	GOCC_LARGE_RIBOSOMAL_SUBUNIT, GOCC_LARGE_RIBOSOMA GOBP_MITOCHONDRIAL_TRANSLATION, GOBP_MITOCHONDRIA GSE34156_NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6 GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_BCELL_DN, GSE296 GSE26030_TH1_VS_TH17_RESTIMULATED_DAY15_POST_POLARIZA GOBP_TRANSLATIONAL_ELONGATION, GOBP_TRANSLATIONAL	JCTURAL_CONSTITUENT_OF_RIBOSOME DRIAL_GENE_EXPRESSION M, GOBP_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM L_SUBUNIT AL_TRANSLATION H_TREATED_MONOCYTE_UP, GSE34156_NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_UP 18_LAIV_VS_TIV_FLU_VACCINE_DAY7_BCELL_DN ATION_DN, GSE26030_TH1_VS_TH17_RESTIMULATED_DAY15_POST_POLARIZATION_DN _ELONGATION
	GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_3H_IFNG_STIM YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_13, Y GSE22886_NAIVE_TCELL_VS_DC_UP, GSE22886_NAIVE_TCELL_VS_	D_MONOCYTE_DN, GSE34156_NOD2_LIGAND_VS_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_DN I_UP, GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_3H_IFNG_STIM_UP YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_13 _DC_UP RACTION_OF_VZV_SPECIFIC_T_CELLS_PEAK_TO_28DYAT_1DY_NEGATIVE, QI_PBMC_ZOSTAVAX_AGE_50_75YO_CORRELATED_WITH_CONTRACTION_OF_VZV_SPECIFIC_T_CELLS_PEAK_TO_28DYA K4_VS_CPG_12H_BMDC_UP CL6_HIGH_TFH_VS_TFH_CD4_TCELL_DN BCELL_DN
	GSE5099_DAY3_VS_DAY7_MCSF_TREATED_MACROPHAGE_UP, GSE17721_0.5H_VS_12H_LFGSE3982_NEUTROPHIL_VS_BCELL_DN, GSE3982_NEUTROPHIL_VS_GSE3982_DC_VS_NEUTROPHIL_LPS_STIM_UP, GSE3982_DC_VS_NEUTROPHIL_LPS_STIM_UP, GSE3982_DC_VS_NEUTROPHIL_LPS_CD34POS_CD10NEG_CD62LPOSGOBP_TRANSLATIONAL_TERMINATION, GOBP_TRANSLATIONALGSE22886_IGM_MEMORY_BCELL_VS_BM_PLASMA_CELL_UP, GSE3286_IGM_MEMORY_BCELL_VS_BM_PLASMA_CELL_UP, GSE32886_IGM_MEMORY_BCELL_VS_BM_PLASMA_CELL_UP, GSE32886_IGM_MEMORY_BCELL_UP, GSE3288_IGM_MEMORY_BCELL_UP, GSE3288_IGM_MEMORY_BCELL_UP, GSE3288_IGM_MEMORY_BCELL_UP, GSE3288_IGM_MEMORY_BCELL_UP, GSE3288_IGM_MEMORY_BCELL_UP, GSE3288_IGM_MEMORY_BCELL_UP, GSE3288_IGM_MEMORY_BCELL_UP, GSE3288_IGM_MEMORY_BC	SE5099_DAY3_VS_DAY7_MCSF_TREATED_MACROPHAGE_UP PS_BMDC_UP S_BMDC_UP S_BCELL_DN SUTROPHIL_LPS_STIM_UP S_BONE_MARROW_UP, GSE35685_CD34POS_CD34POS_CD34POS_CD10NEG_CD62LPOS_BONE_MARROW_UP L_TERMINATION S22886_IGM_MEMORY_BCELL_VS_BM_PLASMA_CELL_UP 4H_TREATED_MONOCYTE_DN, GSE34156_NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_24H_TREATED_MONOCYTE_DN
	GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE27 GSE42088_2H_VS_24H_LEISHMANIA_INF_DC_UP, GSE42088_2H_VS FAN_OVARY_CL12_T_LYMPHOCYTE_NK_CELL_2, FAN_OVARY_C GSE3982_EOSINOPHIL_VS_BASOPHIL_DN, GSE3982_EOSINOPHIL_ GSE10239_NAIVE_VS_DAY4.5_EFF_CD8_TCELL_UP, GSE10239_NAIV GSE5099_CLASSICAL_M1_VS_ALTERNATIVE_M2_MACROPHAGE_ GSE17721_POLYIC_VS_CPG_4H_BMDC_UP, GSE17721_POLYIC_VS_C	TO TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_2H_UP S_24H_LEISHMANIA_INF_DC_UP S_24H_LEISHMANIA_INF_DC_UP VS_BASOPHIL_DN VE_VS_DAY4.5_EFF_CD8_TCELL_UP UP, GSE5099_CLASSICAL_M1_VS_ALTERNATIVE_M2_MACROPHAGE_UP CPG_4H_BMDC_UP GH_MONOCYTE_DN, GSE32034_UNTREATED_VS_ROSIGLIZATONE_TREATED_LY6C_HIGH_MONOCYTE_DN
	KAECH_NAIVE_VS_DAY8_EFF_CD8_TCELL_UP, KAECH_NAIVE_V GSE37605_TREG_VS_TCONV_C57BL6_FOXP3_IRES_GFP_DN, GSE376 GSE17721_0.5H_VS_8H_LPS_BMDC_UP, GSE17721_0.5H_VS_8H_LPS_ GSE22886_NAIVE_CD4_TCELL_VS_48H_ACT_TH2_UP, GSE22886_NA GSE14000_TRANSLATED_RNA_VS_MRNA_16H_LPS_DC_DN, GSE14 AIZARANI_LIVER_C3_NK_NKT_CELLS_2, AIZARANI_LIVER_C3_N GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_DN, GSE42088_UN GOBP_MITOCHONDRIAL_TRANSLATIONAL_TERMINATION, GOE GSE46606_IRF4_KO_VS_WT_UNSTIM_BCELL_UP, GSE46606_IRF4_KO	505_TREG_VS_TCONV_C57BL6_FOXP3_IRES_GFP_DN _BMDC_UP AIVE_CD4_TCELL_VS_48H_ACT_TH2_UP \$000_TRANSLATED_RNA_VS_MRNA_16H_LPS_DC_DN K_NKT_CELLS_2 NINF_VS_LEISHMANIA_INF_DC_4H_DN \$P_MITOCHONDRIAL_TRANSLATIONAL_TERMINATION D_VS_WT_UNSTIM_BCELL_UP
