

	GCM_RAN, GCM_RAN
	GSE13484_UNSTIM_VS_YF17D_VACCINE_STIM_PBMIC_UP, GSE13484_UNSTIM_VS_YF17D_VACCINE_STIM_PBMIC_UP
	GCM_MLL, GCM_MLL
	GCM_GSPT1, GCM_GSPT1
	GCM_RAB10, GCM_RAB10
	GSE39022_LN_VS_SPLEEN_DC_DN, GSE39022_LN_VS_SPLEEN_DC_DN
	GCM_UBE2N, GCM_UBE2N
	GCM_DFFA, GCM_DFFA
	GCM_MAP4K4, GCM_MAP4K4
	CSE37416_0H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GSE37416_0H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN
	GCM_ZNF198, GCM_ZNF198
	CSE34515_CD16_POS_MONOCYTE_VS_DC_UP, CSE34515_CD16_POS_MONOCYTE_VS_DC_UP
	CSE8621_UNSTIM_VS_LPS_PRIMED_UNSTIM_MACROPHAGE_DN, CSE8621_UNSTIM_VS_LPS_PRIMED_UNSTIM_MACROPHAGE_DN
	CSE4176_UNSTIM_VS_ANTI_IGM_STIM_BCELL_24H_DN, GSE4176_UNSTIM_VS_ANTI_IGM_STIM_BCELL_24H_DN
	MIR5484V_3P, MIR5484V_3P
	GOBP_PROTEIN_ACETYLATION, GOBP_PROTEIN_ACETYLATION
	HP_ASTIGMATISM, HP_ASTIGMATISM
	MODULE_182, MODULE_182
	GCM_CHUK, GCM_CHUK
	MIR6768_5P, MIR6768_5P
	GOBP_PEPITIDYL_LYSINE_ACETYLATION, GOBP_PEPITIDYL_LYSINE_ACETYLATION
	GSE7348_UNSTIM_VS_LPS_STIM_MACROPHAGE_UP, GSE7348_UNSTIM_VS_LPS_STIM_MACROPHAGE_UP
	GSE4990_PRE_BCELL_VS_LARGE_PRE_BCELL_DN, GSE4990_PRE_BCELL_VS_LARGE_PRE_BCELL_DN
	GSE40666_NAIVE_VS_EFFECTOR_CD8_TCELL_WITH_IFNA_STIM_90MIN_DN, GSE40666_NAIVE_VS_EFFECTOR_CD8_TCELL_WITH_IFNA_STIM_90MIN_DN
	GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_6H_DN, GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_6H_DN
	GSE29617_CTRL_VS_DAY3_TIV_FLU_VACCINE_PBMIC_2008_UP, GSE29617_CTRL_VS_DAY3_TIV_FLU_VACCINE_PBMIC_2008_UP
	GOCC_PROTEIN_ACETYLTRANSFERASE_COMPLEX, GOCC_PROTEIN_ACETYLTRANSFERASE_COMPLEX
	MIR371A_5P, MIR371A_5P
	MIR221_3P, MIR221_3P
	MIR222_3P, MIR222_3P
	GSE40274_FOXP3_VS_FOXP3_AND_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_FOXP3_VS_FOXP3_AND_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
	GAGCTGG_MIR337, GAGCTGG_MIR337
	MIR187_5P, MIR187_5P
	GCM_PTK2, GCM_PTK2
	MIR19A_5P, MIR19A_5P
	GOBP_MYELOID_CELL_HOMEOSTASIS, GOBP_MYELOID_CELL_HOMEOSTASIS
	MIR6513_3P, MIR6513_3P
	MIR19B_1_5P_MIR19B_2_5P, MIR19B_1_5P_MIR19B_2_5P
	GSE3982_MEMORY_CD4_TCELL_VS_TH1_UP, GSE3982_MEMORY_CD4_TCELL_VS_TH1_UP
	HP_HIP_DYSPLASIA, HP_HIP_DYSPLASIA
	GCM_RBM8A, GCM_RBM8A
	MIR150_5P, MIR150_5P
	TCATCTC_MIR143, TCATCTC_MIR143
	HP_DEEPLY_SET_EYE, HP_DEEPLY_SET_EYE
	GCM_SUFU, GCM_SUFU
	TNCATNTCCYR_UNKNOWN, TNCATNTCCYR_UNKNOWN
