		BCELL_DN, GSE12845_NAIVE_VS_DARKZONE_GC_TONSIL_BCELL_DN
	GSE5679_PPARG_LIGAND_ROSIGLITAZONE_VS GSE5455_EX_VIVO_VS_POST_24H_INCUBATION GSE31082_DP_VS_CD4_SP_THYMOCYTE_UP, GSI	CD4_TCELL_48H_DN, GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_48H_DN 5_RARA_AGONIST_AM580_TREATED_DC_UP, GSE5679_PPARG_LIGAND_ROSIGLITAZONE_VS_RARA_AGONIST_AM580_TREATED_DC_UP 5_MONOCYTES_FROM_TUMOR_BEARING_MOUSE_UP, GSE5455_EX_VIVO_VS_POST_24H_INCUBATION_MONOCYTES_FROM_TUMOR_BEARING_MOUSE_UP
	GOBP_PROTEIN_LOCALIZATION_TO_MITOCHO GSE9239_CTRL_VS_TNF_INHIBITOR_TREATED_I GSE10273_HIGH_IL7_VS_HIGH_IL7_AND_IRF4_I	ONDRION, GOBP_PROTEIN_LOCALIZATION_TO_MITOCHONDRION DC_DN, GSE9239_CTRL_VS_TNF_INHIBITOR_TREATED_DC_DN N_IRF4_8_NULL_PRE_BCELL_UP, GSE10273_HIGH_IL7_VS_HIGH_IL7_AND_IRF4_IN_IRF4_8_NULL_PRE_BCELL_UP ATID_COHESION, REACTOME_RESOLUTION_OF_SISTER_CHROMATID_COHESION
	GSE2770_UNTREATED_VS_IL12_TREATED_ACT_ GSE14413_UNSTIM_VS_IFNB_STIM_NIH3T3_CEL GSE40068_CXCR5NEG_BCL6NEG_CD4_TCELL_V	CD4_TCELL_48H_DN, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_DN LS_DN, GSE14413_UNSTIM_VS_IFNB_STIM_NIH3T3_CELLS_DN S_CXCR5POS_BCL6NEG_TFH_UP, GSE40068_CXCR5NEG_BCL6NEG_CD4_TCELL_VS_CXCR5POS_BCL6NEG_TFH_UP DRION, GOBP_PROTEIN_TARGETING_TO_MITOCHONDRION
	GOBP_TRANSCRIPTION_INITIATION_FROM_RNGOMF_UBIQUITIN_LIKE_PROTEIN_BINDING, GHP_PSYCHOMOTOR_RETARDATION, HP_PSYCHOMOTOR_TREG_DN, GSE24210_TCOGOMF_EXONUCLEASE_ACTIVITY, GOMF_EXONUCLEASE_ACTIVITY, GOMF_EXONUCLEASE_ACTIVITY	HOMOTOR_RETARDATION DNV_VS_TREG_DN
	GSE29614_DAY3_VS_DAY7_TIV_FLU_VACCINE_ GSE17721_CTRL_VS_LPS_12H_BMDC_DN, GSE17 GSE7460_FOXP3_MUT_VS_WT_ACT_WITH_TGFE	PBMC_DN, GSE29614_CTRL_VS_DAY7_TIV_FLU_VACCINE_PBMC_DN PBMC_DN, GSE29614_DAY3_VS_DAY7_TIV_FLU_VACCINE_PBMC_DN 721_CTRL_VS_LPS_12H_BMDC_DN 8_TCONV_UP, GSE7460_FOXP3_MUT_VS_WT_ACT_WITH_TGFB_TCONV_UP DN, GSE21670_TGFB_VS_IL6_TREATED_CD4_TCELL_DN
	REACTOME_HIV_TRANSCRIPTION_INITIATION HP_CLUBBING, HP_CLUBBING GOMF_EXONUCLEASE_ACTIVITY_ACTIVE_WIT	E_6H_DN, GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_6H_DN I, REACTOME_HIV_TRANSCRIPTION_INITIATION TH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS, GOMF_EXONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUC RICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA, GOBP_MATURATION_OF_5_8S_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA
	GOBP_RNA_CAPPING, GOBP_RNA_CAPPING GOMF_3_5_EXONUCLEASE_ACTIVITY, GOMF_3 GOBP_DICARBOXYLIC_ACID_METABOLIC_PRO	S_PLASMA_CELL_DAY7_UP, GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP  _5_EXONUCLEASE_ACTIVITY DCESS, GOBP_DICARBOXYLIC_ACID_METABOLIC_PROCESS N_FACTOR_BINDING, GOMF_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_BINDING
	GOBP_NCRNA_CATABOLIC_PROCESS, GOBP_N GOBP_RNA_PHOSPHODIESTER_BOND_HYDRO	PLEX_ASSEMBLY, GOBP_TRANSCRIPTION_PREINITIATION_COMPLEX_ASSEMBLY
	HP_KINETIC_TREMOR, HP_KINETIC_TREMOR MIR6787_3P, MIR6787_3P HP_NEONATAL_RESPIRATORY_DISTRESS, HP_I REACTOME_SYNTHESIS_OF_ACTIVE_UBIQUITI	
	HP_DRY_SKIN, HP_DRY_SKIN GOCC_EXORIBONUCLEASE_COMPLEX, GOCC_I GOMF_EXORIBONUCLEASE_ACTIVITY, GOMF_I MIR4695_5P, MIR4695_5P	EXORIBONUCLEASE_COMPLEX
	MIR660_3P, MIR660_3P GOBP_RRNA_CATABOLIC_PROCESS, GOBP_RRN	RANSCRIPTION_INITIATION, GOBP_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION  NA_CATABOLIC_PROCESS
	HP_MOTOR_AXONAL_NEUROPATHY, HP_MOT BENITEZ_GBM_PROTEASOME_INHIBITION_RES	MORPHOLOGY, HP_ABNORMAL_PULMONARY_INTERSTITIAL_MORPHOLOGY  FOR_AXONAL_NEUROPATHY  SPONSE, BENITEZ_GBM_PROTEASOME_INHIBITION_RESPONSE  PLEX, GOCC_TRANSCRIPTION_FACTOR_TFIID_COMPLEX
