	GSE41867_DAY6_VS_DAY8_LCMV_CLONE13_EFFE GSE41867_DAY8_EFFECTOR_VS_DAY30_EXHAUST GSE14308_TH2_VS_INDUCED_TREG_DN, GSE14308 GSE18893_CTRL_VS_TNF_TREATED_TCONV_2H_U GSE14350_IL2RB_KO_VS_WT_TREG_UP, GSE14350_ GSE2405_S_AUREUS_VS_UNTREATED_NEUTROPE	UP, GSE18893_CTRL_VS_TNF_TREATED_TCONV_2H_UP _IL2RB_KO_VS_WT_TREG_UP HIL_UP, GSE2405_S_AUREUS_VS_UNTREATED_NEUTROPHIL_UP
	GCM_ING1, GCM_ING1 GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_ HP_POOR_EYE_CONTACT, HP_POOR_EYE_CONT GSE12198_CTRL_VS_HIGH_IL2_STIM_NK_CELL_D	MC_UP, GSE9960_GRAM_NEG_VS_GRAM_POS_SEPSIS_PBMC_UP _DN, GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_DN
	GSE14308_NAIVE_CD4_TCELL_VS_NATURAL_TRE MIR8076, MIR8076 GSE20152_HTNFA_OVERXPRESS_ANKLE_VS_CTR HP_INCREASED_SERUM_PYRUVATE, HP_INCREA GSE18893_TCONV_VS_TREG_2H_TNF_STIM_UP, G MIR297, MIR297	EG_DN, GSE14308_NAIVE_CD4_TCELL_VS_NATURAL_TREG_DN RL_SPHK1_KO_ANKLE_UP, GSE20152_HTNFA_OVERXPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_UP ASED_SERUM_PYRUVATE
	GSE15930_STIM_VS_STIM_AND_IFNAB_24H_CD8_ HP_ABNORMAL_CIRCULATING_CARBOHYDRAT GSE30153_LUPUS_VS_HEALTHY_DONOR_BCELL_ MODULE_195, MODULE_195	O_DOUBLE_POSITIVE_THYMOCYTE_UP, GSE26488_WT_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_UP _T_CELL_UP, GSE15930_STIM_VS_STIM_AND_IFNAB_24H_CD8_T_CELL_UP _TE_CONCENTRATION, HP_ABNORMAL_CIRCULATING_CARBOHYDRATE_CONCENTRATION _UP, GSE30153_LUPUS_VS_HEALTHY_DONOR_BCELL_UP _IG_ENZYME_ACTIVITY, GOMF_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_ACTIVITY _GSE10239_NAIVE_VS_MEMORY_CD8_TCELL_DN
	RAX2_TARGET_GENES, RAX2_TARGET_GENES MODULE_147, MODULE_147	JP, GSE12198_CTRL_VS_HIGH_IL2_STIM_NK_CELL_UP
	HOLLEMAN_VINCRISTINE_RESISTANCE_B_ALL_ MODULE_576, MODULE_576	
	AMIT_SERUM_RESPONSE_480_MCF10A, AMIT_SEROCC_ANAPHASE_PROMOTING_COMPLEX, GOOCC_RNA_POLYMERASE_II_CORE_COMPLEX, GOOCC_RNA_POLYMERASE_II_CORE_COMPLEX	CRUM_RESPONSE_480_MCF10A CC_ANAPHASE_PROMOTING_COMPLEX GOCC_RNA_POLYMERASE_II_CORE_COMPLEX MALLY_FOLDED_HELIX
