	GSE22886_NAIVE_CD4_TCELL_VS_48H_ACT_TH2_DN, GSE2288 GSE24634_NAIVE_CD4_TCELL_VS_DAY7_IL4_CONV_TREG_DN GSE22886_NAIVE_CD4_TCELL_VS_NKCELL_DN, GSE22886_NA	N, GSE24634_NAIVE_CD4_TCELL_VS_DAY7_IL4_CONV_TREG_DN AIVE_CD4_TCELL_VS_NKCELL_DN _STIM_DC_UP, GSE32986_UNSTIM_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_UP
	GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_LOWDOSE_ST GSE22886_NAIVE_BCELL_VS_BLOOD_PLASMA_CELL_DN, GSE GSE7764_IL15_NK_CELL_24H_VS_SPLENOCYTE_UP, GSE7764_IL GSE22033_UNTREATED_VS_MRL24_TREATED_MEF_UP, GSE220 GSE22886_NAIVE_CD4_TCELL_VS_MEMORY_TCELL_DN, GSE220 GSE22886_CD8_VS_CD4_NAIVE_TCELL_UP, GSE22886_CD8_VS_	TIM_DC_UP, GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_LOWDOSE_STIM_DC_UP E22886_NAIVE_BCELL_VS_BLOOD_PLASMA_CELL_DN IL15_NK_CELL_24H_VS_SPLENOCYTE_UP 2033_UNTREATED_VS_MRL24_TREATED_MEF_UP 22886_NAIVE_CD4_TCELL_VS_MEMORY_TCELL_DN
	GSE10325_BCELL_VS_LUPUS_BCELL_DN, GSE10325_BCELL_VS_HALLMARK_DNA_REPAIR, HALLMARK_DNA_REPAIR GSE3982_BASOPHIL_VS_TH1_DN, GSE3982_BASOPHIL_VS_TH1 GSE17974_0H_VS_2H_IN_VITRO_ACT_CD4_TCELL_DN, GSE179 GSE10325_MYELOID_VS_LUPUS_MYELOID_DN, GSE10325_MYE GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_6H_DN, GSE2770_IL4_GSE37416_0H_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN	S_LUPUS_BCELL_DN 1_DN 974_0H_VS_2H_IN_VITRO_ACT_CD4_TCELL_DN ELOID_VS_LUPUS_MYELOID_DN _ACT_VS_ACT_CD4_TCELL_6H_DN N, GSE37416_0H_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN
	GSE24726_WT_VS_E2_2_KO_PDC_DAY4_POST_DELETION_UP, GSE33425_CD161_HIGH_VS_INT_CD8_TCELL_DN, GSE33425_CIGSE27786_LIN_NEG_VS_CD8_TCELL_UP, GSE27786_LIN_NEG_VGSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_EPPERT_PROGENITOR, EPPERT_PROGENITOR GSE6092_IFNG_VS_IFNG_AND_B_BURGDORFERI_INF_ENDOTIGSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_1H_CD4_TCELI	GSE24726_WT_VS_E2_2_KO_PDC_DAY4_POST_DELETION_UP D161_HIGH_VS_INT_CD8_TCELL_DN VS_CD8_TCELL_UP _48H_UP, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_UP THELIAL_CELL_DN, GSE6092_IFNG_VS_IFNG_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_DN L_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_1H_CD4_TCELL_DN
	HALLMARK_IL2_STAT5_SIGNALING, HALLMARK_IL2_STAT5_MTOR_UP.N4.V1_UP, MTOR_UP.N4.V1_UP GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_48H_UP, GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DGSE11864_UNTREATED_VS_CSF1_PAM3CYS_IN_MAC_DN, GSEGSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_	GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_48H_UP L_2H_UP, GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP DN, GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN E11864_UNTREATED_VS_CSF1_PAM3CYS_IN_MAC_DN _UP, GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_UP
	GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD4_TCELL_UP, GSGSE17721_PAM3CSK4_VS_CPG_1H_BMDC_DN, GSE17721_PAM3GSE22611_NOD2_VS_MUTANT_NOD2_TRANSDUCED_HEK293GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNG_KOMODULE_321, MODULE_321 GSE27859_MACROPHAGE_VS_CD11C_INT_F480_INT_DC_DN, GSE27850_INT_DC_DN, GSE27850_I	GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD4_TCELL_UP 13CSK4_VS_CPG_1H_BMDC_DN 13T_CELL_UP, GSE22611_NOD2_VS_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_UP 12T_CELL_UP, GSE22611_NOD2_TRANSDUCED_HEK293T_CELL_UP 12T_CEL
	KEGG_PURINE_METABOLISM, KEGG_PURINE_METABOLISM GSE360_T_GONDII_VS_B_MALAYI_LOW_DOSE_DC_UP, GSE360_GSE10325_CD4_TCELL_VS_LUPUS_CD4_TCELL_DN, GSE10325_0 MARTORIATI_MDM4_TARGETS_NEUROEPITHELIUM_UP, MAGSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IL4_TREATINGSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_TREATED_CD4_T0GSE29949_CD8_NEG_DC_SPLEEN_VS_MONOCYTE_BONE_MAI	60_T_GONDII_VS_B_MALAYI_LOW_DOSE_DC_UP _CD4_TCELL_VS_LUPUS_CD4_TCELL_DN
	GOBP_NUCLEOBASE_CONTAINING_SMALL_MOLECULE_BIOGSE22501_PERIPHERAL_BLOOD_VS_CORD_BLOOD_TREG_DNGSE21033_3H_VS_12H_POLYIC_STIM_DC_UP, GSE21033_3H_VS_JIANG_VHL_TARGETS, JIANG_VHL_TARGETS MIR216B_3P, MIR216B_3P	OSYNTHETIC_PROCESS, GOBP_NUCLEOBASE_CONTAINING_SMALL_MOLECULE_BIOSYNTHETIC_PROCESS N, GSE22501_PERIPHERAL_BLOOD_VS_CORD_BLOOD_TREG_DN S_12H_POLYIC_STIM_DC_UP AMAGE_STIMULUS, GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS HIRSCH_CELLULAR_TRANSFORMATION_SIGNATURE_DN
	GSE360_CTRL_VS_B_MALAYI_HIGH_DOSE_DC_UP, GSE360_CTGSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMC_1MONTH_PGSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_INAKAYA_PBMC_FLUARIX_FLUVIRIN_AGE_18_50YO_CORRELGSE19512_NAUTRAL_VS_INDUCED_TREG_DN, GSE19512_NAUWP_TNF_ALPHA_SIGNALING_PATHWAY, WP_TNF_ALPHA_SIGNALING_PATHWAY, WP_TNF_ALP	TRL_VS_B_MALAYI_HIGH_DOSE_DC_UP POST_DX_DN, GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMC_1MONTH_POST_DX_DN _DN, GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_DN LATED_WITH_HAI_28DY_RESPONSE_AT_7DY_POSITIVE, NAKAYA_PBMC_FLUARIX_FLUVIRIN_AGE_18_50YO_CORRELATED_WITH_HAI_28DY_RESPONSE_AT_7DY_POSITIVE UTRAL_VS_INDUCED_TREG_DN