	GOMF_K63_LINKED_POLYUBIQUITIN_MODIFIC GOMF_RNA_POLYMERASE_II_GENERAL_TRAN GOMF_PEPTIDE_N_ACETYLTRANSFERASE_ACT	JP, GSE17721_LPS_VS_GARDIQUIMOD_1H_BMDC_UP CATION_DEPENDENT_PROTEIN_BINDING, GOMF_K63_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING ISCRIPTION_INITIATION_FACTOR_ACTIVITY, GOMF_RNA_POLYMERASE_II_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY IIVITY, GOMF_PEPTIDE_N_ACETYLTRANSFERASE_ACTIVITY IIATION_FROM_RNA_POLYMERASE_II_PROMOTER, GOBP_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER
	GOBP_MICROTUBULE_POLYMERIZATION, GOB	LLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY
	REACTOME_TRANSCRIPTIONAL_ACTIVITY_OF REACTOME_KSRP_KHSRP_BINDS_AND_DESTA	J, REACTOME_HDR_THROUGH_MMEJ_ALT_NHEJ F_SMAD2_SMAD3_SMAD4_HETEROTRIMER, REACTOME_TRANSCRIPTIONAL_ACTIVITY_OF_SMAD2_SMAD3_SMAD4_HETEROTRIMER BILIZES_MRNA, REACTOME_KSRP_KHSRP_BINDS_AND_DESTABILIZES_MRNA THETIC_PROCESS, GOBP_SERINE_FAMILY_AMINO_ACID_BIOSYNTHETIC_PROCESS TEILLANCE
	REACTOME_BRANCHED_CHAIN_AMINO_ACII	
	HP_ABNORMAL_HAIR_LABORATORY_EXAMINGOBP_POLYOL_BIOSYNTHETIC_PROCESS, GOBKEGG_BASAL_TRANSCRIPTION_FACTORS, KEGGOBP_MICROTUBULE_NUCLEATION, GOBP_MICROTUBULE_NUCLEATION, GOBP_MICROTUBULE_NUCLEAT	NATION, HP_ABNORMAL_HAIR_LABORATORY_EXAMINATION P_POLYOL_BIOSYNTHETIC_PROCESS G_BASAL_TRANSCRIPTION_FACTORS
	NIKOLSKY_BREAST_CANCER_17Q11_Q21_AMPI GOBP_MATERNAL_PROCESS_INVOLVED_IN_FI HUMMEL_BURKITTS_LYMPHOMA_UP, HUMMI GOBP_EXIT_FROM_MITOSIS, GOBP_EXIT_FROM	LICON, NIKOLSKY_BREAST_CANCER_17Q11_Q21_AMPLICON EMALE_PREGNANCY, GOBP_MATERNAL_PROCESS_INVOLVED_IN_FEMALE_PREGNANCY EL_BURKITTS_LYMPHOMA_UP
	MIR1238_5P_MIR4758_5P, MIR1238_5P_MIR4758_5 GOBP_HORMONE_MEDIATED_SIGNALING_PATHP_MOTOR_POLYNEUROPATHY, HP_MOTOR_I	THWAY, GOBP_HORMONE_MEDIATED_SIGNALING_PATHWAY POLYNEUROPATHY
	GOBP_CELLULAR_MODIFIED_AMINO_ACID_BI HP_ABNORMALITY_OF_DNA_REPAIR, HP_ABN GOCC_GAMMA_TUBULIN_COMPLEX, GOCC_G HP_ABSENCE_OF_SUBCUTANEOUS_FAT, HP_A GOCC_SIN3_TYPE_COMPLEX, GOCC_SIN3_TYPE	AMMA_TUBULIN_COMPLEX BSENCE_OF_SUBCUTANEOUS_FAT
	HP_GENERALIZED_HYPERREFLEXIA, HP_GENE	T_PROCESS, GOBP_MODULATION_BY_HOST_OF_SYMBIONT_PROCESS
	HASLINGER_B_CLL_WITH_CHROMOSOME_12_GOBP_FOLIC_ACID_CONTAINING_COMPOUNI HP_ADIPOSE_TISSUE_LOSS, HP_ADIPOSE_TISSU GOMF_RNA_POLYMERASE_II_REPRESSING_TRA	TRISOMY, HASLINGER_B_CLL_WITH_CHROMOSOME_12_TRISOMY D_METABOLIC_PROCESS, GOBP_FOLIC_ACID_CONTAINING_COMPOUND_METABOLIC_PROCESS JE_LOSS ANSCRIPTION_FACTOR_BINDING, GOMF_RNA_POLYMERASE_II_REPRESSING_TRANSCRIPTION_FACTOR_BINDING OGY, HP_ABNORMAL_THORACIC_SPINE_MORPHOLOGY
	HP_BIRD_LIKE_FACIES, HP_BIRD_LIKE_FACIES HP_DIFFUSE_CEREBELLAR_ATROPHY, HP_DIFFUSE_FACIES GOBP_PHOSPHATE_ION_TRANSPORT, GOBP_P CHENG_RESPONSE_TO_NICKEL_ACETATE, CH	FUSE_CEREBELLAR_ATROPHY HOSPHATE_ION_TRANSPORT ENG_RESPONSE_TO_NICKEL_ACETATE
	GOBP_HISTONE_H2A_UBIQUITINATION, GOBP HP_LOW_SET_NIPPLES, HP_LOW_SET_NIPPLES HP_RHINITIS, HP_RHINITIS MIR6728_5P, MIR6728_5P	
	BILANGES_SERUM_SENSITIVE_VIA_TSC2, BILANGES_HP_ABNORMAL_BRONCHUS_PHYSIOLOGY, HE HP_3_METHYLGLUTACONIC_ACIDURIA, HP_3_HP_THORACIC_KYPHOSIS, HP_THORACIC_KYPHOSIS, HP_THORACIC_KYPHOSIS, HP_SPLIT_NAIL	P_ABNORMAL_BRONCHUS_PHYSIOLOGY _METHYLGLUTACONIC_ACIDURIA
	HP_DYSTROPHIC_FINGERNAILS, HP_DYSTROP	ND_HD_MTX_DN, CHEOK_RESPONSE_TO_MERCAPTOPURINE_AND_HD_MTX_DN
	HP_ABNORMAL_VENA_CAVA_MORPHOLOGY, WP_ONE_CARBON_METABOLISM, WP_ONE_CA HP_DEFECTIVE_DNA_REPAIR_AFTER_ULTRAV GOMF_PRENYLTRANSFERASE_ACTIVITY, GOM	, HP_ABNORMAL_VENA_CAVA_MORPHOLOGY ARBON_METABOLISM IOLET_RADIATION_DAMAGE, HP_DEFECTIVE_DNA_REPAIR_AFTER_ULTRAVIOLET_RADIATION_DAMAGE
