

	GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN
	GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_DN, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_DN
	GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN
	GSE17186_BLOOD_VS_CORD_BLOOD_CD21LOW_TRANSITIONAL_BCELL_DN, GSE17186_BLOOD_VS_CORD_BLOOD_CD21LOW_TRANSITIONAL_BCELL_DN
	GSE37416_CTRL_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_CTRL_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_UP
	GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP, GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP
	GSE25088_WT_VS_STAT6_KO_MACROPHAGE_ROSIGLITAZONE_STIM_UP, GSE25088_WT_VS_STAT6_KO_MACROPHAGE_ROSIGLITAZONE_STIM_UP
	KARLSSON_TGFB1_TARGETS_DN, KARLSSON_TGFB1_TARGETS_DN
	GSE13484_12H_VS_3H_YF17D_VACCINE_STIM_PBMCMC_UP, GSE13484_12H_VS_3H_YF17D_VACCINE_STIM_PBMCMC_UP
	GSE13493_CD4INTCD8POS_VS_CD8POS_THYMOCYTE_UP, GSE13493_CD4INTCD8POS_VS_CD8POS_THYMOCYTE_UP
	GSE10273_HIGH_VS_LOW_IL7_TREATED_IRF4_8_NULL_PRE_BCELL_UP, GSE10273_HIGH_VS_LOW_IL7_TREATED_IRF4_8_NULL_PRE_BCELL_UP
	KAECH_NAIVE_VS_DAY15_EFF_CD8_TCELL_DN, KAECH_NAIVE_VS_DAY15_EFF_CD8_TCELL_DN
	MIR298, MIR298
	GSE42021_TCONV_PLN_VS_TREG_PRECURSORS_THYMUS_UP, GSE42021_TCONV_PLN_VS_TREG_PRECURSORS_THYMUS_UP
	HALLMARK_XENOBIOTIC_METABOLISM, HALLMARK_XENOBIOTIC_METABOLISM
	MIR513B_3P, MIR513B_3P
	GSE25087_FETAL_VS_ADULT_TCONV_DN, GSE25087_FETAL_VS_ADULT_TCONV_DN
	MIR1236_3P, MIR1236_3P
	PGF_UP.V1_UP, PGF_UP.V1_UP
	WP_GASTRIN_SIGNALING_PATHWAY, WP_GASTRIN_SIGNALING_PATHWAY
	MIR4277, MIR4277
	REACTOME_RAC2_GTPASE_CYCLE, REACTOME_RAC2_GTPASE_CYCLE
	MIR633, MIR633
	GSE37563_WT_VS_CTLA4_KO_CD4_TCELL_D4_POST_IMMUNIZATION_UP, GSE37563_WT_VS_CTLA4_KO_CD4_TCELL_D4_POST_IMMUNIZATION_UP
	MIR5087, MIR5087
	MIR1208, MIR1208
	PID_FAK_PATHWAY, PID_FAK_PATHWAY
	HP_DICARBOXYLIC_ACIDURIA, HP_DICARBOXYLIC_ACIDURIA
	MIR7153_5P, MIR7153_5P
	MIR4678, MIR4678
	HP_ABNORMAL_CIRCULATING_DICARBOXYLIC_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_DICARBOXYLIC_ACID_CONCENTRATION
	RUAN_RESPONSE_TO_TNF_DN, RUAN_RESPONSE_TO_TNF_DN
	GSE22886_TH1_VS_TH2_12H_ACT_DN, GSE22886_TH1_VS_TH2_12H_ACT_DN
	MIR4797_5P, MIR4797_5P
	GOBP_POLYSACCHARIDE_METABOLIC_PROCESS, GOBP_POLYSACCHARIDE_METABOLIC_PROCESS
	GOBP_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS
	MIR4326, MIR4326
	GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP, GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP
	GOBP_REGULATION_OF_CELLULAR_CARBOHYDRATE_METABOLIC_PROCESS, GOBP_REGULATION_OF_CELLULAR_CARBOHYDRATE_METABOLIC_PROCESS
	MIR3913_5P, MIR3913_5P
	HP_AMAUROSIS_FUGAX, HP_AMAUROSIS_FUGAX
	GOBP_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT, GOBP_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT
	MIR1229_3P, MIR1229_3P
	GSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_UP, GSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_UP