		IAIVE_VS_KLRGIINT_EFF_CD8_TCELL_UP TRL_VS_GARDIQUIMOD_12H_BMDC_UP OCHONDRIAL_TRANSLATION ME _VS_MEMORY_CD8_TCELL_DN _IL2_VS_IL12_TREATED_ACT_CD8_TCELL_DN DMAL_SMALL_SUBUNIT_BIOGENESIS GE_UP, GSE20484_MCSG_VS_CXCL4_MONOCYTE_DERIVED_MACROPHAGE_UP
	GSE1460_DP_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOGSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_UP, GSE17721_CTRL_VHP_ABNORMAL_NEUTROPHIL_COUNT, HP_ABNORMAL_NEUTRGSE39820_TGFBETA1_IL6_VS_TGFBETA1_IL6_IL23A_TREATED_CD4GSE369_PRE_VS_POST_IL6_INJECTION_SOCS3_KO_LIVER_UP, GSE369_PRE_VS_POST_IL6_INJECTION_SOCS3_KO_LIVER_UP, GSE369_PRE_VS_POST_INDEX_POST_I	O_UP, GSE45365_HEALTHY_VS_MCMV_INFECTION_CD8A_DC_IFNAR_KO_UP D_DN, GSE1460_DP_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_DN /S_PAM3CSK4_24H_BMDC_UP OPHIL_COUNT I_TCELL_DN, GSE39820_TGFBETA1_IL6_VS_TGFBETA1_IL6_IL23A_TREATED_CD4_TCELL_DN 369_PRE_VS_POST_IL6_INJECTION_SOCS3_KO_LIVER_UP P, GSE39110_DAY3_VS_DAY6_POST_IMMUNIZATION_CD8_TCELL_UP HEALTHY_VS_RSV_INF_INFANT_PBMC_UP
	PECE_MAMMARY_STEM_CELL_UP, PECE_MAMMARY_STEM_CELGSE21033_1H_VS_24H_POLYIC_STIM_DC_DN, GSE21033_1H_VS_24H_	L_UP H_POLYIC_STIM_DC_DN H_POLYIC_STIM_DC_DN H_POLYIC_STIM_DC_DN H_POLYIC_STIM_DC_DN H_POLYIC_STIM_DC_UP H_POLYIC_STIM_DC_UP H_POLYIC_STIM_MAC_UP H_POLYIC_UP H_PO
	GSE24574_BCL6_HIGH_VS_LOW_TFH_CD4_TCELL_UP, GSE24574_B GSE9960_HEALTHY_VS_GRAM_POS_SEPSIS_PBMC_UP, GSE9960_H GSE26488_WT_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSI HP_TRIANGULAR_SHAPED_PHALANGES_OF_THE_HAND, HP_TFHP_APLASIA_HYPOPLASIA_OF_THE_EAR, HP_APLASIA_HYPOPLHP_ANEMIA_OF_INADEQUATE_PRODUCTION, HP_ANEMIA_OF_	EALTHY_VS_GRAM_POS_SEPSIS_PBMC_UP TIVE_THYMOCYTE_DN, GSE26488_WT_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN XIANGULAR_SHAPED_PHALANGES_OF_THE_HAND ASIA_OF_THE_EAR INADEQUATE_PRODUCTION TION_UP, GSE26030_TH1_VS_TH17_RESTIMULATED_DAY5_POST_POLARIZATION_UP CHERER_PBMC_APSV_WETVAX_AGE_18_32YO_5_TO_7DY_DN
		DD_DN, GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_ADULT_BLOOD_DN NG_UP, GSE36392_TYPE_2_MYELOID_VS_EOSINOPHIL_IL25_TREATED_LUNG_UP 14_VS_CPG_8H_BMDC_UP 121670_IL6_VS_TGFB_AND_IL6_TREATED_CD4_TCELL_UP 122_EOSINOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN IC_MEYLOGENOUS_LEUKEMIA_DN _DC_UP
	GSE39820_TGFBETA3_IL6_VS_TGFBETA3_IL6_IL23A_TREATED_CD4	3920_IFNB_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP
	GSE29618_BCELL_VS_MONOCYTE_UP, GSE29618_BCELL_VS_MONOGOMF_RRNA_BINDING, GOMF_RRNA_BINDING LINDGREN_BLADDER_CANCER_CLUSTER_1_UP, LINDGREN_BLAGSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_DN, GSE7460_TREG_GOCC_MITOCHONDRIAL_LARGE_RIBOSOMAL_SUBUNIT, GOCC_GNF2_TPT1, GNF2_TPT1 GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_TREATED_MACCHOPSETED_NATE	DDER_CANCER_CLUSTER_1_UP _VS_TCONV_ACT_WITH_TGFB_DN
	HP_RENAL_AGENESIS, HP_RENAL_AGENESIS TRAVAGLINI_LUNG_CLUB_CELL, TRAVAGLINI_LUNG_CLUB_CEL GSE25087_TREG_VS_TCONV_ADULT_DN, GSE25087_TREG_VS_TCO AIZARANI_LIVER_C34_MHC_II_POS_B_CELLS, AIZARANI_LIVER_HP_LOW_ANTERIOR_HAIRLINE, HP_LOW_ANTERIOR_HAIRLINE GSE37416_12H_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GS GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_DN, HP_PALLOR, HP_PALLOR GOBP_MATURATION_OF_SSU_RRNA, GOBP_MATURATION_OF_S	ONV_ADULT_DN C34_MHC_II_POS_B_CELLS GE37416_12H_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_DN GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_DN
	GSE12505_WT_VS_E2_2_HET_PDC_DN, GSE12505_WT_VS_E2_2_HET HP_APLASIA_HYPOPLASIA_OF_THE_THUMB, HP_APLASIA_HYPOPLASIA_OF_THE_THUMB, HP_APLASIA_HYPOPLASIA_HYPOPLASIA_HYPOPLASIA_HYPOPLASIA_HYPOPLASIA_HYPOPLASIA_IN_IL6_TREATED_CD4_TCELIC HP_NEOPLASM_OF_THE_LARGE_INTESTINE, HP_NEOPLASM_OF_THE_LARGE_INTESTINE, HP_NEOPLASM_OF_THE_LARGE_INTESTINE, HP_NEOPLASM_OF_THE_LARGE_INTESTINE, HP_NEOPLASM_OF_THE_THEOPLASM_OF_T	DPLASIA_OF_THE_THUMB L_DN, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_TREATED_CD4_TCELL_DN _THE_LARGE_INTESTINE