	GOMF_RNA_POLYMERASE_III_GENERAL_TRAN REACTOME_METABOLISM_OF_FOLATE_AND_I GOBP_NEGATIVE_REGULATION_OF_DNA_DEP GOMF_PROTEIN_DISULFIDE_ISOMERASE_ACTI	NSCRIPTION_INITIATION_FACTOR_ACTIVITY, GOMF_RNA_POLYMERASE_III_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY PTERINES, REACTOME_METABOLISM_OF_FOLATE_AND_PTERINES PENDENT_DNA_REPLICATION, GOBP_NEGATIVE_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION IVITY, GOMF_PROTEIN_DISULFIDE_ISOMERASE_ACTIVITY
	BROWNE_HCMV_INFECTION_8HR_DN, BROWN HP_SUPERNUMERARY_SPLEENS, HP_SUPERNU NKX25_01, NKX25_01 GOBP_HISTONE_SERINE_PHOSPHORYLATION,	JMERARY_SPLEENS  GOBP_HISTONE_SERINE_PHOSPHORYLATION
	HP_SHUFFLING_GAIT, HP_SHUFFLING_GAIT HP_NUMEROUS_PIGMENTED_FRECKLES, HP_N GOCC_RNA_POLYMERASE_III_TRANSCRIPTION GOBP_POSITIVE_REGULATION_OF_MICROTUB	MODIFICATION, GOBP_N_TERMINAL_PROTEIN_AMINO_ACID_MODIFICATION  NUMEROUS_PIGMENTED_FRECKLES  N_REGULATOR_COMPLEX, GOCC_RNA_POLYMERASE_III_TRANSCRIPTION_REGULATOR_COMPLEX  ULE_POLYMERIZATION_OR_DEPOLYMERIZATION, GOBP_POSITIVE_REGULATION_OF_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION
	HP_ERYTHRODERMA, HP_ERYTHRODERMA HP_ATELECTASIS, HP_ATELECTASIS HP_BRITTLE_HAIR, HP_BRITTLE_HAIR MIR382_5P, MIR382_5P GOBP_DIOL_BIOSYNTHETIC_PROCESS, GOBP_D	DIOL_BIOSYNTHETIC_PROCESS
	GOBP_EMBRYONIC_CRANIAL_SKELETON_MOINTENSE HP_BRONCHITIS, HP_BRONCHITIS GOBP_TRANSCRIPTION_INITIATION_FROM_RN	MERIZATION, GOBP_REGULATION_OF_MICROTUBULE_POLYMERIZATION RPHOGENESIS, GOBP_EMBRYONIC_CRANIAL_SKELETON_MORPHOGENESIS NA_POLYMERASE_III_PROMOTER, GOBP_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_III_PROMOTER ATHWAY, REACTOME_NOTCH_HLH_TRANSCRIPTION_PATHWAY
	GOBP_NEGATIVE_REGULATION_OF_ANDROGE GOCC_SIN3_COMPLEX, GOCC_SIN3_COMPLEX GOMF_NUCLEOSOMAL_HISTONE_BINDING, GOBP_CORTICOSTEROID_RECEPTOR_SIGNALING	EN_RECEPTOR_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY
	GOBP_GLYCEROL_3_PHOSPHATE_METABOLIC_GOBP_POSITIVE_REGULATION_OF_TRANSLATE HP_ALOPECIA_OF_SCALP, HP_ALOPECIA_OF_STAIL_BANDING, HP_TIGER_TAIL_BANDING, HP_TIGER_TA	_PROCESS, GOBP_GLYCEROL_3_PHOSPHATE_METABOLIC_PROCESS IONAL_INITIATION, GOBP_POSITIVE_REGULATION_OF_TRANSLATIONAL_INITIATION SCALP ANDING
	GOCC_KINETOCHORE_MICROTUBULE, GOCC_ GOMF_HISTONE_THREONINE_KINASE_ACTIVI HP_WHEEZING, HP_WHEEZING GOBP_PTERIDINE_CONTAINING_COMPOUND_	TION, GOBP_HISTONE_THREONINE_PHOSPHORYLATION KINETOCHORE_MICROTUBULE TY, GOMF_HISTONE_THREONINE_KINASE_ACTIVITY  _BIOSYNTHETIC_PROCESS, GOBP_PTERIDINE_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS
	MIR5186, MIR5186  GOBP_EXTRACELLULAR_TRANSPORT, GOBP_E GOBP_ONE_CARBON_METABOLIC_PROCESS, G HP_FOCAL_CORTICAL_DYSPLASIA, HP_FOCAL HP_ABNORMAL_SPERM_MOTILITY, HP_ABNORMAL_SPERM	COBP_ONE_CARBON_METABOLIC_PROCESS C_CORTICAL_DYSPLASIA
	GOBP_GOLGI_DISASSEMBLY, GOBP_GOLGI_DIS	COUGH IVITY, GOMF_SODIUM_CHANNEL_REGULATOR_ACTIVITY
	GOBP_OUTER_MITOCHONDRIAL_MEMBRANE_GOBP_DNA_CATABOLIC_PROCESS_EXONUCLEGOBP_TRNA_TRANSCRIPTION, GOBP_TRNA_TIGOCC_TRNA_SPLICING_LIGASE_COMPLEX, GO	_ORGANIZATION, GOBP_OUTER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION EOLYTIC, GOBP_DNA_CATABOLIC_PROCESS_EXONUCLEOLYTIC RANSCRIPTION
	HOXD1_TARGET_GENES, HOXD1_TARGET_GENES HP_ABNORMAL_RESPIRATORY_MOTILE_CILIU GOBP_TETRAHYDROFOLATE_METABOLIC_PRO HP_RECURRENT_MYCOBACTERIAL_INFECTION	NES IM_MORPHOLOGY, HP_ABNORMAL_RESPIRATORY_MOTILE_CILIUM_MORPHOLOGY OCESS, GOBP_TETRAHYDROFOLATE_METABOLIC_PROCESS NS, HP_RECURRENT_MYCOBACTERIAL_INFECTIONS
	HP_CHRONIC_RHINITIS, HP_CHRONIC_RHINIT GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ HP_FEMALE_INFERTILITY, HP_FEMALE_INFERT HP_SHYNESS, HP_SHYNESS	_ON_THE_CH_NH_GROUP_OF_DONORS, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH_GROUP_OF_DONORS TILITY