	GSE11961_FOLLICULAR_BCELL_VS_MEMORY_BCELL_DAY40_DN, GSE11961_FOLLICULAR_BCELL_VS_MEMORY_BCELL_DAY40_DN
	MIR4773, MIR4773
	HP_ABNORMALITY_OF_THE_HALLUX, HP_ABNORMALITY_OF_THE_HALLUX
	TTGGAGA_MIR5155P_MIR519E, TTGGAGA_MIR5155P_MIR519E
	GOBP_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_DEPENDENT_DECAY, GOBP_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_DEPENDENT_DECAY
	GCM_CRKL, GCM_CRKL
	MIR512_3P, MIR512_3P
	GSE32255_WT_VS_JMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_UP, GSE32255_WT_VS_JMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_UP
	MIR6740_3P, MIR6740_3P
	GCM_CALM1, GCM_CALM1
	MODULE_323, MODULE_323
	GCM_NUMA1, GCM_NUMA1
	MIR5697, MIR5697
	ATGTTTC_MIR494, ATGTTTC_MIR494
	MIR888_3P, MIR888_3P
	HP_BROAD_LONG_BONES, HP_BROAD_LONG_BONES
	MIR3155A_MIR3155B, MIR3155A_MIR3155B
	HP_THICK_EYEBROW, HP_THICK_EYEBROW
	ACTCCAG_MIR173P, ACTCCAG_MIR173P
	GOBP_ERYTHROCYTE_HOMEOSTASIS, GOBP_ERYTHROCYTE_HOMEOSTASIS
	MIR4499, MIR4499
	GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCTYE_UP, GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCTYE_UP
	GCM_BMPR2, GCM_BMPR2
	HP_PREMATURE_BIRTH, HP_PREMATURE_BIRTH
	CCCAGAG_MIR326, CCCAGAG_MIR326
	MODULE_277, MODULE_277
	GAGCCAG_MIR149, GAGCCAG_MIR149
	MIR484, MIR484
	PANGAS_TUMOR_SUPPRESSION_BY_SMAD1_AND_SMAD5_DN, PANGAS_TUMOR_SUPPRESSION_BY_SMAD1_AND_SMAD5_DN
	GCM_GSTA4, GCM_GSTA4
	AAGGGAT_MIR188, AAGGGAT_MIR188
	GOBP_DENDRITE_MORPHOGENESIS, GOBP_DENDRITE_MORPHOGENESIS
	BILD_CTNNB1_ONCOGENIC_SIGNATURE, BILD_CTNNB1_ONCOGENIC_SIGNATURE
	MIR6891_5P, MIR6891_5P
	MIR6088, MIR6088
	MIR8064, MIR8064
	MIR6762_3P, MIR6762_3P
	HP_BROAD_FINGER, HP_BROAD_FINGER
	HP_ABNORMALITY_OF_THE_DIAPHRAGM, HP_ABNORMALITY_OF_THE_DIAPHRAGM
	HP_DOWNTURNED_CORNERS_OF_MOUTH, HP_DOWNTURNED_CORNERS_OF_MOUTH
	AACTGAC_MIR223, AACTGAC_MIR223
	WP_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY, WP_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY
	GOBP_PROTEIN_K48_LINKED_UBIQUITINATION, GOBP_PROTEIN_K48_LINKED_UBIQUITINATION
	GCM_PTPRD, GCM_PTPRD
	MIR1910_5P, MIR1910_5P
	MIR10526_3P, MIR10526_3P
	HP_CONGENITAL_DIAPHRAGMATIC_HERNIA, HP_CONGENITAL_DIAPHRAGMATIC_HERNIA
	MIR2681_5P, MIR2681_5P
	GCTCTTG_MIR335, GCTCTTG_MIR335
	HP_SPEECH_APRAXIA, HP_SPEECH_APRAXIA
	MIR622, MIR622
	GAGACTG_MIR452, GAGACTG_MIR452
	AGGAAGC_MIR5163P, AGGAAGC_MIR5163P
	GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_IFNAB_CD8_TCELL_UP, GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_IFNAB_CD8_TCELL_UP
	MIR7109_5P, MIR7109_5P
	MIR5690, MIR5690
	MIR1226_3P, MIR1226_3P
	MIR642A_3P_MIR642B_3P, MIR642A_3P_MIR642B_3P
	BOYAUULT_LIVER_CANCER_SUBCLASS_G1_UP, BOYAUULT_LIVER_CANCER_SUBCLASS_G1_UP
	MIR203B_3P, MIR203B_3P
	MIR630, MIR630
	MIR2276_3P, MIR2276_3P
	MIR5681A, MIR5681A
	MIR4700_3P, MIR4700_3P
	MIR1237_3P, MIR1237_3P
	WP_THERMOGENESIS, WP_THERMOGENESIS
	MIR212_5P, MIR212_5P
	MIR6773_3P, MIR6773_3P
	GOBP_HISTONE_H3_ACETYLATION, GOBP_HISTONE_H3_ACETYLATION
	HP_OBSESSIVE_COMPULSIVE_BEHAVIOR, HP_OBSESSIVE_COMPULSIVE_BEHAVIOR
	HP_ABSENT_TOE, HP_ABSENT_TOE
	GSE37301_LYMPHOID_PRIMED_MPP_VS_CD4_TCELL_UP, GSE37301_LYMPHOID_PRIMED_MPP_VS_CD4_TCELL_UP
	ATGTAGC_MIR221_MIR222, ATGTAGC_MIR221_MIR222
	HP_INCOMITANT_STRABISMUS, HP_INCOMITANT_STRABISMUS
	AGTCTTA_MIR499, AGTCTTA_MIR499
	MIR4436B_3P, MIR4436B_3P
	GCM_MAX, GCM_MAX
	HP_MACRODONTIA, HP_MACRODONTIA
	MIR335_5P, MIR335_5P
	GALE_APL_WITH_FLT3_MUTATED_UP, GALE_APL_WITH_FLT3_MUTATED_UP
	HP_SYNOSTOSIS_OF_JOINTS, HP_SYNOSTOSIS_OF_JOINTS
	GOBP_ENDOPLASMIC_RETICULUM_MANNNOSE_TRIMMING, GOBP_ENDOPLASMIC_RETICULUM_MANNNOSE_TRIMMING
	CCAWNWNNGGC_UNKNOWN, CCAWNWNNGGC_UNKNOWN
	MIR4433B_5P, MIR4433B_5P
	MIR6851_5P, MIR6851_5P
	TCTCTCC_MIR185, TCTCTCC_MIR185
	GTAAGAT_MIR200A, GTAAGAT_MIR200A
	HUANG_DASATINIB_RESISTANCE_DN, HUANG_DASATINIB_RESISTANCE_DN
	MIR6879_5P, MIR6879_5P
	GOBP_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION, GOBP_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION
	MIR4696, MIR4696
	MIR3622A_3P_MIR3622B_3P, MIR3622A_3P_MIR3622B_3P
	HP_FRONTAL_CORTICAL_ATROPHY, HP_FRONTAL_CORTICAL_ATROPHY
	HP_RELATIVE_MACROCEPHALY, HP_RELATIVE_MACROCEPHALY
	MIR6735_5P, MIR6735_5P
	GSE11884_WT_VS_FURIN_KO_NAIVE_CD4_TCELL_DN, GSE11884_WT_VS_FURIN_KO_NAIVE_CD4_TCELL_DN
	GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_ROSIGLITAZONE_TREATED_UP, GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_ROSIGLITAZONE_TREATED_UP
	HP_ABNORMALITY_OF_THE_AUDITORY_CANAL, HP_ABNORMALITY_OF_THE_AUDITORY_CANAL
	MIR3689D, MIR3689D
	MIR628_3P, MIR628_3P
	GCM_RAP2A, GCM_RAP2A
	HP_FLAT_OCCIPUT, HP_FLAT_OCCIPUT
	HP_ABNORMAL_LARYNX_MORPHOLOGY, HP_ABNORMAL_LARYNX_MORPHOLOGY
