	GSE39864_WT_VS_GATA3_KO_TREG_DN, GSE39864_WT_VS_GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_DGSE9960_GRAM_POS_VS_GRAM_NEG_AND_POS_SEPSIS_PIGSE339_EX_VIVO_VS_IN_CULTURE_CD4POS_DC_UP, GSE33GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOGGSE17721_CTRL_VS_LPS_6H_BMDC_UP, GSE17721_CTRL_VS_	DN, GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_DN BMC_UP, GSE9960_GRAM_POS_VS_GRAM_NEG_AND_POS_SEPSIS_PBMC_UP 39_EX_VIVO_VS_IN_CULTURE_CD4POS_DC_UP CYTE_UP, GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOCYTE_UP 5_LPS_6H_BMDC_UP
	GSE34515_CD16_NEG_MONOCYTE_VS_DC_DN, GSE34515_C GSE7831_CPG_VS_INFLUENZA_STIM_PDC_1H_UP, GSE7831_ GSE360_L_DONOVANI_VS_B_MALAYI_LOW_DOSE_DC_DN GSE38681_WT_VS_LYL1_KO_LYMPHOID_PRIMED_MULTIPO	21_0.5H_VS_4H_GARDIQUIMOD_BMDC_UP DIETIC_PROGENITORS_UP, GSE26351_UNSTIM_VS_WNT_PATHWAY_STIM_HEMATOPOIETIC_PROGENITORS_UP D16_NEG_MONOCYTE_VS_DC_DN CPG_VS_INFLUENZA_STIM_PDC_1H_UP I, GSE360_L_DONOVANI_VS_B_MALAYI_LOW_DOSE_DC_DN OTENT_PROGENITOR_UP, GSE38681_WT_VS_LYL1_KO_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_UP
		CYTE_UP ED_MACROPHAGE_DN, GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_DN JP, GSE43955_1H_VS_42H_ACT_CD4_TCELL_WITH_TGFB_IL6_UP 721_CTRL_VS_GARDIQUIMOD_6H_BMDC_UP 721_CTRL_VS_GARDIQUIMOD_8H_BMDC_UP
	HALLMARK_P53_PATHWAY, HALLMARK_P53_PATHWAY GSE33374_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIG	GH_TCELL_DN, GSE33374_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_DN N, GSE26030_TH1_VS_TH17_DAY15_POST_POLARIZATION_DN GETS_DN GSE8384_CTRL_VS_B_ABORTUS_4H_MAC_CELL_LINE_UP
	GSE17721_CTRL_VS_GARDIQUIMOD_24H_BMDC_UP, GSE17 GSE17721_POLYIC_VS_PAM3CSK4_24H_BMDC_UP, GSE1772 GSE26488_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP	7721_CTRL_VS_GARDIQUIMOD_24H_BMDC_UP 21_POLYIC_VS_PAM3CSK4_24H_BMDC_UP 2_TG_OT2_THYMOCYTE_DN, GSE26488_CTRL_VS_PEPTIDE_INJECTION_HDAC7_DELTAP_TG_OT2_THYMOCYTE_DN JP, GSE43955_1H_VS_20H_ACT_CD4_TCELL_WITH_TGFB_IL6_UP NULE
	GSE339_CD4POS_VS_CD4CD8DN_DC_UP, GSE339_CD4POS_ HP_ABNORMAL_GLIAL_CELL_MORPHOLOGY, HP_ABNOR GSE13411_NAIVE_VS_IGM_MEMORY_BCELL_DN, GSE13411_ GSE44649_NAIVE_VS_ACTIVATED_CD8_TCELL_UP, GSE44644464646464646464646464646464646464	RMAL_GLIAL_CELL_MORPHOLOGY _NAIVE_VS_IGM_MEMORY_BCELL_DN 49_NAIVE_VS_ACTIVATED_CD8_TCELL_UP GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_24H_DN VS_20H_ACT_CD4_TCELL_UP