	HP_INCREASED_MEAN_CORPUSCULAR_HEMO / MIR12126, MIR12126 / GOBP_POSITIVE_REGULATION_OF_FEEDING_B / HP_MONOTONIC_SPEECH, HP_MONOTONIC_S	
S_CD4_SP_THYMOCYTE_UP, GSE31082_DN_VS_CD4_SP_THYMOCYTE_U	GOBP_PHOSPHORYLATION_OF_RNA_POLYME	
	chr17p11, chr17p11 WP_PRION_DISEASE_PATHWAY, WP_PRION_D	IP_ABNORMALITY_OF_BODY_MASS_INDEX "Y, GOMF_PHOSPHOLIPID_SCRAMBLASE_ACTIVITY ISEASE_PATHWAY
	HP_ABNORMALITY_OF_THE_PULMONARY_VE MIR193A_5P, MIR193A_5P GOBP_REGULATION_OF_CILIUM_MOVEMENT,	N_DEPENDENT_PROTEIN_BINDING, GOMF_UBIQUITINATION_LIKE_MODIFICATION_DEPENDENT_PROTEIN_BINDING INS, HP_ABNORMALITY_OF_THE_PULMONARY_VEINS  GOBP_REGULATION_OF_CILIUM_MOVEMENT RIGHT_HEART, HP_CONGENITAL_MALFORMATION_OF_THE_RIGHT_HEART
	HP_NASAL_OBSTRUCTION, HP_NASAL_OBSTR HP_TETRAPARESIS, HP_TETRAPARESIS GOMF_NEDD8_SPECIFIC_PROTEASE_ACTIVITY HP_CONCAVE_NAIL, HP_CONCAVE_NAIL GOCC_MICROTUBULE_END, GOCC_MICROTUB	, GOMF_NEDD8_SPECIFIC_PROTEASE_ACTIVITY
	PID_PS1_PATHWAY, PID_PS1_PATHWAY HP_METAPHYSEAL_CHONDRODYSPLASIA, HP GOBP_CENTRIOLE_ELONGATION, GOBP_CENT	
	GOMF_SUCCINATE_TRANSMEMBRANE_TRANS HP_GENERALIZED_LYMPHADENOPATHY, HP_ GOBP_NEGATIVE_REGULATION_OF_PROTEIN_ GOMF_KINETOCHORE_BINDING, GOMF_KINET	SPORTER_ACTIVITY, GOMF_SUCCINATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY _GENERALIZED_LYMPHADENOPATHY _FOLDING, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_FOLDING
	GOBP_POSITIVE_REGULATION_OF_EXTRACELIGION GOCC_CENTRIOLAR_SUBDISTAL_APPENDAGE	ROTUBULE_PLUS_END OGY, HP_ABNORMAL_NASAL_MUCOSA_MORPHOLOGY LULAR_MATRIX_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_EXTRACELLULAR_MATRIX_ASSEMBLY I, GOCC_CENTRIOLAR_SUBDISTAL_APPENDAGE MINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_AROMATIC_AMINO_ACID_CONCENTRATION
	MIR6871_5P, MIR6871_5P GOBP_REPLICATION_FORK_PROTECTION, GOE REACTOME_REGULATION_OF_PTEN_LOCALIZE GOMF_PROTEIN_PRENYLTRANSFERASE_ACTIVE	BP_REPLICATION_FORK_PROTECTION ZATION, REACTOME_REGULATION_OF_PTEN_LOCALIZATION VITY, GOMF_PROTEIN_PRENYLTRANSFERASE_ACTIVITY
	REACTOME_PTK6_EXPRESSION, REACTOME_PTGOCC_AP_4_ADAPTOR_COMPLEX, GOCC_AP_4 GOCC_HULC_COMPLEX, GOCC_HULC_COMPLEX ACGCACA_MIR210, ACGCACA_MIR210 GOMF_SODIUM_CHANNEL_INHIBITOR_ACTIV	4_ADAPTOR_COMPLEX
	GOBP_LEUCINE_METABOLIC_PROCESS, GOBP_GOBP_POSITIVE_REGULATION_OF_EXTRACELIGOCC_CYCLIN_CDK_POSITIVE_TRANSCRIPTION	T_IT, WP_LDLRAD4_AND_WHAT_WE_KNOW_ABOUT_IT LEUCINE_METABOLIC_PROCESS LULAR_MATRIX_DISASSEMBLY, GOBP_POSITIVE_REGULATION_OF_EXTRACELLULAR_MATRIX_DISASSEMBLY ON_ELONGATION_FACTOR_COMPLEX, GOCC_CYCLIN_CDK_POSITIVE_TRANSCRIPTION_ELONGATION_FACTOR_COMPLEX IE, GOCC_POSTSYNAPTIC_RECYCLING_ENDOSOME
	EVI1_06, EVI1_06 HP_SUBCORTICAL_HETEROTOPIA, HP_SUBCOI	TON, GOBP_REGULATION_OF_PROTEIN_NEDDYLATION
	GOBP_FOLIC_ACID_CONTAINING_COMPOUNI GOBP_VITAMIN_D_RECEPTOR_SIGNALING_PA REACTOME_ORGANIC_ANION_TRANSPORTER GOMF_NADPH_BINDING, GOMF_NADPH_BINI	D_BIOSYNTHETIC_PROCESS, GOBP_FOLIC_ACID_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS THWAY, GOBP_VITAMIN_D_RECEPTOR_SIGNALING_PATHWAY S, REACTOME_ORGANIC_ANION_TRANSPORTERS
	EHLERS_ANEUPLOIDY_DN, EHLERS_ANEUPLOGOMF_VOLTAGE_GATED_CHLORIDE_CHANNICHP_RECURRENT_CUTANEOUS_ABSCESS_FORMGOBP_5S_CLASS_RRNA_TRANSCRIPTION_BY_RECURRENT_CUTANEOUS_CRIPTION_BY_RECURRENT_CUTANE	DIDY_DN EL_ACTIVITY, GOMF_VOLTAGE_GATED_CHLORIDE_CHANNEL_ACTIVITY MATION, HP_RECURRENT_CUTANEOUS_ABSCESS_FORMATION RNA_POLYMERASE_III, GOBP_5S_CLASS_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_III
	GOBP_DNA_CATABOLIC_PROCESS, GOBP_DNA HP_ABNORMAL_MEAN_CORPUSCULAR_HEMO GOBP_T_HELPER_17_TYPE_IMMUNE_RESPONSO HP_DECREASED_FERTILITY_IN_FEMALES, HP_1	OGLOBIN_CONCENTRATION, HP_ABNORMAL_MEAN_CORPUSCULAR_HEMOGLOBIN_CONCENTRATION E, GOBP_T_HELPER_17_TYPE_IMMUNE_RESPONSE DECREASED_FERTILITY_IN_FEMALES