	GOBP_REGULATION_OF_GLUCOSE_METABOLIC_PROCESS, GOBP_REGULATION_OF_GLUCOSE_METABOLIC_PROCESS
	TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_6HR_UP, TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_6HR_UP
	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
	WP_RAC1PAK1P38MMP2_PATHWAY, WP_RAC1PAK1P38MMP2_PATHWAY
	GOBP_AMINO_ACID_TRANSMEMBRANE_TRANSPORT, GOBP_AMINO_ACID_TRANSMEMBRANE_TRANSPORT
	MIR449C_3P, MIR449C_3P
	GNF2_MAP2K3, GNF2_MAP2K3
	GNF2_BNIP3L, GNF2_BNIP3L
	GOBP_AMINO_ACID_IMPORT, GOBP_AMINO_ACID_IMPORT
	REACTOME_SIGNAL_TRANSDUCTION_BY_L1, REACTOME_SIGNAL_TRANSDUCTION_BY_L1
	MIR5094, MIR5094
	GOBP_AUTOPHAGOSOME_MATURATION, GOBP_AUTOPHAGOSOME_MATURATION
	YAO_HOXA10_TARGETS_VIA_PROGESTERONE_UP, YAO_HOXA10_TARGETS_VIA_PROGESTERONE_UP
	MIR3122, MIR3122
	GOMF_SOLUTE_CATION_ANTIPORTER_ACTIVITY, GOMF_SOLUTE_CATION_ANTIPORTER_ACTIVITY
	GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_CYAN_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_CYAN_UP
	MIR7975, MIR7975
	MIR6816_3P, MIR6816_3P
	HALLMARK_ANGIOGENESIS, HALLMARK_ANGIOGENESIS
	GOBP_ACIDIC_AMINO_ACID_TRANSPORT, GOBP_ACIDIC_AMINO_ACID_TRANSPORT
	GOBP_L_GLUTAMATE_TRANSMEMBRANE_TRANSPORT, GOBP_L_GLUTAMATE_TRANSMEMBRANE_TRANSPORT
	GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY
	GOBP_AMINE_TRANSPORT, GOBP_AMINE_TRANSPORT
	GOCC_GLIAL_CELL_PROJECTION, GOCC_GLIAL_CELL_PROJECTION
	CAHOY_OLIGODENDROCUTIC, CAHOY_OLIGODENDROCUTIC
	GOBP_NEGATIVE_REGULATION_OF_CARBOHYDRATE_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_CARBOHYDRATE_METABOLIC_PROCESS
	WP_CANNABINOID_RECEPTOR_SIGNALING, WP_CANNABINOID_RECEPTOR_SIGNALING
	GSE11818_WT_VS_DICER_KO_TREG_UP, GSE11818_WT_VS_DICER_KO_TREG_UP
	HP_SCAPHOCEPHALY, HP_SCAPHOCEPHALY
	GOBP_RESPONSE_TO_CATECHOLAMINE, GOBP_RESPONSE_TO_CATECHOLAMINE
	GOBP_POSITIVE_REGULATION_OF_LIPID_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_LIPID_TRANSPORT
	GOBP_NEGATIVE_REGULATION_OF_PHAGOCYTOSIS, GOBP_NEGATIVE_REGULATION_OF_PHAGOCYTOSIS
	MODULE_362, MODULE_362
	HP_CURLY_HAIR, HP_CURLY_HAIR
	GOBP_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY, GOBP_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY
	GOUYER_TUMOR_INVASIVENESS, GOUYER_TUMOR_INVASIVENESS
	GOBP_RESPONSE_TO_COBALT_ION, GOBP_RESPONSE_TO_COBALT_ION
	FRASOR_RESPONSE_TO_SERM_OR_FULVESTRANT_UP, FRASOR_RESPONSE_TO_SERM_OR_FULVESTRANT_UP
	GOMF_SOLUTE_PROTON_ANTIPORTER_ACTIVITY, GOMF_SOLUTE_PROTON_ANTIPORTER_ACTIVITY
	PID_CDC42_REG_PATHWAY, PID_CDC42_REG_PATHWAY
	HP_ATRIAL_FLUTTER, HP_ATRIAL_FLUTTER
	WIKMAN_ASBESTOS_LUNG_CANCER_UP, WIKMAN_ASBESTOS_LUNG_CANCER_UP
	GOMF_CELL_MATRIX_ADHESION_MEDIATOR_ACTIVITY, GOMF_CELL_MATRIX_ADHESION_MEDIATOR_ACTIVITY
	GOBP_L_GLUTAMATE_IMPORT_ACROSS_PLASMA_MEMBRANE, GOBP_L_GLUTAMATE_IMPORT_ACROSS_PLASMA_MEMBRANE
	GOCC_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_COMPLEX, GOCC_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_COMPLEX
	IRITANI_MAD1_TARGETS_UP, IRITANI_MAD1_TARGETS_UP
	MIR5191, MIR5191