	GSE360_CTRL_VS_B_MALAYI_HIGH_DOSE_MAC_DN, GSE3	GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_UP H_VS_16H_IFNG_IN_CD8POS_DC_DN E7831_UNSTIM_VS_INFLUENZA_STIM_PDC_4H_DN E_MEDULLA_THYMUS_UP, GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_UP B60_CTRL_VS_B_MALAYI_HIGH_DOSE_MAC_DN
	GSE2826_WT_VS_XID_BCELL_UP, GSE2826_WT_VS_XID_BCE HALLMARK_FATTY_ACID_METABOLISM, HALLMARK_FAT GSE27670_BLIMP1_VS_LMP1_TRANSDUCED_GC_BCELL_UP GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_U GSE3337_CTRL_VS_4H_IFNG_IN_CD8POS_DC_UP, GSE3337_	TTY_ACID_METABOLISM P, GSE27670_BLIMP1_VS_LMP1_TRANSDUCED_GC_BCELL_UP UP, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_UP _CTRL_VS_4H_IFNG_IN_CD8POS_DC_UP
	GSE17721_CTRL_VS_POLYIC_2H_BMDC_UP, GSE17721_CTRI GOBP_REGULATION_OF_PROTEIN_BINDING, GOBP_REGU GSE43955_10H_VS_30H_ACT_CD4_TCELL_UP, GSE43955_10H GSE360_T_GONDII_VS_B_MALAYI_LOW_DOSE_DC_DN, GSE GSE4984_GALECTIN1_VS_LPS_STIM_DC_DN, GSE4984_GALI HP_LIPID_ACCUMULATION_IN_HEPATOCYTES, HP_LIPID HP_IRRITABILITY, HP_IRRITABILITY	JLATION_OF_PROTEIN_BINDING H_VS_30H_ACT_CD4_TCELL_UP JE360_T_GONDII_VS_B_MALAYI_LOW_DOSE_DC_DN ECTIN1_VS_LPS_STIM_DC_DN
	GSE43955_10H_VS_60H_ACT_CD4_TCELL_WITH_TGFB_IL6_ HP_MALABSORPTION, HP_MALABSORPTION	RESPONSE_UP
	GOBP_TEMPERATURE_HOMEOSTASIS, GOBP_TEMPERATU WP_ANGIOPOIETIN_LIKE_PROTEIN_8_REGULATORY_PAT WP_THERMOGENESIS, WP_THERMOGENESIS	YAY VIM_NK_CELL_DN, GSE21774_CD62L_POS_CD56_DIM_VS_CD62L_NEG_CD56_DIM_NK_CELL_DN
	GOBP_POSITIVE_REGULATION_OF_CHROMATIN_ORGANI GSE40274_FOXP3_VS_FOXP3_AND_GATA1_TRANSDUCED_A HP_NEOPLASM_OF_THE_ENDOCRINE_SYSTEM, HP_NEOP GOBP_NEGATIVE_REGULATION_OF_DNA_METABOLIC_PI	T, WP_DNA_DAMAGE_RESPONSE_ONLY_ATM_DEPENDENT IZATION, GOBP_POSITIVE_REGULATION_OF_CHROMATIN_ORGANIZATION ACTIVATED_CD4_TCELL_UP, GSE40274_FOXP3_VS_FOXP3_AND_GATA1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP PLASM_OF_THE_ENDOCRINE_SYSTEM ROCESS, GOBP_NEGATIVE_REGULATION_OF_DNA_METABOLIC_PROCESS ATHWAY, GOBP_INTERFERON_GAMMA_MEDIATED_SIGNALING_PATHWAY
