

	GSE22103_UNSTIM_VS_LPS_STIM_NEUTROPHIL_UP, GSE22103_UNSTIM_VS_LPS_STIM_NEUTROPHIL_UP
	GSE22589_HEALTHY_VS_SIV_INFECTED_DC_UP, GSE22589_HEALTHY_VS_SIV_INFECTED_DC_UP
	GSE40274_XBP1_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_XBP1_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
	GSE17721_0.5H_VS_4H_PAM3CSK4_BMDC_UP, GSE17721_0.5H_VS_4H_PAM3CSK4_BMDC_UP
	GSE22935_UNSTIM_VS_12H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_DN, GSE22935_UNSTIM_VS_12H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_DN
	GSE17721_LPS_VS_CPG_12H_BMDC_UP, GSE17721_LPS_VS_CPG_12H_BMDC_UP
	GSE23398_WT_VS_IL2_KO_CD4_TCELL_SCURFY_MOUSE_UP, GSE23398_WT_VS_IL2_KO_CD4_TCELL_SCURFY_MOUSE_UP
	GOBP_PEPTIDYL_LYSINE_ACETYLATION, GOBP_PEPTIDYL_LYSINE_ACETYLATION
	GSE14415_FOXP3_KO_NATURAL_TREG_VS_TCONV_DN, GSE14415_FOXP3_KO_NATURAL_TREG_VS_TCONV_DN
	GSE7852_TREG_VS_TCONV_THYMUS_DN, GSE7852_TREG_VS_TCONV_THYMUS_DN
	GSE15330_LYMPHOID_MULTIPOTENT_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN, GSE15330_LYMPHOID_MULTIPOTENT_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN
	GSE17721_PAM3CSK4_VS_CPG_1H_BMDC_UP, GSE17721_PAM3CSK4_VS_CPG_1H_BMDC_UP
	GSE18281_PERIMEDULLARY_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_UP, GSE18281_PERIMEDULLARY_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_UP
	GSE40274_FOXP3_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_FOXP3_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
	BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_TRANS, BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_TRANS
	GSE11961_MARGINAL_ZONE_BCELL_VS_PLASMA_CELL_DAY7_DN, GSE11961_MARGINAL_ZONE_BCELL_VS_PLASMA_CELL_DAY7_DN
	MORF_HEAB, MORF_HEAB
	MIR324_3P, MIR324_3P
	GOBP_HISTONE_H4_ACETYLATION, GOBP_HISTONE_H4_ACETYLATION
	MIR1255B_2_3P, MIR1255B_2_3P
	HP_HYPOPLASIA_OF_THE_BRAINSTEM, HP_HYPOPLASIA_OF_THE_BRAINSTEM
	HUTTMANN_B_CLL_POOR_SURVIVAL_DN, HUTTMANN_B_CLL_POOR_SURVIVAL_DN
	GOMF_REPRESSING_TRANSCRIPTION_FACTOR_BINDING, GOMF_REPRESSING_TRANSCRIPTION_FACTOR_BINDING
	MIR190B_5P, MIR190B_5P
	HP_APLASTIC_HYPOPLASTIC_TOENAIL, HP_APLASTIC_HYPOPLASTIC_TOENAIL
	MIR887_5P, MIR887_5P
	HP_OPTIC_NEUROPATHY, HP_OPTIC_NEUROPATHY
	GSE14350_TREG_VS_TEFF_DN, GSE14350_TREG_VS_TEFF_DN
	HP_ABNORMAL_OVARIAN_MORPHOLOGY, HP_ABNORMAL_OVARIAN_MORPHOLOGY
	HP_AGANGLIONIC_MEGACOLON, HP_AGANGLIONIC_MEGACOLON
	MIR190A_5P, MIR190A_5P
	MIR3187_3P, MIR3187_3P
	GSE17721_LPS_VS_PAM3CSK4_2H_BMDC_DN, GSE17721_LPS_VS_PAM3CSK4_2H_BMDC_DN
	HP_CONGENITAL_HEPATIC_FIBROSIS, HP_CONGENITAL_HEPATIC_FIBROSIS
	GSE28237_EARLY_VS_LATE_GC_BCELL_UP, GSE28237_EARLY_VS_LATE_GC_BCELL_UP
	MIR15B_3P, MIR15B_3P