	MODULE_562, MODULE_562
	CTGYNNCTYTAA_UNKNOWN, CTGYNNCTYTAA_UNKNOWN
	GOBP_EPITHELIAL_CELL_FATE_COMMITMENT, GOBP_EPITHELIAL_CELL_FATE_COMMITMENT
	GOMF_ALDITOL_NADPPLUS_1_OXIDOREDUCTASE_ACTIVITY, GOMF_ALDITOL_NADPPLUS_1_OXIDOREDUCTASE_ACTIVITY
	GOBP_MEMBRANE_PROTEIN_INTRACELLULAR_DOMAIN_PROTEOLYSIS, GOBP_MEMBRANE_PROTEIN_INTRACELLULAR_DOMAIN_PROTEOLYSIS
	GOBP_VERY_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE, GOBP_VERY_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE
	GOBP_REGULATION_OF_AMINO_ACID_TRANSPORT, GOBP_REGULATION_OF_AMINO_ACID_TRANSPORT
	GOMF_N_ACETYLGLUCOSAMINE_6_O_SULFOTRANSFERASE_ACTIVITY, GOMF_N_ACETYLGLUCOSAMINE_6_O_SULFOTRANSFERASE_ACTIVITY
	MIR5705, MIR5705
	GOBP_POSITIVE_REGULATION_OF_ARACHIDONIC_ACID_SECRETION, GOBP_POSITIVE_REGULATION_OF_ARACHIDONIC_ACID_SECRETION
	GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT
	REACTOME_FIBRONECTIN_MATRIX_FORMATION, REACTOME_FIBRONECTIN_MATRIX_FORMATION
	GOMF_CHEMOATTRACTANT_ACTIVITY, GOMF_CHEMOATTRACTANT_ACTIVITY
	GOMF_COLLAGEN_BINDING_INVOLVED_IN_CELL_MATRIX_ADHESION, GOMF_COLLAGEN_BINDING_INVOLVED_IN_CELL_MATRIX_ADHESION
	GOBP_COLLECTING_DUCT_DEVELOPMENT, GOBP_COLLECTING_DUCT_DEVELOPMENT
	GOBP_POSITIVE_REGULATION_OF_AMYLOID_BETA_CLEARANCE, GOBP_POSITIVE_REGULATION_OF_AMYLOID_BETA_CLEARANCE
	HOEK_T_CELL_2011_2012_TIV_ADULT_1DY_DN, HOEK_T_CELL_2011_2012_TIV_ADULT_1DY_DN
	GOBP_D_AMINO_ACID_TRANSPORT, GOBP_D_AMINO_ACID_TRANSPORT
	HP_HYPERPHOSPHATURIA, HP_HYPERPHOSPHATURIA
	GOBP_POSITIVE_REGULATION_OF_ORGANIC_ACID_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_ORGANIC_ACID_TRANSPORT
	GOMF_C_X_C_CHEMOKINE_BINDING, GOMF_C_X_C_CHEMOKINE_BINDING
	GOBP_REGULATION_OF_VERY_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE, GOBP_REGULATION_OF_VERY_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE
	PALOMERO_GSI_SENSITIVITY_DN, PALOMERO_GSI_SENSITIVITY_DN
	GOBP_CARBON_CATABOLITE_REGULATION_OF_TRANSCRIPTION, GOBP_CARBON_CATABOLITE_REGULATION_OF_TRANSCRIPTION
	GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_UPTAKE, GOBP_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_UPTAKE
	GOBP_NEGATIVE_REGULATION_OF_MUSCLE_CONTRACTION, GOBP_NEGATIVE_REGULATION_OF_MUSCLE_CONTRACTION
	GOBP_RESPONSE_TO_DOPAMINE, GOBP_RESPONSE_TO_DOPAMINE
	GOBP_GLUTAMATE_SECRETION, GOBP_GLUTAMATE_SECRETION
	GOBP_LIPID_EXPORT_FROM_CELL, GOBP_LIPID_EXPORT_FROM_CELL
	HP_PERSONALITY_DISORDER, HP_PERSONALITY_DISORDER
	GOBP_NEGATIVE_REGULATION_OF_RECEPTOR_SIGNALING_PATHWAY_VIA_JAK_STAT, GOBP_NEGATIVE_REGULATION_OF_RECEPTOR_SIGNALING_PATHWAY_VIA_JAK_STAT
	GOBP_SENSORY_PERCEPTION_OF_TEMPERATURE_STIMULUS, GOBP_SENSORY_PERCEPTION_OF_TEMPERATURE_STIMULUS
	GOBP_POSITIVE_REGULATION_OF_INHIBITORY_POSTSYNAPTIC_POTENTIAL, GOBP_POSITIVE_REGULATION_OF_INHIBITORY_POSTSYNAPTIC_POTENTIAL
	GOMF_GAMMA_AMINO BUTYRIC ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_GAMMA_AMINO BUTYRIC ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY