	GSE22432_MULTIPOTENT_VS_COMMON_D GSE24634_NAIVE_CD4_TCELL_VS_DAY5_IL GSE31082_DN_VS_CD8_SP_THYMOCYTE_US GSE10239_MEMORY_VS_DAY4.5_EFF_CD8_S GSE24210_CTRL_VS_IL35_TREATED_TCONV	_IFNAB_CD8_TCELL_DN, GSE15930_NAIVE_VS_48H_IN_VITRO_STIM_IFNAB_CD8_TCELL_DN DC_PROGENITOR_UNTREATED_UP, GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_UP A_CONV_TREG_DN, GSE24634_NAIVE_CD4_TCELL_VS_DAY5_IL4_CONV_TREG_DN P, GSE31082_DN_VS_CD8_SP_THYMOCYTE_UP TCELL_DN, GSE10239_MEMORY_VS_DAY4.5_EFF_CD8_TCELL_DN V_CD4_TCELL_DN, GSE24210_CTRL_VS_IL35_TREATED_TCONV_CD4_TCELL_DN
	GSE7568_IL4_VS_IL4_AND_TGFB_TREATED GSE9650_GP33_VS_GP276_LCMV_SPECIFIC_ GSE19825_NAIVE_VS_DAY3_EFF_CD8_TCEL GSE36476_CTRL_VS_TSST_ACT_16H_MEMO GSE2770_TGFB_AND_IL4_VS_IL12_TREATEI	CROPHAGE_DN, GSE411_100MIN_VS_400MIN_IL6_STIM_MACROPHAGE_DN _MACROPHAGE_24H_DN, GSE7568_IL4_VS_IL4_AND_TGFB_TREATED_MACROPHAGE_24H_DN _EXHAUSTED_CD8_TCELL_DN, GSE9650_GP33_VS_GP276_LCMV_SPECIFIC_EXHAUSTED_CD8_TCELL_DN _LL_DN, GSE19825_NAIVE_VS_DAY3_EFF_CD8_TCELL_DN DRY_CD4_TCELL_OLD_DN, GSE36476_CTRL_VS_TSST_ACT_16H_MEMORY_CD4_TCELL_OLD_DN D_ACT_CD4_TCELL_6H_DN, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_6H_DN L4_CONVERSION_UP, GSE24634_TREG_VS_TCONV_POST_DAY5_IL4_CONVERSION_UP
	GSE32986_UNSTIM_VS_GMCSF_AND_CURE REACTOME_S_PHASE, REACTOME_S_PHASE GSE12845_IGD_NEG_BLOOD_VS_DARKZON GSE40273_GATA1_KO_VS_WT_TREG_UP, GS REACTOME_DNA_REPLICATION, REACTOR	NE_GC_TONSIL_BCELL_DN, GSE12845_IGD_NEG_BLOOD_VS_DARKZONE_GC_TONSIL_BCELL_DN SE40273_GATA1_KO_VS_WT_TREG_UP ME_DNA_REPLICATION
	GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_II GSE24210_RESTING_TREG_VS_TCONV_DN, GSE7764_IL15_NK_CELL_24H_VS_SPLENOC GSE23505_IL6_IL1_VS_IL6_IL1_TGFB_TREAT GSE32986_CURDLAN_LOWDOSE_VS_GMCS	TCELL_DN, GSE9650_EXHAUSTED_VS_MEMORY_CD8_TCELL_DN L12_72H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_72H_CD4_TCELL_DN , GSE24210_RESTING_TREG_VS_TCONV_DN EYTE_UP, GSE7764_IL15_NK_CELL_24H_VS_SPLENOCYTE_UP TED_CD4_TCELL_DN, GSE23505_IL6_IL1_VS_IL6_IL1_TGFB_TREATED_CD4_TCELL_DN GF_AND_CURDLAN_LOWDOSE_STIM_DC_DN, GSE32986_CURDLAN_LOWDOSE_VS_GMCSF_AND_CURDLAN_LOWDOSE_STIM_DC_DN GMACYTOID_PDC_UP, GSE22432_CONVENTIONAL_CDC_VS_PLASMACYTOID_PDC_UP
	GSE22432_MULTIPOTENT_PROGENITOR_VSGSE6259_FLT3L_INDUCED_33D1_POS_DC_VGSE37532_TREG_VS_TCONV_PPARG_KO_CSE19923_WT_VS_HEB_KO_DP_THYMOCYTGSE36476_CTRL_VS_TSST_ACT_16H_MEMO	Y3_EFF_CD8_TCELL_DN, GSE19825_CD24LOW_VS_IL2RA_HIGH_DAY3_EFF_CD8_TCELL_DN S_CDC_DN, GSE22432_MULTIPOTENT_PROGENITOR_VS_CDC_DN VS_BCELL_UP, GSE6259_FLT3L_INDUCED_33D1_POS_DC_VS_BCELL_UP D4_TCELL_FROM_VISCERAL_ADIPOSE_TISSUE_UP, GSE37532_TREG_VS_TCONV_PPARG_KO_CD4_TCELL_FROM_VISCERAL_ADIPOSE_TISSUE_UP TE_DN, GSE19923_WT_VS_HEB_KO_DP_THYMOCYTE_DN DRY_CD4_TCELL_YOUNG_DN, GSE36476_CTRL_VS_TSST_ACT_16H_MEMORY_CD4_TCELL_YOUNG_DN
	GSE22611_NOD2_TRANSDUCED_VS_CTRL_ GSE19772_HCMV_INFL_VS_HCMV_INF_MC FISCHER_G1_S_CELL_CYCLE, FISCHER_G1_ GSE14308_TH2_VS_NATURAL_TREG_UP, GS GSE17974_0H_VS_4H_IN_VITRO_ACT_CD4_	
	GSE37301_MULTIPOTENT_PROGENITOR_VEGOBP_PROTEIN_IMPORT, GOBP_PROTEIN_GSE17721_POLYIC_VS_CPG_24H_BMDC_DNGSE3337_CTRL_VS_16H_IFNG_IN_CD8POS_GSE27786_LSK_VS_NKTCELL_UP, GSE27786_	N, GSE17721_POLYIC_VS_CPG_24H_BMDC_DN DC_DN, GSE3337_CTRL_VS_16H_IFNG_IN_CD8POS_DC_DN
	GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_ GSE17721_LPS_VS_PAM3CSK4_24H_BMDC_I GSE4748_CTRL_VS_CYANOBACTERIUM_LP GSE19198_1H_VS_6H_IL21_TREATED_TCELI	CD8_TCELL_UP, GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD8_TCELL_UP DN, GSE17721_LPS_VS_PAM3CSK4_24H_BMDC_DN PSLIKE_STIM_DC_3H_DN, GSE4748_CTRL_VS_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_DN L_DN, GSE19198_1H_VS_6H_IL21_TREATED_TCELL_DN DCHONDRION, GOBP_PROTEIN_LOCALIZATION_TO_MITOCHONDRION