	GOBP_REGULATION_OF_MITOCHONDRIAL_GENE_EXPRESSION, GOBP_REGULATION_OF_MITOCHONDRIAL_GENE_EXPRESSION
	HP_ANIRIDIA, HP_ANIRIDIA
	MIR320E, MIR320E
	HP_2_3_TOE_SYNDACTYLY, HP_2_3_TOE_SYNDACTYLY
	HP_MENINGOCELE, HP_MENINGOCELE
	HP_APLASIA_CUTIS_CONGENITA, HP_APLASIA_CUTIS_CONGENITA
	HP_ECTRODACTYLY, HP_ECTRODACTYLY
	WP_NOTCH_SIGNALING, WP_NOTCH_SIGNALING
	HP_METATARSUS_ADDUCTUS, HP_METATARSUS_ADDUCTUS
	QUEREC_MODEL_PBMCI_YF_17D_VACCINE_AGE_18_45_7DY_PREDICTIVE, QUEREC_MODEL_PBMCI_YF_17D_VACCINE_AGE_18_45_7DY_PREDICTIVE
	GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_BALBC_MOUSE_DN, GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_BALBC_MOUSE_DN
	MIR3923, MIR3923
	HP_MEDIAL_FLARING_OF_THE_EYEBROW, HP_MEDIAL_FLARING_OF_THE_EYEBROW
	GOBP_BROWN_FAT_CELL_DIFFERENTIATION, GOBP_BROWN_FAT_CELL_DIFFERENTIATION
	THRAP3_TARGET_GENES, THRAP3_TARGET_GENES
	HP_ABNORMALITY_OF_THE_PONS, HP_ABNORMALITY_OF_THE_PONS
	HP_GENERALIZED_JOINT_LAXITY, HP_GENERALIZED_JOINT_LAXITY
	GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREEN_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREEN_DN
	HP_APLASIA_HYPOPLASIA_OF_THE_OVARY, HP_APLASIA_HYPOPLASIA_OF_THE_OVARY
	GOBP_POSITIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE, GOBP_POSITIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE
	BIOCARTA_ETS_PATHWAY, BIOCARTA_ETS_PATHWAY
	HP_ABSENT_HAND, HP_ABSENT_HAND
	GOMF_PROTEIN_KINASE_A_REGULATORY_SUBUNIT_BINDING, GOMF_PROTEIN_KINASE_A_REGULATORY_SUBUNIT_BINDING
	HP_PROTRUDING_TONGUE, HP_PROTRUDING_TONGUE
	HP_CALVARIAL_SKULL_DEFECT, HP_CALVARIAL_SKULL_DEFECT
	HP_PULMONARY_ARTERY_ATRESIA, HP_PULMONARY_ARTERY_ATRESIA
	WINNEPENINCKX_MELANOMA_METASTASIS_DN, WINNEPENINCKX_MELANOMA_METASTASIS_DN
	XU_HGF_SIGNALING_NOT_VIA_AKT1_6HR, XU_HGF_SIGNALING_NOT_VIA_AKT1_6HR
	PID_INTEGRIN_A9B1_PATHWAY, PID_INTEGRIN_A9B1_PATHWAY
	GOBP_NEUTROPHIL_EXTRAVASATION, GOBP_NEUTROPHIL_EXTRAVASATION
	GOBP_AORTA_DEVELOPMENT, GOBP_AORTA_DEVELOPMENT
	GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_GLYCOPROTEIN, GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_GLYCOPROTEIN
	HP_ABNORMAL_LIBIDO, HP_ABNORMAL_LIBIDO
	HP_ABNORMAL_HAND_BONE_OSSIFICATION, HP_ABNORMAL_HAND_BONE_OSSIFICATION
	HP_DYSPAREUNIA, HP_DYSPAREUNIA
	GOBP_HISTONE_H4_K20_METHYLATION, GOBP_HISTONE_H4_K20_METHYLATION
	HP_11_PAIRS_OF_RIBS, HP_11_PAIRS_OF_RIBS
	TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_SUSTAINED_IN_MONOCYTE_UP, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_SUSTAINED_IN_MONOCYTE_UP
	GOBP_NUCLEOSIDE_TRANSPORT, GOBP_NUCLEOSIDE_TRANSPORT
	HP_HYPOTHALAMIC_GONADOTROPIN_RELEASING_HORMONE_DEFICIENCY, HP_HYPOTHALAMIC_GONADOTROPIN_RELEASING_HORMONE_DEFICIENCY