	MIR218_2_3P, MIR218_2_3P BILANGES_SERUM_SENSITIVE_GENES, BILANGES REACTOME_MICRORNA_MIRNA_BIOGENESIS, RI GOMF_ATPASE_REGULATOR_ACTIVITY, GOMF_A HP_APLASIA_HYPOPLASIA_INVOLVING_THE_VI	S_SERUM_SENSITIVE_GENES EACTOME_MICRORNA_MIRNA_BIOGENESIS
	BIOCARTA_G2_PATHWAY, BIOCARTA_G2_PATH HP_DEEP_PHILTRUM, HP_DEEP_PHILTRUM MIR6731_3P, MIR6731_3P	MAPK_ACTIVATION, REACTOME_ACTIVATED_TAK1_MEDIATES_P38_MAPK_ACTIVATION
	GOBP_FIBROBLAST_MIGRATION, GOBP_FIBROBL GOBP_REGULATION_OF_PEPTIDYL_SERINE_PHO GOBP_SOMATIC_STEM_CELL_POPULATION_MAT GOBP_NEGATIVE_REGULATION_OF_INTRACELL GSE22589_SIV_VS_HIV_AND_SIV_INFECTED_DC_I	_C, REACTOME_PHOSPHORYLATION_OF_THE_APC_C LAST_MIGRATION OSPHORYLATION, GOBP_REGULATION_OF_PEPTIDYL_SERINE_PHOSPHORYLATION INTENANCE, GOBP_SOMATIC_STEM_CELL_POPULATION_MAINTENANCE LULAR_PROTEIN_TRANSPORT, GOBP_NEGATIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT DN, GSE22589_SIV_VS_HIV_AND_SIV_INFECTED_DC_DN
	MIR4693_3P, MIR4693_3P MIR1231, MIR1231	
		P_RESPONSE_TO_INTERFERON_BETA
	GOBP_REGULATION_OF_PROTEIN_EXPORT_FROGOBP_PROTEASOME_ASSEMBLY, GOBP_PROTEALLIAO_HAVE_SOX4_BINDING_SITES, LIAO_HAVE_	_SOX4_BINDING_SITES ON, GOBP_REGULATION_OF_FIBROBLAST_MIGRATION
	GOBP_REGULATION_OF_MAINTENANCE_OF_SISTEM PPARA_01, PPARA_01 BIOCARTA_TOLL_PATHWAY, BIOCARTA_TOLL_I ZNF560_TARGET_GENES, ZNF560_TARGET_GENE	ES
	HP_HEPATIC_ENCEPHALOPATHY, HP_HEPATIC_HP_BEAKING_OF_VERTEBRAL_BODIES, HP_BEAKING_ABNORMALITY_OF_THE_DISTAL_PHALANX_GOBP_MODIFIED_AMINO_ACID_TRANSPORT, GOBP_MODIFIED_AMINO_ACID_TRANSPORT, GOBP_MODIFIED_AMINO_ACID_TRANSPORT	KING_OF_VERTEBRAL_BODIES K_OF_THE_THUMB, HP_ABNORMALITY_OF_THE_DISTAL_PHALANX_OF_THE_THUMB OBP_MODIFIED_AMINO_ACID_TRANSPORT
	GOBP_REGULATION_OF_CHROMATIN_ASSEMBLE GOBP_POSITIVE_REGULATION_OF_PEPTIDYL_SEGOBP_REGULATION_OF_PROTEIN_LOCALIZATION GOBP_POSITIVE_REGULATION_OF_MITOTIC_NUTURE GOCC_GLYCOGEN_GRANULE, GOCC_GLYCOGENWAESCH_ANAPHASE_PROMOTING_COMPLEX, NGOMF_MICROTUBULE_MINUS_END_BINDING, GOODS	JMORS, SNIJDERS_AMPLIFIED_IN_HEAD_AND_NECK_TUMORS LY_OR_DISASSEMBLY, GOBP_REGULATION_OF_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY ERINE_PHOSPHORYLATION, GOBP_POSITIVE_REGULATION_OF_PEPTIDYL_SERINE_PHOSPHORYLATION ON_TO_NUCLEOLUS, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEOLUS JCLEAR_DIVISION, GOBP_POSITIVE_REGULATION_OF_MITOTIC_NUCLEAR_DIVISION IN_GRANULE WAESCH_ANAPHASE_PROMOTING_COMPLEX