	GOBP_POSITIVE_REGULATION_OF_BASEMENT	REASED_BODY_MASS_INDEX  SYNTHETIC_PROCESS, GOBP_BRANCHED_CHAIN_AMINO_ACID_BIOSYNTHETIC_PROCESS  "_MEMBRANE_ASSEMBLY_INVOLVED_IN_EMBRYONIC_BODY_MORPHOGENESIS, GOBP_POSITIVE_REGULATION_OF_BASEMENT_MEMBRANE_ASSEMBLY_INVOLVED_IN_EMBRYONIC_BODY_MORPHOGENESIS  JITINATION, GOBP_REGULATION_OF_HISTONE_H2B_UBIQUITINATION
	GOBP_TETRAHYDROFOLATE_BIOSYNTHETIC_I GOBP_DETERMINATION_OF_DIGESTIVE_TRAC	CTIVITY, GOMF_NAD_P_H_OXIDASE_H2O2_FORMING_ACTIVITY PROCESS, GOBP_TETRAHYDROFOLATE_BIOSYNTHETIC_PROCESS T_LEFT_RIGHT_ASYMMETRY, GOBP_DETERMINATION_OF_DIGESTIVE_TRACT_LEFT_RIGHT_ASYMMETRY ION_IN_LATE_STAGE_BRANCHING_MORPHOGENESIS_PANCREATIC_BUD_PRECURSOR_CELLS, REACTOME_REGULATION_OF_GENE_EXPRESSION_IN_LATE_STAGE_BRANCHING_MORPHOGENESIS_PANCREATI OBP_RETINAL_ROD_CELL_DEVELOPMENT
	HP_LOWER_LIMB_UNDERGROWTH, HP_LOWE HP_PROLONGED_PR_INTERVAL, HP_PROLONGED_PR_INTERVAL	R_LIMB_UNDERGROWTH GED_PR_INTERVAL PROTEIN_TARGETING_TO_MITOCHONDRION, GOBP_PROTEIN_PROCESSING_INVOLVED_IN_PROTEIN_TARGETING_TO_MITOCHONDRION VALINE_METABOLIC_PROCESS
	GOBP_NEGATIVE_REGULATION_OF_DNA_TEM GOBP_MEIOTIC_SPINDLE_ORGANIZATION, GO HP_SHORT_LOWER_LIMBS, HP_SHORT_LOWER HP_HYPOPLASIA_OF_THE_VENTRAL_PONS, HI	IPLATED_TRANSCRIPTION_INITIATION, GOBP_NEGATIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION DBP_MEIOTIC_SPINDLE_ORGANIZATION R_LIMBS
	GOBP_HISTONE_H2A_K63_LINKED_UBIQUITIN REACTOME_SULFIDE_OXIDATION_TO_SULFAT	FT_RIGHT_ASYMMETRY, GOBP_DETERMINATION_OF_PANCREATIC_LEFT_RIGHT_ASYMMETRY ATION, GOBP_HISTONE_H2A_K63_LINKED_UBIQUITINATION TE, REACTOME_SULFIDE_OXIDATION_TO_SULFATE PHATE, REACTOME_SYNTHESIS_OF_DOLICHYL_PHOSPHATE
	HP_ABNORMAL_RESPIRATORY_MOTILE_CILIU GOBP_INTERLEUKIN_17_PRODUCTION, GOBP_ HP_CONGENITAL_NONBULLOUS_ICHTHYOSIF LI_PBMC_MENACTRA_AGE_18_45YO_7DY_UP, I	FORM_ERYTHRODERMA, HP_CONGENITAL_NONBULLOUS_ICHTHYOSIFORM_ERYTHRODERMA LI_PBMC_MENACTRA_AGE_18_45YO_7DY_UP
	HP_SHORT_STEPPED_SHUFFLING_GAIT, HP_SH HP_EVERTED_UPPER_LIP_VERMILION, HP_EVE SCHAEFFER_PROSTATE_DEVELOPMENT_AND_ HP_UPLIFTED_EARLOBE, HP_UPLIFTED_EARLO GOBP_INOSITOL_TRISPHOSPHATE_METABOLIC	HORT_STEPPED_SHUFFLING_GAIT ERTED_UPPER_LIP_VERMILION _CANCER_BOX4_UP, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX4_UP DBE C_PROCESS, GOBP_INOSITOL_TRISPHOSPHATE_METABOLIC_PROCESS
	GOBP_REGULATION_OF_WOUND_HEALING_SIGNATURE GOMF_ARYL_HYDROCARBON_RECEPTOR_BIN	GOBP_EMBRYONIC_BODY_MORPHOGENESIS N, HP_EXCESSIVE_WRINKLING_OF_PALMAR_SKIN PREADING_OF_EPIDERMAL_CELLS, GOBP_REGULATION_OF_WOUND_HEALING_SPREADING_OF_EPIDERMAL_CELLS DING, GOMF_ARYL_HYDROCARBON_RECEPTOR_BINDING IGNALING, REACTOME_DOWNREGULATION_OF_ERBB4_SIGNALING
	GOBP_DOLICHYL_DIPHOSPHATE_BIOSYNTHET	
	GOBP_NEGATIVE_REGULATION_OF_OXIDATIV	YE_PHOSPHORYLATION, GOBP_NEGATIVE_REGULATION_OF_OXIDATIVE_PHOSPHORYLATION TP_SYNTHESIS_COUPLED_ELECTRON_TRANSPORT, GOBP_REGULATION_OF_MITOCHONDRIAL_ATP_SYNTHESIS_COUPLED_ELECTRON_TRANSPORT EXPRESSED_WITH_CYP19A1
	HP_ENLARGEMENT_OF_THE_WRISTS, HP_ENL REACTOME_MECP2_REGULATES_TRANSCRIPT HP_ABNORMAL_ATRIAL_ARRANGEMENT, HP_ chr14q23, chr14q23	ARGEMENT_OF_THE_WRISTS ION_OF_NEURONAL_LIGANDS, REACTOME_MECP2_REGULATES_TRANSCRIPTION_OF_NEURONAL_LIGANDS _ABNORMAL_ATRIAL_ARRANGEMENT
	HP_LEWY_BODIES, HP_LEWY_BODIES GOBP_EPITHELIAL_CILIUM_MOVEMENT_INVO SNAPC2_TARGET_GENES, SNAPC2_TARGET_GE HP_POLYSPLENIA, HP_POLYSPLENIA	
	FARMER_BREAST_CANCER_CLUSTER_6, FARMS GOBP_CALCIUM_ACTIVATED_PHOSPHOLIPID_ LET_7E_3P, LET_7E_3P GOBP_10_FORMYLTETRAHYDROFOLATE_META	SCRAMBLING, GOBP_CALCIUM_ACTIVATED_PHOSPHOLIPID_SCRAMBLING  ABOLIC_PROCESS, GOBP_10_FORMYLTETRAHYDROFOLATE_METABOLIC_PROCESS