	HP_ABNORMALITY_OF_INCISOR_MORPHOLOGY, HP_ABNORMALITY_OF_INCISOR_MORPHOLOGY
	GCM_BAG5, GCM_BAG5
	VERHAAK_GLIOBLASTOMA_PRONEURAL, VERHAAK_GLIOBLASTOMA_PRONEURAL
	ATAACCT_MIR154, ATAACCT_MIR154
	MIR4632_5P, MIR4632_5P
	MIR1251_5P, MIR1251_5P
	MIR370_5P, MIR370_5P
	MIR645, MIR645
	MIR3121_5P, MIR3121_5P
	MIR127_5P, MIR127_5P
	CSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCTYE_RTOC_CULTURE_DN, CSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCTYE_RTOC_CULTURE_DN
	GCM_MAP1B, GCM_MAP1B
	MIR7843_5P, MIR7843_5P
	MIR6783_5P, MIR6783_5P
	MIR8077, MIR8077
	HP_AORTIC_ROOT_ANEURYSM, HP_AORTIC_ROOT_ANEURYSM
	HP_CONGENITAL_NYSTAGMUS, HP_CONGENITAL_NYSTAGMUS
	MIR1285_5P, MIR1285_5P
	GOMF_PEPITIDE_LYSINE_N_ACETYLTRANSFERASE_ACTIVITY, GOMF_PEPITIDE_LYSINE_N_ACETYLTRANSFERASE_ACTIVITY
	MIR647, MIR647
	AATGGAG_MIR136, AATGGAG_MIR136
	MIR892A, MIR892A
	GOBP_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION, GOBP_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION
	MIR3074_3P, MIR3074_3P
	GOMF_VITAMIN_D_RECEPTOR_BINDING, GOMF_VITAMIN_D_RECEPTOR_BINDING
	HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_PYRAMIDAL_TRACT, HP_MORPHOLOGICAL_ABNORMALITY_OF_THE_PYRAMIDAL_TRACT
	CRGAARNNNNGCA_UNKNOWN, CRGAARNNNNGCA_UNKNOWN
	MIR10397_5P, MIR10397_5P
	MIR6873_5P, MIR6873_5P
	HP_ATRESIA_OF_THE_EXTERNAL_AUDITORY_CANAL, HP_ATRESIA_OF_THE_EXTERNAL_AUDITORY_CANAL
	GTACGGA_MIR378, GTACGGA_MIR378
	GOBP_HISTONE_H3_K9_DEMETHYLATION, GOBP_HISTONE_H3_K9_DEMETHYLATION
	HP_CHRONIC_CONSTIPATION, HP_CHRONIC_CONSTIPATION
	MIR6794_3P, MIR6794_3P
	GOMF_HISTONE_DEMETHYLASE_ACTIVITY_H3_K9_SPECIFIC, GOMF_HISTONE_DEMETHYLASE_ACTIVITY_H3_K9_SPECIFIC
	MIR550B_3P, MIR550B_3P
	HP_PROTRUDING_TONGUE, HP_PROTRUDING_TONGUE
	HP_APLASIA_HYPOPLASIA_OF_THE_HALLUX, HP_APLASIA_HYPOPLASIA_OF_THE_HALLUX
	GOBP_LIPID_IMPORT_INTO_CELL, GOBP_LIPID_IMPORT_INTO_CELL
	MIR4327, MIR4327
	GOBP_REGULATION_OF_TYPE_I_INTERFERON_MEDIATED_SIGNALING_PATHWAY, GOBP_REGULATION_OF_TYPE_I_INTERFERON_MEDIATED_SIGNALING_PATHWAY
	MIR652_5P, MIR652_5P
	AR_Q2, AR_Q2
	MIR711, MIR711
	MIR3152_5P, MIR3152_5P
	GOBP_P_BODY_ASSEMBLY, GOBP_P_BODY_ASSEMBLY
	MIR4483, MIR4483
	MIR4710, MIR4710
	MIR6845_5P, MIR6845_5P
	MIR2861, MIR2861
	GTGTCAA_MIR514, GTGTCAA_MIR514
	MIR1293, MIR1293
	HP_TRUNCAL_OBESITY, HP_TRUNCAL_OBESITY
	MIR4301, MIR4301
	MIR6762_5P, MIR6762_5P
