

	GSE22886_NAIVE_BCELL_VS_MONOCYTE_DN, GSE22886_NAIVE_BCELL_VS_MONOCYTE_DN
	GSE45365_WT_VS_IFNAR_KO_CD8A_DC_MCMV_INFECTION_UP, GSE45365_WT_VS_IFNAR_KO_CD8A_DC_MCMV_INFECTION_UP
	GSE22886_NAIVE_TCELL_VS_MONOCYTE_DN, GSE22886_NAIVE_TCELL_VS_MONOCYTE_DN
	GSE29618_BCELL_VS_MONOCYTE_DN, GSE29618_BCELL_VS_MONOCYTE_DN
	GSE29618_BCELL_VS_MONOCYTE_DAY7_FLU_VACCINE_DN, GSE29618_BCELL_VS_MONOCYTE_DAY7_FLU_VACCINE_DN
	GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_KO_2H_DN, GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_KO_2H_DN
	GSE22886_NAIVE_CD4_TCELL_VS_MONOCYTE_DN, GSE22886_NAIVE_CD4_TCELL_VS_MONOCYTE_DN
	GSE29618_PDC_VS_MDC_DN, GSE29618_PDC_VS_MDC_DN
	GSE29618_MONOCYTE_VS_PDC_DAY7_FLU_VACCINE_UP, GSE29618_MONOCYTE_VS_PDC_DAY7_FLU_VACCINE_UP
	GSE42724_NAIVE_BCELL_VS_PLASMABLAST_DN, GSE42724_NAIVE_BCELL_VS_PLASMABLAST_DN
	GSE29618_MONOCYTE_VS_PDC_UP, GSE29618_MONOCYTE_VS_PDC_UP
	HOXA9_DN.V1_UP, HOXA9_DN.V1_UP
	GSE10325_LUPUS_BCELL_VS_LUPUS_MYELOID_DN, GSE10325_LUPUS_BCELL_VS_LUPUS_MYELOID_DN
	GSE22886_NAIVE_CD8_TCELL_VS_MONOCYTE_DN, GSE22886_NAIVE_CD8_TCELL_VS_MONOCYTE_DN
	GSE10325_LUPUS_CD4_TCELL_VS_LUPUS_MYELOID_DN, GSE10325_LUPUS_CD4_TCELL_VS_LUPUS_MYELOID_DN
	GSE24634_IL4_VS_CTRL_TREATED_NAIVE_CD4_TCELL_DAY10_DN, GSE24634_IL4_VS_CTRL_TREATED_NAIVE_CD4_TCELL_DAY10_DN
	GSE45365_HEALTHY_VS_MCMV_INFECTION_CD8A_DC_IFNAR_KO_DN, GSE45365_HEALTHY_VS_MCMV_INFECTION_CD8A_DC_IFNAR_KO_DN
	GSE22886_NAIVE_BCELL_VS_BM_PLASMA_CELL_UP, GSE22886_NAIVE_BCELL_VS_BM_PLASMA_CELL_UP
	GSE3982_MAST_CELL_VS_EFF_MEMORY_CD4_TCELL_UP, GSE3982_MAST_CELL_VS_EFF_MEMORY_CD4_TCELL_UP
	GSE11057_CD4_EFF_MEM_VS_PBMC_DN, GSE11057_CD4_EFF_MEM_VS_PBMC_DN
	GSE22886_NAIVE_BCELL_VS_NEUTROPHIL_DN, GSE22886_NAIVE_BCELL_VS_NEUTROPHIL_DN
	GSE29618_PDC_VS_MDC_DAY7_FLU_VACCINE_DN, GSE29618_PDC_VS_MDC_DAY7_FLU_VACCINE_DN
	GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCTYE_ADULT_UP, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN3_THYMOCTYE_ADULT_UP
	GSE10325_CD4_TCELL_VS_MYELOID_DN, GSE10325_CD4_TCELL_VS_MYELOID_DN
	GSE30971_WBP7_HET_VS_KO_MACROPHAGE_2H_LPS_STIM_UP, GSE30971_WBP7_HET_VS_KO_MACROPHAGE_2H_LPS_STIM_UP
	GSE36476_CTRL_VS_TSST_ACT_16H_MEMORY_CD4_TCELL_YOUNG_UP, GSE36476_CTRL_VS_TSST_ACT_16H_MEMORY_CD4_TCELL_YOUNG_UP