	GSE20727_CTRL_VS_DNFB_ALLERGEN_TREATED_DC_DN, GSE20726 GSE9006_TYPE_1_DIABETES_AT_DX_VS_4MONTH_POST_DX_PBMC HP_SHORT_THUMB, HP_SHORT_THUMB	LLS_FIBRINOGEN_POS_S3, LAKE_ADULT_KIDNEY_C6_PROXIMAL_TUBULE_EPITHELIAL_CELLS_FIBRINOGEN_POS_S3 27_CTRL_VS_DNFB_ALLERGEN_TREATED_DC_DN 22_DN, GSE9006_TYPE_1_DIABETES_AT_DX_VS_4MONTH_POST_DX_PBMC_DN AL_CELL_1, FAN_OVARY_CL2_PUTATIVE_EARLY_ATRETIC_FOLLICLE_THECAL_CELL_1 PAM3CSK4_0.5H_BMDC_UP
	GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_STIM_UP, GSE2 GRUETZMANN_PANCREATIC_CANCER_DN, GRUETZMANN_PAN GSE13485_DAY3_VS_DAY7_YF17D_VACCINE_PBMC_UP, GSE13485_ HP_DEVELOPMENTAL_CATARACT, HP_DEVELOPMENTAL_CATA HP_NEOPLASM_OF_THE_COLON, HP_NEOPLASM_OF_THE_COLO GSE41176_WT_VS_TAK1_KO_UNSTIM_BCELL_UP, GSE41176_WT_VS_ HP_MACROCYTIC_ANEMIA, HP_MACROCYTIC_ANEMIA GOCC_POLYSOMAL_RIBOSOME, GOCC_POLYSOMAL_RIBOSOME GSE24671_CTRL_VS_BAKIMULC_INFECTED_MOUSE_SPLENOCYTE	NCREATIC_CANCER_DN DAY3_VS_DAY7_YF17D_VACCINE_PBMC_UP ARACT DN S_TAK1_KO_UNSTIM_BCELL_UP
	BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_CIS, GSE17721_CPG_VS_GARDIQUIMOD_0.5H_BMDC_UP, GSE17721_CPG GSE22886_IGM_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_UP, GOBP_MATURATION_OF_LSU_RRNA, GOBP_MATURATION_OF_L	MEOSTASIS JLL_UP, GSE22611_MUTANT_NOD2_VS_CTRL_TRANSDUCED_HEK293T_CELL_UP BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_CIS G_VS_GARDIQUIMOD_0.5H_BMDC_UP GSE22886_IGM_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_UP
	GOBP_NEGATIVE_REGULATION_OF_UBIQUITIN_DEPENDENT_PE	THYMIC_DC_VS_THYMIC_MACROPHAGE_DN AGE_DN, GSE25123_CTRL_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DN ROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS TCELL_UP, GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_0.5H_ACT_CD4_TCELL_UP F_THE_SKELETAL_SYSTEM F_THUMB_PHALANX UNSTIM_VS_IFNB_STIM_NIH3T3_CELLS_UP
	HP_ABNORMAL_SCAPULA_MORPHOLOGY, HP_ABNORMAL_SCAPULA_HP_TRIPHALANGEAL_THUMB, HP_TRIPHALANGEAL_THUMB GSE6269_FLU_VS_STREP_PNEUMO_INF_PBMC_DN, GSE6269_FLU_GSE10239_MEMORY_VS_KLRG1INT_EFF_CD8_TCELL_UP, GSE10239_BHATI_G2M_ARREST_BY_2METHOXYESTRADIOL_DN, BHATI_G2M_HP_NONIMMUNE_HYDROPS_FETALIS, HP_NONIMMUNE_HYDROPS_TRAVAGLINI_LUNG_AIRWAY_SMOOTH_MUSCLE_CELL, TRAVAGLINI_LUNG_AIRWAY_SMOOTH_MUSCLE_CELL, TRAVAGLINI_LUNG_AIRWAY_SMOOTH_MUSCLE_CELL, TRAVAGLINI_LUNG_AIRWAY_SMOOTH_MUSCLE_CELL, TRAVAGLINI_LUNG_AIRWAY_SMOOTH_MUSCLE_CELL, TRAVAGLINI_LUNG_AIRWAY_SMOOTH_MUSCLE_CELL, TRAVAGLINI_LUNG_AIRWAY_SMOOTH_MUSCLE_CELL, TRAVAGLINI_LUNG_AIRWAY_SMOOTH_SMUSCLE_CELL, TRAVAGLINI_STATION_OF_AORTA_STA	VS_STREP_PNEUMO_INF_PBMC_DN D_MEMORY_VS_KLRG1INT_EFF_CD8_TCELL_UP M_ARREST_BY_2METHOXYESTRADIOL_DN DPS_FETALIS GLINI_LUNG_AIRWAY_SMOOTH_MUSCLE_CELL
		N_TREATED_DC_UP, GSE20727_DNFB_ALLERGEN_VS_ROS_INH_AND_DNFB_ALLERGEN_TREATED_DC_UP BY_SMALL_PROTEIN_CONJUGATION_OR_REMOVAL, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_CONJUGATION_OR_REMOVAL
	HP_ABNORMAL_ERYTHROCYTE_ENZYME_LEVEL, HP_ABNORMAL HP_APLASIA_OF_THE_FINGERS, HP_APLASIA_OF_THE_FINGERS PTPRA_TARGET_GENES, PTPRA_TARGET_GENES HP_SPRENGEL_ANOMALY, HP_SPRENGEL_ANOMALY HP_MALIGNANT_GENITOURINARY_TRACT_TUMOR, HP_MALIGNANT_GENITOURINARY_TRACT_TUMOR, HP_MALIGNANT_GENITOURINARY_TRACT_TUMOR, HP_MALIGNANT_GENITOURINARY_TRACT_TUMOR, HP_MALIGNANT_ASSEMBLY, GOBP_RIBOSOMAL_LARGE_SUBUNIT_ASSEMBLY, GOBP_RIBOSOM HP_ATRESIA_OF_THE_EXTERNAL_AUDITORY_CANAL, HP_ATRE	NANT_GENITOURINARY_TRACT_TUMOR MAL_LARGE_SUBUNIT_ASSEMBLY