	GSE27859_MACROPHAGE_VS_CD11C_INT_CGOBP_DNA_BIOSYNTHETIC_PROCESS, GOTGOBP_IMPORT_INTO_NUCLEUS, GOBP_IMGSE8685_IL2_ACT_IL2_STARVED_VS_IL15_AGSE33424_CD161_HIGH_VS_INT_CD8_TCEL	IPORT_INTO_NUCLEUS ACT_IL2_STARVED_CD4_TCELL_DN, GSE8685_IL2_ACT_IL2_STARVED_VS_IL15_ACT_IL2_STARVED_CD4_TCELL_DN .L_UP, GSE33424_CD161_HIGH_VS_INT_CD8_TCELL_UP
	GSE4984_UNTREATED_VS_LPS_TREATED_I GSE14308_TH2_VS_INDUCED_TREG_DN, GS GSE3982_EOSINOPHIL_VS_BCELL_DN, GSE3 GSE39820_CTRL_VS_IL1B_IL6_IL23A_CD4_T	LL_DN, GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_DN DC_DN, GSE4984_UNTREATED_VS_LPS_TREATED_DC_DN SE14308_TH2_VS_INDUCED_TREG_DN
	GSE12392_CD8A_POS_VS_NEG_SPLEEN_DC KAECH_NAIVE_VS_DAY15_EFF_CD8_TCEL GSE6259_FLT3L_INDUCED_33D1_POS_DC_V GSE14908_ATOPIC_VS_NONATOPIC_PATIE GSE43863_TFH_VS_LY6C_INT_CXCR5POS_E	C_UP, GSE12392_CD8A_POS_VS_NEG_SPLEEN_DC_UP L_DN, KAECH_NAIVE_VS_DAY15_EFF_CD8_TCELL_DN VS_CD4_TCELL_DN, GSE6259_FLT3L_INDUCED_33D1_POS_DC_VS_CD4_TCELL_DN ENT_HDM_STIM_CD4_TCELL_DN, GSE14908_ATOPIC_VS_NONATOPIC_PATIENT_HDM_STIM_CD4_TCELL_DN EFFECTOR_CD4_TCELL_UP, GSE43863_TFH_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_UP CD8_TCELL_DN, GSE39152_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_DN
	SCHUHMACHER_MYC_TARGETS_UP, SCHUGSE25123_CTRL_VS_ROSIGLITAZONE_STIMGOBP_PROTEIN_TARGETING_TO_MITOCHGSE15930_STIM_VS_STIM_AND_TRICHOSTAGSE42021_CD24LO_TREG_VS_CD24LO_TCO	HYMOCYTE_UP, GSE1448_CTRL_VS_ANTI_VALPHA2_DP_THYMOCYTE_UP UHMACHER_MYC_TARGETS_UP M_MACROPHAGE_DN, GSE25123_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_DN HONDRION, GOBP_PROTEIN_TARGETING_TO_MITOCHONDRION ATINA_48H_CD8_T_CELL_UP, GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_48H_CD8_T_CELL_UP NV_THYMUS_UP, GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP _TREATED_DC_DN, GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_DN
	GSE3039_ALPHAALPHA_VS_ALPHABETA_G GSE15930_STIM_VS_STIM_AND_IL12_48H_C GSE37301_GRANULOCYTE_MONOCYTE_PE GOBP_CARBOHYDRATE_CATABOLIC_PRO GSE9650_NAIVE_VS_MEMORY_CD8_TCELL GSE36392_TYPE_2_MYELOID_VS_EOSINOPE	CD8_TCELL_UP, GSE3039_ALPHAALPHA_VS_ALPHABETA_CD8_TCELL_UP CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_IL12_48H_CD8_T_CELL_DN ROGENITOR_VS_RAG2_KO_NK_CELL_DN, GSE37301_GRANULOCYTE_MONOCYTE_PROGENITOR_VS_RAG2_KO_NK_CELL_DN PCESS, GOBP_CARBOHYDRATE_CATABOLIC_PROCESS PLON, GSE9650_NAIVE_VS_MEMORY_CD8_TCELL_DN HIL_IL25_TREATED_LUNG_UP, GSE36392_TYPE_2_MYELOID_VS_EOSINOPHIL_IL25_TREATED_LUNG_UP
	GSE13493_DP_VS_CD4INTCD8POS_THYMOOK KAECH_NAIVE_VS_MEMORY_CD8_TCELL_ GOCC_ORGANELLE_ENVELOPE_LUMEN, OF GSE10239_KLRG1INT_VS_KLRG1HIGH_EFF_ GSE1432_CTRL_VS_IFNG_1H_MICROGLIA_N	CD4_TCELL_UP, GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD4_TCELL_UP CYTE_DN, GSE13493_DP_VS_CD4INTCD8POS_THYMOCYTE_DN _DN, KAECH_NAIVE_VS_MEMORY_CD8_TCELL_DN GOCC_ORGANELLE_ENVELOPE_LUMEN _CD8_TCELL_UP, GSE10239_KLRG1INT_VS_KLRG1HIGH_EFF_CD8_TCELL_UP UP, GSE1432_CTRL_VS_IFNG_1H_MICROGLIA_UP F_UP, GSE17580_TREG_VS_TEFF_S_MANSONI_INF_UP
	GSE25088_CTRL_VS_ROSIGLITAZONE_STIM GSE339_CD4POS_VS_CD8POS_DC_IN_CULT GSE15330_LYMPHOID_MULTIPOTENT_VS_I GSE37301_COMMON_LYMPHOID_PROGEN GOCC_NUCLEAR_PORE, GOCC_NUCLEAR_	M_MACROPHAGE_UP, GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP FURE_DN, GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_DN MEGAKARYOCYTE_ERYTHROID_PROGENITOR_IKAROS_KO_DN, GSE15330_LYMPHOID_MULTIPOTENT_VS_MEGAKARYOCYTE_ERYTHROID_PROG IITOR_VS_GRAN_MONO_PROGENITOR_UP, GSE37301_COMMON_LYMPHOID_PROGENITOR_VS_GRAN_MONO_PROGENITOR_UP
	GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INGSE17721_PAM3CSK4_VS_CPG_1H_BMDC_INGSE17721_PAM3CSK4_VS_CPG_1H_BMDC_INGSE170_TEMPORAL_RESPONSE_TO_PROGEST GOBP_CELL_CYCLE_DNA_REPLICATION, ON KEGG_PYRIMIDINE_METABOLISM, KEGG_INGSE170_PROGEST RESPONSE_TO_PROGEST RESPONSE_TO_PROGET	
	GSE3720_VD1_VS_VD2_GAMMADELTA_TC GOBP_PYRUVATE_METABOLIC_PROCESS, ZNF595_TARGET_GENES, ZNF595_TARGET_ GSE42021_TCONV_PLN_VS_CD24LO_TCON GSE3691_IFN_PRODUCING_KILLER_DC_VS	ELL_UP, GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_UP GOBP_PYRUVATE_METABOLIC_PROCESS
	HALLMARK_XENOBIOTIC_METABOLISM, IN WP_DNA_REPLICATION, WP_DNA_REPLICATION, WP_DNA_REPLICATION, WP_DNA_REPLICATION, WP_DNA_REPLICATION_OFLOWINGSE25123_WT_VS_PPARG_KO_MACROPHAGROSS_HYPOXIA_VIA_HIF1A_UP, GROSS_INGOBP_POSITIVE_REGULATION_OFLOWINAL_	CATION AGE_IL4_STIM_DN, GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_DN
	SHAFFER_IRF4_TARGETS_IN_ACTIVATED_GSE45365_NK_CELL_VS_CD11B_DC_MCMV_GSE40274_CTRL_VS_HELIOS_TRANSDUCELGSE34217_MIR17_92_OVEREXPRESS_VS_WTRB_DN.V1_UP, RB_DN.V1_UP	D4_TCELL_32H_DIN, G3E43935_THIO_V3_TGFB_IL6_HH7_ACT_CD4_TCELL_32H_DIN _B_LYMPHOCYTE, SHAFFER_IRF4_TARGETS_IN_ACTIVATED_B_LYMPHOCYTE _INFECTION_UP, GSE45365_NK_CELL_VS_CD11B_DC_MCMV_INFECTION_UP D_ACTIVATED_CD4_TCELL_DN, GSE40274_CTRL_VS_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN _ACT_CD8_TCELL_DN, GSE34217_MIR17_92_OVEREXPRESS_VS_WT_ACT_CD8_TCELL_DN DSE_MAC_UP, GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_MAC_UP
	GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCOGOBP_NUCLEOTIDE_PHOSPHORYLATIONOUELLET_CULTURED_OVARIAN_CANCERGOBP_MULTI_ORGANISM_LOCALIZATIONOUP_PYRIMIDINE_METABOLISM, WP_PYRIMIDINE_METABOLISM, WP_PYRIMIDINE_METABOLISM, WP_PYRIMIDINE	
	GSE17721_PAM3CSK4_VS_GADIQUIMOD_4I HALLMARK_CHOLESTEROL_HOMEOSTAS REACTOME_ACTIVATION_OF_ATR_IN_RES SANSOM_APC_TARGETS_UP, SANSOM_AP MORI_SMALL_PRE_BII_LYMPHOCYTE_DN,	REACTOME_EXTENSION_OF_TELOMERES H_BMDC_UP, GSE17721_PAM3CSK4_VS_GADIQUIMOD_4H_BMDC_UP GIS, HALLMARK_CHOLESTEROL_HOMEOSTASIS SPONSE_TO_REPLICATION_STRESS, REACTOME_ACTIVATION_OF_ATR_IN_RESPONSE_TO_REPLICATION_STRESS PC_TARGETS_UP , MORI_SMALL_PRE_BII_LYMPHOCYTE_DN ATED_ALLOGENIC_TCELL_UP, GSE5503_PLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_UP
	REACTOME_TRNA_AMINOACYLATION, REACTOME_TRNA_DEGRADATION, KEGG_RNA_DEGRADATION, KEGG_RNA_DEGRADATION, KEGG_RNA_DEGRADATION_RESPONSE GSE13306_TREG_RA_VS_TCONV_RA_UP, GSTART REACTOME_GLUCOSE_METABOLISM, REACTOME_GLUCOSE_METABOLISM, REACTOME_REACTOME_REPLICATION_FORK, GOOCC_NUCLEAR_REPLICATION_FORK, GOOCC_NUCLEAR_REPLICA	EACTOME_TRNA_AMINOACYLATION DEGRADATION E_SYNERGY, SCHLOSSER_MYC_AND_SERUM_RESPONSE_SYNERGY SE13306_TREG_RA_VS_TCONV_RA_UP ACTOME_GLUCOSE_METABOLISM DCC_NUCLEAR_REPLICATION_FORK
	KEGG_DNA_REPLICATION, KEGG_DNA_REACTOME_ACTIVATION_OF_THE_PRE_REGOCC_HOST_CELLULAR_COMPONENT, GOCSE360_DC_VS_MAC_L_MAJOR_DN, GSE360_BROWN_MYELOID_CELL_DEVELOPMENT_	EPLICATIVE_COMPLEX, REACTOME_ACTIVATION_OF_THE_PRE_REPLICATIVE_COMPLEX OCC_HOST_CELLULAR_COMPONENT
	GOBP_NCRNA_EXPORT_FROM_NUCLEUS, REACTOME_GLYCOLYSIS, REACTOME_GLY HOSHIDA_LIVER_CANCER_SURVIVAL_DN GUO_TARGETS_OF_IRS1_AND_IRS2, GUO_T REACTOME_NS1_MEDIATED_EFFECTS_ON	GOBP_NCRNA_EXPORT_FROM_NUCLEUS YCOLYSIS I, HOSHIDA_LIVER_CANCER_SURVIVAL_DN
	GSE37534_GW1929_VS_ROSIGLITAZONE_TEGSE3720_UNSTIM_VS_LPS_STIM_VD1_GAMGSE17721_CPG_VS_GARDIQUIMOD_12H_BMREACTOME_REGULATION_OF_CHOLESTEGOBP_ENTRY_INTO_HOST, GOBP_ENTRY_	
