABERRANT_MITOSIS_BY_DOCETACEL_4NM_DN, HERNANDEZ_ABERRANT_MITOSIS_BY_DOCETACEL_4NM_DN,
	CHESLER_BRAIN_QIT_TRANS, CHESLER_BRAIN_QIT_TRANS GOBP_MITOCHONDRIAL_RNA_CATABOLIC_PROCESS, GOBP_MITOCHONDRIAL_RNA_CATABOLIC_PROCESS GOBP_PURINE_NUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_PURINE_NUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS GOBP_RIBONUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_RIBONUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS GOBP_DIRECT_OSSIFICATION, GOBP_DIRECT_OSSIFICATION GOBP_CRIBONUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_RIBONUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS GOBP_DIRECT_OSSIFICATION, GOBP_DIRECT_OSSIFICATION GOBP_CRIBONUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS GOBP_CRIBONUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS GOBP_DIRECT_OSSIFICATION, GOBP_DIRECT_OSSIFICATION GOBP_CRIBONUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROCESS GOBP_CRIBONUCLEOSIDE_DIPHOSPHATE_BIOSYNTHETIC_PROC
	GOBP_PROTEIN_MATURATION_BY_4FE_4S_CLUSTER_TRANSFER, GOBP_PROTEIN_MATURATION_BY_4FE_4S_CLUSTER_TRANSFER GOBP_L_LYSINE_TRANSMEMBRANE_TRANSPORT, GOBP_L_LYSINE_TRANSMEMBRANE_TRANSPORT GOMF_L_LYSINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_L_LYSINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY REACTOME_PROTON_COUPLED_MONOCARBOXYLATE_TRANSPORT, REACTOME_PROTON_COUPLED_MONOCARBOXYLATE_TRANSPORT REACTOME_RUNX1_ETEGULATE_ESTROGEN_RECEPTOR_MEDIATED_TRANSCRIPTION, REACTOME_RUNX1_REGULATES_ESTROGEN_RECEPTOR_MEDIATED_TRANSCRIPTION LEL_HOXCS_TARGETS_UP, LEL_HOXCS_TARGETS_UP WP_MIRNA_REGULATION_OF_P53_PATHWAY_IN_PROSTATE_CANCER, WP_MIRNA_REGULATION_OF_P53_PATHWAY_IN_PROSTATE_CANCER MACLACHLAN_BRCA1_TARGETS_DN, MACLACHLAN_BRCA1_TARGETS_DN MIR604, MIR604 MIR604 MIR604
	GOBP_REGULATION_OF_APOPTOTIC_PROCESS_INVOLVED_IN_DEVELOPMENT, GOBP_REGULATION_OF_APOPTOTIC_PROCESS_INVOLVED_IN_DEVELOPMENT GOBP_MAGNESIUM_ION_HOMEOSTASIS, GOBP_MAGNESIUM_ION_HOMEOSTASIS LIU_TOPBP1_TARGETS, LIU_TOPBP1_TARGETS GOMF_INTERLEUKIN_1_BINDING, GOMF_INTERLEUKIN_1_BINDING BIOCARTA_BBCELL_PATHWAY, BIOCARTA_BBCELL_PATHWAY HP_LACK_OF_T_CELL_FUNCTION, HP_LACK_OF_T_CELL_FUNCTION KASLER_HDAC7_TARGETS_2_UP, KASLER_HDAC7_TARGETS_2_UP GOBP_POSITIVE_REGULATION_OF_SECONDARY_METABOLITE_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_SECONDARY_METABOLITE_BIOSYNTHETIC_PROCESS