	GCNP_SHH_UP_LATE.V1_DN, GCNP_SHH_UP_LATE.V1_DN HP_ABNORMALITY_OF_PANCREAS_PHYSIOLOGY, HP_AB HP_ABNORMAL_SHAPE_OF_THE_OCCIPUT, HP_ABNORM HP_FETAL_ULTRASOUND_SOFT_MARKER, HP_FETAL_ULT HP_ABNORMALITY_OF_THE_ANTERIOR_FONTANELLE, H	N SNORMALITY_OF_PANCREAS_PHYSIOLOGY (AL_SHAPE_OF_THE_OCCIPUT TRASOUND_SOFT_MARKER
	WP_APOPTOSIS_MODULATION_AND_SIGNALING, WP_APOSIS_MODULATION_AND_SIGNALING, WP_APOSISE6875_TCONV_VS_TREGORY WP_EGFR_TYROSINE_KINASE_INHIBITOR_RESISTANCE, WORK HP_ABNORMALITY_OF_FONTANELLES, HP_ABNORMALITY_OF_FONTANELLES, HP_ABNORMALITY_OF_SETSON_TOTALLES, HP_ABNORMALITY_TOTALLES, HP_ABNORMALITY_OF_SETSON_TOTALLES, HP_ABNORMALITY_TOTALLES, HP_ABNORMALITY_OF_SETSON_TOTALLES, HP_ABNORMALITY_TOTALLES, HP_ABNORMALITY_TOTALL	POPTOSIS_MODULATION_AND_SIGNALING G_DN WP_EGFR_TYROSINE_KINASE_INHIBITOR_RESISTANCE TY_OF_FONTANELLES 'S_EZH2_KO_LN_TCELL_DN
	WP_SIGNALING_PATHWAYS_IN_GLIOBLASTOMA, WP_SIGNALING_PATHWAYS_IN_GLIOBLASTOMA, WP_SIGNALING_PATHWAYS_IN_GLIOBLASTOMA, WP_SIGNALING_PATHICKENED_INGOBP_ADAPTIVE_THERMOGENESIS, GOBP_ADAPTIVE_THERMOGENESIS, GOBP_ADAPTIVE_THE	NUCHAL_SKIN_FOLD HERMOGENESIS AL_THYROID_MORPHOLOGY PATHWAY EIDENBLAD_AMPLIFIED_IN_PANCREATIC_CANCER
	YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTIVE YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTIVE GSE39152_SPLEEN_CD103_NEG_VS_BRAIN_CD103_POS_METERONE_HOMOLOGOUS_DNA_PAIRING_AND_STRANT	_TRANSCRIPTION_BY_RNA_POLYMERASE_III NG, WP_AMPACTIVATED_PROTEIN_KINASE_AMPK_SIGNALING ER_1, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_1 ER_12, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_12 MORY_CD8_TCELL_DN, GSE39152_SPLEEN_CD103_NEG_VS_BRAIN_CD103_POS_MEMORY_CD8_TCELL_DN ID_EXCHANGE, REACTOME_HOMOLOGOUS_DNA_PAIRING_AND_STRAND_EXCHANGE
	HP_LYMPHOMA, HP_LYMPHOMA MIR6801_3P, MIR6801_3P KEGG_RENAL_CELL_CARCINOMA, KEGG_RENAL_CELL_C BENITEZ_GBM_PROTEASOME_INHIBITION_RESPONSE, BEITHP_CHOREOATHETOSIS, HP_CHOREOATHETOSIS HP_ABNORMALITY_OF_THE_SUPRAORBITAL_RIDGES, HP_GOBP_TELOMERE_CAPPING	NITEZ_GBM_PROTEASOME_INHIBITION_RESPONSE P_ABNORMALITY_OF_THE_SUPRAORBITAL_RIDGES
	GOCC_MLL1_2_COMPLEX, GOCC_MLL1_2_COMPLEX GOBP_POST_EMBRYONIC_DEVELOPMENT, GOBP_POST_EN DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_MA HUMMERICH_SKIN_CANCER_PROGRESSION_DN, HUMME	MBRYONIC_DEVELOPMENT ACROPHAGES, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_MACROPHAGES
