

	GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP
	GSE32533_WT_VS_MIR17_KO_ACT_CD4_TCELL_DN, GSE32533_WT_VS_MIR17_KO_ACT_CD4_TCELL_DN
	GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_WITH_IFNA_STIM_UP
	GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP, GSE3400_UNTREATED_VS_IFNB_TREATED_MEF_UP
	GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_DN, GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_DN
	GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_OT1_VS_HY_CD8AB_THYMOCYTE_RTOC_CULTURE_DN
	GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN, GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_DN
	GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP
	GSE37416_12H_VS_48H_F_TULARENSIS_1VS_NEUTROPHIL_UP, GSE37416_12H_VS_48H_F_TULARENSIS_1VS_NEUTROPHIL_UP
	GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN
	GSE22886_TH1_VS_TH2_48H_ACT_DN, GSE22886_TH1_VS_TH2_48H_ACT_DN
	GSE3982_DC_VS_EFF_MEMORY_CD4_TCELL_UP, GSE3982_DC_VS_EFF_MEMORY_CD4_TCELL_UP
	GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP
	GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_DN, GSE2128_C57BL6_VS_NOD_THYMOCYTE_MIMETOPE_NEGATIVE_SELECTION_DN
	GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_48H_UP
	GSE24142_DN2_VS_DN3_THYMOCYTE_ADULT_DN, GSE24142_DN2_VS_DN3_THYMOCYTE_ADULT_DN
	PECE_MAMMARY_STEM_CELL_DN, PECE_MAMMARY_STEM_CELL_DN
	GSE42021_TCONV_PLN_VS_TREG_PRECURSORS_THYMUS_UP, GSE42021_TCONV_PLN_VS_TREG_PRECURSORS_THYMUS_UP
	GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_6H_UP
	GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_UP, GSE2128_C57BL6_VS_NOD_CD4CD8_DP_THYMOCYTE_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
	STEARMAN_LUNG_CANCER_EARLY_VS_LATE_UP, STEARMAN_LUNG_CANCER_EARLY_VS_LATE_UP
	GSE5679_CTRL_VS_RARA_AGONIST_AM580_TREATED_DC_DN, GSE5679_CTRL_VS_RARA_AGONIST_AM580_TREATED_DC_DN
	GSE15767_MED_VS_SCS_MAC_LN_UP, GSE15767_MED_VS_SCS_MAC_LN_UP
	PUIFFE_INVASION_INHIBITED_BY_ASCITES_DN, PUIFFE_INVASION_INHIBITED_BY_ASCITES_DN
	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
	GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_3H_DN, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_3H_DN
	MIR6768_3P, MIR6768_3P
	MIR6505_5P, MIR6505_5P
	GSE37301_RAG2_KO_VS_RAG2_AND_ETSI_KO_NK_CELL_UP, GSE37301_RAG2_KO_VS_RAG2_AND_ETSI_KO_NK_CELL_UP
	GSE42021_TREG_PLN_VS_TREG_PRECURSORS_THYMUS_UP, GSE42021_TREG_PLN_VS_TREG_PRECURSORS_THYMUS_UP
	GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_UP, GSE21927_BALBC_VS_C57BL6_MONOCYTE_SPLEEN_UP
	MIR8065, MIR8065
	MIR1208, MIR1208
	HP_THORACIC_HYPOPLASIA, HP_THORACIC_HYPOPLASIA