	HP_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_HAND, HE HP_ELEVATED_RED_CELL_ADENOSINE_DEAMINASE_LEVEL, HP_HP_ADENOCARCINOMA_OF_THE_INTESTINES, HP_ADENOCARCINOMAL_NUMBER_OF_ERYTHROID_PRECURSORS, HP_ABHP_ABSENT_THUMB, HP_ABSENT_THUMB HP_ABNORMAL_MEAN_CORPUSCULAR_VOLUME, HP_ABNORMHONMAL_DOCETAXEL_RESISTANCE, HONMAL_DOCETAXEL_RESISTANCE, HONM	P_ABNORMALITY_OF_THE_MUSCULATURE_OF_THE_HAND _ELEVATED_RED_CELL_ADENOSINE_DEAMINASE_LEVEL CINOMA_OF_THE_INTESTINES BNORMAL_NUMBER_OF_ERYTHROID_PRECURSORS AL_MEAN_CORPUSCULAR_VOLUME STANCE
	JAZAG_TGFB1_SIGNALING_VIA_SMAD4_UP, JAZAG_TGFB1_SIGNALING_VIA_SMAD4_UP, JAZAG_TGFB1_SIGNALING_COBP_REGULATION_OF_MITOCHONDRIAL_GENE_EXPRESSION, BUFFA_HYPOXIA_METAGENE GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_UP, GSE17721_POLYIC_ADENOCARCINOMA_OF_THE_LARGE_INTESTINE, HP_ADENOCARCINOMA_OF_THE_LARGE_INTESTINE, HP_ADENOCARCINIC_CD4_MEMORY_EFFECTOR_T_CELL, TRAVAC	ALING_VIA_SMAD4_UP GOBP_REGULATION_OF_MITOCHONDRIAL_GENE_EXPRESSION YIC_VS_PAM3CSK4_0.5H_BMDC_UP OCARCINOMA_OF_THE_LARGE_INTESTINE GLINI_LUNG_CD4_MEMORY_EFFECTOR_T_CELL LL_DN, GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_DN
	GSE13547_CTRL_VS_ANTI_IGM_STIM_BCELL_2H_DN, GSE13547_CTREACTOME_TNF_SIGNALING, REACTOME_TNF_SIGNALING HP_PERSISTENCE_OF_HEMOGLOBIN_F, HP_PERSISTENCE_OF_HE REACTOME_NEGATIVE_REGULATORS_OF_DDX58_IFIH1_SIGNAL NIKOLSKY_BREAST_CANCER_16P13_AMPLICON, NIKOLSKY_BREACTOMA, HP_OSTEOSARCOMA	EMOGLOBIN_F ING, REACTOME_NEGATIVE_REGULATORS_OF_DDX58_IFIH1_SIGNALING AST_CANCER_16P13_AMPLICON IID_SORTED_BCELL_DN, GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4MID_SORTED_BCELL_DN CIN_SENSITIVE_GENES
	GOBP_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGN NUTT_GBM_VS_AO_GLIOMA_INTT_GBM_VS_AO_GLIOMA_INTT_GBM_VS_AO_GLIOMA_INTT_GBM_VS_AO_GLIOMA_INTT_GBM_VS_AO_GLIOMA_INTT_GBM_VS_AO_GLIOMA_INTT_GBM_VS_AO_GLIOMA_INTT_GBM_VS_AO_GLIOMA_INTT_GBM_VS_AO_GLIOMA_INTT_GOCC_HOUGH_ENDOPLASMIC_RETICULUM_MEMBRANE, GOCC_WANG_PROSTATE_CANCER_ANDROGEN_INDEPENDENT, WANG_HP_DEVELOPMENTAL_GLAUCOMA, HP_DEVELOPMENTAL_GLAUCOMA_INTT_GLAUCOMA_INTT_AL_GLAUCOMA_INTT_AL_GLAUCOMA_INT_AL_GL	MITOCHONDRIAL_SMALL_RIBOSOMAL_SUBUNIT PARTIAL_DUPLICATION_OF_THE_PHALANX_OF_HAND ROUGH_ENDOPLASMIC_RETICULUM_MEMBRANE S_PROSTATE_CANCER_ANDROGEN_INDEPENDENT UCOMA
	REACTOME_REGULATION_OF_TNFR1_SIGNALING, REACTOME_IGSE10422_WT_VS_BAFF_TRANSGENIC_LN_BCELL_DN, GSE10422_WT_PRETICULOCYTOPENIA, HP_RETICULOCYTOPENIA LEE_LIVER_CANCER_MYC_UP, LEE_LIVER_CANCER_MYC_UP GSE10325_BCELL_VS_LUPUS_BCELL_UP, GSE10325_BCELL_VS_LUPUS_HOXC11_TARGET_GENES	REGULATION_OF_TNFR1_SIGNALING WT_VS_BAFF_TRANSGENIC_LN_BCELL_DN PUS_BCELL_UP PING, GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_DUPLEX_UNWINDING
OMAL_SUBUNIT, GOCC_RIBOSOMAL_SUBUNIT	HU_GENOTOXIC_DAMAGE_24HR, HU_GENOTOXIC_DAMAGE_24 GOBP_MATURATION_OF_LSU_RRNA_FROM_TRICISTRONIC_RRN REACTOME_TNFR1_INDUCED_NFKAPPAB_SIGNALING_PATHWA MELLMAN_TUT1_TARGETS_UP, MELLMAN_TUT1_TARGETS_UP REACTOME_ER_QUALITY_CONTROL_COMPARTMENT_ERQC, RE GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_TRANSLA BILANGES_SERUM_SENSITIVE_GENES, BILANGES_SERUM_SENSITIVE_CONTROL_COMPARTMENT_CONTROL_COMPA	'A_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA, GOBP_MATURATION_OF_LSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA AY, REACTOME_TNFR1_INDUCED_NFKAPPAB_SIGNALING_PATHWAY ACTOME_ER_QUALITY_CONTROL_COMPARTMENT_ERQC TION, GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_TRANSLATION
OWIAL_SUBUNIT, GOCC_RIBOSOWAL_SUBUNIT	REACTOME_INFECTION_WITH_MYCOBACTERIUM_TUBERCULOS GSE10147_IL3_VS_IL3_AND_HIVP17_STIM_PDC_DN, GSE10147_IL3_ SCHERER_PBMC_APSV_WETVAX_AGE_18_32YO_2_TO_4DY_DN, SO GOBP_POSITIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_BY CDX1_TARGET_GENES, CDX1_TARGET_GENES TERAO_AOX4_TARGETS_SKIN_UP, TERAO_AOX4_TARGETS_SKIN_ GOBP_NEGATIVE_REGULATION_OF_UBIQUITIN_PROTEIN_TRANSOBP_RRNA_METHYLATION	CHERER_PBMC_APSV_WETVAX_AGE_18_32YO_2_TO_4DY_DN _P53_CLASS_MEDIATOR, GOBP_POSITIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR _UP ISFERASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_UBIQUITIN_PROTEIN_TRANSFERASE_ACTIVITY