	GOBP_MODULATION_OF_PROCESS_OF_OTMORI_PLASMA_CELL_UP, MORI_PLASMA_GOBP_NUCLEAR_TRANSCRIBED_MRNA_CDAIRKEE_CANCER_PRONE_RESPONSE_BP	CATABOLIC_PROCESS_EXONUCLEOLYTIC, GOBP_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_EXONUCLEOLYTIC PA, DAIRKEE_CANCER_PRONE_RESPONSE_BPA TION_BY_E2F6, REACTOME_TRANSCRIPTIONAL_REGULATION_BY_E2F6
	PID_RB_1PATHWAY, PID_RB_1PATHWAY ZAMORA_NOS2_TARGETS_UP, ZAMORA_N REACTOME_TELOMERE_C_STRAND_LAGO REACTOME_TRANSPORT_OF_THE_SLBP_D GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GI	TY, GOBP_REGULATION_OF_ATPASE_ACTIVITY NOS2_TARGETS_UP GING_STRAND_SYNTHESIS, REACTOME_TELOMERE_C_STRAND_LAGGING_STRAND_SYNTHESIS DEPENDANT_MATURE_MRNA, REACTOME_TRANSPORT_OF_THE_SLBP_DEPENDANT_MATURE_MRNA FP_TCONV_UP, GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GFP_TCONV_UP
	GOCC_REPLISOME, GOCC_REPLISOME ZHOU_TNF_SIGNALING_30MIN, ZHOU_TN BURTON_ADIPOGENESIS_2, BURTON_ADII XU_GH1_AUTOCRINE_TARGETS_UP, XU_G KEGG_MISMATCH_REPAIR, KEGG_MISMATCH_REPAIR, KEGG_MISMATCH_REPAIR, HUANG_GATA2_TARGETS_DN, HUANG_G KEGG_GLYCEROPHOSPHOLIPID_METABO	POGENESIS_2 GH1_AUTOCRINE_TARGETS_UP TCH_REPAIR
	POMEROY_MEDULLOBLASTOMA_DESMOI	A_TARGETS TO_G1_TRANSITION GOBP_HISTONE_H3_K4_METHYLATION NASE_BY_GLUCOKINASE_REGULATORY_PROTEIN, REACTOME_REGULATION_OF_GLUCOKINASE_BY_GLUCOKINASE_REGULATORY_PROTEIN PLASIC_VS_CLASSIC_UP, POMEROY_MEDULLOBLASTOMA_DESMOPLASIC_VS_CLASSIC_UP
	WP_DNA_MISMATCH_REPAIR, WP_DNA_M GOBP_DNA_STRAND_ELONGATION, GOBI GOBP_TETRAPYRROLE_BIOSYNTHETIC_PR WIERENGA_PML_INTERACTOME, WIEREN WP_TRANSSULFURATION_AND_ONE_CAR	P_DNA_STRAND_ELONGATION ROCESS, GOBP_TETRAPYRROLE_BIOSYNTHETIC_PROCESS
	RIZKI_TUMOR_INVASIVENESS_2D_DN, RIZ GOBP_MONOSACCHARIDE_CATABOLIC_F GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_IN WP_CHOLESTEROL_METABOLISM_INCLUI HP_ABNORMALITY_OF_DNA_REPAIR, HP_	PROCESS, GOBP_MONOSACCHARIDE_CATABOLIC_PROCESS NF_SKIN_IFNG_KO_DN, GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNG_KO_DN DES_BOTH_BLOCH_AND_KANDUTSCHRUSSELL_PATHWAYS, WP_CHOLESTEROL_METABOLISM_INCLUDES_BOTH_BLOCH_AND_KANDUTSCHRU _ABNORMALITY_OF_DNA_REPAIR
	MIR4474_3P, MIR4474_3P PID_HIF1_TFPATHWAY, PID_HIF1_TFPATH GOMF_NUCLEAR_IMPORT_SIGNAL_RECEI	PTOR_ACTIVITY, GOMF_NUCLEAR_IMPORT_SIGNAL_RECEPTOR_ACTIVITY SIS, REACTOME_LAGGING_STRAND_SYNTHESIS , HP_PERIVENTRICULAR_LEUKOMALACIA
	GOBP_POSITIVE_REGULATION_OF_HISTON WP_NUCLEOTIDE_METABOLISM, WP_NUC	ABOLIC_PROCESS, GOBP_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS
	HP_FRONTOTEMPORAL_DEMENTIA, HP_F REACTOME_MISMATCH_REPAIR, REACTO GOBP_NAD_METABOLIC_PROCESS, GOBP_ PID_HIF1A_PATHWAY, PID_HIF1A_PATHW MIR7109_3P, MIR7109_3P	OME_MISMATCH_REPAIR _NAD_METABOLIC_PROCESS
	GOBP_POSITIVE_REGULATION_OF_ATPAS HP_BILATERAL_MICROPHTHALMOS, HP_I GOBP_MODULATION_OF_PROCESS_OF_OT GOBP_MODULATION_BY_SYMBIONT_OF_I GOBP_DNA_UNWINDING_INVOLVED_IN_	DUND_METABOLIC_PROCESS, GOBP_PORPHYRIN_CONTAINING_COMPOUND_METABOLIC_PROCESS SE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_ATPASE_ACTIVITY BILATERAL_MICROPHTHALMOS THER_ORGANISM, GOBP_MODULATION_OF_PROCESS_OF_OTHER_ORGANISM HOST_CELLULAR_PROCESS, GOBP_MODULATION_BY_SYMBIONT_OF_HOST_CELLULAR_PROCESS DNA_REPLICATION, GOBP_DNA_UNWINDING_INVOLVED_IN_DNA_REPLICATION SIONT_PROCESS, GOBP_MODULATION_BY_HOST_OF_SYMBIONT_PROCESS
	BIOCARTA_MCM_PATHWAY, BIOCARTA_N MIR1251_5P, MIR1251_5P HP_FOCAL_SEGMENTAL_GLOMERULOSCI GOBP_HEME_METABOLIC_PROCESS, GOBI GOBP_NEGATIVE_REGULATION_OF_DNA	MCM_PATHWAY LEROSIS, HP_FOCAL_SEGMENTAL_GLOMERULOSCLEROSIS
	GOBP_REGULATION_OF_CHOLESTEROL_N XU_CREBBP_TARGETS_UP, XU_CREBBP_TA HP_ABNORMALITY_OF_THE_INTRINSIC_P	PATHWAY, HP_ABNORMALITY_OF_THE_INTRINSIC_PATHWAY Γ_CELLULAR_PROCESS, GOBP_MODULATION_BY_VIRUS_OF_HOST_CELLULAR_PROCESS
_48H_IN_VITRO_STIM_IL12_CD8_TCELL_DN, GSE15930_NAIVE_VS_48H_IN_VITRO_STIM_IL12_CD8_TCELL_DN	KEGG_PENTOSE_PHOSPHATE_PATHWAY,	OBP_GLUCOSE_CATABOLIC_PROCESS
	BLANCO_MELO_COVID19_SARS_COV_2_IN REACTOME_MRNA_DECAY_BY_3_TO_5_EX WP_CHOLESTEROL_BIOSYNTHESIS_PATHY REACTOME_MRNA_DECAY_BY_5_TO_3_EX	NFECTION_A594_ACE2_EXPRESSING_CELLS_RUXOLITINIB_DN, BLANCO_MELO_COVID19_SARS_COV_2_INFECTION_A594_ACE2_EXPRESSING_CELL KORIBONUCLEASE, REACTOME_MRNA_DECAY_BY_3_TO_5_EXORIBONUCLEASE WAY, WP_CHOLESTEROL_BIOSYNTHESIS_PATHWAY KORIBONUCLEASE, REACTOME_MRNA_DECAY_BY_5_TO_3_EXORIBONUCLEASE E_ACTIVITY, GOMF_INTRAMOLECULAR_TRANSFERASE_ACTIVITY
	GOBP_REGULATION_OF_CHOLESTEROL_E GOMF_CYCLIN_DEPENDENT_PROTEIN_KI WP_INITIATION_OF_TRANSCRIPTION_AN BRUNO_HEMATOPOIESIS, BRUNO_HEMAT GOBP_DNA_STRAND_ELONGATION_INVO	THE_LAGGING_STRAND, REACTOME_PROCESSIVE_SYNTHESIS_ON_THE_LAGGING_STRAND SIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_CHOLESTEROL_BIOSYNTHETIC_PROCESS INASE_ACTIVITY, GOMF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY ID_TRANSLATION_ELONGATION_AT_THE_HIV1_LTR, WP_INITIATION_OF_TRANSCRIPTION_AND_TRANSLATION_ELONGATION_AT_THE_HIV1_L TOPOIESIS DLVED_IN_DNA_REPLICATION, GOBP_DNA_STRAND_ELONGATION_INVOLVED_IN_DNA_REPLICATION Y, HP_ABNORMAL_BRONCHUS_PHYSIOLOGY
	GOBP_RECEPTOR_CLUSTERING, GOBP_REGULAR WALLACE_JAK2_TARGETS_UP, WALLACE_HP_HYPOPLASIA_OF_THE_ODONTOID_PRCAFFAREL_RESPONSE_TO_THC_UP, CAFFAGOBP_GLYCOLYTIC_PROCESS_THROUGH_	CEPTOR_CLUSTERING _JAK2_TARGETS_UP ROCESS, HP_HYPOPLASIA_OF_THE_ODONTOID_PROCESS
	GOMF_PROTEIN_TRANSPORTER_ACTIVITY HP_DIFFUSE_CEREBELLAR_ATROPHY, HP_ WP_GLYCEROPHOSPHOLIPID_BIOSYNTHE	MF_MONOSACCHARIDE_BINDING YTE_CELL, TRAVAGLINI_LUNG_CAPILLARY_AEROCYTE_CELL Y, GOMF_PROTEIN_TRANSPORTER_ACTIVITY
	HP_RETICULOCYTOSIS, HP_RETICULOCYT MARCINIAK_ER_STRESS_RESPONSE_VIA_C WP_ONE_CARBON_METABOLISM, WP_ON GOBP_NLS_BEARING_PROTEIN_IMPORT_II GOMF_UBIQUITIN_LIKE_PROTEIN_SPECIFI	CHOP, MARCINIAK_ER_STRESS_RESPONSE_VIA_CHOP IE_CARBON_METABOLISM NTO_NUCLEUS, GOBP_NLS_BEARING_PROTEIN_IMPORT_INTO_NUCLEUS IC_PROTEASE_ACTIVITY, GOMF_UBIQUITIN_LIKE_PROTEIN_SPECIFIC_PROTEASE_ACTIVITY
	BURTON_ADIPOGENESIS_10, BURTON_ADITERAO_AOX4_TARGETS_SKIN_UP, TERAO_WP_FATTY_ACID_BETA_OXIDATION, WP_IMP_PSEUDOBULBAR_SIGNS, HP_PSEUDOBULBAR_SIGNS, HP_PSEUDOBULBAR_SIGNS, SEMENZA_HIF1_MOOTHA_GLYCOLYSIS, MOOTHA_GLYCOBERENIENO TRANSFORMED BY RHOA R	_AOX4_TARGETS_SKIN_UP FATTY_ACID_BETA_OXIDATION BULBAR_SIGNS 1_TARGETS
	BRACHAT_RESPONSE_TO_METHOTREXAT FONTAINE_THYROID_TUMOR_UNCERTAIN KEGG_GLYCOLYSIS_GLUCONEOGENESIS, HP_TETRAPARESIS HP_CORONAL_CRANIOSYNOSTOSIS, HP_COROP_LIPID_DROPLET_ORGANIZATION, CORONAL_CRANIOSYNOSTOSIS, HP_COROP_LIPID_DROPLET_ORGANIZATION, CORONAL_CRANIOSYNOSTOSIS, HP_COROP_LIPID_DROPLET_ORGANIZATION, CORONAL_CRANIOSYNOSTOSIS, HP_COROP_LIPID_DROPLET_ORGANIZATION, CORONAL_CRANIOSYNOSTOSIS, HP_COROP_LIPID_DROPLET_ORGANIZATION, CORONAL_CRANIOSYNOSTOSIS, HP_CORONAL_CRANIOSYNOSTOSIS, HP_COROP_LIPID_DROPLET_ORGANIZATION, CORONAL_CRANIOSYNOSTOSIS, HP_CORONAL_CRANIOSYNOSTOSIS, HP_CORONAL_CRANIO	TE_DN, BRACHAT_RESPONSE_TO_METHOTREXATE_DN TN_MALIGNANCY_UP, FONTAINE_THYROID_TUMOR_UNCERTAIN_MALIGNANCY_UP KEGG_GLYCOLYSIS_GLUCONEOGENESIS CORONAL_CRANIOSYNOSTOSIS GOBP_LIPID_DROPLET_ORGANIZATION
	HP_ADIPOSE_TISSUE_LOSS, HP_ADIPOSE_T BIOCARTA_NPC_PATHWAY, BIOCARTA_N MIR10395_5P, MIR10395_5P ZNF202_TARGET_GENES, ZNF202_TARGET_	IPC_PATHWAY _GENES _METHYLATION, GOBP_REGULATION_OF_HISTONE_H3_K4_METHYLATION