	REACTOME_RUNX1_REGULATES_GENES_INVOLVED_IN_MEGAKARYOCYTE_DIFFERENTIATION_AND_PLATELET_FUNCTION, REACTOME_RUNX1_REGULATES_GENES_INVOLVED_IN_MEGAKARYOCYTE_DIFFERENTIATION_AND_PLATELET_FUNCTION
	MIR12132, MIR12132
	GOMF_HORMONE_RECEPTOR_BINDING, GOMF_HORMONE_RECEPTOR_BINDING
	MIR558, MIR558
	MIR4727_5P, MIR4727_5P
	MIR4456, MIR4456
	MIR642B_5P, MIR642B_5P
	GOBP_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION
	GOBP_POSITIVE_REGULATION_OF_HEMOPOIESIS, GOBP_POSITIVE_REGULATION_OF_HEMOPOIESIS
	GSE32423_CTRL_VS_IL7_MEMORY_CD8_TCELL_DN, GSE32423_CTRL_VS_IL7_MEMORY_CD8_TCELL_DN
	MIR5001_3P, MIR5001_3P
	MENON_FETAL_KIDNEY_9_ENDOTHELIAL_CELLS, MENON_FETAL_KIDNEY_9_ENDOTHELIAL_CELLS
	MIR7162_5P, MIR7162_5P
	GOBP_AMINOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_AMINOGLYCAN_BIOSYNTHETIC_PROCESS
	GOBP_PROTEOGLYCAN_METABOLIC_PROCESS, GOBP_PROTEOGLYCAN_METABOLIC_PROCESS
	MIR211_3P, MIR211_3P
	GOMF_INTEGRIN_BINDING, GOMF_INTEGRIN_BINDING
	GSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN
	REACTOME_RHOG_GTPASE_CYCLE, REACTOME_RHOG_GTPASE_CYCLE
	MIR495_5P, MIR495_5P
	GOBP_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS, GOBP_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS
	GOBP_RESPONSE_TO_OSMOTIC_STRESS, GOBP_RESPONSE_TO_OSMOTIC_STRESS
	MIR3682_3P, MIR3682_3P
	MIR1912_3P, MIR1912_3P
	GOBP_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
	RELA_DN.V1_UP, RELA_DN.V1_UP
	GRAHAM_NORMAL_QUIESCENT_VS_NORMAL_DIVIDING_UP, GRAHAM_NORMAL_QUIESCENT_VS_NORMAL_DIVIDING_UP
	REACTOME_ELASTIC_FIBRE_FORMATION, REACTOME_ELASTIC_FIBRE_FORMATION
	GOMF_CYTOKINE_BINDING, GOMF_CYTOKINE_BINDING
	GOBP_CELL_ADHESION_MEDIATED_BY_INTEGRIN, GOBP_CELL_ADHESION_MEDIATED_BY_INTEGRIN
	MIR148B_5P, MIR148B_5P
	REACTOME_CA2_PATHWAY, REACTOME_CA2_PATHWAY
	WP_G_PROTEIN_SIGNALING_PATHWAYS, WP_G_PROTEIN_SIGNALING_PATHWAYS
	MIR1295B_5P, MIR1295B_5P
	GOBP_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS
	MIR584_5P, MIR584_5P
	MIR1178_3P, MIR1178_3P
	MIR5588_3P, MIR5588_3P
	GSE34205_RSV_VS_FLU_INF_INFANT_PPMC_UP, GSE34205_RSV_VS_FLU_INF_INFANT_PPMC_UP
	ZHAN_MULTIPLE_MYELOMA_CD1_VS_CD2_DN, ZHAN_MULTIPLE_MYELOMA_CD1_VS_CD2_DN
	GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_POSITIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
	MIR597_5P, MIR597_5P
	TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_6HR_UP, TAKEDA_TARGETS_OF_NUP98_HOXA9_FUSION_6HR_UP
	HP_ANKYLOSIS, HP_ANKYLOSIS
	ZHU_CMV_8_HR_DN, ZHU_CMV_8_HR_DN
	GOBP_REGULATION_OF_WOUND_HEALING, GOBP_REGULATION_OF_WOUND_HEALING
	REACTOME_MOLECULES_ASSOCIATED_WITH_ELASTIC_FIBRES, REACTOME_MOLECULES_ASSOCIATED_WITH_ELASTIC_FIBRES
	CASORELLI_APL_SECONDARY_VS_DE_NOVO_UP, CASORELLI_APL_SECONDARY_VS_DE_NOVO_UP