	GSE41867_NAIVE_VS_DAY8_LCMV_CLONE13_EFFECTOR_CD8_SIRNA_EIF4GI_DN, SIRNA_EIF4GI_DN GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_ITURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_LOBULAR_	B_TCELL_DN, GSE41867_NAIVE_VS_DAY8_LCMV_CLONE13_EFFECTOR_CD8_TCELL_DN _DN, GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_DN _NORMAL_UP, TURASHVILI_BREAST_DUCTAL_CARCINOMA_VS_LOBULAR_NORMAL_UP LPOS_BONE_MARROW_DN, GSE35685_CD34POS_CD38NEG_VS_CD34POS_CD10NEG_CD62LPOS_BONE_MARROW_DN
	GSE2826_XID_VS_BTK_KO_BCELL_DN, GSE2826_XID_VS_BTK_E MARKEY_RB1_CHRONIC_LOF_UP, MARKEY_RB1_CHRONIC_I GSE16450_CTRL_VS_IFNA_12H_STIM_IMMATURE_NEURON_C GOBP_ACTIVATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_A GSE28130_ACTIVATED_VS_INDUCEED_TREG_UP, GSE28130_ACTIVATED_VS_INDUCEED_TREG_UP, GSE28130_ACTIVATED_PHOSPHATE_CATABOLIC_PROCESS, GO GSE5099_DAY3_VS_DAY7_MCSF_TREATED_MACROPHAGE_DE	KO_BCELL_DN LOF_UP CELL_LINE_DN, GSE16450_CTRL_VS_IFNA_12H_STIM_IMMATURE_NEURON_CELL_LINE_DN ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS, GOBP_ACTIVATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS ACTIVATED_VS_INDUCEED_TREG_UP
	MIR6837_3P, MIR6837_3P GOBP_POSITIVE_REGULATION_OF_CELL_GROWTH, GOBP_POSITIVE_REGULATION_OF_CELL_GROWTH, GOBP_POSITIVE_VS_PTPRC_NEG_CD4_TCELL_DN, GSE15659_GSE22886_TH1_VS_TH2_12H_ACT_DN, GSE22886_TH1_VS_TH2_KEGG_P53_SIGNALING_PATHWAY, KEGG_P53_SIGNALING_PHP_ABNORMAL_CONJUNCTIVA_MORPHOLOGY, HP_ABNORGOBP_PROTEIN_HOMOOLIGOMERIZATION, GOBP_PROTEIN_	OSITIVE_REGULATION_OF_CELL_GROWTH _NAIVE_VS_PTPRC_NEG_CD4_TCELL_DN 2_12H_ACT_DN PATHWAY RMAL_CONJUNCTIVA_MORPHOLOGY I_HOMOOLIGOMERIZATION
	GSE25846_IL10_POS_VS_NEG_CD8_TCELL_DAY7_POST_CORON BUFFA_HYPOXIA_METAGENE, BUFFA_HYPOXIA_METAGENE HP_CUTANEOUS_PHOTOSENSITIVITY, HP_RED_EYE, HP_RED_EYE SCHEIDEREIT_IKK_INTERACTING_PROTEINS, SCHEIDEREIT_I	NAVIRUS_BRAIN_UP, GSE25846_IL10_POS_VS_NEG_CD8_TCELL_DAY7_POST_CORONAVIRUS_BRAIN_UP E OTOSENSITIVITY ISEASES_OF_GLYCOSYLATION RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS JIKK_INTERACTING_PROTEINS
	GOBP_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTOR HP_HYDROURETER, HP_HYDROURETER GOBP_SOMATIC_DIVERSIFICATION_OF_IMMUNOGLOBULING GOBP_MONOSACCHARIDE_CATABOLIC_PROCESS, GOBP_MOTE GOCC_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE, IGARASHI_ATF4_TARGETS_DN, IGARASHI_ATF4_TARGETS_DX KIM_MYCN_AMPLIFICATION_TARGETS_UP, KIM_MYCN_TARGETS_UP,	RS, GOBP_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS IS, GOBP_SOMATIC_DIVERSIFICATION_OF_IMMUNOGLOBULINS ONOSACCHARIDE_CATABOLIC_PROCESS IS, GOCC_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE ON IPLIFICATION_TARGETS_UP
	GSE2585_CD80_HIGH_VS_LOW_MTEC_DN, GSE2585_CD80_HIC WP_SMALL_CELL_LUNG_CANCER, WP_SMALL_CELL_LUNG_ GOBP_POST_EMBRYONIC_DEVELOPMENT, GOBP_POST_EMBI GOBP_SOMATIC_RECOMBINATION_OF_IMMUNOGLOBULIN_ REACTOME_INTEGRATION_OF_ENERGY_METABOLISM, REACTOME_INTEGRATION_OF_ENERGY_METABOLISM, REACTOME_IMMATURE_VS_LISTERIA_INF_MATURE_DC_DN, GSGOBP_RIBONUCLEOTIDE_CATABOLIC_PROCESS, GOBP_RIBO	GH_VS_LOW_MTEC_DN _CANCER BRYONIC_DEVELOPMENT I_GENE_SEGMENTS, GOBP_SOMATIC_RECOMBINATION_OF_IMMUNOGLOBULIN_GENE_SEGMENTS ACTOME_INTEGRATION_OF_ENERGY_METABOLISM SE9946_IMMATURE_VS_LISTERIA_INF_MATURE_DC_DN DNUCLEOTIDE_CATABOLIC_PROCESS
	HP_DUPLICATION_OF_THUMB_PHALANX, HP_DUPLICATION HP_SLENDER_LONG_BONE, HP_SLENDER_LONG_BONE MIR4686, MIR4686 GSE46606_IRF4HIGH_VS_IRF4MID_CD40L_IL2_IL5_DAY1_STIM WARTERS_IR_RESPONSE_5GY, WARTERS_IR_RESPONSE_5GY GOCC_GOLGI_CISTERNA_MEMBRANE, GOCC_GOLGI_CISTERWP_P53_TRANSCRIPTIONAL_GENE_NETWORK, WP_P53_TRANSCRIPTIONAL_GENE_NETWORK, WP_P53_TRANSCRIPTIONAL_GENE_MP\$	ON_OF_THUMB_PHALANX IULATED_BCELL_DN, GSE46606_IRF4HIGH_VS_IRF4MID_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_DN RNA_MEMBRANE ANSCRIPTIONAL_GENE_NETWORK
	VANASSE_BCL2_TARGETS_DN, VANASSE_BCL2_TARGETS_DN MIR6814_5P, MIR6814_5P GOMF_THYROID_HORMONE_RECEPTOR_BINDING, GOMF_TI AMUNDSON_DNA_DAMAGE_RESPONSE_TP53, AMUNDSON_GSE25147_UNSTIM_VS_HELIOBACTER_PYLORI_LPS_STIM_MK HP_ABNORMAL_THORACIC_SPINE_MORPHOLOGY, HP_ABN GOMF_NUCLEOBASE_CONTAINING_COMPOUND_KINASE_A	N THYROID_HORMONE_RECEPTOR_BINDING _DNA_DAMAGE_RESPONSE_TP53 KN45_CELL_DN, GSE25147_UNSTIM_VS_HELIOBACTER_PYLORI_LPS_STIM_MKN45_CELL_DN NORMAL_THORACIC_SPINE_MORPHOLOGY ACTIVITY, GOMF_NUCLEOBASE_CONTAINING_COMPOUND_KINASE_ACTIVITY
	GSE13762_CTRL_VS_125_VITAMIND_DAY12_DC_DN, GSE13762 KEGG_FRUCTOSE_AND_MANNOSE_METABOLISM, KEGG_FRUBIOCARTA_CERAMIDE_PATHWAY, BIOCARTA_CERAMIDE_P	2_CTRL_VS_125_VITAMIND_DAY12_DC_DN RUCTOSE_AND_MANNOSE_METABOLISM PATHWAY OCESS, GOBP_PURINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS 5_CTRL_VS_LAIV_FLU_VACCINE_PBMC_UP L4_ISOFORM_A
	MIR101_5P, MIR101_5P GNF2_CDC27, GNF2_CDC27	RESPONSE_TO_OXIDATIVE_STRESS, GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_OXIDATIVE_STRESS _LIVER_CANCER_PROGRESSION_G1_G2_DN
