

	GSE2270_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP
	GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN
	GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN, GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN
	GSE41176_UNSTIM_VS_ANTI_IGM_STIM_BCELL_6H_UP, GSE41176_UNSTIM_VS_ANTI_IGM_STIM_BCELL_6H_UP
	GSE40666_UNTREATED_VS_IFNA_STIM_STAT4_KO_EFFECTOR_CD8_TCELL_90MIN_DN, GSE40666_UNTREATED_VS_IFNA_STIM_STAT4_KO_EFFECTOR_CD8_TCELL_90MIN_DN
	GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP, GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP
	MIR194_5P, MIR194_5P
	GSE3982_CTRL_VS_IGE_STIM_MAST_CELL_UP, GSE3982_CTRL_VS_IGE_STIM_MAST_CELL_UP
	GSE360_L_DONOVANI_VS_B_MALAYI_HIGH_DOSE_MAC_UP, GSE360_L_DONOVANI_VS_B_MALAYI_HIGH_DOSE_MAC_UP
	GSE18804_SPLEEN_MACROPHAGE_VS_TUMORAL_MACROPHAGE_DN, GSE18804_SPLEEN_MACROPHAGE_VS_TUMORAL_MACROPHAGE_DN
	REACTOME_ESTROGEN_DEPENDENT_GENE_EXPRESSION, REACTOME_ESTROGEN_DEPENDENT_GENE_EXPRESSION
	GSE17721_CTRL_VS_LPS_2H_BMDC_DN, GSE17721_CTRL_VS_LPS_2H_BMDC_DN
	MIR12132, MIR12132
	GOBP_REGULATION_OF_GENE_SILENCING, GOBP_REGULATION_OF_GENE_SILENCING
	MIR6768_3P, MIR6768_3P
	GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP
	GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_UP, GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_UP
	REACTOME_RUNX1_REGULATES_GENES_INVOLVED_IN_MEGAKARYOCYTE_DIFFERENTIATION_AND_PLATELET_FUNCTION, REACTOME_RUNX1_REGULATES_GENES_INVOLVED_IN_MEGAKARYOCYTE_DIFFERENTIATION_AND_PLATELET_FUNCTION
	GOBP_REGULATION_OF_GENE_SILENCING_BY_RNA, GOBP_REGULATION_OF_GENE_SILENCING_BY_RNA
	MIR500A_5P, MIR500A_5P
	GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_UP, GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_UP
	MIR4286, MIR4286
	MIR1208, MIR1208
	MIR558, MIR558
	MIR891A_3P, MIR891A_3P
	MIR6770_5P, MIR6770_5P
	GOBP_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC, GOBP_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC
	MIR488_3P, MIR488_3P
	GSE40685_NAIVE_CD4_TCELL_VS_FOXP3_KO_TREG_PRECURSOR_DN, GSE40685_NAIVE_CD4_TCELL_VS_FOXP3_KO_TREG_PRECURSOR_DN
	MIR4456, MIR4456
	MIR4302, MIR4302
	GOBP_AMINOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_AMINOGLYCAN_BIOSYNTHETIC_PROCESS
	GRAHAM_CML_DIVIDING_VS_NORMAL_QUIESCENT_DN, GRAHAM_CML_DIVIDING_VS_NORMAL_QUIESCENT_DN
	GOBP_NEPHRON_DEVELOPMENT, GOBP_NEPHRON_DEVELOPMENT
	GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_DN