	KEGG_FRUCTOSE_AND_MANNOSE_META HP_IMPAIRED_ANTIGEN_SPECIFIC_RESPO GOBP_PROTEIN_IMPORT_INTO_MITOCHO MIR4453, MIR4453 HP_ABNORMALITY_OF_IMMUNE_SERUM_	HEMOGLOBIN_CONCENTRATION, HP_ABNORMAL_MEAN_CORPUSCULAR_HEMOGLOBIN_CONCENTRATION ABOLISM, KEGG_FRUCTOSE_AND_MANNOSE_METABOLISM ONSE, HP_IMPAIRED_ANTIGEN_SPECIFIC_RESPONSE ONDRIAL_MATRIX, GOBP_PROTEIN_IMPORT_INTO_MITOCHONDRIAL_MATRIX _PROTEIN_PHYSIOLOGY, HP_ABNORMALITY_OF_IMMUNE_SERUM_PROTEIN_PHYSIOLOGY
	MUELLER_COMMON_TARGETS_OF_AML_GOBP_CHAPERONE_MEDIATED_PROTEINGAKL_HTLV1_INFECTION_DN, AKL_HTLV1_HP_PANHYPOGAMMAGLOBULINEMIA, HTML	P_PANHYPOGAMMAGLOBULINEMIA _BY_SIRT1, WANG_ADIPOGENIC_GENES_REPRESSED_BY_SIRT1
	MIR4538, MIR4538 KEGG_GALACTOSE_METABOLISM, KEGG_ GOBP_NEGATIVE_REGULATION_OF_ESTA LEIN_OLIGODENDROCYTE_MARKERS, LEI REACTOME_SLC_TRANSPORTER_DISORDE	BLISHMENT_OF_PROTEIN_LOCALIZATION, GOBP_NEGATIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION IN_OLIGODENDROCYTE_MARKERS ERS, REACTOME_SLC_TRANSPORTER_DISORDERS
	HP_ABNORMALITY_OF_THE_EXTRINSIC_F GOMF_PROTEIN_TRANSMEMBRANE_TRAI GOBP_ALDITOL_METABOLIC_PROCESS, GO	F_SPECIFIC_ANTIBODY, HP_DECREASED_CIRCULATING_LEVEL_OF_SPECIFIC_ANTIBODY FORMED_SELLA_TURCICA
	HP_ABNORMALITY_OF_PROTHROMBIN, H HP_RIDGED_NAIL, HP_RIDGED_NAIL GOMF_5_3_EXONUCLEASE_ACTIVITY, GOM GOBP_DOUBLE_STRAND_BREAK_REPAIR_ HP_CONCAVE_NAIL, HP_CONCAVE_NAIL	MF_5_3_EXONUCLEASE_ACTIVITY VIA_BREAK_INDUCED_REPLICATION, GOBP_DOUBLE_STRAND_BREAK_REPAIR_VIA_BREAK_INDUCED_REPLICATION
	BRACHAT_RESPONSE_TO_CISPLATIN, BRACHAT_RESPONSE_TO_CISPLATIN, BRACHAT_SMALL_EARLOID GOMF_PEPTIDE_TRANSMEMBRANE_TRANGOMF_METALLOENDOPEPTIDASE_ACTIVE	TE_UP, BRACHAT_RESPONSE_TO_METHOTREXATE_UP ACHAT_RESPONSE_TO_CISPLATIN
	GOBP_INTRA_S_DNA_DAMAGE_CHECKPO GOBP_POSITIVE_REGULATION_OF_HISTON MIR6859_3P, MIR6859_3P	RANE_TRANSPORTER_ACTIVITY, GOMF_MACROMOLECULE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY DINT, GOBP_INTRA_S_DNA_DAMAGE_CHECKPOINT NE_H3_K4_METHYLATION, GOBP_POSITIVE_REGULATION_OF_HISTONE_H3_K4_METHYLATION
	HP_LOW_SET_NIPPLES, HP_LOW	ANIZATION, GOBP_POSTSYNAPTIC_MEMBRANE_ORGANIZATION PLES TA_AMYLOID_PROTEIN, HP_DEPOSITS_IMMUNOREACTIVE_TO_BETA_AMYLOID_PROTEIN LUM, GOCC_SMOOTH_ENDOPLASMIC_RETICULUM P_3_UP, VANDESLUIS_COMMD1_TARGETS_GROUP_3_UP AGULATOR_COMPLEX
	HP_CEREBRAL_DYSMYELINATION, HP_CEGOBP_PLASMA_MEMBRANE_PHOSPHOLIIMING MIR6861_3P, MIR6861_3P GOMF_RNA_CAP_BINDING, GOMF_RNA_CGOBP_MITOTIC_DNA_REPLICATION, GOBTGOBP_DNA_REPLICATION_CHECKPOINT, GOMF_MANNOSE_BINDING, GOMF_MA	PID_SCRAMBLING, GOBP_PLASMA_MEMBRANE_PHOSPHOLIPID_SCRAMBLING CAP_BINDING P_MITOTIC_DNA_REPLICATION GOBP_DNA_REPLICATION_CHECKPOINT
	BIOCARTA_SKP2E2F_PATHWAY, BIOCARTA LASTOWSKA_COAMPLIFIED_WITH_MYCN HP_J_SHAPED_SELLA_TURCICA, HP_J_SHA HP_GENERALIZED_HYPERREFLEXIA, HP_C	A_SKP2E2F_PATHWAY I, LASTOWSKA_COAMPLIFIED_WITH_MYCN APED_SELLA_TURCICA GENERALIZED_HYPERREFLEXIA N_UP, BRACHAT_RESPONSE_TO_CAMPTOTHECIN_UP
		_GENES ON, GOBP_PROTEIN_HETEROOLIGOMERIZATION HEMOGLOBIN_CONCENTRATION, HP_INCREASED_MEAN_CORPUSCULAR_HEMOGLOBIN_CONCENTRATION OLIOSIS LACE_PROSTATE_CANCER_UP
	HP_BASAL_GANGLIA_GLIOSIS, HP_BASAL REACTOME_ABERRANT_REGULATION_OF WENG_POR_TARGETS_LIVER_UP, WENG_F GOBP_NUCLEOTIDE_SUGAR_BIOSYNTHET	,_GANGLIA_GLIOSIS F_MITOTIC_G1_S_TRANSITION_IN_CANCER_DUE_TO_RB1_DEFECTS, REACTOME_ABERRANT_REGULATION_OF_MITOTIC_G1_S_TRANSITION_IN_C POR_TARGETS_LIVER_UP TIC_PROCESS, GOBP_NUCLEOTIDE_SUGAR_BIOSYNTHETIC_PROCESS CICA, HP_ABNORMALITY_OF_THE_SELLA_TURCICA
		OSIA E_ROBIN_SEQUENCE
	YOKOE_CANCER_TESTIS_ANTIGENS, YOKO GOCC_ENDOPLASMIC_RETICULUM_EXIT_	OE_CANCER_TESTIS_ANTIGENS _SITE, GOCC_ENDOPLASMIC_RETICULUM_EXIT_SITE APPEARANCE, HP_ABNORMAL_MACROSCOPIC_URINE_APPEARANCE IS 5_PATHWAY