	LAKE_ADULT_KIDNEY_C25_ENDOTHELIAL_CELLS_UNASSIGNED, LAKE_ADULT_KIDNEY_C25_ENDOTHELIAL_CELLS_UNASSIGNED
	GOBP_NEPHRON_DEVELOPMENT, GOBP_NEPHRON_DEVELOPMENT
	MIR4638_3P, MIR4638_3P
	MIR601, MIR601
	HP_BILATERAL_TONIC_CLONIC_SEIZURE_WITH_GENERALIZED_ONSET, HP_BILATERAL_TONIC_CLONIC_SEIZURE_WITH_GENERALIZED_ONSET
	GOBP_RESPONSE_TO_PROGESTERONE, GOBP_RESPONSE_TO_PROGESTERONE
	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
	HP_BROAD_EYEBROW, HP_BROAD_EYEBROW
	WP_MECHANOREGULATION_AND_PATHOLOGY_OF_YAPTAZ_VIA_HIPPO_AND_NONHIPPO_MECHANISMS, WP_MECHANOREGULATION_AND_PATHOLOGY_OF_YAPTAZ_VIA_HIPPO_AND_NONHIPPO_MECHANISMS
	WP_EXTRACELLULAR_MATRIX_BINDING, GOMF_EXTRACELLULAR_MATRIX_BINDING
	WP_NANOPARTICLEMEDIATED_ACTIVATION_OF_RECEPTOR_SIGNALING, WP_NANOPARTICLEMEDIATED_ACTIVATION_OF_RECEPTOR_SIGNALING
	HP_PLEURAL_EFFUSION, HP_PLEURAL_EFFUSION
	GOBP_MESODERM_DEVELOPMENT, GOBP_MESODERM_DEVELOPMENT
	GOMF_CORECEPTOR_ACTIVITY, GOMF_CORECEPTOR_ACTIVITY
	GUENTHER_GROWTH_SPHERICAL_VS_ADHERENT_DN, GUENTHER_GROWTH_SPHERICAL_VS_ADHERENT_DN
	HP_SHORT_RIBS, HP_SHORT_RIBS
	GOMF_MANNOSIDASE_ACTIVITY, GOMF_MANNOSIDASE_ACTIVITY
	GOBP_POSITIVE_REGULATION_OF_WOUND_HEALING, GOBP_POSITIVE_REGULATION_OF_WOUND_HEALING
	MIR7112_3P, MIR7112_3P
	GOMF_SOLUTE_PROTON_ANTIPORTER_ACTIVITY, GOMF_SOLUTE_PROTON_ANTIPORTER_ACTIVITY
	MOREAUX_B_LYMPHOCYTE_MATURATION_BY_TAC1_UP, MOREAUX_B_LYMPHOCYTE_MATURATION_BY_TAC1_UP
	GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS, GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS
	GOMF_COLLAGEN_BINDING, GOMF_COLLAGEN_BINDING
	GOBP_INDOLALKYLAMINE_METABOLIC_PROCESS, GOBP_INDOLALKYLAMINE_METABOLIC_PROCESS
	GOBP_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION
	HP_ABNORMAL_PLATELET_AGGREGATION, HP_ABNORMAL_PLATELET_AGGREGATION
	GOBP_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS, GOBP_REGULATION_OF_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS
	WP_MFAP5_EFFECT_ON_PERMEABILITY_AND_MOTILITY_OF_ENDOTHELIAL_CELLS_VIA_CYTOSKELETON_REARRANGEMENT, WP_MFAP5_EFFECT_ON_PERMEABILITY_AND_MOTILITY_OF_ENDOTHELIAL_CELLS_VIA_CYTOSKELETON_REARRANGEMENT
	MAHAJAN_RESPONSE_TO_IL1A_UP, MAHAJAN_RESPONSE_TO_IL1A_UP
	GOBP_INDOLE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_INDOLE_CONTAINING_COMPOUND_METABOLIC_PROCESS
	RIZ_ERYTHROID_DIFFERENTIATION_HEMGN, RIZ_ERYTHROID_DIFFERENTIATION_HEMGN
	GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS
	TACGGGT_MIR99A_MIR100_MIR99B, TACGGGT_MIR99A_MIR100_MIR99B
	GOBP_TRYPTOPHAN_METABOLIC_PROCESS, GOBP_TRYPTOPHAN_METABOLIC_PROCESS
	WP_OVARIAN_INFERTILITY_GENES, WP_OVARIAN_INFERTILITY_GENES
	PID_ATE2_PATHWAY, PID_ATE2_PATHWAY
	GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS, GOBP_CHONDROITIN_SULFATE_PROTEOGLYCAN_BIOSYNTHETIC_PROCESS
	GOBP_AMINE_CATABOLIC_PROCESS, GOBP_AMINE_CATABOLIC_PROCESS
	MIR135A_3P, MIR135A_3P
	HP_INAPPROPRIATE_LAUGHTER, HP_INAPPROPRIATE_LAUGHTER
	KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE, KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE
	HP_U_SHAPED_UPPER_LIP_VERMILION, HP_U_SHAPED_UPPER_LIP_VERMILION
	WP_NAD_BIOSYNTHESIS_II_FROM_TRYPTOPHAN, WP_NAD_BIOSYNTHESIS_II_FROM_TRYPTOPHAN
	REACTOME_GLUCAGON_SIGNALING_IN_METABOLIC_REGULATION, REACTOME_GLUCAGON_SIGNALING_IN_METABOLIC_REGULATION
	GOBP_DICARBOXYLIC_ACID_TRANSPORT, GOBP_DICARBOXYLIC_ACID_TRANSPORT
	MIR3677_5P, MIR3677_5P
	HARALAMBIEVA_PPMC_M_M_R_ILAGE_13_16YO_STIMULATED_VS_UNSTIMULATED_3_TO_7YR_POST_TOP_DEG_UP, HARALAMBIEVA_PPMC_M_M_R_ILAGE_13_16YO_STIMULATED_VS_UNSTIMULATED_3_TO_7YR_POST_TOP_DEG_UP
	GOBP_PHOSPHATIDYLSERINE_METABOLIC_PROCESS, GOBP_PHOSPHATIDYLSERINE_METABOLIC_PROCESS
	GOBP_NEGATIVE_REGULATION_OF_PROTEIN_MATURATION, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_MATURATION
	GOCC_ANCHORED_COMPONENT_OF_SYNAPTIC_VESICLE_MEMBRANE, GOCC_ANCHORED_COMPONENT_OF_SYNAPTIC_VESICLE_MEMBRANE
	GOBP_DE_NOVO_NAD_BIOSYNTHETIC_PROCESS, GOBP_DE_NOVO_NAD_BIOSYNTHETIC_PROCESS
	REACTOME_SYNDECAN_INTERACTIONS, REACTOME_SYNDECAN_INTERACTIONS
	HP_REFLEX_SEIZURE, HP_REFLEX_SEIZURE
	HP_ABNORMAL_PLACENTA_MORPHOLOGY, HP_ABNORMAL_PLACENTA_MORPHOLOGY
	GOCC_GLIAL_CELL_PROJECTION, GOCC_GLIAL_CELL_PROJECTION
	SERVITJA_ISLET_HNF1A_TARGETS_DN, SERVITJA_ISLET_HNF1A_TARGETS_DN
	HP_ABNORMAL_SPERM_MOTILITY, HP_ABNORMAL_SPERM_MOTILITY
	REACTOME_SYNTHESIS_OF_BILE_ACIDS_AND_BILE_SALTS_VIA_7ALPHA_HYDROXYCHOLESTEROL, REACTOME_SYNTHESIS_OF_BILE_ACIDS_AND_BILE_SALTS_VIA_7ALPHA_HYDROXYCHOLESTEROL
	GOBP_EXTRACELLULAR_TRANSPORT, GOBP_EXTRACELLULAR_TRANSPORT
	GOBP_METANEPHRIC_S_SHAPED_BODY_MORPHOGENESIS, GOBP_METANEPHRIC_S_SHAPED_BODY_MORPHOGENESIS
	GOBP_CELL_CELL_ADHESION_MEDIATED_BY_INTEGRIN, GOBP_CELL_CELL_ADHESION_MEDIATED_BY_INTEGRIN
	GOBP_ACIDIC_AMINO_ACID_TRANSPORT, GOBP_ACIDIC_AMINO_ACID_TRANSPORT
	FERRARI_RESPONSE_TO_FENRETINIDE_UP, FERRARI_RESPONSE_TO_FENRETINIDE_UP
	GOBP_AMINE_TRANSPORT, GOBP_AMINE_TRANSPORT
	PID_VEGF_VEGFR_PATHWAY, PID_VEGF_VEGFR_PATHWAY
	HP_REDUCED_SPERM_MOTILITY, HP_REDUCED_SPERM_MOTILITY
	HP_ABSENT_TOENAIL, HP_ABSENT_TOENAIL
	GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_NEGATIVE_REGULATION_OF_ALPHA_BETA_T_CELL_ACTIVATION
	WP_MFAP5MEDIATED_OVARIAN_CANCER_CELL_MOTILITY_AND_INVASIVENESS, WP_MFAP5MEDIATED_OVARIAN_CANCER_CELL_MOTILITY_AND_INVASIVENESS
	GOBP_REGULATION_OF_EPIDERMIS_DEVELOPMENT, GOBP_REGULATION_OF_EPIDERMIS_DEVELOPMENT