	GOBP_CORTICOSTERONE_SECRETION, GOBP_C GOBP_NEGATIVE_REGULATION_OF_TRANSCR MIR10396A_3P_MIR10396B_3P, MIR10396A_3P_MI HP_HYPERPHENYLALANINEMIA, HP_HYPERPI HP_BLUE_NEVUS, HP_BLUE_NEVUS	IPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER, GOBP_NEGATIVE_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER IR10396B_3P
	GOBP_UBIQUITIN_DEPENDENT_ENDOCYTOSIS	
	GOBP_POSITIVE_REGULATION_OF_PROTEIN_N GOCC_TRANSCRIPTION_FACTOR_TFIIA_COMP GOCC_POLAR_MICROTUBULE, GOCC_POLAR_I GOCC_BASAL_CORTEX, GOCC_BASAL_CORTEX GOMF_CYCLOHYDROLASE_ACTIVITY, GOMF_C	IEDDYLATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_NEDDYLATION PLEX, GOCC_TRANSCRIPTION_FACTOR_TFIIA_COMPLEX MICROTUBULE ( CYCLOHYDROLASE_ACTIVITY
	GOMF_BETA_2_ADRENERGIC_RECEPTOR_BINE GOMF_HISTONE_ACETYLTRANSFERASE_ACTIV	DING, GOMF_BETA_2_ADRENERGIC_RECEPTOR_BINDING VITY_H3_K23_SPECIFIC, GOMF_HISTONE_ACETYLTRANSFERASE_ACTIVITY_H3_K23_SPECIFIC CTIVITY, GOMF_INOSITOL_TRISPHOSPHATE_KINASE_ACTIVITY NEAL_MUSCLE_ATROPHY
	REACTOME_ROLE_OF_ABL_IN_ROBO_SLIT_SIG	NALING, REACTOME_ROLE_OF_ABL_IN_ROBO_SLIT_SIGNALING BIOGENESIS, WP_EXRNA_MECHANISM_OF_ACTION_AND_BIOGENESIS
	HP_BREATHING_DYSREGULATION, HP_BREAT HP_PROMINENT_ANTIHELIX, HP_PROMINENT GOBP_EMBRYONIC_VISCEROCRANIUM_MORP KRISHNAN_FURIN_TARGETS_DN, KRISHNAN_	_ANTIHELIX HOGENESIS, GOBP_EMBRYONIC_VISCEROCRANIUM_MORPHOGENESIS FURIN_TARGETS_DN
	GOBP_GLUCOCORTICOID_SECRETION, GOBP_G HP_ABNORMAL_THYROID_HORMONE_LEVEL, PID_LPA4_PATHWAY, PID_LPA4_PATHWAY chr2q14, chr2q14 HP_SPOTTY_HYPOPIGMENTATION, HP_SPOTTY HP_LONG_LIPPER_LIP_HP_LONG_LIPPER_LIP	HP_ABNORMAL_THYROID_HORMONE_LEVEL
	HP_LONG_UPPER_LIP, HP_LONG_UPPER_LIP HP_INTRAVENTRICULAR_HEMORRHAGE, HP_ BIOCARTA_PELP1_PATHWAY, BIOCARTA_PELI	INTRAVENTRICULAR_HEMORRHAGE P1_PATHWAY LEUCINE_CATABOLIC_PROCESS _RESPONSE_TO_AMMONIUM_ION
	GOBP_RESPONSE_TO_AMMONIUM_ION, GOBP	DIAC_MUSCLE_CELL_MEMBRANE_DEPOLARIZATION, GOBP_REGULATION_OF_VENTRICULAR_CARDIAC_MUSCLE_CELL_MEMBRANE_DEPOLARIZATION
	GOBP_RESPONSE_TO_AMMONIUM_ION, GOBP_GOBP_REGULATION_OF_VENTRICULAR_CARD HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTANTIVE_REGULATION_OF_MAST_CELL GOBP_CORTICOSTEROID_HORMONE_SECRETICAL LOPEZ_MESOTHELIOMA_SURVIVAL_OVERALL	L_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_POSITIVE_REGULATION_OF_MAST_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE ON, GOBP_CORTICOSTEROID_HORMONE_SECRETION _DN, LOPEZ_MESOTHELIOMA_SURVIVAL_OVERALL_DN
	GOBP_RESPONSE_TO_AMMONIUM_ION, GOBP_GOBP_REGULATION_OF_VENTRICULAR_CARE HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_PERCUSSION_MYOTONIA, HP_PERCUSSION_BIOCARTA_RARRXR_PATHWAY, BIOCARTA_RARRXR_PATHWAY, BIOCARTA_PATHWAY,	L_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_POSITIVE_REGULATION_OF_MAST_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE ON, GOBP_CORTICOSTEROID_HORMONE_SECRETION ,_DN, LOPEZ_MESOTHELIOMA_SURVIVAL_OVERALL_DN HOEK_B_CELL_2011_2012_TIV_ADULT_7DY_UP OGENESIS, GOBP_EMBRYONIC_NEUROCRANIUM_MORPHOGENESIS N_MYOTONIA
	GOBP_RESPONSE_TO_AMMONIUM_ION, GOBP_GOBP_REGULATION_OF_VENTRICULAR_CARE HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTANTIA_CINCULATING_PHENYLALANTIA_GOBP_THYROID_HORMONE_MEDIATED_SIGN_GRAPANTIA_SIGN_GLIOSIS, HP_SUBSTANTIA_GLIOSIS, HP_SUBSTANTIA_GLIOSIS, HP_SUBSTANTIA_GLIOSIS, HP_SUBSTANTIA_CINCULATING_PHENYLALANTIA_GOBP_THYROID_HORMONE_MEDIATED_SIGN_GRAPANTIA_GLIOSIS AND AMBOLIATED_SIGN_GOBP_THYROID_HORMONE_MEDIATED_SIGN_GRAPANTIA_GLIOSIS AND AMBOLIATED_SIGN_GOBP_THYROID_HORMONE_MEDIATED_THYROID_HORMONE_MEDIATED_SIGN_GOBP_THYROID_HORMONE_MEDIATED_THYROID_HORMONE_MEDIATED_THYROID_HORMONE_MEDIATED_THYROID_HORMONE_MEDIATED_THYROID_HORMONE_MEDIATED_THYROID_H	