	GSE12392_CD8A_POS_VS_NEG_SPLEEN_IFNB_KO_DC_DN, GSE12392_CD8A_POS_VS_NEG_SPLEEN_IFNB_KO_DC_DN
	RPS14_DN.V1_UP, RPS14_DN.V1_UP
	GSE34156_UNTREATED_VS_24H_NOD2_LIGAND_TREATED_MONOCYTE_DN, GSE34156_UNTREATED_VS_24H_NOD2_LIGAND_TREATED_MONOCYTE_DN
	GSE3982_MAST_CELL_VS_MAC_DN, GSE3982_MAST_CELL_VS_MAC_DN
	GSE24634_TREG_VS_TCONV_POST_DAY10_IL4_CONVERSION_DN, GSE24634_TREG_VS_TCONV_POST_DAY10_IL4_CONVERSION_DN
	GSE2706_2H_VS_8H_R848_STIM_DC_UP, GSE2706_2H_VS_8H_R848_STIM_DC_UP
	AIZARANI_LIVER_C2_KUPFFER_CELLS_1, AIZARANI_LIVER_C2_KUPFFER_CELLS_1
	GSE32255_WT_UNSTIM_VS_JMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_DN, GSE32255_WT_UNSTIM_VS_JMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_DN
	GSE34156_UNTREATED_VS_6H_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP, GSE34156_UNTREATED_VS_6H_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP
	GSE13306_RA_VS_UNTREATED_MEM_CD4_TCELL_UP, GSE13306_RA_VS_UNTREATED_MEM_CD4_TCELL_UP
	HUANG_GATA2_TARGETS_UP, HUANG_GATA2_TARGETS_UP
	RAF_UP.V1_UP, RAF_UP.V1_UP
	GSE3982_BASOPHIL_VS_TH2_UP, GSE3982_BASOPHIL_VS_TH2_UP
	GSE12845_NAIVE_VS_DARKZONE_GC_TONSIL_BCELL_UP, GSE12845_NAIVE_VS_DARKZONE_GC_TONSIL_BCELL_UP
	AIZARANI_LIVER_C25_KUPFFER_CELLS_4, AIZARANI_LIVER_C25_KUPFFER_CELLS_4
	GSE32901_THI7_EMRICHD_VS_THI7_NEG_CD4_TCELL_DN, GSE32901_THI7_EMRICHD_VS_THI7_NEG_CD4_TCELL_DN
	VERHAAK_AML_WITH_NPM1_MUTATED_UP, VERHAAK_AML_WITH_NPM1_MUTATED_UP
	TONKS_TARGETS_OF_RUNX1_RUNXIT1_FUSION_HSC_DN, TONKS_TARGETS_OF_RUNX1_RUNXIT1_FUSION_HSC_DN
	GSE6269_HEALTHY_VS_STAPH_PNEUMO_INF_PBMC_DN, GSE6269_HEALTHY_VS_STAPH_PNEUMO_INF_PBMC_DN
	GSE3982_CTRL_VS_LPS_48H_DC_UP, GSE3982_CTRL_VS_LPS_48H_DC_UP
	GSE32533_MIR17_KO_VS_MIR17_OVEREXPRESS_ACT_CD4_TCELL_UP, GSE32533_MIR17_KO_VS_MIR17_OVEREXPRESS_ACT_CD4_TCELL_UP
	GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_CORTEX_THYMUS_UP, GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_CORTEX_THYMUS_UP
	LTE2_UP.V1_UP, LTE2_UP.V1_UP
	GSE3982_MAC_VS_TH1_UP, GSE3982_MAC_VS_TH1_UP
	GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_MONOCYTE_UP, GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_MONOCYTE_UP
	GSE7460_TCONV_VS_TREG_THYMUS_UP, GSE7460_TCONV_VS_TREG_THYMUS_UP