	REACTOME_NON_INTEGRIN_MEMBRANE_ECM_INTERACTIONS, REACTOME_NON_INTEGRIN_MEMBRANE_ECM_INTERACTIONS
	REACTOME_SIGNAL_ATTENUATION, REACTOME_SIGNAL_ATTENUATION
	GOBP_NEGATIVE_REGULATION_OF_VASOCONSTRICTION, GOBP_NEGATIVE_REGULATION_OF_VASOCONSTRICTION
	HINATA_NFKB_TARGETS_KERATINOCYTE_DN, HINATA_NFKB_TARGETS_KERATINOCYTE_DN
	REACTOME_BILE_ACID_AND_BILE_SALT_METABOLISM, REACTOME_BILE_ACID_AND_BILE_SALT_METABOLISM
	GOCC_INFLAMMASOME_COMPLEX, GOCC_INFLAMMASOME_COMPLEX
	GOBP_TRYPTOPHAN_CATABOLIC_PROCESS_TO_KYNURENINE, GOBP_TRYPTOPHAN_CATABOLIC_PROCESS_TO_KYNURENINE
	GOBP_S_SHAPED_BODY_MORPHOGENESIS, GOBP_S_SHAPED_BODY_MORPHOGENESIS
	REACTOME_CYTOCHROME_C_MEDIATED_APOPTOTIC_RESPONSE, REACTOME_CYTOCHROME_C_MEDIATED_APOPTOTIC_RESPONSE
	HP_HYPOPLASTIC_VERTEBRAL_BODIES, HP_HYPOPLASTIC_VERTEBRAL_BODIES
	WP_SEROTONIN_TRANSPORTER_ACTIVITY, WP_SEROTONIN_TRANSPORTER_ACTIVITY
	HP_ABNORMAL_PELVIS_BONE_OSSIFICATION, HP_ABNORMAL_PELVIS_BONE_OSSIFICATION
	HP_MULTIPLE_PRENATAL_FRACTURES, HP_MULTIPLE_PRENATAL_FRACTURES
	HP_IMPAIRED_RISTOCETIN_INDUCED_PLATELET_AGGREGATION, HP_IMPAIRED_RISTOCETIN_INDUCED_PLATELET_AGGREGATION
	HP_ABNORMAL_FOOT_BONE_OSSIFICATION, HP_ABNORMAL_FOOT_BONE_OSSIFICATION
	GSE3720_UNSTIM_VS_PMA_STIM_VD1_GAMMADELTA_TCELL_DN, GSE3720_UNSTIM_VS_PMA_STIM_VD1_GAMMADELTA_TCELL_DN
	GOBP_INTERLEUKIN_1_PRODUCTION, GOBP_INTERLEUKIN_1_PRODUCTION
	GOBP_NEUROTRANSMITTER_REUPTAKE, GOBP_NEUROTRANSMITTER_REUPTAKE
	REACTOME_CHONDROITIN_SULFATE_BIOSYNTHESIS, REACTOME_CHONDROITIN_SULFATE_BIOSYNTHESIS
	HALLMARK_ANGIOGENESIS, HALLMARK_ANGIOGENESIS
	HP_ABNORMALITY_OF_THE_4TH_METACARPAL, HP_ABNORMALITY_OF_THE_4TH_METACARPAL
	GAO_SMALL_INTESTINE_24W_C3_ENTEROCYTE_PROGENITOR_SUBTYPE_1, GAO_SMALL_INTESTINE_24W_C3_ENTEROCYTE_PROGENITOR_SUBTYPE_1
	HP_VERTEBRAL_HYPOPLASIA, HP_VERTEBRAL_HYPOPLASIA
	GOBP_NEGATIVE_REGULATION_OF_ANTIGEN_PROCESSING_AND_PRESENTATION, GOBP_NEGATIVE_REGULATION_OF_ANTIGEN_PROCESSING_AND_PRESENTATION
	GOBP_REGULATION_OF_SEQUESTERING_OF_TRIGLYCERIDE, GOBP_REGULATION_OF_SEQUESTERING_OF_TRIGLYCERIDE
	GOBP_NEPHRIC_DUCT_DEVELOPMENT, GOBP_NEPHRIC_DUCT_DEVELOPMENT
	HP_APLASIA_HYPOPLASIA_OF_THE_4TH_METACARPAL, HP_APLASIA_HYPOPLASIA_OF_THE_4TH_METACARPAL
	GOBP_POSITIVE_REGULATION_OF_AXON_EXTENSION_INVOLVED_IN_AXON_GUIDANCE, GOBP_POSITIVE_REGULATION_OF_AXON_EXTENSION_INVOLVED_IN_AXON_GUIDANCE
	GOBP_MESONEPHRIC_DUCT_DEVELOPMENT, GOBP_MESONEPHRIC_DUCT_DEVELOPMENT
	GOMF_VIRION_BINDING, GOMF_VIRION_BINDING
	GOBP_LABYRINTHINE_LAYER_MORPHOGENESIS, GOBP_LABYRINTHINE_LAYER_MORPHOGENESIS
	DESCARTES_MAIN_FETAL_INHIBITORY_INTERNEURONS, DESCARTES_MAIN_FETAL_INHIBITORY_INTERNEURONS
	GOBP_MONOCYTE_CHEMOTACTIC_PROTEIN_1_PRODUCTION, GOBP_MONOCYTE_CHEMOTACTIC_PROTEIN_1_PRODUCTION
	chr4p15, chr4p15