L_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_POSITIVE_REGULATION_OF_MAST_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE ON, GOBP_CORTICOSTEROID_HORMONE_SECRETION ,_DN, LOPEZ_MESOTHELIOMA_SURVIVAL_OVERALL_DN HOEK_B_CELL_2011_2012_TIV_ADULT_7DY_UP OGENESIS, GOBP_EMBRYONIC_NEUROCRANIUM_MORPHOGENESIS N_MYOTONIA ARRXR_PATHWAY NT, HP_HIGH_FREQUENCY_HEARING_IMPAIRMENT JENCY_INVOLVED_IN_CILIARY_MOTILITY, GOBP_REGULATION_OF_CILIUM_BEAT_FREQUENCY_INVOLVED_IN_CILIARY_MOTILITY
	GOBP_RESPONSE_TO_AMMONIUM_ION, GOBP_GOBP_REGULATION_OF_VENTRICULAR_CARE HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTAGOBP_POSITIVE_REGULATION_OF_MAST_CELICOBP_CORTICOSTEROID_HORMONE_SECRETICE LOPEZ_MESOTHELIOMA_SURVIVAL_OVERALE HOEK_B_CELL_2011_2012_TIV_ADULT_7DY_UP, GOBP_EMBRYONIC_NEUROCRANIUM_MORPH HP_PERCUSSION_MYOTONIA, HP_PERCUSSION_BIOCARTA_RARRXR_PATHWAY, BIOCARTA_RAY HP_HIGH_FREQUENCY_HEARING_IMPAIRMENGOBP_REGULATION_OF_CILIUM_BEAT_FREQUENCY_HEARING_PHENYLALANGOBP_THYROID_HORMONE_MEDIATED_SIGN_GAO_SMALL_INTESTINE_24W_C3_ENTEROCYTGOBP_REGULATION_OF_CILIUM_BEAT_FREQUENCY_HISTONE_H3_K14_ACETYLATION, GOBP_NICK_RESPONSE_TO_PROC_TREATMENT_UP, Nothryq24, chryq24	L_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_POSITIVE_REGULATION_OF_MAST_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE  DN, GOBP_CORTICOSTEROID_HORMONE_SECRETION DN, LOPEZ_MESOTHELIOMA_SURVIVAL_OVERALL_DN  HOEK B CELL_2011_2012_TIV_ADULT_7DY_UP  OGENESIS, GOBP_EMBRYONIC_NEUROCRANIUM_MORPHOGENESIS  N_MYOTONIA  ARRXR_PATHWAY  TT, HP_HIGH_PREQUENCY_HEARING_IMPAIRMENT  JENCY_INVOLVED_IN_CILLIARY_MOTILITY, GOBP_REGULATION_OF_CILLIUM_BEAT_FREQUENCY_INVOLVED_IN_CILLIARY_MOTILITY  ALT_ERRIOR_FONTANELLE  JINE_CONCENTRATION, HP_ABNORMAL_CIRCULATING_PHENYLALANINE_CONCENTRATION  ALING_PATHWAY, GOBP_THYROID_HORMONE_MEDIATED_SIGNALING_PATHWAY  E_PROGENITIOR_SUBTYPE_1, GAO_SMALL_INTESTINE_24W_C3_ENTEROCYTE_PROGENITOR_SUBTYPE_1  JENCY_GOBP_REGULATION_OF_CILLIUM_BEAT_FREQUENCY  _*HISTONE_H3_K14_ACETYLATION  NICK_RESPONSE_TO_PROC_TREATMENT_UP
	GOBP_RESPONSE_TO_AMMONIUM_ION, GOBP_GOBP_REGULATION_OF_VENTRICULAR_CARE HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTA GOBP_POSITIVE_REGULATION_OF_MAST_CELIGOBP_CORTICOSTEROID_HORMONE_SECRETICELOPEZ_MESOTHELIOMA_SURVIVAL_OVERALICHOEK_B_CELL_2011_2012_TIV_ADULT_7DY_UP, GOBP_EMBRYONIC_NEUROCRANIUM_MORPHHP_PERCUSSION_MYOTONIA, HP_PERCUSSION_BIOCARTA_RARRXR_PATHWAY, BIOCARTA_RAHP_HIGH_FREQUENCY_HEARING_IMPAIRMENGOBP_REGULATION_OF_CILIUM_BEAT_FREQUENCY_HEARING_PHENYLALANGOBP_THYROID_HORMONE_MEDIATED_SIGN_GAO_SMALL_INTESTINE_24W_C3_ENTEROCYTGOBP_REGULATION_OF_CILIUM_BEAT_FREQUENCY_COBP_HISTONE_H3_K14_ACETYLATION, GOBP_NICK_RESPONSE_TO_PROC_TREATMENT_UP, ChrXq24, chrXq24 GOBP_NEGATIVE_REGULATION_OF_FATTY_ACMIR3197, MIR3197 GOCC_NUCLEAR_COHESIN_COMPLEX, GOCC_MIR3683, MIR3683 GOBP_REGULATION_OF_THYROID_HORMONE	L_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, COBP_POSITIVE_REGULATION_OF_MAST_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE  DN, GOBP_CORTICOSTEROID_HORMONE_SECRETION