	HP_HYDROURETER, HP_HYDROURETER chr15q22, chr15q22 GOBP_POSITIVE_REGULATION_OF_NUCLEAR_DEGUBP_POSITIVE_REGULATION_OF_MAINTENANGOBP_FLAVIN_CONTAINING_COMPOUND_MET	DIVISION, GOBP_POSITIVE_REGULATION_OF_NUCLEAR_DIVISION NCE_OF_SISTER_CHROMATID_COHESION, GOBP_POSITIVE_REGULATION_OF_MAINTENANCE_OF_SISTER_CHROMATID_COHESION TABOLIC_PROCESS, GOBP_FLAVIN_CONTAINING_COMPOUND_METABOLIC_PROCESS TT_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_FIBROBLAST_APOPTOTIC_PROCESS
	WRN_TARGET_GENES, WRN_TARGET_GENES GOBP_NUCLEAR_ENVELOPE_REASSEMBLY, GOB GOMF_MODIFIED_AMINO_ACID_TRANSMEMBR. chr17q22, chr17q22 GOBP_POSITIVE_REGULATION_OF_ACTION_POT	
	GOBP_NEGATIVE_REGULATION_OF_FIBROBLAST REACTOME_TRAF6_MEDIATED_IRF7_ACTIVATION ZNF404_TARGET_GENES, ZNF404_TARGET_GENE GOBP_CARTILAGE_MORPHOGENESIS, GOBP_CARTILAGE	RANSPORT, GOBP_PLASMA_MEMBRANE_TO_ENDOSOME_TRANSPORT T_MIGRATION, GOBP_NEGATIVE_REGULATION_OF_FIBROBLAST_MIGRATION DN_IN_TLR7_8_OR_9_SIGNALING, REACTOME_TRAF6_MEDIATED_IRF7_ACTIVATION_IN_TLR7_8_OR_9_SIGNALING ES LRTILAGE_MORPHOGENESIS VE_T_CELLS, HP_DECREASED_PROPORTION_OF_CD3_POSITIVE_T_CELLS ACIAL_DYSOSTOSIS
	GOBP_TRACHEA_CARTILAGE_DEVELOPMENT, CGOMF_ALCOHOL_DEHYDROGENASE_NADPPLUHP_MIXED_HEARING_IMPAIRMENT, HP_MIXED_REACTOME_IRAK2_MEDIATED_ACTIVATION_OF	JS_ACTIVITY, GOMF_ALCOHOL_DEHYDROGENASE_NADPPLUS_ACTIVITY _HEARING_IMPAIRMENT F_TAK1_COMPLEX, REACTOME_IRAK2_MEDIATED_ACTIVATION_OF_TAK1_COMPLEX
	GOMF_ADENYL_NUCLEOTIDE_EXCHANGE_FACTOR GOBP_CENTROMERIC_SISTER_CHROMATID_COME HP_UNUSUAL_SKIN_INFECTION, HP_UNUSUAL_GOBP_RNA_GUANINE_N7_METHYLATION, GOBD	INDLE_POLE_CENTROSOME OT_BY_UV_IN_WS, KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_UV_IN_WS CTOR_ACTIVITY, GOMF_ADENYL_NUCLEOTIDE_EXCHANGE_FACTOR_ACTIVITY HESION, GOBP_CENTROMERIC_SISTER_CHROMATID_COHESION _SKIN_INFECTION