	REACTOME_RELEASE_OF_APOPTOTIC_FACTORS_FROM_THE_MITOCHONDRIA, REACTOME_RELEASE_OF_APOPTOTIC_FACTORS_FROM_THE_MITOCHONDRIA HP_THORACOLUMBAR_KYPHOSCOLIOSIS, HP_THORACOLUMBAR_KYPHOSCOLIOSIS HP_PROGRESSIVE_VISUAL_FIELD_DEFECTS, HP_PROGRESSIVE_VISUAL_FIELD_DEFECTS CAO_BLOOD_FLUMIST_AGE_05_14VO_1DV_UP, CAO_BLOOD_FLUMIST_AGE_05_14VO_1DV_UP GOBP_CORTICOTROPIN_SECRETION, GOBP_CORTICOTROPIN_SECRETION GOBP_NEGATIVE_REGULATION_OF_GLYCOGEN_METABOLIC_PROCESS, GOBP_MEGATIVE_REGULATION_OF_GLYCOGEN_METABOLIC_PROCESS GOBP_LARGININE_TRANSMEMBRANE_TRANSPORT, GOBP_L_ARGININE_TRANSMEMBRANE_TRANSPORT GOBP_NEGATIVE_REGULATION_OF_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY GOMF_RIBOSOMAL_PROTEIN_S6_KINASE_ACTIVITY, GOMF_RIBOSOMAL_PROTEIN_S6_KINASE_ACTIVITY
	SHIN_B_CELL_LYMPHOMA_CLUSTER_9, SHIN_B_CELL_LYMPHOMA_CLUSTER_9 HP_ANOMALOUS_TRICHROMACY, HP_ANOMALOUS_TRICHROMACY GOMF_CHONDROITIN_SULFATE_BINDING, GOMF_CHONDROITIN_SULFATE_BINDING ZHENG_RESPONSE_TO_ARSENITE_UP, ZHENG_RESPONSE_TO_ARSENITE_UP GOBP_PROGRAMMED_CELL_DEVELOPMENT, GOBP_PROGRAMMED_CELL_DEATH_INVOLVED_IN_CELL_DEVELOPMENT GOBP_RESPONSE_TO_REDOX_STATE, GOBP_RESPONSE_TO_REDOX_STATE cht12q23, cht12q23 GOBP_RESPONSE_TO_PROTOZOAN, GOBP_RESPONSE_TO_PROTOZOAN
	HP_EXERCISE_INDUCED_MUSCLE_STIFFNESS, HP_EXERCISE_INDUCED_MUSCLE_STIFFNESS HP_ABNORMAL_PERIOSTEUM_MORPHOLOGY, HP_ABNORMAL_PERIOSTEUM_MORPHOLOGY GOBP_POST_EMBRYONIC_EYE_MORPHOGENESIS, GOBP_POST_EMBRYONIC_EYE_MORPHOGENESIS GGCGGCA_MIR371, GGCCGGCA_MIR371 VERRECCHIA_RESPONSE_TO_TGFBL_C6, VERRECCHIA_RESPONSE_TO_TGFBL_C6 HP_PERIPHERAL_HYPERMYELINATION, HP_PERIPHERAL_HYPERMYELINATION GOCC_DENSE_BODY, GOCC_DENSE_BODY HP_HYPOPLASIA_OF_THE_FEMORAL_HEAD, HP_HYPOPLASIA_OF_THE_FEMORAL_HEAD HP_HYPOPLASIA_OF_THE_FEMORAL_HEAD, HP_HYPOPLASIA_OF_THE_FEMORAL_HEAD
	HP_SPONTANEOUS_PAIN_SENSATION, HP_SPONTANEOUS_PAIN_SENSATION HP_CHRONIC_PAIN, HP_CHRONIC_PAIN HP_CHRONIC_PAIN, HP_CHRONIC_PAIN HP_TRUNCAL_TITUBATION HP_SHOULDER_PAIN, HP_SHOULDER_PAIN HP_SHOULDER_PAIN, HP_SHOULDER_PAIN GOBP_REGULATION_OF_MITOCHONDRIAL_RNA_CATABOLIC_PROCESS, GOBP_REGULATION_OF_MITOCHONDRIAL_RNA_CATABOLIC_PROCESS GOBP_RNA_IMPORT_INTO_MITOCHONDRION, GOBP_RNA_IMPORT_INTO_MITOCHONDRION GOBP_POSITIVE_REGULATION_OF_CORTICOTROPIN_SECRETION, GOBP_POSITIVE_REGULATION_OF_CORTICOTROPIN_SECRETION GOBP_NEGATIVE_REGULATION_OF_RESPIRATORY_BURST_INVOLVED_IN_INFLAMMATORY_RESPONSE, GOBP_NEGATIVE_REGULATION_OF_RESPIRATORY_BURST_INVOLVED_IN_INFLAMMATORY_RESPONSE GOBP_L_ARGININE_IMPORT_ACROSS_PLASMA_MEMBRANE, GOBP_L_ARGININE_IMPORT_ACROSS_PLASMA_MEMBRANE
