

	GSE37534_UNTREATED_VS_GW1929_TREATED_CD4_TCELL_PPARG1_AND_FOXP3_TRASDUCED_DN
	GSE14000_UNSTIM_VS_16H_LPS_DC_TRANSLATED_RNA_UP, GSE14000_UNSTIM_VS_16H_LPS_DC_TRANSLATED_RNA_UP
	GSE14000_UNSTIM_VS_4H_LPS_DC_TRANSLATED_RNA_UP, GSE14000_UNSTIM_VS_4H_LPS_DC_TRANSLATED_RNA_UP
	GSE37533_PPARG1_FOXP3_VS_FOXP3_TRANSDUCED_CD4_TCELL_PIOGLITAZONE_TREATED_DN, GSE37533_PPARG1_FOXP3_VS_FOXP3_TRANSDUCED_CD4_TCELL_PIOGLITAZONE_TREATED_DN
	GSE39022_LN_VS_SPLEEN_DC_DN, GSE39022_LN_VS_SPLEEN_DC_DN
	GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_2H_UP
	GSE22886_CD8_VS_CD4_NAIVE_TCELL_UP, GSE22886_CD8_VS_CD4_NAIVE_TCELL_UP
	GSE42021_TREG_VS_TCONV_PLN_DN, GSE42021_TREG_VS_TCONV_PLN_DN
	GSE3982_EOSINOPHIL_VS_NKCELL_DN, GSE3982_EOSINOPHIL_VS_NKCELL_DN
	GSE1432_CTRL_VS_IFNG_6H_MICROGLIA_UP, GSE1432_CTRL_VS_IFNG_6H_MICROGLIA_UP
	GSE5589_LPS_VS_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_DN, GSE5589_LPS_VS_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_DN
	GSE42021_CD24HL_VS_CD24INT_TREG_THYMUS_UP, GSE42021_CD24HL_VS_CD24INT_TREG_THYMUS_UP
	GSE1432_1H_VS_6H_IFNG_MICROGLIA_UP, GSE1432_1H_VS_6H_IFNG_MICROGLIA_UP
	GSE42021_CD24HL_VS_CD24INT_TCONV_THYMUS_UP, GSE42021_CD24HL_VS_CD24INT_TCONV_THYMUS_UP
	GSE42021_CD24HL_VS_CD24LOW_TCONV_THYMUS_UP, GSE42021_CD24HL_VS_CD24LOW_TCONV_THYMUS_UP
	GSE21360_NAIVE_VS_TERTIARY_MEMORY_CD8_TCELL_UP, GSE21360_NAIVE_VS_TERTIARY_MEMORY_CD8_TCELL_UP
	GSE41978_ID2_KO_AND_BIM_KO_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN, GSE41978_ID2_KO_AND_BIM_KO_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN
	GSE26495_PD1HIGH_VS_PD1LOW_CD8_TCELL_DN, GSE26495_PD1HIGH_VS_PD1LOW_CD8_TCELL_DN
	GSE37533_PPARG1_FOXP3_VS_PPARG2_FOXP3_TRANSDUCED_CD4_TCELL_PIOGLITAZONE_TREATED_UP, GSE37533_PPARG1_FOXP3_VS_PPARG2_FOXP3_TRANSDUCED_CD4_TCELL_PIOGLITAZONE_TREATED_UP
	DESERT_PERIVENOUS_HEPATOCELLULAR_CARCINOMA_SUBCLASS_UP, DESERT_PERIVENOUS_HEPATOCELLULAR_CARCINOMA_SUBCLASS_UP
	GSE19401_PLN_VS_PEYERS_PATCH_FOLLICULAR_DC_UP, GSE19401_PLN_VS_PEYERS_PATCH_FOLLICULAR_DC_UP
	GSE42021_TREG_PLN_VS_TREG_PRECURSORS_THYMUS_UP, GSE42021_TREG_PLN_VS_TREG_PRECURSORS_THYMUS_UP
	GSE3982_CTRL_VS_LPS_48H_DC_UP, GSE3982_CTRL_VS_LPS_48H_DC_UP
	ACAWNRNRSRCGG_UNKNOWN, ACAWNRNRSRCGG_UNKNOWN
	GSE42021_TREG_PLN_VS_CD24LO_TREG_THYMUS_UP, GSE42021_TREG_PLN_VS_CD24LO_TREG_THYMUS_UP
	GALE_APL_WITH_FLT3_MUTATED_UP, GALE_APL_WITH_FLT3_MUTATED_UP