	GOMF_INTEGRIN_BINDING, GOMF_INTEGRIN_BINDING
	MIR198, MIR198
	MIR4638_3P, MIR4638_3P
	ZHAN_MULTIPLE_MYELOMA_CD1_VS_CD2_DN, ZHAN_MULTIPLE_MYELOMA_CD1_VS_CD2_DN
	GOBP_CHROMATIN_SILENCING, GOBP_CHROMATIN_SILENCING
	GOBP_ZYMOGEN_ACTIVATION, GOBP_ZYMOGEN_ACTIVATION
	HALLMARK_COAGULATION, HALLMARK_COAGULATION
	MIR584_5P, MIR584_5P
	MIR597_5P, MIR597_5P
	HP_ABNORMAL_ANATOMIC_LOCATION_OF_THE_HEART, HP_ABNORMAL_ANATOMIC_LOCATION_OF_THE_HEART
	MIR1912_3P, MIR1912_3P
	GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMCDN, GSE29615_DAY3_VS_DAY7_LAIV_FLU_VACCINE_PBMCDN
	GOBP_EMBRYONIC_PLACENTA_DEVELOPMENT, GOBP_EMBRYONIC_PLACENTA_DEVELOPMENT
	GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_WOUNDING, GOBP_POSITIVE_REGULATION_OF_RESPONSE_TO_WOUNDING
	GOMF_CORECEPTOR_ACTIVITY, GOMF_CORECEPTOR_ACTIVITY
	GOBP_AROMATIC_AMINO_ACID_FAMILY_CATABOLIC_PROCESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_CATABOLIC_PROCESS
ROMEDINU_IN_T_CELL_LINE_6H_DN, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_DN	WP_MECHANOREGULATION_AND_PATHOLOGY_OF_YAPTAZ_VIA_HIPPO_AND_NONHIPPO_MECHANISMS, WP_MECHANOREGULATION_AND_PATHOLOGY_OF_YAPTAZ_VIA_HIPPO_AND_NONHIPPO_MECHANISMS
	HP_SHORT_RIBS, HP_SHORT_RIBS
	GOBP_WNT_SIGNALING_PATHWAY_CALCMIUM_MODULATING_PATHWAY, GOBP_WNT_SIGNALING_PATHWAY_CALCMIUM_MODULATING_PATHWAY
	GOBP_LABYRINTHINE_LAYER_DEVELOPMENT, GOBP_LABYRINTHINE_LAYER_DEVELOPMENT
	GOBP_CONNECTIVE_TISSUE_REPLACEMENT, GOBP_CONNECTIVE_TISSUE_REPLACEMENT
	GOMF_EXTRACELLULAR_MATRIX_BINDING, GOMF_EXTRACELLULAR_MATRIX_BINDING
	GOBP_APOPTOTIC_CELL_CLEARANCE, GOBP_APOPTOTIC_CELL_CLEARANCE
	HARALAMBIEVA_PBMCDN, HARALAMBIEVA_PBMCDN
	HP_ABNORMALITY_OF_THE_CHOROID_PLEXUS, HP_ABNORMALITY_OF_THE_CHOROID_PLEXUS
	GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS
	GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY
	GOBP_REGULATION_OF_CONNECTIVE_TISSUE_REPLACEMENT, GOBP_REGULATION_OF_CONNECTIVE_TISSUE_REPLACEMENT
	GOBP_REGULATION_OF_GOLGI_ORGANIZATION, GOBP_REGULATION_OF_GOLGI_ORGANIZATION
	GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_DN, GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_DN
	COMP1_01, COMP1_01
	GOBP_LIPID_IMPORT_INTO_CELL, GOBP_LIPID_IMPORT_INTO_CELL
	GOBP_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT, GOBP_REGULATION_OF_ANION_TRANSMEMBRANE_TRANSPORT
	HP_ATYPICAL_ABSENCE_SEIZURE, HP_ATYPICAL_ABSENCE_SEIZURE
	GOBP_CHROMATIN_SILENCING_AT_TELOMERE, GOBP_CHROMATIN_SILENCING_AT_TELOMERE
	REACTOME_CA2_PATHWAY, REACTOME_CA2_PATHWAY
	BECKER_TAMOXIFEN_RESISTANCE_DN, BECKER_TAMOXIFEN_RESISTANCE_DN
	GOBP_LABYRINTHINE_LAYER_MORPHOGENESIS, GOBP_LABYRINTHINE_LAYER_MORPHOGENESIS
	GOBP_SEQUESTERING_OF_TRIGLYCERIDE, GOBP_SEQUESTERING_OF_TRIGLYCERIDE
	GOBP_AMINE_CATABOLIC_PROCESS, GOBP_AMINE_CATABOLIC_PROCESS
	HP_REDUCED_SPERM_MOTILITY, HP_REDUCED_SPERM_MOTILITY
	HP_ABNORMALITY_OF_THE_4TH_METACARPAL, HP_ABNORMALITY_OF_THE_4TH_METACARPAL
	WP_OSTEOPONTIN_SIGNALING, WP_OSTEOPONTIN_SIGNALING
	GOMF_H4_HISTONE_ACETYLTRANSFERASE_ACTIVITY, GOMF_H4_HISTONE_ACETYLTRANSFERASE_ACTIVITY
	REACTOME_NON_INTEGRIN_MEMBRANE_ECM_INTERACTIONS, REACTOME_NON_INTEGRIN_MEMBRANE_ECM_INTERACTIONS
	HP_APLASIA_HYPOPLASIA_OF_THE_4TH_METACARPAL, HP_APLASIA_HYPOPLASIA_OF_THE_4TH_METACARPAL
	HP_BONE_FRACTURE, HP_BONE_FRACTURE
	HP_ANKYLOSIS, HP_ANKYLOSIS
	GOBP_REGULATION_OF_SEQUESTERING_OF_TRIGLYCERIDE, GOBP_REGULATION_OF_SEQUESTERING_OF_TRIGLYCERIDE
	GOBP_KETONE_CATABOLIC_PROCESS, GOBP_KETONE_CATABOLIC_PROCESS
	GOMF_ACETYLGALACTOSAMINYLTRANSFERASE_ACTIVITY, GOMF_ACETYLGALACTOSAMINYLTRANSFERASE_ACTIVITY
	GOBP_COLLAGEN_METABOLIC_PROCESS, GOBP_COLLAGEN_METABOLIC_PROCESS
	HASLINGER_B_CLL_WITH_MUTATED_VH_GENES, HASLINGER_B_CLL_WITH_MUTATED_VH_GENES
	GOBP_INOSITOL_PHOSPHATE_MEDIATED_SIGNALING, GOBP_INOSITOL_PHOSPHATE_MEDIATED_SIGNALING
	WP_PREGNANE_X_RECEPTOR_PATHWAY, WP_PREGNANE_X_RECEPTOR_PATHWAY
	GOBP_DENDRITE_EXTENSION, GOBP_DENDRITE_EXTENSION
	MATZUK_EARLY_ANTRAL_FOLLICLE, MATZUK_EARLY_ANTRAL_FOLLICLE
	GOBP_POSTSYNAPTIC_MODULATION_OF_CHEMICAL_SYNAPTIC_TRANSMISSION, GOBP_POSTSYNAPTIC_MODULATION_OF_CHEMICAL_SYNAPTIC_TRANSMISSION
	GOBP_SYMPATHETIC_GANGLION_DEVELOPMENT, GOBP_SYMPATHETIC_GANGLION_DEVELOPMENT
	GOBP_TRYPTOPHAN_METABOLIC_PROCESS, GOBP_TRYPTOPHAN_METABOLIC_PROCESS
	REACTOME_G_ALPHA_S_SIGNALING_EVENTS, REACTOME_G_ALPHA_S_SIGNALING_EVENTS
	HP_ABNORMAL_PLACENTA_MORPHOLOGY, HP_ABNORMAL_PLACENTA_MORPHOLOGY
	GOBP_POSITIVE_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS, GOBP_POSITIVE_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS
	HP_DIAPHYSEAL_THICKENING, HP_DIAPHYSEAL_THICKENING
	GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_DN, GSE7459_UNTREATED_VS_IL6_TREATED_ACT_CD4_TCELL_DN
	GOCC_NUCLEAR_CYCLIN_DEPENDENT_PROTEIN_KINASE_HOLOENZYME_COMPLEX, GOCC_NUCLEAR_CYCLIN_DEPENDENT_PROTEIN_KINASE_HOLOENZYME_COMPLEX
	HINATA_NFKB_TARGETS_KERATINOCYTE_DN, HINATA_NFKB_TARGETS_KERATINOCYTE_DN