	GOBP_POSITIVE_REGULATION_OF_APOPTOTIC_PROCESS_INVOLVED_IN_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_APOPTOTIC_PROCESS_INVOLVED_IN_DEVELOPMENT GOCC_NUCLEOLAR_RIBONUCLEASE_P_COMPLEX, GOCC_NUCLEOLAR_RIBONUCLEASE_P_COMPLEX GOCC_TRNA_METHYLTRANSFERASE_COMPLEX, GOCC_TRNA_METHYLTRANSFERASE_COMPLEX GOMF_NUCLEOBASE_BINDING, GOMF_NUCLEOBASE_BINDING GOMF_NUCLEOBASE_BINDING, GOMF_NUCLEOBASE_BINDING GOMF_MITOGEN_ACTIVATED_PROTEIN_KINASE_P38_BINDING, GOMF_MITOGEN_ACTIVATED_PROTEIN_KINASE_P38_BINDING REACTOME_ABACAVIR_METABOLISM, REACTOME_ABACAVIR_METABOLISM GOBP_CALMODULIN_DEPENDENT_KINASE_ISGNALING_PATHWAY, GOBP_CALMODULIN_DEPENDENT_KINASE_SIGNALING_PATHWAY FUJIWARA_PARK2_IN_LIVER_CANCER_UP, FUJIWARA_PARK2_IN_LIVER_CANCER_UP GOBP_ADENOSINE_METABOLIC_PROCESS, GOBP_ADENOSINE_METABOLIC_PROCESS
	HP_ABNORMAL_GLOMERULAR_BASEMENT_MEMBRANE_MORPHOLOGY, HP_ABNORMAL_GLOMERULAR_BASEMENT_MEMBRANE_MORPHOLOGY COBP_PROTEIN_SULFATION, GOBP_PROTEIN_SULFATION GALL_TP53_TARGETS_APOPTOTIC_UP, GALL_TP53_TARGETS_APOPTOTIC_UP MODULE_501 MODULE_501 GOBP_RESPONSE_TO_MACNESIUM_ION, GOBP_RESPONSE_TO_MACNESIUM_ION WEBER_METHYLATED_IN_COLON_CANCER, WEBER_METHYLATED_IN_COLON_CANCER GOBP_NEGATIVE_REGULATION_OF_MYELOID_CELL_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_MYELOID_CELL_APOPTOTIC_PROCESS GOBP_REGULATION_OF_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY, GOBP_REGULATION_OF_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY, GOBP_REGULATION_OF_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY,
	GOME_DYSTROGLYCAN_BINDING, GOME_DYSTROGLYCAN_BINDING HP_PERISYLVIAN_POLYMICROGYRIA, HP_PERISYLVIAN_POLYMICROGYRIA GOBP_NEGATIVE_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLATION, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLATION MIR1973, MIR1973 HP_ABNORMALITY_OF_THE_PERIOSTEUM, HP_ABNORMALITY_OF_THE_PERIOSTEUM GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_INTERFERON_GAMMA, GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_INTERFERON_GAMMA HP_CARDIAC_SHUNT, HP_CARDIAC_SHUNT TERAO_AOX4_TARGETS_HG_DN, TERAO_AOX4_TARGETS_HG_DN MIR10392 3P MIR10392 3P MIR10392 3P MIR10392 3P MIR10392 3P