	GOBP_RESPONSE_TO_AMMONIUM_ION, GOBP_GOBP_REGULATION_OF_VENTRICULAR_CARE HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTANTIA_NIGRA_GOBP_CORTICOSTEROID_HORMONE_SECRETICAL LOPEZ_MESOTHELIOMA_SURVIVAL_OVERALLANTERIOMA_SURVIVAL_OVERALLANTERIOMA_SURVIVAL_OVERALLANTERIOMA_SURVIVAL_OVERALLANTERIOMA_SURVIVAL_OVERALLANTERIOMA_HP_PERCUSSION_MYOTONIA, HP_PERCUSSION_BIOCARTA_RARRXR_PATHWAY, BIOCARTA_RAMP_HIGH_FREQUENCY_HEARING_IMPAIRMENTAMENTAMENTAMENTAMENTAMENTAMENTAMENTA	LACTIVATION INVOLVED IN LIMININE, RESPONSE, GOBE_POSITIVE_REGULATION_OF_MAST_CELL_ACTIVATION_INVOLVED_IN_LIMINUTE_RESPONSE ON, COBP_CORTICOSTEROID HORMONE SECRETION LON, LOPEZ_MISOTHELIONA_SURVIVAL_COURALL_DN HOFE B_CELL_2011_2012_TV_ADULT_TOY_UP OSCENESS, GOBP_EMBRYONIC_NEUROCRANIUM_MORPHOGENESIS S. MYOTONIA ARRER, PATHIWAY IT, HP_HIGH_FREQUENCY_HEARING_IMPAIRMENT ENCY_INVOLVED_IN_CILLARY_MOILLITY, COBP_REGULATION_OF_CILLUM_BEAT_FREQUENCY_INVOLVED_IN_CILLARY_MOTILLITY ENCY_INVOLVED_IN_CILLARY_MOILLITY, COBP_REGULATION_OF_CILLUM_BEAT_FREQUENCY_INVOLVED_IN_CILLARY_MOTILLITY ENCY_INVOLVED_IN_TILTARY_MORE_INTEROULLITY INC_PHENYLALANINE_CONCENTRATION ALING_PATHIWAY, GOBE_INTROID_HORMONE_MEDIATED_SIGNALING_PATHIWAY E_PROCENITOR_SUBTYPE_I_GAO_SMALL_INTESTINE_24W_C3_ENTEROCYTE_PROGENITOR_SUBTYPE_I ENCY_COBP_RECULATION_OF_CILLUM_BEAT_FREQUENCY _INSTONE_ID_ENCY_COBP_RECULATION_OF_CILLUM_BEAT_FREQUENCY _INSTONE_ID_ENCY_COBP_RECULATION_OF_CILLUM_BEAT_FREQUENCY _INSTONE_ID_ENCY_COBP_RECULATION_OF_THY ACID_BIOSYNTHETIC_PROCESS  NUCLEAR_COHESIN_COMPLEX  **MEDIATED_SIGNALING_PATHWAY, GOBP_REGULATION_OF_THYROID_HORMONE_MEDIATED_SIGNALING_PATHWAY RESPONSE_TO_ALUMINUM_ION TICL_PROTEIN_FOLDING, GOBP_REGULATION_OF_CHAPERONE_MEDIATED_PROTEIN_FOLDING  **PORT_ACROSS_PLASMA_MEMBRANE_COBP_REGULATION OF_POTASSIUM_ION_EXPORT_ACROSS_PLASMA_MEMBRANE V, COBP_REGULATION_OF_FEMALE_RECEPTIVITY
	GOBP_RESPONSE_TO_AMMONIUM_ION, GOBP_GOBP_REGULATION_OF_VENTRICULAR_CARE HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTANTIA_NIGRAL_AND HELD HELD HP_SECUSSION_MYOTONIA, HP_PERCUSSION_MYOTONIA, HP_PERCUSSION_BIOCARTA_RARRXR_PATHWAY, BIOCARTA_RAMD HP_HIGH_FREQUENCY_HEARING_IMPAIRMENTAMENTAMENTAMENTAMENTAMENTAMENTAMENTA	LACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE_CORP_POSITIVE_REGULATION_OF_MAST_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE  OR, CORP_CORPICOSETRODI IDORMONE_SERVERALL_DN  HORK_B_CFLI_2011_2012_IV_ADALIT_ZDY_UP  OCERPSIS, CORP_FMIRRYCNIC_NEUROCRANIUM_MORPHOGENESIS  NAYOTORIA  REROR PATHWAY  IV_IP_IRIGIL_REGUENCY_HEARING_IMPAIRMENT  INFOC_INCOLOR_BLANY_MORITITY_CORP_REGULATION_OF_CRIJUM_BEAT_EREQUENCY_INVOLVED_IN_CRIJARY_MORITITY ALL_ANTERIOR_FONTANELLE  INFOC_INCOLOR_BLANY_MORITITY_CORP_REGULATION_OF_CRIJUM_BEAT_EREQUENCY_INVOLVED_IN_CRIJARY_MORITITY ALL_ANTERIOR_FONTANELLE  INFOC_INCOLOR_BLANY_MORITITY_CORP_REGULATION_OF_CRIJUM_BEAT_EREQUENCY_INVOLVED_IN_CRIJARY_MORITITY ALL_ANTERIOR_FONTANELLE  INFOC_INCOLOR_BLANY_MORITITY_CORP_REGULATION_OF_CRIJUM_BEAT_EREQUENCY_INVOLVED_IN_CRIJARY_MORITITY ALL_ANTERIOR_FONTANELLE  INFOC_INCOLOR_BLANY_MORITITY_CORP_REGULATION_OF_CRIJUM_BEAT_EREQUENCY_INVOLVED_IN_CRIJARY_MORITITY ALL_ANTERIOR_FONTANELLE  INFOC_INCOLOR_BLANY_MORITITY ALL_ANTERIOR_FORTANELLE  INFORM_FORTANELLE
	GOBP_RESPONSE_TO_AMMONIUM_ION, GOBP_GOBP_REGULATION_OF_VENTRICULAR_CARE_HP_SUBSTANTIA_NIGRA_GLIOSIS, HP_SUBSTA_GOBP_POSITIVE_REGULATION_OF_MAST_CELIGOBP_CORTICOSTEROID_HORMONE_SECRETIC_LOPEZ_MESOTHELIOMA_SURVIVAL_OVERALL_HOEK_B_CELL_2011_2012_TIV_ADULT_7DY_UP, GOBP_EMBRYONIC_NEUROCRANIUM_MORPH_HP_PERCUSSION_MYOTONIA, HP_PERCUSSION_BIOCARTA_RARRXR_PATHWAY, BIOCARTA_RAMP_HIGH_FREQUENCY_HEARING_IMPAIRMENGOBP_REGULATION_OF_CILLIUM_BEAT_FREQUENCY_HEARING_PHENYLALANGOBP_THYROID_HORMONE_MEDIATED_SIGN_GAO_SMALL_INTESTINE_24W_C3_ENTEROCYTGOBP_REGULATION_OF_CILIUM_BEAT_FREQUENCY_PROC_TREATMENT_UP, Not	LACTIVATION INVOLVED IN IMMUNE RESPONSE CODE POSITIVE REGULATION OF MAST CELL ACTIVATION INVOLVED IN IMMUNE RESPONSE NO, CODE CONTECTSTED HORMOND SPERTS OF THE NOTION OF MAST CELL ACTIVATION INVOLVED IN IMMUNE RESPONSE NO, CODE CONTENT OF THE NOTION OF T