	LI_WILMS_TUMOR_VS_FETAL_KIDNEY_1_UP, LI_WILMS_TUMOR_VS_FETAL_KIDNEY_1_UP
	GSE5679_PPARG_LIGAND_ROSIGLITAZONE_VS_RARA_AGONIST_AM580_TREATED_DC_DN, GSE5679_PPARG_LIGAND_ROSIGLITAZONE_VS_RARA_AGONIST_AM580_TREATED_DC_DN
	REACTOME_FATTY_ACID_METABOLISM, REACTOME_FATTY_ACID_METABOLISM
	GSE12392_IFNAR_KO_VS_IFNB_KO_CD8_NEG_SPLEEN_DC_DN, GSE12392_IFNAR_KO_VS_IFNB_KO_CD8_NEG_SPLEEN_DC_DN
	AIZARANI_LIVER_C18_NK_NKT_CELLS_5, AIZARANI_LIVER_C18_NK_NKT_CELLS_5
	GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_INF_UP, GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_INF_UP
	REACTOME_RAB_GEPF_EXCHANGE_GTP_FOR_GDP_ON_RABS, REACTOME_RAB_GEPF_EXCHANGE_GTP_FOR_GDP_ON_RABS
	GSE3982_MAST_CELL_VS_DC_DN, GSE3982_MAST_CELL_VS_DC_DN
	ESC_J1_UP_EARLY.V1_DN, ESC_J1_UP_EARLY.V1_DN
	DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_DENDRITIC_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_DENDRITIC_CELLS
	GSE2706_2H_VS_8H_LPS_STIM_DC_UP, GSE2706_2H_VS_8H_LPS_STIM_DC_UP
	ALTEMEIER_RESPONSE_TO_LPS_WITH_MECHANICAL_VENTILATION, ALTEMEIER_RESPONSE_TO_LPS_WITH_MECHANICAL_VENTILATION
	GSE34156_NOD2_LIGAND_VS_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_DN, GSE34156_NOD2_LIGAND_VS_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_DN
	HUANG_DASATINIB_SENSITIVITY_UP, HUANG_DASATINIB_SENSITIVITY_UP
	GSE22611_UNSTIM_VS_6H_MDP_STIM_NOD2_TRANSDUCED_HEK293T_CELL_DN, GSE22611_UNSTIM_VS_6H_MDP_STIM_NOD2_TRANSDUCED_HEK293T_CELL_DN
	GNF2_HCK, GNF2_HCK
	GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_WITH_EGR2_DELETED_DN, GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_WITH_EGR2_DELETED_DN
	GSE5455_EX_VIVO_VS_POST_24H_INCUBATION_MONOCYTES_FROM_TUMOR_BEARING_MOUSE_DN, GSE5455_EX_VIVO_VS_POST_24H_INCUBATION_MONOCYTES_FROM_TUMOR_BEARING_MOUSE_DN
	MIR4728_3P, MIR4728_3P
	GSE24574_BCL6_HIGH_TFH_VS_TCONV_CD4_TCELL_DN, GSE24574_BCL6_HIGH_TFH_VS_TCONV_CD4_TCELL_DN
	CHIARADONNA_NEOPLASTIC_TRANSFORMATION_CDC25_UP, CHIARADONNA_NEOPLASTIC_TRANSFORMATION_CDC25_UP
	DESCARTES_FETAL_HEART_MYELOID_CELLS, DESCARTES_FETAL_HEART_MYELOID_CELLS
	TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_10D_DN, TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_10D_DN
	GSE2706_R848_VS_R848_AND_LPS_8H_STIM_DC_UP, GSE2706_R848_VS_R848_AND_LPS_8H_STIM_DC_UP