	GSE45365_CD8A_DC_VS_CD11B_DC_IFNAR_KO_MCMV_INFECTION_UP, GSE45365_CD8A_DC_VS_CD11B_DC_IFNAR_KO_MCMV_INFECTION_UP
	MORF_RFC5, MORF_RFC5
	GSE3982_BASOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN, GSE3982_BASOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN
	DROSHA_TARGET_GENES, DROSHA_TARGET_GENES
	GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG2_AND_FOXP3 TRASDUCED_UP, GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG2_AND_FOXP3 TRASDUCED_UP
	GOBP_PALLIUM_DEVELOPMENT, GOBP_PALLIUM_DEVELOPMENT
	FREAC7_01, FREAC7_01
	MIR3121_5P, MIR3121_5P
	GSE37301_COMMON_LYMPHOID_PROGENITOR_VS_RAG2_KO_NK_CELL_UP, GSE37301_COMMON_LYMPHOID_PROGENITOR_VS_RAG2_KO_NK_CELL_UP
	COATES_MACROPHAGE_M1_VS_M2_DN, COATES_MACROPHAGE_M1_VS_M2_DN
	GUTIERREZ_CHRONIC_LYMPHOCYTIC_LEUKEMIA_DN, GUTIERREZ_CHRONIC_LYMPHOCYTIC_LEUKEMIA_DN
	SHETH_LIVER_CANCER_VS_TXNIP_LOSS_PAM6, SHETH_LIVER_CANCER_VS_TXNIP_LOSS_PAM6
	MIR3929, MIR3929
	GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BLUE_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_BLUE_DN
	GSE37319_WT_VS_RC3H1_KO_CD44LOW_CD8_TCELL_DN, GSE37319_WT_VS_RC3H1_KO_CD44LOW_CD8_TCELL_DN
	BERNARD_PPAPDC1B_TARGETS_UP, BERNARD_PPAPDC1B_TARGETS_UP
	MORF_RUNX1, MORF_RUNX1
	HP_GAIT_IMBALANCE, HP_GAIT_IMBALANCE
	GOBP_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
	GOBP_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION, GOBP_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION
	ROYLANCE_BREAST_CANCER_16Q_COPY_NUMBER_UP, ROYLANCE_BREAST_CANCER_16Q_COPY_NUMBER_UP
	MIR655_5P, MIR655_5P
	IIZUKA_LIVER_CANCER_PROGRESSION_G1_G2_DN, IIZUKA_LIVER_CANCER_PROGRESSION_G1_G2_DN
	DESCARTES_FETAL_INTESTINE_ERYTHROBLASTS, DESCARTES_FETAL_INTESTINE_ERYTHROBLASTS
	KANNAN_TP53_TARGETS_UP, KANNAN_TP53_TARGETS_UP
	JAZAERI_BREAST_CANCER_BRCA1_VS_BRCA2_DN, JAZAERI_BREAST_CANCER_BRCA1_VS_BRCA2_DN
	BILBAN_B_CLL_LPL_UP, BILBAN_B_CLL_LPL_UP
	GOBP_NEGATIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
	GOBP_RECEPTOR_CLUSTERING, GOBP_RECEPTOR_CLUSTERING
	GOBP_POSITIVE_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA, GOBP_POSITIVE_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA
	GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE
	YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_16, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_16