	GSE14769_UNSTIM_VS_40MIN_LPS_BMDM_UP, GSE14769_UAR_01, AR_01 LANDIS_ERBB2_BREAST_PRENEOPLASTIC_DN, LANDIS_EFKEGG_GLIOMA, KEGG_GLIOMA HP_WIDE_ANTERIOR_FONTANEL, HP_WIDE_ANTERIOR_FGOBP_MULTICELLULAR_ORGANISM_GROWTH, GOBP_MU	RBB2_BREAST_PRENEOPLASTIC_DN FONTANEL
		NTO_CELL DEVELOPMENT IENT DONESIA_05_2005_H5N1_AGE_18_49YO_28DY_DN, HOWARD_NK_CELL_INACT_MONOV_INFLUENZA_A_INDONESIA_05_2005_H5N1_AGE_18_49YO_28DY_DN
	HP_NEOPLASM_OF_THE_THYROID_GLAND, HP_NEOPLASG GOBP_POSITIVE_REGULATION_OF_NEURON_DEATH, GOE GSE5589_LPS_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHA MODULE_20, MODULE_20 NADLER_OBESITY_UP, NADLER_OBESITY_UP	
	BIOCARTA_FCER1_PATHWAY, BIOCARTA_FCER1_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_BINDING, GOBP_REGULATION_OF_DOUBLE_STRAND_BREAK_REPATE	
	BIOCARTA_PYK2_PATHWAY, BIOCARTA_PYK2_PATHWAY GSE12505_WT_VS_E2_2_HET_PDC_UP, GSE12505_WT_VS_E2 HP_ABNORMAL_ASTROCYTE_MORPHOLOGY, HP_ABNOR BIOCARTA_TCR_PATHWAY, BIOCARTA_TCR_PATHWAY ZHAN_MULTIPLE_MYELOMA_MS_DN, ZHAN_MULTIPLE_ HP_CONSTRICTION_OF_PERIPHERAL_VISUAL_FIELD, HP_	2_1HET_PDC_UP RMAL_ASTROCYTE_MORPHOLOGY MYELOMA_MS_DN
	KEGG_GNRH_SIGNALING_PATHWAY, KEGG_GNRH_SIGN HP_PANCREATITIS, HP_PANCREATITIS REACTOME_REGULATION_OF_TP53_EXPRESSION_AND_D	IALING_PATHWAY DEGRADATION, REACTOME_REGULATION_OF_TP53_EXPRESSION_AND_DEGRADATION NANCE, GOBP_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE ANSPORT
	WP_SPINAL_CORD_INJURY, WP_SPINAL_CORD_INJURY BIOCARTA_INTEGRIN_PATHWAY, BIOCARTA_INTEGRIN_	CROPHAGE_UP, GSE8621_LPS_STIM_VS_LPS_PRIMED_AND_LPS_STIM_MACROPHAGE_UP PATHWAY
	RYCACNNRNNRNCAG_UNKNOWN, RYCACNNRNNRNCAG_GNF2_CD14, GNF2_CD14 GOBP_THYMUS_DEVELOPMENT, GOBP_THYMUS_DEVELOF REACTOME_SIGNALING_BY_INSULIN_RECEPTOR, REACTOME_SIGNALING_BY_INSULIN_RECEPTOR, REACTOME_SAME TARGET_GENES WP_OXIDATIVE_DAMAGE, WP_OXIDATIVE_DAMAGE WP_PDGFRBETA_PATHWAY, WP_PDGFRBETA_PATHWAY	DPMENT OME_SIGNALING_BY_INSULIN_RECEPTOR
	BLANCO_MELO_COVID19_SARS_COV_2_INFECTION_A594_ GOCC_NON_MOTILE_CILIUM, GOCC_NON_MOTILE_CILIU GOBP_POSITIVE_REGULATION_OF_LIPID_METABOLIC_PR HP_MACULAR_DEGENERATION, HP_MACULAR_DEGENE	_ACE2_EXPRESSING_CELLS_DN, BLANCO_MELO_COVID19_SARS_COV_2_INFECTION_A594_ACE2_EXPRESSING_CELLS_DN JM OCESS, GOBP_POSITIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS
G_VS_CD24HI_TCONV_THYMUS_UP, GSE42021_CD24HI_TREG_VS_CD24HI_TCONV_THYMUS_UP	WP_IL5_SIGNALING_PATHWAY, WP_IL5_SIGNALING_PAT GOBP_NEUTRAL_LIPID_BIOSYNTHETIC_PROCESS, GOBP_N HP_ABSCESS, HP_ABSCESS WP_BLADDER_CANCER, WP_BLADDER_CANCER WP_IL2_SIGNALING_PATHWAY, WP_IL2_SIGNALING_PAT BROWNE_HCMV_INFECTION_16HR_DN, BROWNE_HCMV_	NEUTRAL_LIPID_BIOSYNTHETIC_PROCESS THWAY _INFECTION_16HR_DN
	HOWARD_NK_CELL_INACT_MONOV_INFLUENZA_A_IND HOFFMANN_IMMATURE_TO_MATURE_B_LYMPHOCYTE_I	
	MISHRA_CARCINOMA_ASSOCIATED_FIBROBLAST_DN, MIGOBP_LONG_CHAIN_FATTY_ACID_IMPORT_INTO_CELL, GOBP_MACROPHAGE_ACTIVATION, GOBP_MACROPHAGGOBP_DNA_CATABOLIC_PROCESS, GOBP_DNA_CATABOLIC_KAYO_CALORIE_RESTRICTION_MUSCLE_UP, KAYO_CALORIG_LUNG_CANCER_MUTATED_SIGNIFICANTLY, DING_	ISHRA_CARCINOMA_ASSOCIATED_FIBROBLAST_DN GOBP_LONG_CHAIN_FATTY_ACID_IMPORT_INTO_CELL GE_ACTIVATION LIC_PROCESS DRIE_RESTRICTION_MUSCLE_UP
	GOBP_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLY	MENT
	GOBP_SEGMENTATION, GOBP_SEGMENTATION GOBP_FOREBRAIN_NEURON_DEVELOPMENT, GOBP_FORE REACTOME_ESTROGEN_DEPENDENT_NUCLEAR_EVENTS HP_ABNORMALITY_OF_THE_CHOROID_PLEXUS, HP_ABNORMALITY_OF_THE_CHOROID_PLEXUS, HP_ABNORMALITY_OF_THE_PLAQUES KEGG_PPAR_SIGNALING_PATHWAY, KEGG_PPAR_SIGNATION, GOBP_VACUOLAR_A	S_DOWNSTREAM_OF_ESR_MEMBRANE_SIGNALING, REACTOME_ESTROGEN_DEPENDENT_NUCLEAR_EVENTS_DOWNSTREAM_OF_ESR_MEMBRANE_SIGNALING FORMALITY_OF_THE_CHOROID_PLEXUS SLING_PATHWAY
	WP_BMP_SIGNALING_PATHWAY_IN_EYELID_DEVELOPMINGORY GOBP_REGULATION_OF_SYNAPTIC_TRANSMISSION_GLU	ENT, WP_BMP_SIGNALING_PATHWAY_IN_EYELID_DEVELOPMENT TAMATERGIC, GOBP_REGULATION_OF_SYNAPTIC_TRANSMISSION_GLUTAMATERGIC DF_OSTEOGENESIS_IMPERFECTA, WP_TYPE_I_COLLAGEN_SYNTHESIS_IN_THE_CONTEXT_OF_OSTEOGENESIS_IMPERFECTA BP_MITOCHONDRIAL_GENOME_MAINTENANCE P_POSITIVE_REGULATION_OF_PHAGOCYTOSIS
	HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_INNER_EGOBP_ASTROCYTE_DIFFERENTIATION, GOBP_ASTROCYTEGOBP_CELLULAR_RESPONSE_TO_ARSENIC_CONTAININGGOBP_NEGATIVE_REGULATION_OF_TORC1_SIGNALING, GOBP_ABNORMAL_PROPORTION_OF_CD4_POSITIVE_T_CELI	EAR, HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_INNER_EAR E_DIFFERENTIATION G_SUBSTANCE, GOBP_CELLULAR_RESPONSE_TO_ARSENIC_CONTAINING_SUBSTANCE
	REACTOME_VLDLR_INTERNALISATION_AND_DEGRADAT	GOBP_AMYLOID_BETA_CLEARANCE_BY_TRANSCYTOSIS HELPER_1_TYPE_IMMUNE_RESPONSE IDING, GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_WOUNDING TION, REACTOME_VLDLR_INTERNALISATION_AND_DEGRADATION
	HP_EEG_WITH_GENERALIZED_SLOW_ACTIVITY, HP_EEG_ HP_BROAD_ALVEOLAR_RIDGES, HP_BROAD_ALVEOLAR_ ZNF404_TARGET_GENES, ZNF404_TARGET_GENES HP_POLYPHAGIA, HP_POLYPHAGIA MIR6759_5P, MIR6759_5P WP_GLYCOLYSIS_IN_SENESCENCE, WP_GLYCOLYSIS_IN_S GOBP_T_CELL_APOPTOTIC_PROCESS, GOBP_T_CELL_APOS	RIDGES SENESCENCE
	GOBP_SOMITOGENESIS, GOBP_SOMITOGENESIS WP_PPAR_SIGNALING_PATHWAY, WP_PPAR_SIGNALING_	ENT, GOBP_CEREBELLAR_PURKINJE_CELL_LAYER_DEVELOPMENT G_PATHWAY
	HP_TEMPORAL_CORTICAL_ATROPHY, HP_TEMPORAL_COGOBP_MAINTENANCE_OF_BLOOD_BRAIN_BARRIER, GOBEGOBP_CELLULAR_RESPONSE_TO_CADMIUM_ION, GOBP_CHP_OSTEOMALACIA, HP_OSTEOMALACIA WANG_IMMORTALIZED_BY_HOXA9_AND_MEIS1_DN, WAGOBP_REGULATION_OF_ORGANIC_ACID_TRANSPORT, GOBEGORD	SP_MAINTENANCE_OF_BLOOD_BRAIN_BARRIER CELLULAR_RESPONSE_TO_CADMIUM_ION ANG_IMMORTALIZED_BY_HOXA9_AND_MEIS1_DN
	GOBP_RESPONSE_TO_UV_C, GOBP_RESPONSE_TO_UV_C GOBP_POSITIVE_REGULATION_OF_LIPID_BIOSYNTHETIC_ GOBP_REGULATION_OF_AMINO_ACID_TRANSPORT, GOB	IX, GOBP_POSITIVE_REGULATION_OF_CHOLESTEROL_EFFLUX _PROCESS, GOBP_POSITIVE_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS BP_REGULATION_OF_AMINO_ACID_TRANSPORT ANIZATION, GOBP_REGULATION_OF_ACTIN_CYTOSKELETON_REORGANIZATION
	GOBP_IRON_SULFUR_CLUSTER_ASSEMBLY, GOBP_IRON_S HP_LACK_OF_INSIGHT, HP_LACK_OF_INSIGHT HP_BROAD_PHALANX_OF_THE_TOES, HP_BROAD_PHALA HP_INCREASED_CIRCULATING_CORTISOL_LEVEL, HP_INCREGG_LIMONENE_AND_PINENE_DEGRADATION, KEGG_D BIOCARTA_SPPA_PATHWAY, BIOCARTA_SPPA_PATHWAY	SULFUR_CLUSTER_ASSEMBLY ANX_OF_THE_TOES CREASED_CIRCULATING_CORTISOL_LEVEL LIMONENE_AND_PINENE_DEGRADATION
	HP_ADRENAL_OVERACTIVITY, HP_ADRENAL_OVERACTI REACTOME_SHC1_EVENTS_IN_ERBB2_SIGNALING, REACT GOBP_SYNAPTIC_TRANSMISSION_GLUTAMATERGIC, GOE HP_THICK_HAIR, HP_THICK_HAIR HP_COLLECTIONISM, HP_COLLECTIONISM HP_RESTRICTIVE_BEHAVIOR, HP_RESTRICTIVE_BEHAVIOR	VITY FOME_SHC1_EVENTS_IN_ERBB2_SIGNALING BP_SYNAPTIC_TRANSMISSION_GLUTAMATERGIC