	GOBP_RETINAL_METABOLIC_PROCESS, GOBP_RI HP_ABNORMALITY_OF_THE_STAPES, HP_ABNOR	PRIMITIVE_STREAK_FORMATION N, HP_ABNORMALITY_OF_THE_ACHILLES_TENDON ETINAL_METABOLIC_PROCESS RMALITY_OF_THE_STAPES
	HP_ABNORMALITY_OF_FRONTALIS_MUSCLE_BE GOMF_CORECEPTOR_ACTIVITY_INVOLVED_IN_ GOMF_INTERLEUKIN_1_BINDING, GOMF_INTER	NTOAXIAL_ABNORMALITY T_PROLIFERATION, GOBP_NEGATIVE_REGULATION_OF_FIBROBLAST_PROLIFERATION ELLY, HP_ABNORMALITY_OF_FRONTALIS_MUSCLE_BELLY WNT_SIGNALING_PATHWAY, GOMF_CORECEPTOR_ACTIVITY_INVOLVED_IN_WNT_SIGNALING_PATHWAY LEUKIN_1_BINDING
		MINERGIC_SYNAPSE METHYLATION ITHESIS, WP_MOLYBDENUM_COFACTOR_MOCO_BIOSYNTHESIS RPHOLOGY, HP_ABNORMAL_PLACENTAL_MEMBRANE_MORPHOLOGY TOMA
ARMSTRONG_MEMORY_CD8_TCELL_DN, GSE41867_NAIVE_VS_DAY30_LCMV_ARMSTRONG_MEMORY_CD8_TCELL_DN	CAO_BLOOD_FLUMIST_AGE_05_14YO_7DY_UP, COME GOBP_POSITIVE_REGULATION_OF_BLOOD_PRESE GOBP_REGULATION_OF_TYPE_B_PANCREATIC_OMP_ABNORMAL_GREATER_SCIATIC_NOTCH_MOME HP_FEMORAL_BOWING, HP_FEMORAL_BOWING	CAO_BLOOD_FLUMIST_AGE_05_14YO_7DY_UP SSURE_BY_EPINEPHRINE_NOREPINEPHRINE, GOBP_POSITIVE_REGULATION_OF_BLOOD_PRESSURE_BY_EPINEPHRINE_NOREPINEPHRINE CELL_PROLIFERATION, GOBP_REGULATION_OF_TYPE_B_PANCREATIC_CELL_PROLIFERATION ORPHOLOGY, HP_ABNORMAL_GREATER_SCIATIC_NOTCH_MORPHOLOGY
	GOBP_NEGATIVE_REGULATION_OF_CAMP_MED GOBP_UDP_N_ACETYLGLUCOSAMINE_BIOSYNT REACTOME_ALPHA_PROTEIN_KINASE_1_SIGNA GOBP_NEGATIVE_REGULATION_OF_GTPASE_AC HP_RHIZOMELIC_ARM_SHORTENING, HP_RHIZO	DIATED_SIGNALING, GOBP_NEGATIVE_REGULATION_OF_CAMP_MEDIATED_SIGNALING THETIC_PROCESS, GOBP_UDP_N_ACETYLGLUCOSAMINE_BIOSYNTHETIC_PROCESS ALING_PATHWAY, REACTOME_ALPHA_PROTEIN_KINASE_1_SIGNALING_PATHWAY CTIVITY, GOBP_NEGATIVE_REGULATION_OF_GTPASE_ACTIVITY
	HP_LINEAR_ARRAYS_OF_MACULAR_HYPERKER HP_MOLLUSCUM_CONTAGIOSUM, HP_MOLLUSCONTAGIOSUM, HP_MOLLUSCONTAGIOSUM, HP_MOLLUSCONTAGETS, OHASHI_AURKA_TAGETS, OHASHI_AURKA_TAGETS_B_DIRECT	RATOSES_IN_FLEXURAL_AREAS, HP_LINEAR_ARRAYS_OF_MACULAR_HYPERKERATOSES_IN_FLEXURAL_AREAS CUM_CONTAGIOSUM
	REACTOME_SIGNALING_BY_RNF43_MUTANTS, F	NT_EARLY_DN, RAMPON_ENRICHED_LEARNING_ENVIRONMENT_EARLY_DN DMF_SPHINGOLIPID_TRANSFER_ACTIVITY . CAO_BLOOD_FLUZONE_AGE_05_14YO_1DY_DN