	GOBP_POSITIVE_REGULATION_OF_PROTEIN_POLYMERIZATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_POLYMERIZATION
	GSE34006_WT_VS_A2AR_KO_TREG_UP, GSE34006_WT_VS_A2AR_KO_TREG_UP
	MODULE_170, MODULE_170
	GNF2_TNFRSF1B, GNF2_TNFRSF1B
	GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_DN, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_DN
	GSE3982_EFF_MEMORY_CD4_TCELL_VS_TH1_UP, GSE3982_EFF_MEMORY_CD4_TCELL_VS_TH1_UP
	HP_VERTIGO, HP_VERTIGO
	GOBP_POSITIVE_REGULATION_OF_ACTIN_FILAMENT_POLYMERIZATION, GOBP_POSITIVE_REGULATION_OF_ACTIN_FILAMENT_POLYMERIZATION
	GSE28783_ANTI_MIR33_VS_UNTREATED_ATHEROSCLEROSIS_MACROPHAGE_DN, GSE28783_ANTI_MIR33_VS_UNTREATED_ATHEROSCLEROSIS_MACROPHAGE_DN
	LIU_IL4_SIGNALING, LIU_IL4_SIGNALING
	GOBP_ACTIN_CYTOSKELETON_REORGANIZATION, GOBP_ACTIN_CYTOSKELETON_REORGANIZATION
	GNF2_CARD15, GNF2_CARD15
	GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_CH_GROUP_OF_DONORS, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_CH_GROUP_OF_DONORS
	DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_FIBROBLASTS_STROMAL_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_FIBROBLASTS_STROMAL_CELLS
	MODULE_292, MODULE_292
	GSE16385_UNTREATED_VS_12H_IL4_TREATED_MACROPHAGE_UP, GSE16385_UNTREATED_VS_12H_IL4_TREATED_MACROPHAGE_UP
	FAELT_B_CLL_WITH_VH_REARRANGEMENTS_UP, FAELT_B_CLL_WITH_VH_REARRANGEMENTS_UP
	KIM_MYC_AMPLIFICATION_TARGETS_DN, KIM_MYC_AMPLIFICATION_TARGETS_DN
	GNF2_S100A4, GNF2_S100A4
	GNF2_PECAM1, GNF2_PECAM1
	RORA2_01, RORA2_01
	DUNNE_TARGETS_OF_AML1_MTG8_FUSION_UP, DUNNE_TARGETS_OF_AML1_MTG8_FUSION_UP
	GOBP_ACTIN_NUCLEATION, GOBP_ACTIN_NUCLEATION
	GSE37319_WT_VS_RC3H1_KO_CD4LOW_CD8_TCELL_DN, GSE37319_WT_VS_RC3H1_KO_CD4LOW_CD8_TCELL_DN
	GSE41978_WT_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_UP, GSE41978_WT_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_UP
	GNF2_CD33, GNF2_CD33
	GOBP_POSITIVE_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION
	GOBP_POSITIVE_REGULATION_OF_T_CELL_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_T_CELL_PROLIFERATION
	MODULE_208, MODULE_208
	GSE43955_1H_VS_42H_ACT_CD4_TCELL_DN, GSE43955_1H_VS_42H_ACT_CD4_TCELL_DN
	REACTOME_FLT3_SIGNALING, REACTOME_FLT3_SIGNALING