	GOBP_MACROPHAGE_ACTIVATION_INVOLVED_IN_IMMU HP_ABNORMAL_EATING_BEHAVIOR, HP_ABNORMAL_EA REACTOME_REGULATION_OF_TP53_ACTIVITY_THROUGH REACTOME_NUCLEAR_SIGNALING_BY_ERBB4, REACTOM	G_HORMONE_LEVEL, HP_ABNORMAL_CIRCULATING_FOLLICLE_STIMULATING_HORMONE_LEVEL UNE_RESPONSE, GOBP_MACROPHAGE_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE ATING_BEHAVIOR H_ASSOCIATION_WITH_CO_FACTORS, REACTOME_REGULATION_OF_TP53_ACTIVITY_THROUGH_ASSOCIATION_WITH_CO_FACTORS IE_NUCLEAR_SIGNALING_BY_ERBB4
	HP_URETEROCELE, HP_URETEROCELE	OBP_NEGATIVE_REGULATION_OF_ERAD_PATHWAY _T_CELL_RESPONSES_1D_INFORMATIVE_OF_LONG_TERM_RESPONSES_NEGATIVE, QI_PBMC_ZOSTAVAX_AGE_50_75YO_CORRELATED_WITH_T_CELL_RESPONSES_1D_INFORMATIVE_OF_LON ABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_PHOSPHOLIPID_METABOLIC_PROCESS
		IERNANDEZ_MITOTIC_ARREST_BY_DOCETAXEL_1_UP OCHONDRIA_PATHWAY
	GOCC_PROTON_TRANSPORTING_V_TYPE_ATPASE_V0_DCGOMF_AMIDINE_LYASE_ACTIVITY, GOMF_AMIDINE_LYAGOBP_REGULATION_OF_MACROPHAGE_ACTIVATION, GOODE ACTIVATION, GOODE A	ARIDE_ANTIBODY_CORRELATION_PROFILE_3DY_UP, LI_PBMC_MENACTRA_AGE_18_45YO_ANTI_POLYSACCHARIDE_ANTIBODY_CORRELATION_PROFILE_3DY_UP OMAIN, GOCC_PROTON_TRANSPORTING_V_TYPE_ATPASE_V0_DOMAIN ISE_ACTIVITY OBP_REGULATION_OF_MACROPHAGE_ACTIVATION
	WP_PPAR_ALPHA_PATHWAY, WP_PPAR_ALPHA_PATHWAY, HP_RETINAL_FLECKS	P_GLOMERULAR_EPITHELIUM_DEVELOPMENT LIAL_CELL_ACTIVATION OMYOCYTE, CUI_DEVELOPING_HEART_RIGHT_VENTRICULAR_CARDIOMYOCYTE
	HP_ANAL_STENOSIS, HP_ANAL_STENOSIS GCM_HMGA2, GCM_HMGA2 WP_EXTRACELLULAR_VESICLEMEDIATED_SIGNALING_IN WP_SEROTONIN_RECEPTOR_2_AND_ELKSRFGATA4_SIGN_GOBP_LONG_CHAIN_FATTY_ACID_IMPORT_ACROSS_PLA	N_RECIPIENT_CELLS, WP_EXTRACELLULAR_VESICLEMEDIATED_SIGNALING_IN_RECIPIENT_CELLS IALING, WP_SEROTONIN_RECEPTOR_2_AND_ELKSRFGATA4_SIGNALING ASMA_MEMBRANE, GOBP_LONG_CHAIN_FATTY_ACID_IMPORT_ACROSS_PLASMA_MEMBRANE IS_SPECIFIC, GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H3_K9_SPECIFIC
	REACTOME_CREB1_PHOSPHORYLATION_THROUGH_NMI HP_PERIPHERAL_VISUAL_FIELD_LOSS, HP_PERIPHERAL_V GOBP_REPLICATIVE_SENESCENCE, GOBP_REPLICATIVE_S HP_CONE_CONE_ROD_DYSTROPHY, HP_CONE_CONE_RO HP_EPIPHORA, HP_EPIPHORA PPARA_01, PPARA_01	DA_RECEPTOR_MEDIATED_ACTIVATION_OF_RAS_SIGNALING, REACTOME_CREB1_PHOSPHORYLATION_THROUGH_NMDA_RECEPTOR_MEDIATED_ACTIVATION_OF_RAS_SIGNALING VISUAL_FIELD_LOSS SENESCENCE OD_DYSTROPHY