	BANDRES_RESPONSE_TO_CARMUSTIN_MGMT_48HR_DN, BANDRES_RESPONSE_TO_CARMUSTIN_MGMT_48HR_DN
	GOCC_DENDRITE_CYTOPLASM, GOCC_DENDRITE_CYTOPLASM
	MIR3122, MIR3122
	AMIT_SERUM_RESPONSE_480_MCF10A, AMIT_SERUM_RESPONSE_480_MCF10A
	REACTOME_POLB_DEPENDENT_LONG_PATCH_BASE_EXCISION_REPAIR, REACTOME_POLB_DEPENDENT_LONG_PATCH_BASE_EXCISION_REPAIR
	RYAAAKNNNNNNTTGW_UNKNOWN, RYAAAKNNNNNNTTGW_UNKNOWN
	GOBP_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION
	HP_HYPOSMIA, HP_HYPOSMIA
	SEITZ_NEOPLASTIC_TRANSFORMATION_BY_8P_DELETION_DN, SEITZ_NEOPLASTIC_TRANSFORMATION_BY_8P_DELETION_DN
	MIR5007_5P, MIR5007_5P
	REN_ALVEOLAR_RHABDOMYOSARCOMA_UP, REN_ALVEOLAR_RHABDOMYOSARCOMA_UP
	BROWNE_HCMV_INFECTION_8HR_DN, BROWNE_HCMV_INFECTION_8HR_DN
	GOBP_NEGATIVE_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS, GOBP_NEGATIVE_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS
	GOCC_NPBAF_COMPLEX, GOCC_NPBAF_COMPLEX
	GOCC_MLL3_4_COMPLEX, GOCC_MLL3_4_COMPLEX
	HP_INSOMNIA, HP_INSOMNIA
	YORDY_RECIPROCAL_REGULATION_BY_ETS1_AND_SP100_UP, YORDY_RECIPROCAL_REGULATION_BY_ETS1_AND_SP100_UP
	GOBP_KERATAN_SULFATE_BIOSYNTHETIC_PROCESS, GOBP_KERATAN_SULFATE_BIOSYNTHETIC_PROCESS
	REACTOME_KERATAN_SULFATE_BIOSYNTHESIS, REACTOME_KERATAN_SULFATE_BIOSYNTHESIS
	HP_NON_MIDLINE_CLEFT_PALATE, HP_NON_MIDLINE_CLEFT_PALATE
	GOCC_CORTICAL_ACTIN_CYTOSKELETON, GOCC_CORTICAL_ACTIN_CYTOSKELETON
	MODULE_200, MODULE_200
	GOBP_REGULATION_OF_NMDA_RECEPTOR_ACTIVITY, GOBP_REGULATION_OF_NMDA_RECEPTOR_ACTIVITY
	MIR3657, MIR3657
	GOBP_REGULATION_OF_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL, GOBP_REGULATION_OF_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL
	GOBP_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
	REACTOME_CHAPERONE_MEDIATED_AUTOPHAGY, REACTOME_CHAPERONE_MEDIATED_AUTOPHAGY
	GOCC_GLIAL_CELL_PROJECTION, GOCC_GLIAL_CELL_PROJECTION
	YAP1_DN, YAP1_DN
	HP_FEMALE_SEXUAL_DYSFUNCTION, HP_FEMALE_SEXUAL_DYSFUNCTION
	ASTON_MAJOR_DEPRESSIVE_DISORDER_UP, ASTON_MAJOR_DEPRESSIVE_DISORDER_UP
	chr7q36, chr7q36
	MIR6742_5P, MIR6742_5P
	HP_LIMITED_ELBOW_FLEXION, HP_LIMITED_ELBOW_FLEXION
	GOMF_PROTEIN_TYROSINE_KINASE_ACTIVATOR_ACTIVITY, GOMF_PROTEIN_TYROSINE_KINASE_ACTIVATOR_ACTIVITY
	CROMER_TUMORIGENESIS_UP, CROMER_TUMORIGENESIS_UP
	GOBP_POSITIVE_REGULATION_OF_NEURAL_PRECURSOR_CELL_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_NEURAL_PRECURSOR_CELL_PROLIFERATION