	HP_ABNORMAL_CSF_CARBOXYLIC_ACID_REACTOME_MUCOPOLYSACCHARIDOSES_MIR6802_3P, MIR6802_3P GOBP_ALDITOL_PHOSPHATE_METABOLIC	CURCUMIN_SULINDAC_TSA_1, MARIADASON_RESPONSE_TO_BUTYRATE_CURCUMIN_SULINDAC_TSA_1 CONCENTRATION, HP_ABNORMAL_CSF_CARBOXYLIC_ACID_CONCENTRATION REACTOME_MUCOPOLYSACCHARIDOSES C_PROCESS, GOBP_ALDITOL_PHOSPHATE_METABOLIC_PROCESS ON, GOBP_PROTEIN_HETEROTETRAMERIZATION TERAMOTO_OPN_TARGETS_CLUSTER_7
	HP_ABNORMALITY_OF_THE_ODONTOID_ HP_FRAGILE_NAILS, HP_FRAGILE_NAILS GOBP_REGULATION_OF_NUCLEAR_CELL_ GOBP_DEADENYLATION_DEPENDENT_DE ROVERSI_GLIOMA_LOH_REGIONS, ROVER chr7p13, chr7p13	PROCESS, HP_ABNORMALITY_OF_THE_ODONTOID_PROCESS _CYCLE_DNA_REPLICATION, GOBP_REGULATION_OF_NUCLEAR_CELL_CYCLE_DNA_REPLICATION ECAPPING_OF_NUCLEAR_TRANSCRIBED_MRNA, GOBP_DEADENYLATION_DEPENDENT_DECAPPING_OF_NUCLEAR_TRANSCRIBED_MRNA
	HP_DILATED_THIRD_VENTRICLE, HP_DILA GOBP_LEUCINE_METABOLIC_PROCESS, GO GOBP_NEGATIVE_REGULATION_OF_TRAN GOBP_ANTIVIRAL_INNATE_IMMUNE_RES	
	HP_MICROTIA_THIRD_DEGREE, HP_MICROGOBP_RRNA_PSEUDOURIDINE_SYNTHESIS HP_ABSENCE_OF_SUBCUTANEOUS_FAT, HHP_ABNORMAL_CAUDATE_NUCLEUS_MOGOBP_REGULATION_OF_PHOSPHOLIPID_TREACTOME_SYNTHESIS_OF_PA, REACTOME	OTIA_THIRD_DEGREE S, GOBP_RRNA_PSEUDOURIDINE_SYNTHESIS HP_ABSENCE_OF_SUBCUTANEOUS_FAT DRPHOLOGY, HP_ABNORMAL_CAUDATE_NUCLEUS_MORPHOLOGY TRANSPORT, GOBP_REGULATION_OF_PHOSPHOLIPID_TRANSPORT
	WP_CYTOSINE_METHYLATION, WP_CYTO HP_ABNORMAL_HAIR_LABORATORY_EXAMINE_RETICULAR_HYPERPIGMENTATION, HE GOBP_HISTONE_THREONINE_PHOSPHORY GOMF_HISTONE_THREONINE_KINASE_ACGROSS_HIF1A_TARGETS_DN, GROSS_HIF1A	OSINE_METHYLATION AMINATION, HP_ABNORMAL_HAIR_LABORATORY_EXAMINATION HP_RETICULAR_HYPERPIGMENTATION YLATION, GOBP_HISTONE_THREONINE_PHOSPHORYLATION CTIVITY, GOMF_HISTONE_THREONINE_KINASE_ACTIVITY A_TARGETS_DN
	HE_PTEN_TARGETS_UP, HE_PTE	TS_UP OBP_U4_SNRNA_3_END_PROCESSING ME_SYNTHESIS_OF_PG ON, REACTOME_DNA_REPLICATION_INITIATION EACTOME_DNA_DAMAGE_REVERSAL ESS, GOBP_ONE_CARBON_METABOLIC_PROCESS
	GOMF_OXIDOREDUCTASE_ACTIVITY_ACT GOMF_INTRAMOLECULAR_TRANSFERASE HP_ABNORMALITY_OF_THE_TRAGUS, HP_ GOBP_RRNA_3_END_PROCESSING, GOBP_I HP_SQUARED_ILIAC_BONES, HP_SQUARE REACTOME_ACTIVATION_OF_PUMA_AND	RRNA_3_END_PROCESSING D_ILIAC_BONES D_TRANSLOCATION_TO_MITOCHONDRIA, REACTOME_ACTIVATION_OF_PUMA_AND_TRANSLOCATION_TO_MITOCHONDRIA
	GOBP_NUCLEAR_PORE_COMPLEX_ASSEM GOMF_CYTIDYLYLTRANSFERASE_ACTIVIT CAFFAREL_RESPONSE_TO_THC_8HR_5_UF HP_GENERALIZED_DYSTONIA, HP_GENER HP_CATAPLEXY, HP_CATAPLEXY HP_CAUDATE_ATROPHY, HP_CAUDATE_A	IBLY, GOBP_NUCLEAR_PORE_COMPLEX_ASSEMBLY IY, GOMF_CYTIDYLYLTRANSFERASE_ACTIVITY P, CAFFAREL_RESPONSE_TO_THC_8HR_5_UP RALIZED_DYSTONIA ATROPHY
	HP_ABNORMAL_POSTURING, HP_ABNORM HP_DECREASED_CIRCULATING_IGG2_LEV ZHONG_PFC_C2_ASCL1_POS_ASTROCYTE, GOBP_NUCLEAR_NCRNA_SURVEILLANCE GOMF_DNA_METHYLTRANSFERASE_ACTI GOMF_FRUCTOSE_BINDING, GOMF_FRUCT	MAL_POSTURING VEL, HP_DECREASED_CIRCULATING_IGG2_LEVEL , ZHONG_PFC_C2_ASCL1_POS_ASTROCYTE E, GOBP_NUCLEAR_NCRNA_SURVEILLANCE IVITY, GOMF_DNA_METHYLTRANSFERASE_ACTIVITY
	GOBP_REGULATION_OF_DNA_METHYLATGOMF_CARBON_CARBON_LYASE_ACTIVITY MARIADASON_RESPONSE_TO_CURCUMIN WP_SREBF_AND_MIR33_IN_CHOLESTEROLGOBP_PYRIMIDINE_NUCLEOBASE_BIOSYNGOBP_PRE_REPLICATIVE_COMPLEX_ASSE	TION_DEPENDENT_HETEROCHROMATIN_ASSEMBLY, GOBP_REGULATION_OF_DNA_METHYLATION_DEPENDENT_HETEROCHROMATIN_ASSEME TY, GOMF_CARBON_CARBON_LYASE_ACTIVITY N_SULINDAC_5, MARIADASON_RESPONSE_TO_CURCUMIN_SULINDAC_5 L_AND_LIPID_HOMEOSTASIS, WP_SREBF_AND_MIR33_IN_CHOLESTEROL_AND_LIPID_HOMEOSTASIS NTHETIC_PROCESS, GOBP_PYRIMIDINE_NUCLEOBASE_BIOSYNTHETIC_PROCESS MBLY_INVOLVED_IN_CELL_CYCLE_DNA_REPLICATION, GOBP_PRE_REPLICATIVE_COMPLEX_ASSEMBLY_INVOLVED_IN_CELL_CYCLE_DNA_REPI