	GOBP_REGULATION_OF_ATP_BIOSYNTHETIC_PROCESS, GZHANG_ADIPOGENESS BY_BMP7, ZHANG_ADIPOGENESS HAY_BONE_MARROW_CD34_POS_MULTILIN, HAY_BONE_HP_EEG_WITH_BURST_SUPPRESSION, HP_EEG_WITH_BURGOBP_POSITIVE_REGULATION_OF_AMYLOID_BETA_CLEAGOBP_NEGATIVE_REGULATION_OF_SODIUM_ION_TRANS	SIS_BY_BMP7 _MARROW_CD34_POS_MULTILIN RST_SUPPRESSION ARANCE, GOBP_POSITIVE_REGULATION_OF_AMYLOID_BETA_CLEARANCE SMEMBRANE_TRANSPORT, GOBP_NEGATIVE_REGULATION_OF_SODIUM_ION_TRANSMEMBRANE_TRANSPORT
	WP_TP53_NETWORK, WP_TP53_NETWORK REACTOME_INSULIN_RECEPTOR_SIGNALLING_CASCADE GOBP_GABAERGIC_NEURON_DIFFERENTIATION, GOBP_G HP_ABNORMAL_HUMERUS_MORPHOLOGY, HP_ABNORM	VITY, GOMF_LONG_CHAIN_FATTY_ACID_TRANSPORTER_ACTIVITY E, REACTOME_INSULIN_RECEPTOR_SIGNALLING_CASCADE GABAERGIC_NEURON_DIFFERENTIATION MAL_HUMERUS_MORPHOLOGY
	REACTOME_VEGFR2_MEDIATED_CELL_PROLIFERATION, I GOBP_POSITIVE_REGULATION_OF_MEMBRANE_POTENTI REACTOME_FOXO_MEDIATED_TRANSCRIPTION_OF_CELL HP_ANOMIA, HP_ANOMIA	IAL, GOBP_POSITIVE_REGULATION_OF_MEMBRANE_POTENTIAL L_CYCLE_GENES, REACTOME_FOXO_MEDIATED_TRANSCRIPTION_OF_CELL_CYCLE_GENES WTH_FACTOR_1_RECEPTOR_IGF1R, REACTOME_SIGNALING_BY_TYPE_1_INSULIN_LIKE_GROWTH_FACTOR_1_RECEPTOR_IGF1R
	GOBP_PANCREATIC_JUICE_SECRETION, GOBP_PANCREAT GOBP_CARDIAC_NEURAL_CREST_CELL_DIFFERENTIATIO	TIC_JUICE_SECRETION ON_INVOLVED_IN_HEART_DEVELOPMENT, GOBP_CARDIAC_NEURAL_CREST_CELL_DIFFERENTIATION_INVOLVED_IN_HEART_DEVELOPMENT ORATION, GOBP_REGULATION_OF_NEURON_PROJECTION_REGENERATION
	HP_RECURRENT_BACTERIAL_SKIN_INFECTIONS, HP_RECOMPLEMENT_FACTOR_I, HP_DECREASED_SERUM_COMPLEMENT_FACTOR_I, HP_DECREASED_SERUM_COMPLEMENT_FACTOR_I, HP_DECREASED_SEROTONIN_FACTOR_ACTION_ACTOR_ACTION_OF_MACROPHAGE_ACTION_RECKETS, HP_RICKETS MODULE_367, MODULE_367	CURRENT_BACTERIAL_SKIN_INFECTIONS
	GOBP_GLOMERULAR_EPITHELIAL_CELL_DIFFERENTIATION HP_WOOLLY_HAIR, HP_WOOLLY_HAIR GOBP_POSITIVE_REGULATION_OF_RECEPTOR_RECYCLING BIOCARTA_CCR3_PATHWAY, BIOCARTA_CCR3_PATHWAY	MOTAXIS, GOBP_POSITIVE_REGULATION_OF_MACROPHAGE_CHEMOTAXIS ON, GOBP_GLOMERULAR_EPITHELIAL_CELL_DIFFERENTIATION IG, GOBP_POSITIVE_REGULATION_OF_RECEPTOR_RECYCLING Y
	HP_VISUAL_AGNOSIA, HP_VISUAL_AGNOSIA JIANG_TIP30_TARGETS_DN, JIANG_TIP30_TARGETS_DN HP_BULL_S_EYE_MACULOPATHY, HP_BULL_S_EYE_MACU HP_TESTICULAR_DYSGENESIS, HP_TESTICULAR_DYSGENI	

GSE42021_TREG_PLN_VS_CD24HI_TREG_THYMUS_DN, GSE42021_TREG_PLN_VS_CD24HI_TREG_THYMUS_DN