	GOBP_REGULATION_OF_UBIQUITIN_PROTEIN_LIGASE_ACTIVITY REACTOME_SMAD2_SMAD3_SMAD4_HETEROTRIMER_REGULATIVE REACTOME_ACTIVATED_TAK1_MEDIATES_P38_MAPK_ACTIVATED_TAK1_MEDIATES_P38_MAPK_ACTIVATED_REACTOME_FLT3_SIGNALING_IN_DISEASE, REACTOME_FLT3_SIGNALING, REACTOME_INACTIVATION_OF_CSF3_G_CSF_SIGNALING, REACTOME_ASSEMBLY_OF_THE_HIV_VIRION, REACTOME_ASSEMBLY_OF_THE_HIV_VIRION_AL_INITIATION_REACTOME_NEGATIVE_REGULATION_OF_FGFR3_SIGNALING, REACTOME_NEGATIVE_REGULATION_OF_FGFR3_SIGNALING, REACTOME_NEGATIVE_REGULATION_OF_FGFR3_SIGNALING.	TOME_INACTIVATION_OF_CSF3_G_CSF_SIGNALING MBLY_OF_THE_HIV_VIRION N, GOBP_POSITIVE_REGULATION_OF_TRANSLATIONAL_INITIATION
	GSE13485_DAY7_VS_DAY21_YF17D_VACCINE_PBMC_DN, GSE13489 REACTOME_MITOPHAGY, REACTOME_MITOPHAGY REACTOME_REGULATION_OF_SIGNALING_BY_CBL, REACTOME_ REACTOME_CONSTITUTIVE_SIGNALING_BY_LIGAND_RESPONSI REACTOME_DOWNREGULATION_OF_TGF_BETA_RECEPTOR_SIGNALING_PAREACTOME_TAK1_ACTIVATES_NFKB_BY_PHOSPHORYLATION_AREACTOME_SPRY_REGULATION_OF_FGF_SIGNALING, REACTOME_	_REGULATION_OF_SIGNALING_BY_CBL VE_EGFR_CANCER_VARIANTS, REACTOME_CONSTITUTIVE_SIGNALING_BY_LIGAND_RESPONSIVE_EGFR_CANCER_VARIANTS NALING, REACTOME_DOWNREGULATION_OF_TGF_BETA_RECEPTOR_SIGNALING NTHWAY_BY_P53_CLASS_MEDIATOR, GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR ND_ACTIVATION_OF_IKKS_COMPLEX, REACTOME_TAK1_ACTIVATES_NFKB_BY_PHOSPHORYLATION_AND_ACTIVATION_OF_IKKS_COMPLEX
	LEE_LIVER_CANCER_MYC_TGFA_UP, LEE_LIVER_CANCER_MYC_ REACTOME_ACTIVATED_NOTCH1_TRANSMITS_SIGNAL_TO_THE	TGFA_UP E_NUCLEUS, REACTOME_ACTIVATED_NOTCH1_TRANSMITS_SIGNAL_TO_THE_NUCLEUS CTIVATING_COMPLEX, REACTOME_DEACTIVATION_OF_THE_BETA_CATENIN_TRANSACTIVATING_COMPLEX _METABOLISM _BREAST_CANCER_17Q11_Q21_AMPLICON
	GOBP_MYD88_DEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_REACTOME_NEGATIVE_REGULATION_OF_FGFR1_SIGNALING, REACTOME_SIGNALING_BY_EGFR_IN_CANCER, REACTOME_SIGNALING_BY_EGFR_IN_CANCER, REACTOME_SIGNALING_BY_EGFR_IN_CHIEF_CELLS, DESCARTES_MAIN_FETAL_PARIETAL_AND_CHIEF_CELLS, DESCARTES_MAIN_FETAL_PARIET	NALING_BY_EGFR_IN_CANCER ARTES_MAIN_FETAL_PARIETAL_AND_CHIEF_CELLS MENT, REACTOME_TICAM1_RIP1_MEDIATED_IKK_COMPLEX_RECRUITMENT BY_NOTCH2
	CUI_DEVELOPING_HEART_COMPACT_ATRIAL_CARDIOMYOCYT REACTOME_AGGREPHAGY, REACTOME_AGGREPHAGY REACTOME_NEGATIVE_REGULATION_OF_MET_ACTIVITY, REACTOME_PBMC_YF_VAX_AGE_18_40YO_ANYD_DN, SCHERER_PERACTOME_LISTERIA_MONOCYTOGENES_ENTRY_INTO_HOST_CREACTOME_SUPPRESSION_OF_PHAGOSOMAL_MATURATION, RIREACTOME_JOSEPHIN_DOMAIN_DUBS, REACTOME_JOSEPHIN_D	E, CUI_DEVELOPING_HEART_COMPACT_ATRIAL_CARDIOMYOCYTE TOME_NEGATIVE_REGULATION_OF_MET_ACTIVITY BMC_YF_VAX_AGE_18_40YO_ANYD_DN CELLS, REACTOME_LISTERIA_MONOCYTOGENES_ENTRY_INTO_HOST_CELLS EACTOME_SUPPRESSION_OF_PHAGOSOMAL_MATURATION DOMAIN_DUBS I, REACTOME_MAP3K8_TPL2_DEPENDENT_MAPK1_3_ACTIVATION
	MIR6864_3P, MIR6864_3P REACTOME_NOTCH2_ACTIVATION_AND_TRANSMISSION_OF_SI	BLAST_GROWTH_FACTOR_BINDING E_DIFFERENTIATION_GENES_DN EACTOME_NEGATIVE_REGULATION_OF_FGFR2_SIGNALING
	REACTOME_NRIF_SIGNALS_CELL_DEATH_FROM_THE_NUCLEUS HP_ABNORMAL_HEMOGLOBIN, HP_ABNORMAL_HEMOGLOBIN REACTOME_INLB_MEDIATED_ENTRY_OF_LISTERIA_MONOCYTOGOMF_RIBOSOMAL_LARGE_SUBUNIT_BINDING, GOMF_RIBOSOM	AKT1_AND_DOK1, REACTOME_PTK6_REGULATES_RTKS_AND_THEIR_EFFECTORS_AKT1_AND_DOK1 G, REACTOME_NRIF_SIGNALS_CELL_DEATH_FROM_THE_NUCLEUS GENES_INTO_HOST_CELL, REACTOME_INLB_MEDIATED_ENTRY_OF_LISTERIA_MONOCYTOGENES_INTO_HOST_CELL MAL_LARGE_SUBUNIT_BINDING TY, GOMF_UBIQUITIN_PROTEIN_TRANSFERASE_INHIBITOR_ACTIVITY