	HP_ELEVATED_URINARY_CARBOXYLIC_A HP_ABNORMAL_CD4_CD8_RATIO, HP_ABN GOBP_NEGATIVE_REGULATION_OF_MEM MTHFD1_TARGET_GENES, MTHFD1_TARG	ACID, HP_ELEVATED_URINARY_CARBOXYLIC_ACID NORMAL_CD4_CD8_RATIO IBRANE_POTENTIAL, GOBP_NEGATIVE_REGULATION_OF_MEMBRANE_POTENTIAL SET_GENES NG_THE_SPINAL_CORD, HP_ATROPHY_DEGENERATION_INVOLVING_THE_SPINAL_CORD P_LIPID_DROPLET_FORMATION
	HP_URETHRAL_STRICTURE, HP_URETHRA GOBP_SKELETAL_MUSCLE_ACETYLCHOLI	
	HP_ANTENATAL_ONSET, HP_ANTENATAL GOBP_GLYCEROL_3_PHOSPHATE_METABOGOBP_NEGATIVE_REGULATION_OF_MRN.	INE_GATED_CHANNEL_CLUSTERING, GOBP_SKELETAL_MUSCLE_ACETYLCHOLINE_GATED_CHANNEL_CLUSTERING NG_BEHAVIOR, GOBP_POSITIVE_REGULATION_OF_FEEDING_BEHAVIOR L_ONSET OLIC_PROCESS, GOBP_GLYCEROL_3_PHOSPHATE_METABOLIC_PROCESS A_3_END_PROCESSING, GOBP_NEGATIVE_REGULATION_OF_MRNA_3_END_PROCESSING OW_WAVES, HP_EEG_WITH_GENERALIZED_SHARP_SLOW_WAVES
	HP_ANTENATAL_ONSET, HP_ANTENATAL GOBP_GLYCEROL_3_PHOSPHATE_METABL GOBP_NEGATIVE_REGULATION_OF_MRN. HP_EEG_WITH_GENERALIZED_SHARP_SLOTE TO BE COMP_VALINE_METABOLIC_PROCESS, GOOD GOBP_MAINTENANCE_OF_DNA_METHYL GOMF_TRANSLATION_TERMINATION_FACTOR GOBP_REGULATION_OF_HELICASE_ACTIVE	NG_BEHAVIOR, GOBP_POSITIVE_REGULATION_OF_FEEDING_BEHAVIOR L_ONSET OLIC_PROCESS, GOBP_GLYCEROL_3_PHOSPHATE_METABOLIC_PROCESS A_3_END_PROCESSING, GOBP_NEGATIVE_REGULATION_OF_MRNA_3_END_PROCESSING OW_WAVES, HP_EEG_WITH_GENERALIZED_SHARP_SLOW_WAVES XP3_TARGETS_DN OBP_VALINE_METABOLIC_PROCESS ATION, GOBP_MAINTENANCE_OF_DNA_METHYLATION CTOR_ACTIVITY, GOMF_TRANSLATION_TERMINATION_FACTOR_ACTIVITY ACTIVITY, GOMF_PHOSPHATIDIC_ACID_TRANSFER_ACTIVITY VITY, GOBP_REGULATION_OF_HELICASE_ACTIVITY
	HP_ANTENATAL_ONSET, HP_ANTENATAL GOBP_GLYCEROL_3_PHOSPHATE_METABL GOBP_NEGATIVE_REGULATION_OF_MRN. HP_EEG_WITH_GENERALIZED_SHARP_SLOTED SHARP_SLOTED SHARP	NG_BEHAVIOR, GOBP_POSITIVE_REGULATION_OF_FEEDING_BEHAVIOR L_ONSET OLIC_PROCESS, GOBP_GLYCEROL_3_PHOSPHATE_METABOLIC_PROCESS A_3_END_PROCESSING, GOBP_NEGATIVE_REGULATION_OF_MRNA_3_END_PROCESSING OW_WAVES, HP_EEG_WITH_GENERALIZED_SHARP_SLOW_WAVES XP3_TARGETS_DN OBP_VALINE_METABOLIC_PROCESS ATION, GOBP_MAINTENANCE_OF_DNA_METHYLATION CTOR_ACTIVITY, GOMF_TRANSLATION_TERMINATION_FACTOR_ACTIVITY ACTIVITY, GOMF_PHOSPHATIDIC_ACID_TRANSFER_ACTIVITY VITY, GOBP_REGULATION_OF_HELICASE_ACTIVITY _ABNORMAL_ECHOCARDIOGRAM NANCE_OF_TRANSCRIPTION, GOBP_CHROMATIN_MEDIATED_MAINTENANCE_OF_TRANSCRIPTION EOSTASIS
	HP_ANTENATAL_ONSET, HP_ANTENATAL GOBP_GLYCEROL_3_PHOSPHATE_METABG GOBP_NEGATIVE_REGULATION_OF_MRN. HP_EEG_WITH_GENERALIZED_SHARP_SLG ZHENG_FOXP3_TARGETS_DN, ZHENG_FOX GOBP_VALINE_METABOLIC_PROCESS, GO GOBP_MAINTENANCE_OF_DNA_METHYL GOMF_TRANSLATION_TERMINATION_FAGOMF_PHOSPHATIDIC_ACID_TRANSFER_GOBP_REGULATION_OF_HELICASE_ACTIVENTED_CHOMEOSTASIS, WP_ZINC_HOMEOSTASIS, WP_ZINC_HOMEOSTASI	NC_BEHAVIOR, GOBP_POSITIVE_REGULATION_OF_FEEDING_BEHAVIOR L_ONSET DLIC_PROCESS, GOBP_GLYCEROL_3_PHOSPHATE_METABOLIC_PROCESS A_3_END_PROCESSING, GOBP_NEGATIVE_REGULATION_OF_MRNA_3_END_PROCESSING OW_WAVES, HP_EEG_WITH_GENERALIZED_SHARP_SLOW_WAVES XP3_TARGETS_DN IBP_VALINE_METABOLIC_PROCESS ATION, GOBP_MAINTENANCE_OF_DNA_METHYLATION CTOR_ACTIVITY, GOMF_TRANSLATION_TERMINATION_FACTOR_ACTIVITY ACTIVITY, GOMF_PHOSPHATIDIC_ACID_TRANSFER_ACTIVITY VITY, GOBP_REGULATION_OF_HELICASE_ACTIVITY _ABNORMAL_ECHOCARDIOGRAM NANCE_OF_TRANSCRIPTION, GOBP_CHROMATIN_MEDIATED_MAINTENANCE_OF_TRANSCRIPTION EOSTASIS _MIDLINE_DEFECT_OF_THE_NOSE METALS, REACTOME_METALLOTHIONEINS_BIND_METALS AL_NYSTAGMUS A_FEEDER_PATHWAY IBX_PATHWAY