	GOBP_PROSTHETIC_GROUP_METABOLIC_PROCE	ESS, GOBP_PROSTHETIC_GROUP_METABOLIC_PROCESS NTHESIS, REACTOME_MOLYBDENUM_COFACTOR_BIOSYNTHESIS ETS
	GOMF_CHROMATIN_INSULATOR_SEQUENCE_BIGORP_FOLIC_ACID_TRANSPORT, GOBP_FOLIC_A	BREAK_REPAIR_VIA_SISTER_CHROMATID_EXCHANGE, GOBP_REPLICATION_BORN_DOUBLE_STRAND_BREAK_REPAIR_VIA_SISTER_CHROMATID_EXCHANGE
	GOBP_REGULATION_OF_ER_ASSOCIATED_UBIQUED HP_ABNORMAL_GLOMERULAR_FILTRATION_RA	
	GOBP_REGULATION_OF_DNA_METHYLATION, C	NTERLEUKIN_23_PRODUCTION SEVERELY_REDUCED_VISUAL_ACUITY ITY, GOMF_ALDO_KETO_REDUCTASE_NADP_ACTIVITY
	HP_DECREASED_HIP_ABDUCTION, HP_DECREASED_HIP_HIGH_PITCHED_CRY, HP_HIGH_PITCHED_CRY GOMF_RNA_POLYMERASE_II_ACTIVITY, GOMF_IRONA_POLYMERASE_I_GENERAL_TRANSC	RY RNA_POLYMERASE_II_ACTIVITY CRIPTION_INITIATION_FACTOR_ACTIVITY, GOMF_RNA_POLYMERASE_I_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY
		ATRIOVENTRICULAR_CANAL_DEFECT
	GOBP_NEGATIVE_REGULATION_OF_HEART_RATGOBP_REGULATION_OF_HEXOKINASE_ACTIVITERACTOME_TICAM1_TRAF6_DEPENDENT_INDUSTREACTOME_TICAM1_TRAF6_DEPENDENT_INDUSTREACTOME_TICAM1_TRAF6_DEPENDENT_HANDHP_ABSENT_HA	TE, GOBP_NEGATIVE_REGULATION_OF_HEART_RATE TY, GOBP_REGULATION_OF_HEXOKINASE_ACTIVITY ICTION_OF_TAK1_COMPLEX, REACTOME_TICAM1_TRAF6_DEPENDENT_INDUCTION_OF_TAK1_COMPLEX
	GOBP_POSITIVE_REGULATION_OF_PROTEIN_KIN WP_NLR_PROTEINS, WP_NLR_PROTEINS HP_ENLARGEMENT_OF_THE_COSTOCHONDRAI HP_BILIARY_ATRESIA, HP_BILIARY_ATRESIA	IN_SILENCING, GOBP_NEGATIVE_REGULATION_OF_CHROMATIN_SILENCING NASE_C_SIGNALING, GOBP_POSITIVE_REGULATION_OF_PROTEIN_KINASE_C_SIGNALING L_JUNCTION, HP_ENLARGEMENT_OF_THE_COSTOCHONDRAL_JUNCTION HALANX_OF_THE_THUMB, HP_APLASIA_HYPOPLASIA_OF_THE_DISTAL_PHALANX_OF_THE_THUMB
	GOMF_CYTIDYLYLTRANSFERASE_ACTIVITY, GOTHP_FOOD_ALLERGY, HP_FOOD_ALLERGY GOCC_IKAPPAB_KINASE_COMPLEX, GOCC_IKAPGOMF_ADRENERGIC_RECEPTOR_ACTIVITY, GOME REACTOME_ADRENOCEPTORS, REACTOME_ADE HP_PARATHYROID_CARCINOMA, HP_PARATHY	PPAB_KINASE_COMPLEX MF_ADRENERGIC_RECEPTOR_ACTIVITY RENOCEPTORS VROID_CARCINOMA