	GOBP_PROTEIN_LOCALIZATION_TO_LYSOSOME, GOBP_PROTEIN_LOCALIZATION_TO_LYSOSOME
	GOBP_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION, GOBP_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION
	GOBP_PLATELET_DEGRANULATION, GOBP_PLATELET_DEGRANULATION
	KIM_GLI32_TARGETS_UP, KIM_GLI32_TARGETS_UP
	LEE_EARLY_T_LYMPHOCYTE_DN, LEE_EARLY_T_LYMPHOCYTE_DN
	GOBP_PHAGOSOME_MATURATION, GOBP_PHAGOSOME_MATURATION
	MOR1_LARGE_PRE_BIL_LYMPHOCYTE_DN, MOR1_LARGE_PRE_BIL_LYMPHOCYTE_DN
	GSE25677_R848_VS_MPL_AND_R848_STIM_BCELL_UP, GSE25677_R848_VS_MPL_AND_R848_STIM_BCELL_UP
	GOBP_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_POTENTIAL, GOBP_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_POTENTIAL
	GOBP_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY, GOBP_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY
	GOMF_AMYLOID_BETA_BINDING, GOMF_AMYLOID_BETA_BINDING
	BUSSLINGER_DUODENAL_LATE_IMMATURE_ENTEROCYTES, BUSSLINGER_DUODENAL_LATE_IMMATURE_ENTEROCYTES
	GSE32423_MEMORY_VS_NAIVE_CD8_TCELL_IL7_IL4_UP, GSE32423_MEMORY_VS_NAIVE_CD8_TCELL_IL7_IL4_UP
	HP_NEURONAL_LOSS_IN_CENTRAL_NERVOUS_SYSTEM, HP_NEURONAL_LOSS_IN_CENTRAL_NERVOUS_SYSTEM
	TONKS_TARGETS_OF_RUNX1_RUNXIT1_FUSION_SUSTAINED_IN_ERYTHROCYTE_UP, TONKS_TARGETS_OF_RUNX1_RUNXIT1_FUSION_SUSTAINED_IN_ERYTHROCYTE_UP
	LIEN_BREAST_CARCINOMA_METAPLASTIC_VS_DUCTAL_UP, LIEN_BREAST_CARCINOMA_METAPLASTIC_VS_DUCTAL_UP
	REACTOME_TRANSPORT_OF_INORGANIC_CATIONS_ANIONS_AND_AMINO_ACIDS_OLIGOPEPTIDES, REACTOME_TRANSPORT_OF_INORGANIC_CATIONS_ANIONS_AND_AMINO_ACIDS_OLIGOPEPTIDES
	HP_PALPITATIONS, HP_PALPITATIONS
	MIR4301, MIR4301
	GSE35543_IN_VIVO_NTREG_VS_IN_VITRO_ITREG_UP, GSE35543_IN_VIVO_NTREG_VS_IN_VITRO_ITREG_UP
	MODULE_248, MODULE_248
	REACTOME_FCGR3A_MEDIATED_IL10_SYNTHESIS, REACTOME_FCGR3A_MEDIATED_IL10_SYNTHESIS
	KEGG_MELANOGENESIS, KEGG_MELANOGENESIS
	GOCC_PHAGOCYTIC_CUP, GOCC_PHAGOCYTIC_CUP
	GOCC_CAVEOLA, GOCC_CAVEOLA
	PETROVA_PROX1_TARGETS_DN, PETROVA_PROX1_TARGETS_DN
	GOMF_ORGANIC_ANION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_ORGANIC_ANION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
	GSE36392_EOSINOPHIL_VS_MAC_IL25_TREATED_LUNG_UP, GSE36392_EOSINOPHIL_VS_MAC_IL25_TREATED_LUNG_UP
	HP_DEHYDRATION, HP_DEHYDRATION
	GOBP_ACTIVATED_T_CELL_PROLIFERATION, GOBP_ACTIVATED_T_CELL_PROLIFERATION
	GOMF_STEROL_TRANSPORTER_ACTIVITY, GOMF_STEROL_TRANSPORTER_ACTIVITY
	MIR656_5P, MIR656_5P
	LIU_VAV3_PROSTATE_CARCINOGENESIS_UP, LIU_VAV3_PROSTATE_CARCINOGENESIS_UP
	MODULE_173, MODULE_173
	HP_FOCAL_MYOCLONIC_SEIZURE, HP_FOCAL_MYOCLONIC_SEIZURE
	GOBP_REGULATION_OF_ANTIGEN_PROCESSING_AND_PRESENTATION, GOBP_REGULATION_OF_ANTIGEN_PROCESSING_AND_PRESENTATION
	GAUSSMANN_MLL_AF4_FUSION_TARGETS_D_UP, GAUSSMANN_MLL_AF4_FUSION_TARGETS_D_UP
	REACTOME_TETRAHYDROBIOPTERIN_BH4_SYNTHESIS_RECYCLING_SALVAGE_AND_REGULATION, REACTOME_TETRAHYDROBIOPTERIN_BH4_SYNTHESIS_RECYCLING_SALVAGE_AND_REGULATION
	GOBP_REGULATION_OF_METALLOENDOPEPTIDASE_ACTIVITY, GOBP_REGULATION_OF_METALLOENDOPEPTIDASE_ACTIVITY
	GOBP_MECHANORECEPTOR_DIFFERENTIATION, GOBP_MECHANORECEPTOR_DIFFERENTIATION
	TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_DUCTAL_NORMAL_UP, TURASHVILI_BREAST_LOBULAR_CARCINOMA_VS_DUCTAL_NORMAL_UP
	GOMF_DIPEPTIDASE_ACTIVITY, GOMF_DIPEPTIDASE_ACTIVITY
	GOCC_CILIARY_BASE, GOCC_CILIARY_BASE
	TRAVAGLINI_LUNG_CAPILLARY_CELL, TRAVAGLINI_LUNG_CAPILLARY_CELL
	GOBP_INNER_EAR_RECEPTOR_CELL_STEREOCILUM_ORGANIZATION, GOBP_INNER_EAR_RECEPTOR_CELL_STEREOCILUM_ORGANIZATION
	GOBP_PTERIDINE_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS, GOBP_PTERIDINE_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS
	REACTOME_METABOLISM_OF_NITRIC_OXIDE_NOS3_ACTIVATION_AND_REGULATION, REACTOME_METABOLISM_OF_NITRIC_OXIDE_NOS3_ACTIVATION_AND_REGULATION
	GOBP_RESPONSE_TO_MACROPHAGE_COLONY_STIMULATING_FACTOR, GOBP_RESPONSE_TO_MACROPHAGE_COLONY_STIMULATING_FACTOR
	GOBP_POSITIVE_REGULATION_OF_PHAGOCYTOSIS_ENGULFMENT, GOBP_POSITIVE_REGULATION_OF_PHAGOCYTOSIS_ENGULFMENT
	REACTOME_PKA_MEDIATED_PHOSPHORYLATION_OF_CREB, REACTOME_PKA_MEDIATED_PHOSPHORYLATION_OF_CREB
	GOMF_COA_HYDROLASE_ACTIVITY, GOMF_COA_HYDROLASE_ACTIVITY
	GOBP_INNER_EAR_RECEPTOR_CELL_DEVELOPMENT, GOBP_INNER_EAR_RECEPTOR_CELL_DEVELOPMENT
	GOMF_ARP2_3_COMPLEX_BINDING, GOMF_ARP2_3_COMPLEX_BINDING
	GOBP_RESPONSE_TO_ATP, GOBP_RESPONSE_TO_ATP
	CTGRYYYNATT_UNKNOWN, CTGRYYYNATT_UNKNOWN
	GOBP_PHAGOLYSOSOME_ASSEMBLY, GOBP_PHAGOLYSOSOME_ASSEMBLY
	REACTOME_ENOS_ACTIVATION, REACTOME_ENOS_ACTIVATION