	HP_BROAD_NECK, HP_BROAD_NECK REACTOME_DOWNREGULATION_OF_ERBB2_ERBB3_SIGNALING, BOGNI_TREATMENT_RELATED_MYELOID_LEUKEMIA_UP, BOGN GOBP_PROTEIN_NEDDYLATION, GOBP_PROTEIN_NEDDYLATION GOBP_NEGATIVE_REGULATION_OF_UBIQUITIN_PROTEIN_LIGAS HEDENFALK_BREAST_CANCER_BRACX_UP, HEDENFALK_BREAS NEBEN_AML_WITH_FLT3_OR_NRAS_DN, NEBEN_AML_WITH_FLT3	N SE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_UBIQUITIN_PROTEIN_LIGASE_ACTIVITY T_CANCER_BRACX_UP I3_OR_NRAS_DN
	REACTOME_RRNA_PROCESSING_IN_THE_MITOCHONDRION, RE REACTOME_DOWNREGULATION_OF_ERBB2_SIGNALING, REACT REACTOME_TICAM1_TRAF6_DEPENDENT_INDUCTION_OF_TAK1 REACTOME_ALPHA_PROTEIN_KINASE_1_SIGNALING_PATHWAY TRAVAGLINI_LUNG_CD8_MEMORY_EFFECTOR_T_CELL, TRAVAGGOBP_MITOCHONDRIAL_RIBOSOME_ASSEMBLY, GOBP_MITOCH REACTOME_IRAK1_RECRUITS_IKK_COMPLEX, REACTOME_IRAK1_RECRUITS_IKK_COMPLEX, REACTOME_IRAK1_RECRUITS_IKK_COMPLEX.	OME_DOWNREGULATION_OF_ERBB2_SIGNALING COMPLEX, REACTOME_TICAM1_TRAF6_DEPENDENT_INDUCTION_OF_TAK1_COMPLEX Y, REACTOME_ALPHA_PROTEIN_KINASE_1_SIGNALING_PATHWAY GLINI_LUNG_CD8_MEMORY_EFFECTOR_T_CELL ONDRIAL_RIBOSOME_ASSEMBLY I_RECRUITS_IKK_COMPLEX XX, REACTOME_IRAK2_MEDIATED_ACTIVATION_OF_TAK1_COMPLEX
	WILLIAMS_ESR2_TARGETS_DN, WILLIAMS_ESR2_TARGETS_DN REACTOME_REGULATION_OF_BACH1_ACTIVITY, REACTOME_RE YANAGISAWA_LUNG_CANCER_RECURRENCE, YANAGISAWA_LGOBP_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_RESEMP_REDUNDANT_NECK_SKIN, HP_REDUNDANT_NECK_SKIN_HP_LOW_LEVELS_OF_VITAMIN_D, HP_LOW_LEVELS_OF_VITAMIN_D, HP_LOW_LEVELS_OF_VITAMIN_D	I_VINCRISTINE_RESISTANCE_B_ALL_DN EMBRANE, GOCC_CYTOPLASMIC_SIDE_OF_ENDOPLASMIC_RETICULUM_MEMBRANE EGULATION_OF_BACH1_ACTIVITY UNG_CANCER_RECURRENCE EULTING_IN_TRANSCRIPTION, GOBP_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_RESULTING_IN_TRANSCRIPTION N_D
	GOBP_POSITIVE_REGULATION_OF_MEMBRANE_DEPOLARIZATION REACTOME_NOTCH3_ACTIVATION_AND_TRANSMISSION_OF_SIGOMF_UBIQUITIN_LIGASE_INHIBITOR_ACTIVITY, GOMF_UBIQUITIN_LIGASE_INHIBITOR_ACTIVITY, GOMF_UBIQUITIN_REACTOME_DISEASES_OF_CARBOHYDRATE_METABOLISM, REACTOME_DISEASES_OF_CARBOHYDRATE_METABOLISM, REACTOME_REACTOME_STRUCTURE_UNWINDING, GOBP_RNADDISTECHE_ESCAPED_FROM_X_INACTIVATION, DISTECHE_ESCAPED_FROM_X_INACTIVATION, DISTECHE_ESCAPED_FROM_STENDEDISEASSEMBLY, GOBP_RIBOSOME_DISASSEMBLY, GOBP_RIBOSOME_DISASSEMBLY	CTOME_DISEASES_OF_CARBOHYDRATE_METABOLISM _SECONDARY_STRUCTURE_UNWINDING PED_FROM_X_INACTIVATION M, REACTOME_MODULATION_BY_MTB_OF_HOST_IMMUNE_SYSTEM Y
	HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VAC REACTOME_REGULATION_OF_FZD_BY_UBIQUITINATION, REACT GOBP_TRANSLATIONAL_READTHROUGH, GOBP_TRANSLATION REACTOME_REGULATION_OF_PTEN_LOCALIZATION, REACTOME GOBP_POSITIVE_REGULATION_OF_TRANSLATION_IN_RESPONSING GOBP_POSITIVE_REGULATION_OF_TRANSLATION_IN_RESPONSING REACTOME_RRNA_MODIFICATION_IN_THE_MITOCHONDRION,	X_WITHIN_TLR4_COMPLEX, REACTOME_TRAF6_MEDIATED_INDUCTION_OF_TAK1_COMPLEX_WITHIN_TLR4_COMPLEX CCINE_AGE_18_45YO_ID_7DY_TOP_100_DEG_EX_VIVO_DN, HOFT_CD4_POSITIVE_ALPHA_BETA_MEMORY_T_CELL_BCG_VACCINE_AGE_18_45YO_ID_7DY_TOP_100_DEG_EX_VIVO_DN TOME_REGULATION_OF_FZD_BY_UBIQUITINATION AL_READTHROUGH TE_REGULATION_OF_PTEN_LOCALIZATION E_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_POSITIVE_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS E_TO_STRESS, GOBP_POSITIVE_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS REACTOME_RRNA_MODIFICATION_IN_THE_MITOCHONDRION
		RMALITY_OF_NEUTROPHIL_MORPHOLOGY IRAS_AND_NRAS_TARGETS_UP SEAL_DYSPLASIA IX R_9_SIGNALING, REACTOME_TRAF6_MEDIATED_IRF7_ACTIVATION_IN_TLR7_8_OR_9_SIGNALING