	HP_MEMBRANOPROLIFERATIVE_GLOMERULON HP_LUMBAR_SCOLIOSIS, HP_LUMBAR_SCOLIOSIS HP_DECREASED_CERVICAL_SPINE_FLEXION_DUHP_KELOIDS, HP_KELOIDS CAO_BLOOD_FLUZONE_AGE_05_14YO_30DY_UP,	JE_TO_CONTRACTURES_OF_POSTERIOR_CERVICAL_MUSCLES, HP_DECREASED_CERVICAL_SPINE_FLEXION_DUE_TO_CONTRACTURES_OF_POSTERIOR_CERVICAL_MUSCLES , CAO_BLOOD_FLUZONE_AGE_05_14YO_30DY_UP
	GOBP_REGULATION_OF_RAB_PROTEIN_SIGNAL	ATE_BINDING E_TYPE_I_IFN_AND_INFLAMMATORY_CYTOKINES_PRODUCTION, REACTOME_DEX_H_BOX_HELICASES_ACTIVATE_TYPE_I_IFN_AND_INFLAMMATORY_CYTOKINES_PRODUCTIO TRANSDUCTION, GOBP_REGULATION_OF_RAB_PROTEIN_SIGNAL_TRANSDUCTION TED_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_ER_ASSOCIATED_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
	CHANG_POU5F1_TARGETS_DN, CHANG_POU5F1 GOBP_VITAMIN_TRANSPORT, GOBP_VITAMIN_T HP_ABNORMALITY_OF_PINEAL_MORPHOLOGY, GOBP_CERAMIDE_1_PHOSPHATE_TRANSPORT, G GOMF_CERAMIDE_1_PHOSPHATE_TRANSFER_AG	TRANSPORT 7, HP_ABNORMALITY_OF_PINEAL_MORPHOLOGY
	GOBP_THALAMUS_DEVELOPMENT, GOBP_THALE HP_CONE_CONE_ROD_DYSTROPHY, HP_CONE_C BUSSLINGER_DUODENAL_K_CELLS, BUSSLINGER BIOCARTA_GABA_PATHWAY, BIOCARTA_GABA_ HP_MOOD_SWINGS, HP_MOOD_SWINGS	LAMUS_DEVELOPMENT CONE_ROD_DYSTROPHY R_DUODENAL_K_CELLS
	REACTOME_NEGATIVE_REGULATION_OF_TCF_IGOBP_NEURON_GLIAL_CELL_SIGNALING, GOBFHP_APLASIA_HYPOPLASIA_INVOLVING_THE_MHP_PAROXYSMAL_SUPRAVENTRICULAR_TACHY	DS, WP_KENNEDY_PATHWAY_FROM_SPHINGOLIPIDS DEPENDENT_SIGNALING_BY_WNT_LIGAND_ANTAGONISTS, REACTOME_NEGATIVE_REGULATION_OF_TCF_DEPENDENT_SIGNALING_BY_WNT_LIGAND_ANTAGONISTS P_NEURON_GLIAL_CELL_SIGNALING IUSCULATURE_OF_THE_UPPER_LIMBS, HP_APLASIA_HYPOPLASIA_INVOLVING_THE_MUSCULATURE_OF_THE_UPPER_LIMBS YCARDIA, HP_PAROXYSMAL_SUPRAVENTRICULAR_TACHYCARDIA ATE_CONCENTRATION, HP_ABNORMAL_URINE_ALPHA_KETOGLUTARATE_CONCENTRATION
	GOBP_NEGATIVE_REGULATION_OF_SISTER_CHI GOBP_NEGATIVE_REGULATION_OF_HEXOKINA	ALATE RANSPORT N_23_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_23_PRODUCTION ROMATID_COHESION, GOBP_NEGATIVE_REGULATION_OF_SISTER_CHROMATID_COHESION .SE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_HEXOKINASE_ACTIVITY