	MIR10392_3P, MIR10392_3P HP_CENTRAL_PRIMITIVE_NEUROECTODERMAL_TUMOR, HP_CENTRAL_PRIMITIVE_NEUROECTODERMAL_TUMOR HP_RECURRENT_PHARYNGITIS, HP_RECURRENT_PHARYNGITIS CAO_BLOOD_FLUMIST_AGE_05_14YO_30DY_UP, CAO_BLOOD_FLUMIST_AGE_05_14YO_30DY_UP GOBP_C_TERMINAL_PROTEIN_LIPIDATION, GOBP_C_TERMINAL_PROTEIN_LIPIDATION GOBP_POSITIVE_REGULATION_OF_AUTOPHAGOSOME_MATURATION, GOBP_POSITIVE_REGULATION_OF_AUTOPHAGOSOME_MATURATION GOBP_REGULATION_OF_NEUROMUSCULAR_JUNCTION_DEVELOPMENT, GOBP_REGULATION_OF_NEUROMUSCULAR_JUNCTION_DEVELOPMENT HP_SUBSARCOLEMMAL_ACCUMULATIONS_OF_ABNORMALLY_SHAPED_MITOCHONDRIA, HP_SUBSARCOLEMMAL_ACCUMULATIONS_OF_ABNORMALLY_SHAPED_MITOCHONDRIA HP_MUSCULAR_VENTRICULAR_SEPTAL_DESECT, HP_MUSCULAR_SEPTAL_DEFECT CORP_MATUR_SCULIAR_VENTRICULAR_SEPTAL_DESECT, HP_MUSCULAR_SEPTAL_DEFECT CORP_MATUR_SCULIAR_VENTRICULAR_SEPTAL_DESCRIPE_CORP_MATUR_SEPTAL_DEFECT CORP_MATUR_SCULINE_VERAMIN_BIOCOMETICATION RECORD_MATUR_SEPTAL_DEFECT CORP_MATUR_SCULINE_VERAMIN_BIOCOMETICATION RECORD_MATUR_SEPTAL_DEFECT CORP_MATUR_SCULINE_VERAMIN_BIOCOMETICATION RECORD_MATUR_SEPTAL_DEFECT CORP_MATUR_SCULINE_VERAMIN_BIOCOMETICATION RECORD MATUR_SEPTAL_DESCRIPE_CORP_MATUR_SCULINE_VERAMIN_BIOCOMETICATION RECORD MATUR_SCULINE_VERAMIN_BIOCOMETICATION RECORD
	GOBP_WATER_SOLUBLE_VITAMIN_BIOSYNTHETIC_PROCESS, GOBP_WATER_SOLUBLE_VITAMIN_BIOSYNTHETIC_PROCESS GOBP_MOTOR_LEARNING, GOBP_MOTOR_LEARNING HP_SUDDEN_EPISODIC_APNEA, HP_SUDDEN_EPISODIC_APNEA GOBP_ESTABLISHMENT_OR_MAINTENANCE_OF_TRANSMEMBRANE_ELECTROCHEMICAL_GRADIENT, GOBP_ESTABLISHMENT_OR_MAINTENANCE_OF_TRANSMEMBRANE_ELECTROCHEMICAL_GRADIENT BIOCARTA_ASBCELL_PATHWAY, BIOCARTA_ASBCELL_PATHWAY HP_CENTRAL_SLEEP_APNEA, HP_CENTRAL_SLEEP_APNEA GOBP_EMBRYONIC_DIGESTIVE_TRACT_MORPHOGENESIS, GOBP_EMBRYONIC_DIGESTIVE_TRACT_MORPHOGENESIS HP_ANKLE_WEAKNESS, HP_ANKLE_WEAKNESS
	GOBP_NUCLEOSIDE_TRANSPORT, GOBP_NUCLEOSIDE_TRANSPORT GOBP_SYNAPTIC_GROWTH_AT_NEUROMUSCULAR_JUNCTION, GOBP_SYNAPTIC_GROWTH_AT_NEUROMUSCULAR_JUNCTION GOMF_PHOSPHATIDYLINOSITOL_4_5_BISPHOSPHATE_5_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_4_5_BISPHOSPHATE_5_PHOSPHATASE_ACTIVITY THUM_MIR21_TARGETS_HEART_DISEASE_DN, THUM_MIR21_TARGETS_HEART_DISEASE_DN REACTOME_MYOCLONIC_EPILEPSY_OF_LAFORA, REACTOME_MYOCLONIC_EPILEPSY_OF_LAFORA GOBP_SHORT_CHAIN_FATTY_ACID_CATABOLIC_PROCESS, GOBP_SHORT_CHAIN_FATTY_ACID_CATABOLIC_PROCESS GOBP_POSITIVE_REGULATION_OF_ISOTYPE_SWITCHING_TO_IGG_ISOTYPES, GOBP_POSITIVE_REGULATION_OF_ISOTYPE_SWITCHING_TO_IGG_ISOTYPES GOMF_UBIQUINOL_CYTOCHROME_C_REDUCTASE_ACTIVITY, GOMF_UBIQUINOL_CYTOCHROME_C_REDUCTASE_ACTIVITY GOBP_MAMMARY_GLAND_INVOLUTION, GOBP_MAMMARY_GLAND_INVOLUTION
	TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_SUSTAINED_IN_GRANULOCYTE_DN, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_SUSTAINED_IN_GRANULOCYTE_DN GOBP_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORT, GOBP_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORT GOBP_RESPIRATORY_BURST_INVOLVED_IN_INFLAMMATORY_RESPONSE, GOBP_RESPIRATORY_BURST_INVOLVED_IN_INFLAMMATORY_RESPONSE GOBP_CELLULAR_RESPONSE_TO_MAGNESIUM_ION, GOBP_CELLULAR_RESPONSE_TO_MAGNESIUM_ION GOMF_DNA_TOPOISOMERASE_ACTIVITY, GOMF_DNA_TOPOISOMERASE_ACTIVITY MODULE_313, MODULE_313 HOLLEMAN_PREDNISOLONE_RESISTANCE_B_ALL_DN, HOLLEMAN_PREDNISOLONE_RESISTANCE_B_ALL_DN chr17p12, chr17p12
	MODULE_156, MODULE_156 MODULE_240, MODULE_240 GOBP_NEGATIVE_REGULATION_OF_OLIGODENDROCYTE_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_OLIGODENDROCYTE_DIFFERENTIATION REACTOME_DEFECTS_OF_CONTACT_ACTIVATION_SYSTEM_CAS_AND_KALLIKREIN_KININ_SYSTEM_KKS, REACTOME_DEFECTS_OF_CONTACT_ACTIVATION_SYSTEM_CAS_AND_KALLIKREIN_KININ_SYSTEM_KKS GOBP_REGULATION_OF_WATER_LOSS_VIA_SKIN, GOBP_REGULATION_OF_WATER_LOSS_VIA_SKIN KEGG_RENIN_ANGIOTENSIN_SYSTEM, KEGG_RENIN_ANGIOTENSIN_SYSTEM HP_HIP_FLEXOR_WEAKNESS, HP_HIP_FLEXOR_WEAKNESS chr12q12, chr12q12
	HP_UPPER_EYELID_EDEMA, HP_UPPER_EYELID_EDEMA HP_CHOROID_PLEXUS_CARCINOMA, HP_CHOROID_PLEXUS_CARCINOMA GOBP_APOPTOTIC_CHROMOSOME_CONDENSATION, GOBP_APOPTOTIC_CHROMOSOME_CONDENSATION GOBP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS, GOBP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS GOBP_UPPER_EYELID_EDEMA, HP_UPPER_EYELID_EDEMA GOBP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS, GOBP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS GOBP_UPPER_EYELID_EDEMA, HP_UPPER_EYELID_EDEMA GOBP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS, GOBP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS GOBP_UPPER_EYELID_EDEMA, HP_UPPER_EYELID_EDEMA