	GOBP_GLYCOSPHINGOLIPID_BIOSYNTHETIC_PROCESS, GOBP_GLYCOSPHINGOLIPID_BIOSYNTHETIC_PROCESS
	GOBP_POSITIVE_REGULATION_OF_OSTEOBLAST_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_OSTEOBLAST_DIFFERENTIATION
	GOBP_SUBSTRATE_DEPENDENT_CELL_MIGRATION, GOBP_SUBSTRATE_DEPENDENT_CELL_MIGRATION
	HP_ABNORMALITY_OF_CARPAL_BONE_OSSIFICATION, HP_ABNORMALITY_OF_CARPAL_BONE_OSSIFICATION
	GOBP_REGULATION_OF_STRIATED_MUSCLE_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_STRIATED_MUSCLE_CELL_DIFFERENTIATION
	HP_MICROSCOPIC_HEMATURIA, HP_MICROSCOPIC_HEMATURIA
	HP_JOINT_SUBLUXATION, HP_JOINT_SUBLUXATION
	MATZUK_IMPLANTATION_AND_UTERINE, MATZUK_IMPLANTATION_AND_UTERINE
	REACTOME_SEMA3A_PLEXIN_REPULSION_SIGNALING_BY_INHIBITING_INTEGRIN_ADHESION, REACTOME_SEMA3A_PLEXIN_REPULSION_SIGNALING_BY_INHIBITING_INTEGRIN_ADHESION
	GOBP_SIALYLATION, GOBP_SIALYLATION
	HOEK_MYELOID_DENDRITIC_CELL_2011_2012_TIV_ADULT_7DY_DN, HOEK_MYELOID_DENDRITIC_CELL_2011_2012_TIV_ADULT_7DY_DN
	GAO_LARGE_INTESTINE_ADULT_CF_GOBLET_CELL_SUBTYPE_1, GAO_LARGE_INTESTINE_ADULT_CF_GOBLET_CELL_SUBTYPE_1
	MIR6787_5P, MIR6787_5P
	HP_BREAST_HYPOPLASIA, HP_BREAST_HYPOPLASIA
	BROWNE_HCMV_INFECTION_30MIN_UP, BROWNE_HCMV_INFECTION_30MIN_UP
	MODULE_465, MODULE_465
	HP_APLASIA_CUTIS_CONGENITA_OF_SCALP, HP_APLASIA_CUTIS_CONGENITA_OF_SCALP
	MIR6767_3P, MIR6767_3P
	GOBP_REGULATION_OF_SOMITOGENESIS, GOBP_REGULATION_OF_SOMITOGENESIS
	GOMF_DOUBLE_STRANDED_METHYLATED_DNA_BINDING, GOMF_DOUBLE_STRANDED_METHYLATED_DNA_BINDING
	GOMF_ESTROGEN_16_ALPHA_HYDROXYLASE_ACTIVITY, GOMF_ESTROGEN_16_ALPHA_HYDROXYLASE_ACTIVITY
	HP_POSTERIOR_PITUITARY_DYSGENESIS, HP_POSTERIOR_PITUITARY_DYSGENESIS
	REACTOME_POU5F1_OCT4_SOX2_NANOG_REPRESS_GENES_RELATED_TO_DIFFERENTIATION, REACTOME_POU5F1_OCT4_SOX2_NANOG_REPRESS_GENES_RELATED_TO_DIFFERENTIATION
	GRAHAM_CML_DIVIDING_VS_NORMAL_DIVIDING_DN, GRAHAM_CML_DIVIDING_VS_NORMAL_DIVIDING_DN
	WP_SOMITOGENESIS_IN_THE_CONTEXT_OF_SPONDYLOCOSTAL_DYSOSTOSIS, WP_SOMITOGENESIS_IN_THE_CONTEXT_OF_SPONDYLOCOSTAL_DYSOSTOSIS
	GOBP_OMEGA_HYDROXYLASE_P450_PATHWAY, GOBP_OMEGA_HYDROXYLASE_P450_PATHWAY
	REACTOME_SYNTHESIS_OF_16_20_HYDROXYEICOSATETRAENOIC_ACIDS_HETE, REACTOME_SYNTHESIS_OF_16_20_HYDROXYEICOSATETRAENOIC_ACIDS_HETE
	GOBP_REGULATION_OF_ODONTOGENESIS_OF_DENTIN_CONTAINING_TOOTH, GOBP_REGULATION_OF_ODONTOGENESIS_OF_DENTIN_CONTAINING_TOOTH
	GOMF_POTASSIUM_CHLORIDE_SYMPORTER_ACTIVITY, GOMF_POTASSIUM_CHLORIDE_SYMPORTER_ACTIVITY
	GOBP_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORT, GOBP_NUCLEOSIDE_TRANSMEMBRANE_TRANSPORT
	HP_EUNUCHOID_HABITUS, HP_EUNUCHOID_HABITUS
	HP_SPARSE_BODY_HAIR, HP_SPARSE_BODY_HAIR