	GOMF_TOXIN_TRANSMEMBRANE_TRANSPORTE BIOCARTA_PARKIN_PATHWAY, BIOCARTA_PAR	PARESIS 2_PAPILLARY_RENAL_CELL_CARCINOMA 5, HP_REDUCED_NATURAL_KILLER_CELL_COUNT ER_ACTIVITY, GOMF_TOXIN_TRANSMEMBRANE_TRANSPORTER_ACTIVITY RKIN_PATHWAY
	GOBP_RAB_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_REGULATION_OF_PEPTIDYL_CYSTEINE_S WP_GAMMAGLUTAMYL_CYCLE_FOR_THE_BIOS	AF1_COMPLEX LUCOSAMINE, REACTOME_SYNTHESIS_OF_UDP_N_ACETYL_GLUCOSAMINE
	SMID_BREAST_CANCER_RELAPSE_IN_BRAIN_UPGOBP_GENITALIA_MORPHOGENESIS, GOBP_GENchr16q11, chr16q11 HP_CARPAL_BONE_HYPOPLASIA, HP_CARPAL_IGOBP_POSITIVE_REGULATION_OF_SYSTEMIC_AIGOBP_NEGATIVE_REGULATION_OF_PROTEIN_T	P, SMID_BREAST_CANCER_RELAPSE_IN_BRAIN_UP NITALIA_MORPHOGENESIS BONE_HYPOPLASIA RTERIAL_BLOOD_PRESSURE, GOBP_POSITIVE_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE CARGETING_TO_MITOCHONDRION, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION
	GOBP_POSITIVE_REGULATION_OF_SYNAPTIC_TI HP_PERSISTENT_EBV_VIREMIA, HP_PERSISTENT_ GOBP_CANONICAL_WNT_SIGNALING_PATHWA GOBP_NEGATIVE_REGULATION_OF_ERAD_PATH GOMF_ALPHA_ADRENERGIC_RECEPTOR_ACTIV HP_HYPOURICEMIA, HP_HYPOURICEMIA	RANSMISSION_GABAERGIC, GOBP_POSITIVE_REGULATION_OF_SYNAPTIC_TRANSMISSION_GABAERGIC LEBV_VIREMIA AY_INVOLVED_IN_REGULATION_OF_CELL_PROLIFERATION, GOBP_CANONICAL_WNT_SIGNALING_PATHWAY_INVOLVED_IN_REGULATION_OF_CELL_PROLIFERATION HWAY, GOBP_NEGATIVE_REGULATION_OF_ERAD_PATHWAY ITTY, GOMF_ALPHA_ADRENERGIC_RECEPTOR_ACTIVITY
	GOMF_DYNEIN_HEAVY_CHAIN_BINDING, GOME GOBP_MIDBRAIN_HINDBRAIN_BOUNDARY_DEV KASLER_HDAC7_TARGETS_2_UP, KASLER_HDAC HP_ABNORMALITY_OF_THE_GLENOID_FOSSA, H GOBP_NEUROTRANSMITTER_GATED_ION_CHAN GOMF_DIACYLGLYCEROL_KINASE_ACTIVITY, GA	VELOPMENT, GOBP_MIDBRAIN_HINDBRAIN_BOUNDARY_DEVELOPMENT C7_TARGETS_2_UP HP_ABNORMALITY_OF_THE_GLENOID_FOSSA NNEL_CLUSTERING, GOBP_NEUROTRANSMITTER_GATED_ION_CHANNEL_CLUSTERING GOMF_DIACYLGLYCEROL_KINASE_ACTIVITY
	GOBP_MICTURITION, GOBP_MICTURITION GOBP_AXIS_ELONGATION_INVOLVED_IN_SOME GOBP_REGULATION_OF_NADP_METABOLIC_PRO GOCC_ER_MEMBRANE_INSERTION_COMPLEX, COMPLEX,	T_CONTRACTION_BY_CHEMICAL_SIGNAL, GOBP_REGULATION_OF_THE_FORCE_OF_HEART_CONTRACTION_BY_CHEMICAL_SIGNAL TOGENESIS, GOBP_AXIS_ELONGATION_INVOLVED_IN_SOMITOGENESIS COCESS, GOBP_REGULATION_OF_NADP_METABOLIC_PROCESS GOCC_ER_MEMBRANE_INSERTION_COMPLEX
	GOBP_REGULATION_OF_NON_MEMBRANE_SPAHP_LONG_THORAX, HP_LONG_THORAX GOBP_POSITIVE_REGULATION_OF_GRANULOCY GOBP_VITAMIN_TRANSMEMBRANE_TRANSPORT REACTOME_REGULATION_OF_FZD_BY_UBIQUIT	SM, REACTOME_VITAMIN_B1_THIAMIN_METABOLISM INNING_PROTEIN_TYROSINE_KINASE_ACTIVITY, GOBP_REGULATION_OF_NON_MEMBRANE_SPANNING_PROTEIN_TYROSINE_KINASE_ACTIVITY YTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_DIFFERENTIATION AT, GOBP_VITAMIN_TRANSMEMBRANE_TRANSPORT INATION, REACTOME_REGULATION_OF_FZD_BY_UBIQUITINATION EAL_JOINT, HP_ABNORMALITY_OF_METACARPOPHALANGEAL_JOINT
	GOBP_TYROSINE_METABOLIC_PROCESS, GOBP_TGOBP_NEGATIVE_REGULATION_OF_SMOOTH_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_SMOOTH_METAMBOLSKY_BOUND_BY_MUTATED_TP53, STANGOBP_RECEPTOR_DIFFUSION_TRAPPING, GOBP_HP_IMPAIRED_LYMPHOCYTE_TRANSFORMATICGOBP_THIAMINE_CONTAINING_COMPOUND_METAGOBP_THIAMINE_CONTAINING_COMPOUND_THIAMINE_CONTAINING_COMPOUND_THIAMINE_CONTAINING_COMPOUND_THIAMINE_CONTAINING_COMPOUND_THIAMINE_CONTAINING_COMPOUND_THIAMINE_CONTAINING_COMPOUND_THIAMINE_CONTAINING_COMPOUND_THIAMINE_CONTAINING_COMPOUND_THIAMINE_CONTAINING_COMPOUND_THIAMINE_CONTAINING_CONTAINING_CONTAINING_CONTAINING_CON	TYROSINE_METABOLIC_PROCESS MUSCLE_CELL_APOPTOTIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_APOPTOTIC_PROCESS MBOLSKY_BOUND_BY_MUTATED_TP53 _RECEPTOR_DIFFUSION_TRAPPING DN_WITH_PHYTOHEMAGGLUTININ, HP_IMPAIRED_LYMPHOCYTE_TRANSFORMATION_WITH_PHYTOHEMAGGLUTININ METABOLIC_PROCESS, GOBP_THIAMINE_CONTAINING_COMPOUND_METABOLIC_PROCESS
	GOBP_EMBRYONIC_RETINA_MORPHOGENESIS_I GOBP_CELLULAR_RESPONSE_TO_CHOLESTEROI GOBP_RECEPTOR_MEDIATED_ENDOCYTOSIS_IN	IN_CAMERA_TYPE_EYE, GOBP_EMBRYONIC_RETINA_MORPHOGENESIS_IN_CAMERA_TYPE_EYE L, GOBP_CELLULAR_RESPONSE_TO_CHOLESTEROL IVOLVED_IN_CHOLESTEROL_TRANSPORT, GOBP_RECEPTOR_MEDIATED_ENDOCYTOSIS_INVOLVED_IN_CHOLESTEROL_TRANSPORT SSURE, GOBP_POSITIVE_REGULATION_OF_BLOOD_PRESSURE