	REACTOME_PEXOPHAGY, REACTOME_PEXOPHAGY ABE_INNER_EAR, ABE_INNER_EAR GOBP_CHEMOKINE_C_C_MOTIF_LIGAND_5_PRODUCTION, GOBE HP_PARIETAL_FORAMINA, HP_PARIETAL_FORAMINA REACTOME_DOWNREGULATION_OF_ERBB4_SIGNALING, REACT HP_ANTENATAL_ONSET, HP_ANTENATAL_ONSET GOCC_DENDRITIC_SPINE_NECK, GOCC_DENDRITIC_SPINE_NECK GOBP_RESPIRATORY_BURST_INVOLVED_IN_INFLAMMATORY_REACT REACTOME_PEXOPHAGY ABE_INNER_EAR GOBP_RESPIRATORY_BURST_INVOLVED_IN_INFLAMMATORY_REACT ABE_INNER_EAR ABE_INNER_EAR GOBP_RESPIRATORY_BURST_INVOLVED_IN_INFLAMMATORY_REACT ABE_INNER_EAR	OME_DOWNREGULATION_OF_ERBB4_SIGNALING K ESPONSE, GOBP_RESPIRATORY_BURST_INVOLVED_IN_INFLAMMATORY_RESPONSE
	OUELLET_OVARIAN_CANCER_INVASIVE_VS_LMP_DN, OUELLET GOBP_REGULATION_OF_PROTEIN_NEDDYLATION, GOBP_REGUL GOBP_RESPIRATORY_BURST_INVOLVED_IN_DEFENSE_RESPONSING GOBP_REGULATION_OF_PROTEIN_K63_LINKED_UBIQUITINATION GOMF_PROTEIN_TAG, GOMF_PROTEIN_TAG REACTOME_GLYCOGEN_STORAGE_DISEASES, REACTOME_GLYCOBCAT_BILD_ET_AL_UP QIU_PBMC_HEPTATITIS_B_SURFACE_ANTIGEN_AGE_UNDER50_N	LATION_OF_PROTEIN_NEDDYLATION E, GOBP_RESPIRATORY_BURST_INVOLVED_IN_DEFENSE_RESPONSE EN, GOBP_RESPIRATORY_BURST_INVOLVED_IN_DEFENSE_RESPONSE EN, GOBP_REGULATION_OF_PROTEIN_K63_LINKED_UBIQUITINATION EOGEN_STORAGE_DISEASES ENON_RESPONDERS_VS_RESPONDERS_28DY_DN, QIU_PBMC_HEPTATITIS_B_SURFACE_ANTIGEN_AGE_UNDER50_NON_RESPONDERS_VS_RESPONDERS_28DY_DN
	GOBP_POSITIVE_REGULATION_OF_CHEMOKINE_C_C_MOTIF_LIGGORD_NEGATIVE_REGULATION_OF_RIBOSOME_BIOGENESIS, GO	ANX_OF_FINGER, HP_ABNORMALITY_OF_THE_EPIPHYSES_OF_THE_DISTAL_PHALANX_OF_FINGER GAND_5_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_CHEMOKINE_C_C_MOTIF_LIGAND_5_PRODUCTION BP_NEGATIVE_REGULATION_OF_RIBOSOME_BIOGENESIS ILUM_MEMBRANE, GOCC_CYTOPLASMIC_SIDE_OF_ROUGH_ENDOPLASMIC_RETICULUM_MEMBRANE MF_SELENOCYSTEINE_INSERTION_SEQUENCE_BINDING CTOME_P75NTR_RECRUITS_SIGNALLING_COMPLEXES EACTOME_NF_KB_IS_ACTIVATED_AND_SIGNALS_SURVIVAL SIGNALS_VIA_NF_KB
	RECHTIEN_PBMC_RVSV_ZEBOV_AGE_18_55YO_1DY_UP, RECHTIEN HP_CRANIOFACIAL_DYSOSTOSIS, HP_CRANIOFACIAL_DYSOSTO HP_LOWER_LIMB_METAPHYSEAL_IRREGULARITY, HP_LOWER_LIMB_METAPHYSEAL_IRREGULARITY, HP_LOWER_LIMB_HAPTOGLOBIN_BINDING, GOMF_HAPTOGLOBIN_BINDING CAIRO_HEPATOBLASTOMA_POOR_SURVIVAL, CAIRO_HEPATOBLASTOMA_POOR_SURVIVAL, CAIRO_HEPATOBLASTOMA_COMPLEX_ASSEMBLY, GOBP_NICTION_GOCC_INFLAMMASOME_COMPLEX, GOCC_INFLAMMASOME_COMPLEX, GOCC_INFLAMMASOME_COMPLEX, GOCC_INFLAMMASOME_COMPLEX_C	IN_PBMC_RVSV_ZEBOV_AGE_18_55YO_1DY_UP SIS IMB_METAPHYSEAL_IRREGULARITY IG LASTOMA_POOR_SURVIVAL LRP3_INFLAMMASOME_COMPLEX_ASSEMBLY DMPLEX
	GOBP_POSITIVE_REGULATION_OF_PROTEIN_POLYUBIQUITINAT LU_TUMOR_VASCULATURE_UP, LU_TUMOR_VASCULATURE_UP BERTUCCI_INVASIVE_CARCINOMA_DUCTAL_VS_LOBULAR_DN, GOBP_ANTIMICROBIAL_HUMORAL_IMMUNE_RESPONSE_MEDIA HP_TIBIALIS_MUSCLE_WEAKNESS, HP_TIBIALIS_MUSCLE_WEAK GOCC_MITOCHONDRIAL_ALPHA_KETOGLUTARATE_DEHYDROG	BERTUCCI_INVASIVE_CARCINOMA_DUCTAL_VS_LOBULAR_DN ATED_BY_ANTIMICROBIAL_PEPTIDE, GOBP_ANTIMICROBIAL_HUMORAL_IMMUNE_RESPONSE_MEDIATED_BY_ANTIMICROBIAL_PEPTIDE NESS GENASE_COMPLEX, GOCC_MITOCHONDRIAL_ALPHA_KETOGLUTARATE_DEHYDROGENASE_COMPLEX AGE_18_45YO_1DY_UP, ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_1DY_UP
	GOBP_CELLULAR_RESPONSE_TO_ANTIBIOTIC, GOBP_CELLULAR GOBP_KILLING_OF_CELLS_OF_OTHER_ORGANISM, GOBP_KILLIN HP_ERYTHROCYTE_INCLUSION_BODIES, HP_ERYTHROCYTE_INC REACTOME_MYOCLONIC_EPILEPSY_OF_LAFORA, REACTOME_M HP_WIDE_NASAL_RIDGE, HP_WIDE_NASAL_RIDGE HP_ANISOCYTOSIS, HP_ANISOCYTOSIS HP_HYPOCHROMIC_MICROCYTIC_ANEMIA, HP_HYPOCHROMIC HP_ENLARGED_METAPHYSES	_RESPONSE_TO_ANTIBIOTIC NG_OF_CELLS_OF_OTHER_ORGANISM CLUSION_BODIES YOCLONIC_EPILEPSY_OF_LAFORA
	GOCC_MITOCHONDRIAL_TRICARBOXYLIC_ACID_CYCLE_ENZYNchr15q25, chr15q25 HP_TOXEMIA_OF_PREGNANCY, HP_TOXEMIA_OF_PREGNANCY BIOCARTA_AHSP_PATHWAY, BIOCARTA_AHSP_PATHWAY GOCC_HAPTOGLOBIN_HEMOGLOBIN_COMPLEX, GOCC_HAPTOGOMF_OXYGEN_CARRIER_	GLOBIN_HEMOGLOBIN_COMPLEX ACTIVITY C_CARBON_DIOXIDE, REACTOME_ERYTHROCYTES_TAKE_UP_OXYGEN_AND_RELEASE_CARBON_DIOXIDE POSITIVE_REGULATION_OF_RESPIRATORY_BURST
	GOCC_NLRP3_INFLAMMASOME_COMPLEX, GOCC_NLRP3_INFLAMP_HYPOPLASTIC_ANEMIA, HP_HYPOPLASTIC_ANEMIA HP_IRREGULAR_FEMORAL_EPIPHYSIS, HP_IRREGULAR_FEMORATE GOCC_POLE_PLASM, GOCC_POLE_PLASM GOMF_EUKARYOTIC_INITIATION_FACTOR_4E_BINDING, GOMF_GOBP_OPTIC_NERVE_DEVELOPMENT, GOBP_OPTIC_NERVE_DEVELOPMENT, GOBP_CONTIGUOUS_GENE_SUBP_CONTIGUOUS_SUBP_CONTIGUOUS_SUBP_CONTIGUOUS_SUBP_CONTIGUOUS_S	EUKARYOTIC_INITIATION_FACTOR_4E_BINDING ELOPMENT ENITAL_GLAUCOMA SYNDROME
	GOBP_POSITIVE_REGULATION_OF_TRANSLATIONAL_ELONGATE GOBP_NEGATIVE_REGULATION_OF_RESPIRATORY_BURST_INVO GOBP_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGN	ION, GOBP_POSITIVE_REGULATION_OF_TRANSLATIONAL_ELONGATION OLVED_IN_INFLAMMATORY_RESPONSE, GOBP_NEGATIVE_REGULATION_OF_RESPIRATORY_BURST_INVOLVED_IN_INFLAMMATORY_RESPONSE VALING_PATHWAY_BY_P53_CLASS_MEDIATOR, GOBP_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR VATION, GOBP_REGULATION_OF_CYTOPLASMIC_TRANSLATIONAL_INITIATION RANSFERASE_ACTIVITY VION_OF_THE_VENTRICULAR_CAVITY
	STAMBOLSKY_RESPONSE_TO_VITAMIN_D3_DN, STAMBOLSKY_R chrXq24, chrXq24 GOBP_POSITIVE_REGULATION_OF_NLRP3_INFLAMMASOME_CO WANG_RESPONSE_TO_FORSKOLIN_DN, WANG_RESPONSE_TO_F FOXF1_TARGET_GENES, FOXF1_TARGET_GENES HP_SECUNDUM_ATRIAL_SEPTAL_DEFECT, HP_SECUNDUM_ATR chr14q21, chr14q21 LU_TUMOR_ENDOTHELIAL_MARKERS_UP, LU_TUMOR_ENDOTHE	ESPONSE_TO_VITAMIN_D3_DN MPLEX_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_NLRP3_INFLAMMASOME_COMPLEX_ASSEMBLY ORSKOLIN_DN IAL_SEPTAL_DEFECT IELIAL_MARKERS_UP
	HP_ABNORMALITY_OF_UPPER_LIMB_METAPHYSIS, HP_ABNORM HP_APLASIA_HYPOPLASIA_OF_THE_SACRUM, HP_APLASIA_HYPOBP_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY, GOBP_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY, GOBP_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY, GOBP_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY, GOBP_TOLL_CANCER_HEAD_AND_NECK_VS_CERVICAL_DN, PYEON_SMIRNOV_CIRCULATING_ENDOTHELIOCYTES_IN_CANCER_DN, HP_BRANCHIAL_CYST_MIR1203, MIR1203 HP_BRANCHIAL_CYST, HP_BRANCHIAL_CYST_MIR1203, MIR1203 HP_TOTAL_ANOMALOUS_PULMONARY_VENOUS_RETURN, HP_HP_IMBALANCED_HEMOGLOBIN_SYNTHESIS, HP_IMBALANCED_HEMOGLOBIN_SYNTHESIS, HP_IMBALANCED_HEMOGLOBIN_SYNTHESIS_HEMOGLOBIN_SYNTHESIS_HEMOGLOBIN_SYNTHESIS_HEMOGLOBIN_SYNTHESIS_HEMOGLOBIN_SYNTHESIS_H	MALITY_OF_UPPER_LIMB_METAPHYSIS POPLASIA_OF_THE_SACRUM OLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY CANCER_HEAD_AND_NECK_VS_CERVICAL_DN SMIRNOV_CIRCULATING_ENDOTHELIOCYTES_IN_CANCER_DN TOTAL_ANOMALOUS_PULMONARY_VENOUS_RETURN O_HEMOGLOBIN_SYNTHESIS
	GOMF_RRNA_CYTOSINE_METHYLTRANSFERASE_ACTIVITY, GONGOBP_NEGATIVE_REGULATION_OF_CYTOPLASMIC_TRANSLATIVE_NONSPHEROCYTIC_HEMOLYTIC_ANEMIA, HP_NONSPHEROCYTIC_HEMOLYTIC_ANEMIA, HP_NONSPHEROCYTES_TOLL_LIKE_RECEPTOR_7_SIGNALING_FERMS_COBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_7_SIGNALING_FERMS_CORPS_	MF_RRNA_CYTOSINE_METHYLTRANSFERASE_ACTIVITY ON, GOBP_NEGATIVE_REGULATION_OF_CYTOPLASMIC_TRANSLATION CYTIC_HEMOLYTIC_ANEMIA PATHWAY, GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_7_SIGNALING_PATHWAY PATHWAY, GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_T_SIGNALING_PATHWAY PATHWAY, GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_T_SIGNALING_PATHWAY PATHWAY, GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_T_SIGNALING_PATHWAY PATHWAY, GOBP_REGULATION_OF_TOLL_LIKE_RECEPTOR_T_SIGNALING_PATHWAY
	TANAKA_METHYLATED_IN_ESOPHAGEAL_CARCINOMA, TANAGOBP_NEGATIVE_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING chr2q11, chr2q11 GRANDVAUX_IRF3_TARGETS_DN, GRANDVAUX_IRF3_TARGETS_HP_DYSCHROMATOPSIA, HP_DYSCHROMATOPSIA chr5q33, chr5q33 chr6q22, chr6q22 chr2q14, chr2q14 chr6q21, chr6q21	KA_METHYLATED_IN_ESOPHAGEAL_CARCINOMA IG, GOBP_NEGATIVE_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING
	chr17q22, chr17q22	