	REACTOME_MITOCHONDRIAL_FATTY_ACID_BETA_OXIDATION_OF_SATURATED_FATTY_ACIDS, REACTOME_MITOCHONDRIAL_FATTY_ACID_BETA_OXIDATION_OF_SATURATED_FATTY_ACIDS
	HOFFMANN_LARGE_TO_SMALL_PRE_BIL_LYMPHOCYTE_DN, HOFFMANN_LARGE_TO_SMALL_PRE_BIL_LYMPHOCYTE_DN
	MODULE_444, MODULE_444
	GOBP_REGULATION_OF_METALLOPEPTIDASE_ACTIVITY, GOBP_REGULATION_OF_METALLOPEPTIDASE_ACTIVITY
	REACTOME_PKA_ACTIVATION_IN_GLUcAGON_SIGNALLING, REACTOME_PKA_ACTIVATION_IN_GLUcAGON_SIGNALLING
	REACTOME_ENDOGENOUS_STEROLS, REACTOME_ENDOGENOUS_STEROLS
	HOWLIN_PUBERTAL_MAMMARY_GLAND, HOWLIN_PUBERTAL_MAMMARY_GLAND
	SCHAEFFER_SOX9_TARGETS_IN_PROSTATE_DEVELOPMENT_UP, SCHAEFFER_SOX9_TARGETS_IN_PROSTATE_DEVELOPMENT_UP
	HP_ESOPHAGEAL_NEOPLASM, HP_ESOPHAGEAL_NEOPLASM
	GOBP_ACTIVATION_OF_PROTEIN_KINASE_A_ACTIVITY, GOBP_ACTIVATION_OF_PROTEIN_KINASE_A_ACTIVITY
	GOBP_POSITIVE_REGULATION_OF_ACTIVATED_T_CELL_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_ACTIVATED_T_CELL_PROLIFERATION
	GOBP_FATTY_ACID_DERIVATIVE_BIOSYNTHETIC_PROCESS, GOBP_FATTY_ACID_DERIVATIVE_BIOSYNTHETIC_PROCESS
	GOBP_CELLULAR_RESPONSE_TO_GLUcAGON_STIMULUS, GOBP_CELLULAR_RESPONSE_TO_GLUcAGON_STIMULUS
	GOBP_REGULATION_OF_MONOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_MONOCYTE_DIFFERENTIATION
	JL_METASTASIS_REPRESSED_BY_STK11, JL_METASTASIS_REPRESSED_BY_STK11
	HP_NEOPLASM_OF_HEAD_AND_NECK, HP_NEOPLASM_OF_HEAD_AND_NECK
	GOBP_INTERLEUKIN_10_PRODUCTION, GOBP_INTERLEUKIN_10_PRODUCTION
	GOBP_CELL_PROLIFERATION_INVOLVED_IN_METANEPHROS_DEVELOPMENT, GOBP_CELL_PROLIFERATION_INVOLVED_IN_METANEPHROS_DEVELOPMENT
	GOBP_SYNAPTIC_VESICLE_MATURATION, GOBP_SYNAPTIC_VESICLE_MATURATION
	GOBP_POSITIVE_REGULATION_OF_MESENCHYMAL_CELL_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_MESENCHYMAL_CELL_PROLIFERATION
	GOBP_REGULATION_OF_MESENCHYMAL_CELL_PROLIFERATION, GOBP_REGULATION_OF_MESENCHYMAL_CELL_PROLIFERATION
	SAKAI_CHRONIC_HEPATITIS_VS_LIVER_CANCER_DN, SAKAI_CHRONIC_HEPATITIS_VS_LIVER_CANCER_DN
	SCHRAETS_MLL_TARGETS_DN, SCHRAETS_MLL_TARGETS_DN
	HP_UNDETECTABLE_ELECTRORETINOGRAM, HP_UNDETECTABLE_ELECTRORETINOGRAM
	GOBP_DORSAL_VENTRAL_PATTERN_FORMATION, GOBP_DORSAL_VENTRAL_PATTERN_FORMATION
	GOBP_ENDOTHELIAL_CELL_MATRIX_ADHESION, GOBP_ENDOTHELIAL_CELL_MATRIX_ADHESION