	HP ABDOMINAL_WALL_MUSCLE_WEAKNESS, HP ABDOMINAL_WALL_MUSCLE_WEAKNESS
	SASAKI_TARGETS_OF_TP73_AND_TP63, SASAKI_TARGETS_OF_TP73_AND_TP63
	GOBP_OXALATE_TRANSPORT, GOBP_OXALATE_TRANSPORT
	GOMF_OXALATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_OXALATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
	KEGG_REGULATION_OF_AUTOPHAGY, KEGG_REGULATION_OF_AUTOPHAGY
	GOBP_POSITIVE_REGULATION_OF_ORGAN_GROWTH, GOBP_POSITIVE_REGULATION_OF_ORGAN_GROWTH
	GOBP_CARDIAC_MUSCLE_FIBER_DEVELOPMENT, GOBP_CARDIAC_MUSCLE_FIBER_DEVELOPMENT
	GOCC_ASTROCYTE_PROJECTION, GOCC_ASTROCYTE_PROJECTION
	WP_1Q211_COPY_NUMBER_VARIATION_SYNDROME, WP_1Q211_COPY_NUMBER_VARIATION_SYNDROME
	GOBP_RESPONSE_TO_IRON_ION, GOBP_RESPONSE_TO_IRON_ION
	GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS
	REACTOME_LONG_TERM_POTENTIATION, REACTOME_LONG_TERM_POTENTIATION
	RNTCANNRNNYNATTW_UNKNOWN, RNTCANNRNNYNATTW_UNKNOWN
	GOBP_L_ASPARTATE_TRANSMEMBRANE_TRANSPORT, GOBP_L_ASPARTATE_TRANSMEMBRANE_TRANSPORT
	GOMF_SMALL_MOLECULE_SENSOR_ACTIVITY, GOMF_SMALL_MOLECULE_SENSOR_ACTIVITY
	HP_PHOTOSENSITIVE_MYOCLONIC_SEIZURE, HP_PHOTOSENSITIVE_MYOCLONIC_SEIZURE
	GOBP_PROTEIN_LOCALIZATION_TO_POSTSYNAPSE, GOBP_PROTEIN_LOCALIZATION_TO_POSTSYNAPSE
	GOBP_CARDIAC_MUSCLE_CELL_CARDIAC_MUSCLE_CELL_ADHESION, GOBP_CARDIAC_MUSCLE_CELL_CARDIAC_MUSCLE_CELL_ADHESION
	HP_LETHAL_SKELETAL_DYSPLASIA, HP_LETHAL_SKELETAL_DYSPLASIA
	GOBP_CARDIOCYTE_DIFFERENTIATION, GOBP_CARDIOCYTE_DIFFERENTIATION
	GARCIA_PINERES_PBMC_HPV_16_L1_VLP_AGE_18_25YO_12MO_CORRELATED_WITH_ANTIBODY_RESPONSE_NEGATIVE, GARCIA_PINERES_PBMC_HPV_16_L1_VLP_AGE_18_25YO_12MO_CORRELATED_WITH_ANTIBODY_RESPONSE_NEGATIVE
	GOBP_NEGATIVE_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT, GOBP_NEGATIVE_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT
	GOBP_RESPONSE_TO_MERCURY_ION, GOBP_RESPONSE_TO_MERCURY_ION
	HP_PEG_SHAPED_MAXILLARY_LATERAL_INCISORS, HP_PEG_SHAPED_MAXILLARY_LATERAL_INCISORS
	GOBP_POSITIVE_REGULATION_OF_HEART_GROWTH, GOBP_POSITIVE_REGULATION_OF_HEART_GROWTH
	VALK_AML_CLUSTER_8, VALK_AML_CLUSTER_8
	GOBP_POSITIVE_REGULATION_OF_FERTILIZATION, GOBP_POSITIVE_REGULATION_OF_FERTILIZATION
	GOBP_REGULATION_OF_UBIQUITIN_SPECIFIC_PROTEASE_ACTIVITY, GOBP_REGULATION_OF_UBIQUITIN_SPECIFIC_PROTEASE_ACTIVITY
	HP_PSEUDOHYPALDOSTERONISM, HP_PSEUDOHYPALDOSTERONISM
VISCERAL_ADIPOSE_TISSUE_TCONV_DN, GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TCONV_DN	