	MODULE_284, MODULE_284 GOBP_PHOSPHATIDYLCHOLINE_BIOSYNTHETIC_PROCESS, C MIR6085, MIR6085	GOBP_PHOSPHATIDYLCHOLINE_BIOSYNTHETIC_PROCESS N, HOFFMANN_IMMATURE_TO_MATURE_B_LYMPHOCYTE_DN _ADIPOGENIC_GENES_REPRESSED_BY_SIRT1
	HP_THORACIC_KYPHOSIS, HP_THORACIC_KYPHOSIS MIR147A, MIR147A GOBP_GLYCOLYTIC_PROCESS_THROUGH_FRUCTOSE_6_PHO HP_LIMB_GIRDLE_MUSCLE_WEAKNESS, HP_LIMB_GIRDLE_M GOBP_CELLULAR_MODIFIED_AMINO_ACID_BIOSYNTHETIC_ GOBP_GLUCOSE_CATABOLIC_PROCESS, GOBP_GLUCOSE_CA GOBP_MISMATCH_REPAIR, GOBP_MISMATCH_REPAIR	OSPHATE, GOBP_GLYCOLYTIC_PROCESS_THROUGH_FRUCTOSE_6_PHOSPHATE MUSCLE_WEAKNESS 2_PROCESS, GOBP_CELLULAR_MODIFIED_AMINO_ACID_BIOSYNTHETIC_PROCESS
	HP_PARKINSONISM_WITH_FAVORABLE_RESPONSE_TO_DOF GOMF_MONOSACCHARIDE_BINDING, GOMF_MONOSACCHARIDE_SALVAGE, REACTOME_NUCLEOT GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_MREACTOME_REGULATION_OF_INSULIN_SECRETION, REACTOGOBP_REGULATION_OF_PROTEIN_DEUBIQUITINATION, GORDON	TARIDE_BINDING TIDE_SALVAGE MEDULLA_THYMUS_DN, GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_DN TOME_REGULATION_OF_INSULIN_SECRETION TOME_REGULATION_OF_PROTEIN_DEUBIQUITINATION PP_REGULATION, GOBP_NUCLEOTIDE_EXCISION_REPAIR_PREINCISION_COMPLEX_STABILIZATION
	NICK_RESPONSE_TO_PROC_TREATMENT_DN, NICK_RESPONDED HP_ARTERIAL_THROMBOSIS, HP_ARTERIAL_THROMBOSIS GOCC_COPII_VESICLE_COAT HP_DIMINISHED_MOVEMENT, HP_DIMINISHED_MOVEMENT GEISS_RESPONSE_TO_DSRNA_UP, G	NSE_TO_PROC_TREATMENT_DN T NA_UP ASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_PHOSPHATASE_ACTIVITY ETAL_LIVER_ERYTHROBLASTS
	GOBP_HISTONE_H2A_MONOUBIQUITINATION, GOBP_HISTOHASLINGER_B_CLL_WITH_11Q23_DELETION, HASLINGER_B_GOMF_CARBOHYDRATE_KINASE_ACTIVITY, GOMF_CARBOHHP_COMPLETE_DUPLICATION_OF_PHALANX_OF_HAND, HEBIOCARTA_TPO_PATHWAY, BIOCARTA_TPO_PATHWAY KIM_LRRC3B_TARGETS, KIM_LRRC3B_TARGETS chr1q22, chr1q22 BIOCARTA_IL6_PATHWAY, BIOCARTA_IL6_PATHWAY	_CLL_WITH_11Q23_DELETION HYDRATE_KINASE_ACTIVITY
	FARMER_BREAST_CANCER_CLUSTER_5, FARMER_BREAST_CANCER_CLUSTER_5, FARMER_BREAST_CANCER_CLUSTER_5, FARMER_BREAST_CANCER_CLUSTER_5, FARMER_BREAST_CANCER_	SIBILITY_OF_THE_KNEE SIDE_CATABOLIC_PROCESS VASCULATURE, HP_ABNORMAL_MORPHOLOGY_OF_THE_CONJUNCTIVAL_VASCULATURE TION_OF_PROTEINS, REACTOME_DISEASES_ASSOCIATED_WITH_N_GLYCOSYLATION_OF_PROTEINS PPONSE_TO_FENRETINIDE_UP
	GOBP_PHOSPHATIDYLINOSITOL_DEPHOSPHORYLATION, GO KEGG_GALACTOSE_METABOLISM, KEGG_GALACTOSE_META MODULE_272, MODULE_272 ZNF433_TARGET_GENES, ZNF433_TARGET_GENES MIR1296_5P, MIR1296_5P SCHEIDEREIT_IKK_TARGETS, SCHEIDEREIT_IKK_TARGETS YANG_BREAST_CANCER_ESR1_BULK_UP, YANG_BREAST_CA	ANCER_ESR1_BULK_UP
	GOBP_NEGATIVE_REGULATION_OF_NEURON_APOPTOTIC_E WU_HBX_TARGETS_1_DN, WU_HBX_TARGETS_1_DN GOMF_FIBRONECTIN_BINDING, GOMF_FIBRONECTIN_BINDI GOBP_CELL_DEATH_IN_RESPONSE_TO_OXIDATIVE_STRESS, WP_EPO_RECEPTOR_SIGNALING, WP_EPO_RECEPTOR_SIGNAL GOMF_MISMATCHED_DNA_BINDING, GOMF_MISMATCHED_ HP_ABNORMALITY_OF_ENDOCRINE_PANCREAS_PHYSIOLOGICAL GOMF_MISMATCHED_DNA_BINDING_COMF_MISMATCHED_ HP_ABNORMALITY_OF_ENDOCRINE_PANCREAS_PHYSIOLOGICAL GOMF_MISMATCHED_DNA_BINDING_COMF_MISMATCHED_ HP_ABNORMALITY_OF_ENDOCRINE_PANCREAS_PHYSIOLOGICAL GOMF_MISMATCHED_DNA_BINDING_COMF_MISMATCHED_ HP_ABNORMALITY_OF_ENDOCRINE_PANCREAS_PHYSIOLOGICAL GOMF_MISMATCHED_DNA_BINDING_COMF_MISMATCHED_ HP_ABNORMALITY_OF_ENDOCRINE_PANCREAS_PHYSIOLOGICAL GOMF_MISMATCHED_DNA_BINDING_COMF_MISMATCHED_ HP_ABNORMALITY_OF_ENDOCRINE_PANCREAS_PHYSIOLOGICAL GOMF_MISMATCHED_DNA_BINDING_COMF_	PROCESS, GOBP_NEGATIVE_REGULATION_OF_NEURON_APOPTOTIC_PROCESS ING GOBP_CELL_DEATH_IN_RESPONSE_TO_OXIDATIVE_STRESS ALING D_DNA_BINDING DGY, HP_ABNORMALITY_OF_ENDOCRINE_PANCREAS_PHYSIOLOGY
	BIOCARTA_TNFR2_PATHWAY, BIOCARTA_TNFR2_PATHWAY GARCIA_PINERES_PBMC_HPV_16_L1_VLP_AGE_18_25YO_7MC GOBP_EXTRACELLULAR_VESICLE_BIOGENESIS, GOBP_EXTRALLEE_CALORIE_RESTRICTION_MUSCLE_UP, LEE_CALORIE_RESTRICTION_MUSCLE_UP, LEE_CALORIE_RESTRICTION_PHATE_KINASE_ACTIVITY, GOMEN_DUPLICATION_OF_PHALANX_OF_HALLUX, HP_DUPLICATION_OF_PHALANX_OF_HALLUX, HP_COMMUNICATING_HYDROCEPHALUS, HP_COMMUNICATION_PHALANX_OF_PHALUS, HP_COMMUNICATION_PHALANX_PERMISSION_PATHWAY	Y O_CORRELATED_WITH_ANTIBODY_RESPONSE_POSITIVE, GARCIA_PINERES_PBMC_HPV_16_L1_VLP_AGE_18_25YO_7MO_CORRELATED_WITH_ANTIBODY_RESPONSE_POSITIVE ACELLULAR_VESICLE_BIOGENESIS ESTRICTION_MUSCLE_UP MF_NUCLEOSIDE_DIPHOSPHATE_KINASE_ACTIVITY ATION_OF_PHALANX_OF_HALLUX ATING_HYDROCEPHALUS
	HP_ABNORMALITY_OF_THE_PHALANGES_OF_THE_HALLUX GOBP_PURINE_CONTAINING_COMPOUND_SALVAGE, GOBP_HU_ANGIOGENESIS_UP, HU_ANGIOGENESIS_UP HP_HYPOKINESIA, HP_HYPOKINESIA HP_CHOLECYSTITIS, HP_CHOLECYSTITIS REACTOME_SYNTHESIS_OF_PIPS_AT_THE_ER_MEMBRANE, RHP_LAMBDOIDAL_CRANIOSYNOSTOSIS, HP_LAMBDOIDAL_CRANIOSYNOSTOSIS.	X, HP_ABNORMALITY_OF_THE_PHALANGES_OF_THE_HALLUX P_PURINE_CONTAINING_COMPOUND_SALVAGE REACTOME_SYNTHESIS_OF_PIPS_AT_THE_ER_MEMBRANE CRANIOSYNOSTOSIS
	REACTOME_PURINE_SALVAGE, REACTOME_PURINE_SALVA HP_PULMONARY_EMBOLISM, HP_PULMONARY_EMBOLISM HP_LIMB_GIRDLE_MUSCLE_ATROPHY, HP_LIMB_GIRDLE_MU GOBP_ALDITOL_METABOLIC_PROCESS, GOBP_ALDITOL_MET TIAN_TNF_SIGNALING_VIA_NFKB, TIAN_TNF_SIGNALING_VIA_BNORMALITY_OF_DNA_REPAIR, HP_ABNORMALITY_O GOBP_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_E	AGE USCLE_ATROPHY ITABOLIC_PROCESS VIA_NFKB DF_DNA_REPAIR ENDOPLASMIC_RETICULUM_STRESS, GOBP_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
	GOMF_MISMATCH_REPAIR_COMPLEX_BINDING, GOMF_MISMONE_CARBON_METABOLISM_AND_RELATED_PATHWAY SMID_BREAST_CANCER_ERBB2_UP, SMID_BREAST_CANCER_WP_PURINE_METABOLISM, WP_PURINE_METABOLISM GOMF_C_C_CHEMOKINE_BINDING, GOMF_C_C_CHEMOKINE WP_THE_HUMAN_IMMUNE_RESPONSE_TO_TUBERCULOSIS, HP_SEROSITIS, HP_SEROSITIS	SMATCH_REPAIR_COMPLEX_BINDING AYS, WP_ONE_CARBON_METABOLISM_AND_RELATED_PATHWAYS _ERBB2_UP
	MIR6870_3P, MIR6870_3P HP_ABNORMALITY_OF_THE_WING_OF_THE_ILIUM, HP_ABNOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROPERTIES HP_SEVERE_COMBINED_IMMUNODEFICIENCY, HP_SEVERE_BIOCARTA_EPO_PATHWAY, BIOCARTA_EPO_PATHWAY MARSON_FOXP3_CORE_DIRECT_TARGETS, MARSON_FOXP3_HP_DEVIATION_OF_THE_2ND_FINGER, HP_DEVIATION_OF_T	CESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS _COMBINED_IMMUNODEFICIENCY _CORE_DIRECT_TARGETS
	GOBP_NEGATIVE_REGULATION_OF_INTRACELLULAR_PROT GOBP_PROTEASOME_ASSEMBLY, GOBP_PROTEASOME_ASSEMBLY. REACTOME_CLEC7A_DECTIN_1_INDUCES_NFAT_ACTIVATION	TEIN_TRANSPORT, GOBP_NEGATIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT EMBLY ON, REACTOME_CLEC7A_DECTIN_1_INDUCES_NFAT_ACTIVATION D_336HR_UP, FULLER_PBMC_F_TULARENSIS_VACCINE_LVS_AGE_22_54YO_336HR_UP TER_LIMB_AMYOTROPHY
	GOBP_PYRIDINE_CONTAINING_COMPOUND_CATABOLIC_PIGOBP_POSITIVE_REGULATION_OF_MIRNA_MEDIATED_INHICGCTGCT_MIR503, CGCTGCT_MIR503 REACTOME_ASSEMBLY_OF_THE_HIV_VIRION, REACTOME_AMORSON_FOXP3_TARGETS_STIMULATED_UP, MARSON_FOXE	PROCESS, GOBP_PYRIDINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS IBITION_OF_TRANSLATION, GOBP_POSITIVE_REGULATION_OF_MIRNA_MEDIATED_INHIBITION_OF_TRANSLATION ASSEMBLY_OF_THE_HIV_VIRION ASSEMBLY_OF_THE_HIV_UP ATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_DEUBIQUITINATION
	YORDY_RECIPROCAL_REGULATION_BY_ETS1_AND_SP100_UIMIR6894_5P, MIR6894_5P GOBP_RIBONUCLEOSIDE_CATABOLIC_PROCESS, GOBP_RIBOURDED_CHEMOATTRACTANT_RECEPT GOBP_POSITIVE_REGULATION_OF_SMAD_PROTEIN_SIGNAL_HP_DEFECTIVE_DNA_REPAIR_AFTER_ULTRAVIOLET_RADIATED_BP_APOPTOTIC_PROCESS_INVOLVED_IN_DEVELOPMENT	P, YORDY_RECIPROCAL_REGULATION_BY_ETS1_AND_SP100_UP
	PID_IL3_PATHWAY, PID_IL3_PATHWAY GOBP_EMBRYONIC_VISCEROCRANIUM_MORPHOGENESIS, G GOCC_ENDOPLASMIC_RETICULUM_PLASMA_MEMBRANE_C HP_BROAD_PALM, HP_BROAD_PALM GOMF_PROTEIN_CARBOXYL_O_METHYLTRANSFERASE_ACT GOBP_TRANSEPITHELIAL_TRANSPORT, GOBP_TRANSEPITHE	GOBP_EMBRYONIC_VISCEROCRANIUM_MORPHOGENESIS CONTACT_SITE, GOCC_ENDOPLASMIC_RETICULUM_PLASMA_MEMBRANE_CONTACT_SITE TIVITY, GOMF_PROTEIN_CARBOXYL_O_METHYLTRANSFERASE_ACTIVITY
	GOBP_NEGATIVE_REGULATION_OF_TELOMERE_CAPPING, CHECK HP_LIMB_TREMOR, HP_LIMB_TREMOR MIR1251_5P, MIR1251_5P GOBP_PROTEIN_TRIMERIZATION, GOBP_PRO	GOBP_NEGATIVE_REGULATION_OF_TELOMERE_CAPPING ZATION IL_UP, WUNDER_INFLAMMATORY_RESPONSE_AND_CHOLESTEROL_UP REACTOME_GROWTH_HORMONE_RECEPTOR_SIGNALING ACTIVE_BLADDER
	GOBP_HISTIDINE_METABOLIC_PROCESS, GOBP_HISTIDINE_N GOCC_PLATELET_DENSE_TUBULAR_NETWORK, GOCC_PLATER REACTOME_INTERLEUKIN_6_FAMILY_SIGNALING, REACTOME GOCC_FILOPODIUM_TIP, GOCC_FILOPODIUM_TIP HP_ABNORMALITY_OF_THE_ACHILLES_TENDON, HP_ABNORMIR219A_1_3P, MIR219A_1_3P	METABOLIC_PROCESS TELET_DENSE_TUBULAR_NETWORK ME_INTERLEUKIN_6_FAMILY_SIGNALING
	HP_TYPE_II_TRANSFERRIN_ISOFORM_PROFILE, HP_TYPE_II_T GOBP_GLUTAMINE_TRANSPORT, GOBP_GLUTAMINE_TRANS GOBP_EMBRYONIC_NEUROCRANIUM_MORPHOGENESIS, GO HP_APLASIA_HYPOPLASIA_OF_THE_EPIGLOTTIS, HP_APLAS GOBP_L_SERINE_TRANSPORT, GOBP_L_SERINE_TRANSPORT GOMF_L_SERINE_TRANSMEMBRANE_TRANSPORTER_ACTIVE HP_INTERMITTENT_CLAUDICATION, HP_INTERMITTENT_CL	TRANSFERRIN_ISOFORM_PROFILE SPORT OBP_EMBRYONIC_NEUROCRANIUM_MORPHOGENESIS SIA_HYPOPLASIA_OF_THE_EPIGLOTTIS TITY, GOMF_L_SERINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY LAUDICATION
CD4_TCELL_VS_12H_ACT_TH2_DN, GSE22886_NAIVE_CD4_TCELL_VS_12H_ACT_TH2_DN	REACTOME_VEGFR2_MEDIATED_CELL_PROLIFERATION, REACTOME_INTERLEUKIN_10_SIGNALING, REACTOME_INTERLEUKIN_10_SIGNALING, REACTOME_INTERLEUKIN_10_SIGNALING, REACTOME_INTERLEUKIN_10_SIGNALING, REACTOME_INTERLEUKIN_10_SIGNALING, REACTOME_INTERLEUKIN_10_SIGNALING, REACTOME_INTERLEUKIN_10_SIGNALING, REACTOME_INTERLEUKIN_10_SIGNALING, REACTOME_INTERLEUKIN_10_SIGNALING, REACTOM, REA	ACTOME_VEGFR2_MEDIATED_CELL_PROLIFERATION ERLEUKIN_10_SIGNALING E, GOCC_PLATELET_DENSE_TUBULAR_NETWORK_MEMBRANE MIA MAL_2ND_FINGER_MORPHOLOGY HCC_TRANS_GOLGI_NETWORK_TRANSPORT_VESICLE HP_DECREASED_PROPORTION_OF_CD3_POSITIVE_T_CELLS
	GOBP_NATURAL_KILLER_CELL_MEDIATED_IMMUNE_RESPO GOBP_TRANSLATIONAL_READTHROUGH, GOBP_TRANSLAT HP_NONPROGRESSIVE, HP_NONPROGRESSIVE GOBP_NEGATIVE_REGULATION_OF_OSSIFICATION, GOBP_N REACTOME_ERYTHROPOIETIN_ACTIVATES_PHOSPHOINOSIT HP_VIRAL_INFECTION_INDUCED_RHABDOMYOLYSIS, HP_VIRAL_INFECTION_INDUCED_RHABDOMYOLYSIS, HP_VIRAL_INFECTION_OF_NEUROTROPHIN_TRK_RE	ONSE_TO_TUMOR_CELL, GOBP_NATURAL_KILLER_CELL_MEDIATED_IMMUNE_RESPONSE_TO_TUMOR_CELL FIONAL_READTHROUGH NEGATIVE_REGULATION_OF_OSSIFICATION ITIDE_3_KINASE_PI3K, REACTOME_ERYTHROPOIETIN_ACTIVATES_PHOSPHOINOSITIDE_3_KINASE_PI3K VIRAL_INFECTION_INDUCED_RHABDOMYOLYSIS ECEPTOR_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_NEUROTROPHIN_TRK_RECEPTOR_SIGNALING_PATHWAY
	GOBP_REGULATION_OF_TRAIL_ACTIVATED_APOPTOTIC_SIGNED GOBP_L_ASPARTATE_TRANSMEMBRANE_TRANSPORT, GOBIT GOBP_REGULATION_OF_POSTSYNAPTIC_CYTOSOLIC_CALCITED HP_PERSISTENCE_OF_PRIMARY_TEETH, HP_PERSISTENCE_OF_WP_IL17_SIGNALING_PATHWAY, WP_IL17_SIGNALING_PATHWAY, WP_IL17_SIGNALING_PATHWAY, SAMOLS_TARGETS_OF_KHSV_MIRNAS_UP, SAMOLS_TARGETARGETARGETARGETARGETARGETARGETARGE	GNALING_PATHWAY, GOBP_REGULATION_OF_TRAIL_ACTIVATED_APOPTOTIC_SIGNALING_PATHWAY IP_L_ASPARTATE_TRANSMEMBRANE_TRANSPORT IUM_ION_CONCENTRATION, GOBP_REGULATION_OF_POSTSYNAPTIC_CYTOSOLIC_CALCIUM_ION_CONCENTRATION IF_PRIMARY_TEETH HWAY TS_OF_KHSV_MIRNAS_UP
	HP_PERITONITIS, HP_PERITONITIS GOMF_L_GLUTAMINE_TRANSMEMBRANE_TRANSPORTER_A	ACTIVITY, GOMF_L_GLUTAMINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY ROCESS, GOBP_REGULATION_OF_CATECHOLAMINE_METABOLIC_PROCESS TOE FLUID_TRANSPORT
	XU_AKT1_TARGETS_48HR, XU_AKT1_TARGETS_48HR GOMF_PHOSPHATIDYLINOSITOL_MONOPHOSPHATE_PHOSE GOBP_LUNG_LOBE_DEVELOPMENT, GOBP_LUNG_LOBE_DEVELOPMENT, GOBP_LUNG_LOBE_DEVELOPMENT, GOBP_EXTRACELLULAR_EXOSOME_ASSEMBLY, GOBP_EXTRACELLULAR_EXOSOME_ASSEMBLY, GOBP_EXTRACELLULAR_EXOSOME_TUBULE_EPITHELIAL_CELLULAR_EXOSOME_ASSEMBLY, GOBP_EXTRACELLULAR_EXOSOME_ASSEMBLY, GOBP_EXOSOME_ASSEMBLY, GOBP_EXOSOME_ASSEMBLY, GOBP_EXOSOME_ASS	SPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_MONOPHOSPHATE_PHOSPHATASE_ACTIVITY VELOPMENT ACELLULAR_EXOSOME_ASSEMBLY LL_DIFFERENTIATION, GOBP_METANEPHRIC_NEPHRON_TUBULE_EPITHELIAL_CELL_DIFFERENTIATION IBIN IUMB, HP_DUPLICATION_OF_THE_DISTAL_PHALANX_OF_THE_THUMB
	HP_RECURRENT_ACUTE_RESPIRATORY_TRACT_INFECTION, HP_VISUAL_FIXATION_INSTABILITY, HP_VISUAL_FIXATION_ HP_ABNORMALITY_OF_VENOUS_PHYSIOLOGY, HP_ABNORM GOBP_ALDITOL_CATABOLIC_PROCESS, GOBP_ALDITOL_CAT GOMF_7S_RNA_BINDING, GOMF_7S_RNA_BINDING HP_MUSCLE_FIBER_INCLUSION_BODIES, HP_MUSCLE_FIBER_ VANDESLUIS_COMMD1_TARGETS_GROUP_2_UP, VANDESLU	, HP_RECURRENT_ACUTE_RESPIRATORY_TRACT_INFECTION _INSTABILITY MALITY_OF_VENOUS_PHYSIOLOGY TABOLIC_PROCESS R_INCLUSION_BODIES JIS_COMMD1_TARGETS_GROUP_2_UP
	GOBP_CATECHOLAMINE_UPTAKE_INVOLVED_IN_SYNAPTIC HP_LARGE_FORAMEN_MAGNUM, HP_LARGE_FORAMEN_HADDAGUM, HP_	C_TRANSMISSION, GOBP_CATECHOLAMINE_UPTAKE_INVOLVED_IN_SYNAPTIC_TRANSMISSION IAGNUM IBP_MODIFICATION_OF_POSTSYNAPTIC_STRUCTURE LIGOSACCHARIDE_METABOLIC_PROCESS GULATION_OF_EXOSOMAL_SECRETION F_CALCIUM_CHANNEL_REGULATOR_ACTIVITY
	KEGG_ONE_CARBON_POOL_BY_FOLATE, KEGG_ONE_CARBON_YAMANAKA_GLIOBLASTOMA_SURVIVAL_UP, YAMANAKA_GLIOBLASTOMA_SURVIVAL_UP, YAMANAKA_GDAZARD_UV_RESPONGOMF_NUCLEOSIDE_DIPHOSPHATASE_ACTIVITY, GOMF_NUROETH_TERT_TARGETS_DN_ROETH_TERT_TARGETS_DN_LET_7D_3P, LET_7D_3P WP_OVARIAN_INFERTILITY_GENES, WP_OVARIAN_INFERTILITY_GENES, WP_OVARIAN_INFERTILITY_UDWICZEK_TR	ON_POOL_BY_FOLATE _GLIOBLASTOMA_SURVIVAL_UP NSE_CLUSTER_G3 UCLEOSIDE_DIPHOSPHATASE_ACTIVITY LITY_GENES
	LUDWICZEK_TREATING_IRON_OVERLOAD, LUDWICZEK_TR GOBP_PANTOTHENATE_METABOLIC_PROCESS, GOBP_PANT	REATING_IRON_OVERLOAD FOTHENATE_METABOLIC_PROCESS IN_MAINTENANCE, GOBP_POSITIVE_REGULATION_OF_STEM_CELL_POPULATION_MAINTENANCE HA_COMPLEX_BINDING LGROWTH_FACTOR_BINDING ACHMENT_AND_ENTRY
	HP_NEONATAL_ASPHYXIA, HP_NEONATAL_ASPHYXIA GOMF_TAP_BINDING, GOMF_TAP_BINDING KOINUMA_COLON_CANCER_MSI_DN, KOINUMA_COLON_C GOBP_PURINE_DEOXYRIBONUCLEOSIDE_TRIPHOSPHATE_M GOCC_VCP_NPL4_UFD1_AAA_ATPASE_COMPLEX, GOCC_VC GSE14413_UNSTIM_VS_IFNB_STIM_L929_CELLS_DN, GSE14413 GNF2_PRDX2, GNF2_PRDX2	CANCER_MSI_DN METABOLIC_PROCESS, GOBP_PURINE_DEOXYRIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS CP_NPL4_UFD1_AAA_ATPASE_COMPLEX
	HP_RECURRENT_OPPORTUNISTIC_INFECTIONS, HP_RECURR HOFFMAN_CLOCK_TARGETS_UP, HOFFMAN_CLOCK_TARGE GUTIERREZ_WALDENSTROEMS_MACROGLOBULINEMIA_1_UGOBP_REGULATION_OF_SNARE_COMPLEX_ASSEMBLY, GOBT REACTOME_REACTIONS_SPECIFIC_TO_THE_COMPLEX_N_GI	RENT_OPPORTUNISTIC_INFECTIONS ETS_UP UP, GUTIERREZ_WALDENSTROEMS_MACROGLOBULINEMIA_1_UP
	GOMF_ACIDIC_AMINO_ACID_TRANSMEMBRANE_TRANSPORT HOWARD_DENDRITIC_CELL_INACT_MONOV_INFLUENZA_A FIGUEROA_AML_METHYLATION_CLUSTER_5_UP, FIGUEROA HP_ABNORMAL_FOOT_BONE_OSSIFICATION, HP_ABNORMA MIR7706, MIR7706 BAUS_TFF2_TARGETS_DN, BAUS_TFF2_TARGETS_DN GOBP_REGULATION_OF_MITOTIC_RECOMBINATION, GOBP_	AL_FOOT_BONE_OSSIFICATION _REGULATION_OF_MITOTIC_RECOMBINATION
	HP_AGENESIS_OF_CEREBELLAR_VERMIS, HP_AGENESIS_OF_THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_70PLS_VS_GOBP_TYROSINE_CATABOLIC_PROCESS, GOBP_TYROSINE_CATABOLISM, REACTOME_TYROSINITHP_HYPERGALACTOSEMIA, HP_HYPERGALACTOSEMIAGOBP_MEIOTIC_MISMATCH_REPAIR, GOBP_MEIOTIC_MISMATCH_REPAIR, GOBP_MEIOTIC_MISMATCH_RESPONSITIVE_REGULATION_OF_TRANSLATION_IN_RESPONSI	CEREBELLAR_VERMIS 1-21_30YO_0DY_UP, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_70PLS_VS_21_30YO_0DY_UP 2-30YO_0DY_UP, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_70PLS_VS_21_30YO_0DY_UP 2-30YO_0DY_UP, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_70PLS_VS_21_30YO_0DY_UP 2-30YO_0DY_UP, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_70PLS_VS_21_30YO_0DY_UP 3-40YO_0DY_UP, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_70PLS_VS_21_30YO_0DY_UP 3-40YO_0DY_UP 3-40YO_0DY_UP, THAKAR_PBMC_INACTIVATED_INFLUENZA_AGE_70PLS_VS_21_30YO_0DY_UP 3-40YO_0DY_UP 3-40YO_0DY_UP, THAKAR_PBMC_INFLUENZA_AGE_70PLS_VS_21_30YO_0DY_UP 3-40YO_0DY_UP 3-40YO_
	GOBP_REGULATION_OF_MODIFICATION_OF_POSTSYNAPTION HP_NON_MEDULLARY_THYROID_CARCINOMA, HP_NON_MEDULLARY_THYROID_CARCINOMA, HP_NON_MEDULLARY_RESPONSE_TO_FOLLICLE_STIMULATING_NEDULE_PELVIC_GIRDLE_MUSCLE_WEAKNESS, HP_PELVIC_GIRDLE_HP_ABNORMAL_TARSAL_OSSIFICATION, HP_ABNORMAL_TARSAL_OSSIFICATIONAL_PROTEIN_TARGORY_NEUROTROPHIN_BINDING, GOMF_NEUROTROPHIN_	C_ACTIN_CYTOSKELETON, GOBP_REGULATION_OF_MODIFICATION_OF_POSTSYNAPTIC_ACTIN_CYTOSKELETON MEDULLARY_THYROID_CARCINOMA _HORMONE_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_FOLLICLE_STIMULATING_HORMONE_STIMULUS DLE_MUSCLE_WEAKNESS ARSAL_OSSIFICATION RGETING_TO_MEMBRANE_SIGNAL_SEQUENCE_RECOGNITION, GOBP_SRP_DEPENDENT_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE_SIGNAL_SEQUENCE_RECOGNITI _BINDING
	JOHANSSON_BRAIN_CANCER_EARLY_VS_LATE_UP, JOHANS GOBP_PROTEIN_REPAIR, GOBP_PROTEIN_REPAIR GOBP_CYTOPLASMIC_SEQUESTERING_OF_TRANSCRIPTION_GOBP_TETRAHYDROFOLATE_METABOLIC_PROCESS, GOBP_TGOBP_MATERNAL_PLACENTA_DEVELOPMENT, GOBP_MATERNAL_HYPOSPADIAS, HP_PENOSCROTAL_HYPOSPADIAS, HP_PENOSCROTAS, HP_PENOSCROTAS, HP_PENOSCROTAS, HP_PENOSCROTAS,	SSON_BRAIN_CANCER_EARLY_VS_LATE_UP _FACTOR, GOBP_CYTOPLASMIC_SEQUESTERING_OF_TRANSCRIPTION_FACTOR TETRAHYDROFOLATE_METABOLIC_PROCESS ERNAL_PLACENTA_DEVELOPMENT POSPADIAS R5_CHEMOKINE_RECEPTOR_BINDING
	GOBP_NEGATIVE_REGULATION_OF_CYSTEINE_TYPE_ENDOING HP_CHRONIC_HEMOLYTIC_ANEMIA, HP_CHRONIC_HEMOL HP_ABNORMALITY_OF_THE_DISTAL_PHALANX_OF_THE_HA GOBP_NEGATIVE_REGULATION_OF_MACROPHAGE_CHEMOL GOBP_ACTOMYOSIN_CONTRACTILE_RING_ORGANIZATION	PEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOT LYTIC_ANEMIA ALLUX, HP_ABNORMALITY_OF_THE_DISTAL_PHALANX_OF_THE_HALLUX OTAXIS, GOBP_NEGATIVE_REGULATION_OF_MACROPHAGE_CHEMOTAXIS I, GOBP_ACTOMYOSIN_CONTRACTILE_RING_ORGANIZATION UCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY , GOBP_REGULATION_OF_PROTEIN_MONOUBIQUITINATION
	GOMF_COPPER_ION_BINDING, GOMF_COPPER_ION_BINDING GOBP_BLOOD_VESSEL_REMODELING, GOBP_BLOOD_VESSEL HP_FLATTENED_EPIPHYSIS, HP_FLATTENED_EPIPHYSIS MAEKAWA_ATF2_TARGETS, MAEKAWA_ATF2_TARGETS HP_SCANNING_SPEECH, HP_SCANNING_SPEECH REACTOME_CGMP_EFFECTS, REACTOME_CGMP_EFFECTS	IG L_REMODELING ACTIVITY, GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY
	HP_CALF_MUSCLE_PSEUDOHYPERTROPHY, HP_CALF_MUSC GOBP_NEGATIVE_REGULATION_OF_BIOMINERALIZATION, OF HP_CUTANEOUS_SYNDACTYLY_OF_TOES, HP_CUTANEOUS_ GOBP_NEGATIVE_REGULATION_OF_VIRAL_TRANSCRIPTION	CLE_PSEUDOHYPERTROPHY GOBP_NEGATIVE_REGULATION_OF_BIOMINERALIZATION _SYNDACTYLY_OF_TOES N, GOBP_NEGATIVE_REGULATION_OF_VIRAL_TRANSCRIPTION CNALING, GOBP_POSITIVE_REGULATION_OF_PROTEIN_KINASE_A_SIGNALING LL_MOLECULE_SENSOR_ACTIVITY
	HP_INSIDIOUS_ONSET, HP_INSIDIOUS_ONSET HP_SMALL_EPIPHYSES, HP_SMALL_EPIPHYSES	NINE, GOBP_PROTEIN_O_LINKED_GLYCOSYLATION_VIA_THREONINE VAY JANOSINE_DIPHOSPHATASE_ACTIVITY MBRYONIC_HEART_TUBE_DEVELOPMENT HP_ABNORMAL_PLACENTAL_MEMBRANE_MORPHOLOGY
	GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_GOBP_NEGATIVE_REGULATION_OF_TRANSFORMING_GROWGOBP_REGULATION_OF_MODIFICATION_OF_POSTSYNAPTICGOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_4_PHOSPHATEGOMF_INOSITOL_TETRAKISPHOSPHATE_PHOSPHATASE_ACREACTOME_ERYTHROPOIETIN_ACTIVATES_STAT5, REACTOMEGOBP_POSITIVE_REGULATION_OF_ACUTE_INFLAMMATORY	, MEDIATED_IMMUNE_RESPONSE_TO_TUMOR_CELL, GOBP_POSITIVE_REGULATION_OF_NATURAL_KILLER_CELL_MEDIATED_IMMUNE_RESPONSE_TO_TUMOR_CELL NTH_FACTOR_BETA1_PRODUCTION, GOBP_NEGATIVE_REGULATION_OF_TRANSFORMING_GROWTH_FACTOR_BETA1_PRODUCTION C_STRUCTURE, GOBP_REGULATION_OF_MODIFICATION_OF_POSTSYNAPTIC_STRUCTURE TASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_4_PHOSPHATASE_ACTIVITY CTIVITY, GOMF_INOSITOL_TETRAKISPHOSPHATE_PHOSPHATASE_ACTIVITY OME_ERYTHROPOIETIN_ACTIVATES_STAT5 Y_RESPONSE_TO_ANTIGENIC_STIMULUS, GOBP_POSITIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE_TO_ANTIGENIC_STIMULUS
	GOBP_POSITIVE_REGULATION_OF_ACUTE_INFLAMMATORY GOBP_REGULATION_OF_MESENCHYMAL_TO_EPITHELIAL_T GOBP_LYMPHOCYTE_MIGRATION_INTO_LYMPHOID_ORGAN GOMF_PHOSPHATIDYLINOSITOL_3_5_BISPHOSPHATE_3_PHO REACTOME_WAX_AND_PLASMALOGEN_BIOSYNTHESIS, REACTOME_ERYTHROPOIETIN_ACTIVATES_PHOSPHOLIPASE	Y_RESPONSE_TO_ANTIGENIC_STIMULUS, GOBP_POSITIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE_TO_ANTIGENIC_STIMULUS TRANSITION_INVOLVED_IN_METANEPHROS_MORPHOGENESIS, GOBP_REGULATION_OF_MESENCHYMAL_TO_EPITHELIAL_TRANSITION_INVOLVED_IN_METANEPHROS_MORPHOGEN NS, GOBP_LYMPHOCYTE_MIGRATION_INTO_LYMPHOID_ORGANS OSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_3_5_BISPHOSPHATE_3_PHOSPHATASE_ACTIVITY ACTOME_WAX_AND_PLASMALOGEN_BIOSYNTHESIS SE_C_GAMMA_PLCG, REACTOME_ERYTHROPOIETIN_ACTIVATES_PHOSPHOLIPASE_C_GAMMA_PLCG IATION, GOBP_POSITIVE_REGULATION_OF_ASTROCYTE_DIFFERENTIATION
	WP_PRADERWILLI_AND_ANGELMAN_SYNDROME, WP_PRADER GOBP_TETRAHYDROFOLATE_BIOSYNTHETIC_PROCESS, GOBTORY GOBP_CELLULAR_DETOXIFICATION_OF_ALDEHYDE, GOBP_CELLTYPE_DEPENDENT_SELECTIVITY_OF_CCK2R_SIGNATE REACTOME_CHEMOKINE_RECEPTORS_BIND_CHEMOKINES, LOPES_METHYLATED_IN_COLON_CANCER_DN, LOPES_METTORS_BOBP_POSITIVE_REGULATION_OF_B_CELL_PROLIFERATION_COLON_CANCER_DN	DERWILLI_AND_ANGELMAN_SYNDROME BP_TETRAHYDROFOLATE_BIOSYNTHETIC_PROCESS _CELLULAR_DETOXIFICATION_OF_ALDEHYDE ALING, WP_CELLTYPE_DEPENDENT_SELECTIVITY_OF_CCK2R_SIGNALING , REACTOME_CHEMOKINE_RECEPTORS_BIND_CHEMOKINES
	HP_SEPTATE_VAGINA, HP_SEPTATE_VAGINA WP_TRANSSULFURATION_PATHWAY, WP_TRANSSULFURAT GOMF_CCR_CHEMOKINE_RECEPTOR_BINDING, GOMF_CCR_ HP_CONGENITAL_ICHTHYOSIFORM_ERYTHRODERMA, HP_C GOBP_OLIGOSACCHARIDE_BIOSYNTHETIC_PROCESS, GOBP_ GOMF_RETINOID_X_RECEPTOR_BINDING, GOMF_RETINOID_ GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_DIFFE	TION_PATHWAY _CHEMOKINE_RECEPTOR_BINDING CONGENITAL_ICHTHYOSIFORM_ERYTHRODERMA _OLIGOSACCHARIDE_BIOSYNTHETIC_PROCESS _X_RECEPTOR_BINDING ERENTIATION, GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION
	GOBP_MEMBRANE_HYPERPOLARIZATION, GOBP_MEMBRANE HP_HYPOPLASIA_OF_THE_EPIGLOTTIS, HP_HYPOPLASIA_OF HP_ADULT_ONSET_SENSORINEURAL_HEARING_IMPAIRMENT HP_ABNORMALITY_OF_THE_3RD_TOE, HP_ABNORMALITY_OR RICHERT_PBMC_HIV_LIPO_5_AGE_37_48YO_STIMULATED_VEHAY_BONE_MARROW_PRE_DENDRITIC, HAY_BONE_MARROW_GOBP_PURINE_DEOXYRIBONUCLEOSIDE_MONOPHOSPHATE	NE_HYPERPOLARIZATION F_THE_EPIGLOTTIS NT, HP_ADULT_ONSET_SENSORINEURAL_HEARING_IMPAIRMENT OF_THE_3RD_TOE S_UNSTIMULATED_14W_INTERFERON_SUBSET_UP, RICHERT_PBMC_HIV_LIPO_5_AGE_37_48YO_STIMULATED_VS_UNSTIMULATED_14W_INTERFERON_SUBSET_UP OW_PRE_DENDRITIC TE_METABOLIC_PROCESS, GOBP_PURINE_DEOXYRIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS
	GOBP_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_RECEINGOBP_REGULATION_OF_GROWTH_HORMONE_RECEPTOR_SEGOBP_UBIQUITIN_DEPENDENT_ENDOCYTOSIS, GOBP_UBIQUEGOMF_TAP1_BINDING, GOMF_TAP1_BINDING HP_BUDD_CHIARI_SYNDROME, HP_BUDD_CHIARI_SYNDROME, HP_REDUCED_FACTOR_XII_ACTIVITY, HP_REDUCED_FACTOR_WIDELY_SPACEING	EPTOR_2_SIGNALING_PATHWAY, GOBP_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_RECEPTOR_2_SIGNALING_PATHWAY SIGNALING_PATHWAY, GOBP_REGULATION_OF_GROWTH_HORMONE_RECEPTOR_SIGNALING_PATHWAY UITIN_DEPENDENT_ENDOCYTOSIS OME OR_XII_ACTIVITY
	HP_LIMB_FASCICULATIONS, HP_LIMB_FASCICULATIONS GOBP_ASSEMBLY_OF_ACTOMYOSIN_APPARATUS_INVOLVED GOBP_PURINE_DEOXYRIBONUCLEOTIDE_BIOSYNTHETIC_PR GOBP_10_FORMYLTETRAHYDROFOLATE_METABOLIC_PROCE GOBP_POSITIVE_REGULATION_OF_PROTEIN_MONOUBIQUIT GOBP_REGULATION_OF_UBIQUITIN_SPECIFIC_PROTEASE_AGGOCC_CALCINEURIN_COMPLEX, GOCC_CALCINEURIN_COMPLEX	ED_IN_CYTOKINESIS, GOBP_ASSEMBLY_OF_ACTOMYOSIN_APPARATUS_INVOLVED_IN_CYTOKINESIS ROCESS, GOBP_PURINE_DEOXYRIBONUCLEOTIDE_BIOSYNTHETIC_PROCESS CESS, GOBP_10_FORMYLTETRAHYDROFOLATE_METABOLIC_PROCESS TINATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_MONOUBIQUITINATION ACTIVITY, GOBP_REGULATION_OF_UBIQUITIN_SPECIFIC_PROTEASE_ACTIVITY MPLEX
	GOMF_GLUTATHIONE_DISULFIDE_OXIDOREDUCTASE_ACTI GOMF_CYCLOHYDROLASE_ACTIVITY, GOMF_CYCLOHYDRO REACTOME_GALACTOSE_CATABOLISM, REACTOME_GALAC CLIMENT_BREAST_CANCER_COPY_NUMBER_DN, CLIMENT_ chr6p11, chr6p11 HP_LONG_THORAX, HP_LONG_THORAX CAO_BLOOD_FLUMIST_AGE_05_14YO_1DY_DN, CAO_BLOOD_	IVITY, GOMF_GLUTATHIONE_DISULFIDE_OXIDOREDUCTASE_ACTIVITY DLASE_ACTIVITY CTOSE_CATABOLISM _BREAST_CANCER_COPY_NUMBER_DN
	HP_NARROW_ILIAC_WINGS, HP_NARROW_ILIAC_WINGS GOBP_CYTOPLASMIC_SEQUESTERING_OF_NF_KAPPAB, GOBS GOCC_HFE_TRANSFERRIN_RECEPTOR_COMPLEX, GOCC_HFS GOBP_EMBRYONIC_HEART_TUBE_MORPHOGENESIS, GOBP_S chr17q23, chr17q23 chr10p15, chr10p15 GOBP_PRIMARY_ALCOHOL_BIOSYNTHETIC_PROCESS, GOBP_GOBP_CRANIOFACIAL_SUTURE_MORPHOGENESIS, GOBP_CF	BP_CYTOPLASMIC_SEQUESTERING_OF_NF_KAPPAB FE_TRANSFERRIN_RECEPTOR_COMPLEX EMBRYONIC_HEART_TUBE_MORPHOGENESIS P_PRIMARY_ALCOHOL_BIOSYNTHETIC_PROCESS
	GOBP_CRANIOFACIAL_SUTURE_MORPHOGENESIS, GOBP_CFGOBP_NEGATIVE_REGULATION_OF_MACROPHAGE_MIGRATED GOMF_COBALAMIN_BINDING, GOMF_COBALAMIN_BINDING GOBP_REGULATION_OF_NAD_P_H_OXIDASE_ACTIVITY, GOBOBP_POSITIVE_REGULATION_OF_PHOSPHOLIPID_BIOSYNTGOBP_REGULATION_OF_STEM_CELL_DIVISION, GOBP_REGULATION_GOBP_REGUL	RANIOFACIAL_SUTURE_MORPHOGENESIS LTION, GOBP_NEGATIVE_REGULATION_OF_MACROPHAGE_MIGRATION IG UBP_REGULATION_OF_NAD_P_H_OXIDASE_ACTIVITY THETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_PHOSPHOLIPID_BIOSYNTHETIC_PROCESS JLATION_OF_STEM_CELL_DIVISION CC_G_PROTEIN_BETA_GAMMA_SUBUNIT_COMPLEX
	GOBP_NEGATIVE_REGULATION_OF_PROTEIN_IMPORT, GOBE HP_OXYCEPHALY, HP_OXYCEPHALY HP_DELAYED_SOCIAL_DEVELOPMENT, HP_DELAYED_SOCIAL GOBP_POLYPHOSPHATE_METABOLIC_PROCESS, GOBP_POLYGOBP_REGULATION_OF_PHOSPHATIDYLINOSITOL_DEPHOSE GOBP_OXYGEN_METABOLIC_PROCESS, GOBP_OXYGEN_METABOLIC_PROC	BP_NEGATIVE_REGULATION_OF_PROTEIN_IMPORT AL_DEVELOPMENT YPHOSPHATE_METABOLIC_PROCESS SPHORYLATION, GOBP_REGULATION_OF_PHOSPHATIDYLINOSITOL_DEPHOSPHORYLATION TABOLIC_PROCESS S_ACTIVITY
	GOMF_INTERLEUKIN_12_RECEPTOR_BINDING, GOMF_INTERGOMF_LIPASE_BINDING, GOMF_LIPASE_BINDING CASTELLANO_HRAS_TARGETS_DN, CASTELLANO_HRAS_TAGERHOLD_RESPONSE_TO_TZD_UP, GERHOLD_RESPONSE_TOBIOCARTA_IFNG_PATHWAY, BIOCARTA_IFNG_PATHWAYHP_REMNANTS_OF_THE_HYALOID_VASCULAR_SYSTEM, HPH_ULNAR_DEVIATION_OF_THE_2ND_FINGER, HP_ULNAR_	RLEUKIN_12_RECEPTOR_BINDING ARGETS_DN O_TZD_UP P_REMNANTS_OF_THE_HYALOID_VASCULAR_SYSTEM _DEVIATION_OF_THE_2ND_FINGER
	HP_PARTIAL_DUPLICATION_OF_THE_DISTAL_PHALANGES_ HP_CEREBELLAR_HEMORRHAGE, HP_CEREBELLAR_HEMORE GOBP_PURINE_NUCLEOSIDE_TRIPHOSPHATE_CATABOLIC_F GOBP_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PART	_OF_THE_HAND, HP_PARTIAL_DUPLICATION_OF_THE_DISTAL_PHALANGES_OF_THE_HAND RRHAGE PROCESS, GOBP_PURINE_NUCLEOSIDE_TRIPHOSPHATE_CATABOLIC_PROCESS TICLE_RECEPTOR_CATABOLIC_PROCESS, GOBP_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_RECEPTOR_CATABOLIC_PROCESS TIVITY, GOBP_POSITIVE_REGULATION_OF_NAD_P_H_OXIDASE_ACTIVITY
	GOCC_MYOSIN_FILAMENT, GOCC_MYOSIN_FILAMENT chr5q22, chr5q22 GOBP_REPLICATIVE_SENESCENCE, GOBP_REPLICATIVE_SEN MATSUMIYA_BLOOD_MODIFIED_VACCINIA_ANKARA_VACCIOBP_URATE_METABOLIC_PROCESS, GOBP_URATE_METABOLIC_PROCESS, GOBP_URATE_PROCESS, GOBP_URATE_	CINE_AGE_4_6MO_VACCINATED_VS_CANDIN_PLACEBO_BCG_PRIMED_1DY_UP, MATSUMIYA_BLOOD_MODIFIED_VACCINIA_ANKARA_VACCINE_AGE_4_6MO_VACCINATED_VS_CANI OLIC_PROCESS Y_RESPONSE, REACTOME_CD163_MEDIATING_AN_ANTI_INFLAMMATORY_RESPONSE
	GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_DIFFER GOMF_G_PROTEIN_COUPLED_RECEPTOR_KINASE_ACTIVITY GOBP_RESPONSE_TO_MERCURY_ION, GOBP_RESPONSE_TO_MERCURY_ION, GOBP_RESPONSE_TO_MERCURY_ION, GOBP_CATECHOL_CONTAINING_COMPOUND_BIOSYNTHET	ERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT, GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT Y, GOMF_G_PROTEIN_COUPLED_RECEPTOR_KINASE_ACTIVITY _MERCURY_ION
	HP_ABNORMALITY_OF_THE_MIDDLE_PHALANGES_OF_THE_GOBP_POSITIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUGOBP_DOPAMINE_BIOSYNTHETIC_PROCESS, GOBP_DOPAMINE_HP_GLANDULAR_HYPOSPADIAS, HP_GLANDULAR_HYPOSPHP_LINEAR_ARRAYS_OF_MACULAR_HYPERKERATOSES_IN_HP_APLASIA_HYPOPLASIA_OF_THE_MIDDLE_PHALANGES_OF_COMESS_OMES	
	GOCC_MUTLALPHA_COMPLEX, GOCC_MUTLALPHA_COMPLEX, GOCC_MUTLALPHA_COMPLEX, GOCC_MUTLALPHA_COMPLEX, GOCC_MUTLALPHA_COMPLEX, GOCC_MUTLALPHA_COMPLEX, GOTTON_COMPLEX, GOTTON_COMPLEX	IGAS_AMPLIFICATION_HOT_SPOT_2 SIA_OF_THE_PRIMARY_TEETH SIA RANSACTIVATION _VASCULATURE_DEVELOPMENT THETIC_PROCESS, GOBP_REGULATION_OF_PHOSPHATIDYLCHOLINE_BIOSYNTHETIC_PROCESS
	GOMF_DTTP_DIPHOSPHATASE_ACTIVITY, GOMF_DTTP_DIPH HP_PERONEAL_MUSCLE_ATROPHY, HP_PERONEAL_MUSCLI GAUSSMANN_MLL_AF4_FUSION_TARGETS_B_DN, GAUSSMA	HOSPHATASE_ACTIVITY JE_ATROPHY
	GOMF_MHC_CLASS_IB_PROTEIN_BINDING, GOMF_MHC_CLAGOBP_ESTABLISHMENT_OF_LYMPHOCYTE_POLARITY, GOBP	ASS_IB_PROTEIN_BINDING P_ESTABLISHMENT_OF_LYMPHOCYTE_POLARITY
	GOMF_MHC_CLASS_IB_PROTEIN_BINDING, GOMF_MHC_CLAGOBP_ESTABLISHMENT_OF_LYMPHOCYTE_POLARITY, GOBPGOMF_INOSITOL_1_4_5_TRISPHOSPHATE_BINDING, GOMF_INGOBP_PROTEIN_O_LINKED_GLYCOSYLATION_VIA_SERINE, GOBP_PROTEIN_O_LINKED_GLYCOSYLATION_VIA_SERINE, GOBP_REGULATION_OF_TOOTH_MINERALIZATION, GOBP_FOLESLER_BRAIN_D6MIT150_QTL_CIS, CHESLER_BRAIN_D6MIT150_QTL_CIS,	ASS_IB_PROTEIN_BINDING P_ESTABLISHMENT_OF_LYMPHOCYTE_POLARITY INOSITOL_1_4_5_TRISPHOSPHATE_BINDING GOBP_PROTEIN_O_LINKED_GLYCOSYLATION_VIA_SERINE GA REGULATION_OF_TOOTH_MINERALIZATION IIT150_QTL_CIS
	GOMF_MHC_CLASS_IB_PROTEIN_BINDING, GOMF_MHC_CLAGOBP_ESTABLISHMENT_OF_LYMPHOCYTE_POLARITY, GOBPGOMF_INOSITOL_1_4_5_TRISPHOSPHATE_BINDING, GOMF_INGOBP_PROTEIN_O_LINKED_GLYCOSYLATION_VIA_SERINE, GOBP_PROTEIN_O_LINKED_GLYCOSYLATION_VIA_SERINE, GOBP_REGULATION_OF_TOOTH_MINERALIZATION, GOBP_FCCHESLER_BRAIN_D6MIT150_QTL_CIS, CHESLER_BRAIN_D6MIT150_QTL_CIS, CHESLER_BRAIN_D6MIT150_QTL_CIS	ASS_IB_PROTEIN_BINDING P_ESTABLISHMENT_OF_LYMPHOCYTE_POLARITY INOSITOL_1_4_5_TRISPHOSPHATE_BINDING GOBP_PROTEIN_O_LINKED_GLYCOSYLATION_VIA_SERINE GA REGULATION_OF_TOOTH_MINERALIZATION IIT150_QTL_CIS INE_RECEPTOR_BINDING _PROCESS, GOBP_CATECHOL_CONTAINING_COMPOUND_METABOLIC_PROCESS WAY, GOBP_LEUKEMIA_INHIBITORY_FACTOR_SIGNALING_PATHWAY HOEK_PBMC_INACTIVATED_INFLUENZA_ADULT_3DY_DN