GOBP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS, GOBP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS GOBP_UPPER_EYELID_EDEMA, HP_UPPER_EYELID_EDEMA GOBP_TRANSCRIPTION_OF_NTICL_OXIDE_SYNTHASCRIPTION_FACTOR_CATABOLIC_PROCESS GOBP_TRANSCRIPTION_OF_TRANSCRIPTION_GORP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS INVOLVED_IN_BLOOD_VESSEL_MORPHOGENESIS, GOBP_APOPTOTIC_PROCESS_INVOLVED_IN_BLOOD_VESSEL_MORPHOGENESIS, GOBP_APOPTOTIC_PROCESS_INVOLVED_IN_BLOOD_VESSEL_MORPHOGENESIS GOBP_POSITIVE_REGULATION_OF_APOPTOTIC_DNA_FRAGMENTATION, GOBP_POSITIVE_REGULATION_OF_APOPTOTIC_DNA_FRAGMENTATION GOMF_TRNA_ADENINE_METHYLTRANSFERASE_ACTIVITY, GOMF_TRNA_ADENINE_METHYLTRANSFERASE_ACTIVITY
	GOMF_LARGININE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_LARGININE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY REACTOME_PROPIONYL_COA_CATABOLISM, REACTOME_PROPIONYL_COA_CATABOLISM GOMF_PHOSPHOLIPID_SCRAMBILASE_ACTIVITY, GOMF_PHOSPHOLIPID_SCRAMBILASE_ACTIVITY HP_VENTRICULAR_SEPTAL_HYPERTROPHY, HP_VENTRICULAR_SEPTAL_HYPERTROPHY REACTOME_SYNTHESIS_OF_PG, REACTOME_SYNTHESIS_OF_PG GOMF_MYOSIN_LIGHT_CHAIN_BINDING, GOMF_MYOSIN_LIGHT_CHAIN_BINDING GOBF_NEGATIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE, GOBP_NEGATIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE MODULE_547, MODULE_547
	chrXq22, chrXq22 GOBP_SULFATION, GOBP_SULFATION DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_RESPIRATORY_SECRETORY_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_RESPIRATORY_SECRETORY_CELLS GOBP_EMBRYONIC_HEMOPOIESIS, GOBP_EMBRYONIC_HEMOPOIESS HP_WEAKNESS_OF_LONG_FINGER_EXTENSOR_MUSCLES, DURANTE_STREETINGOR_EXTENSOR_MUSCLES ZIRN_TRETINGOR_RESPONSE_WTI_DN, ZIRN_TRETINGOR_RESPONSE_WTI_DN GOBP_POSITIVE_REGULATION_OF_DNA_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_DNA_CATABOLIC_PROCESS HP_DECREASED_MINIATURE_ENDPLATE_POTENTIALS, HP_DECREASED_MINIATURE_ENDPLATE_POTENTIALS
	GOBP_POSITIVE_REGULATION_OF_LYMPHOCYTE_APOPTOTIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_LYMPHOCYTE_APOPTOTIC_PROCESS HP_TRICEPS_WEAKNESS, HP_TRICEPS_WEAKNESS HP_WEAKNESS_OF_THE_INTRINSIC_HAND_MUSCLES, HP_WEAKNESS_OF_THE_INTRINSIC_HAND_MUSCLES GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_LIGHTGREEN_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_LIGHTGREEN_DN