	ZHAN_MULTIPLE_MYELOMA_LB_DN, ZHAN_MULTIPLE_MYELOMA_LB_DN
	MIR6810_5P, MIR6810_5P
	MIR4663, MIR4663
	GOBP_POSITIVE_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION
	GOMF_POLY_A_SPECIFIC_RIBONUCLEASE_ACTIVITY, GOMF_POLY_A_SPECIFIC_RIBONUCLEASE_ACTIVITY
	HP_URINARY_URGENCY, HP_URINARY_URGENCY
	MIR4694_5P, MIR4694_5P
	BIOCARTA_ALK_PATHWAY, BIOCARTA_ALK_PATHWAY
	GOBP_PEPITIDYL_METHIONINE_MODIFICATION, GOBP_PEPITIDYL_METHIONINE_MODIFICATION
	MIR6864_3P, MIR6864_3P
	MIR6859_3P, MIR6859_3P
	HP_NARROW_PALPEBRAL_FISSURE, HP_NARROW_PALPEBRAL_FISSURE
	HP_BILATERAL_PTOSIS, HP_BILATERAL_PTOSIS
	SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX1_UP, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX1_UP
	HP_BRAIN_NEOPLASM, HP_BRAIN_NEOPLASM
	MIR4790_3P, MIR4790_3P
	HP_LARYNGOMALACIA, HP_LARYNGOMALACIA
	IKEDA_MIR30_TARGETS_DN, IKEDA_MIR30_TARGETS_DN
	MIR708_5P, MIR708_5P
	GOBP_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING, GOBP_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING
	GOMF_PEPITIDE_ALPHA_N_ACETYLTRANSFERASE_ACTIVITY, GOMF_PEPITIDE_ALPHA_N_ACETYLTRANSFERASE_ACTIVITY
	GOMF_METHYL_CPG_BINDING, GOMF_METHYL_CPG_BINDING
	HP_PERIAURICULAR_SKIN_PITS, HP_PERIAURICULAR_SKIN_PITS
	MIR3139, MIR3139
	GOBP_AMINOPHOSPHOLIPID_TRANSPORT, GOBP_AMINOPHOSPHOLIPID_TRANSPORT
	MIR8056, MIR8056
	GSE12707_AT16L1_HYPOMORPH_VS_WT_THYMUS_DN, GSE12707_AT16L1_HYPOMORPH_VS_WT_THYMUS_DN
	GOBP_RNA_DECAPPING, GOBP_RNA_DECAPPING
	MIR28_5P, MIR28_5P
	GOMF_ARACHIDONATE_COA_LIGASE_ACTIVITY, GOMF_ARACHIDONATE_COA_LIGASE_ACTIVITY
	REACTOME_FOXC_MEDIATED_TRANSCRIPTION_OF_CELL_DEATH_GENES, REACTOME_FOXC_MEDIATED_TRANSCRIPTION_OF_CELL_DEATH_GENES
	chr1p35, chr1p35
	HP_HIGH_ANTERIOR_HAIRLINE, HP_HIGH_ANTERIOR_HAIRLINE
	BYSTRYKH_HEMATOPOIESIS_STEM_CELL_FGF3, BYSTRYKH_HEMATOPOIESIS_STEM_CELL_FGF3
	MIR5586_3P, MIR5586_3P
	GOBP_REGULATION_OF_HEMATOPOIETIC_STEM_CELL_PROLIFERATION, GOBP_REGULATION_OF_HEMATOPOIETIC_STEM_CELL_PROLIFERATION
	MIR5195_5P, MIR5195_5P
	GOBP_N_TERMINAL_PROTEIN_AMINO_ACID_ACETYLATION, GOBP_N_TERMINAL_PROTEIN_AMINO_ACID_ACETYLATION
	GOBP_HISTONE_H3_K36_METHYLATION, GOBP_HISTONE_H3_K36_METHYLATION
	GOBP_REGULATION_OF_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS, GOBP_REGULATION_OF_CELLULAR_CARBOHYDRATE_CATABOLIC_PROCESS
	HP_DEGENERATION_OF_THE_LATERAL_CORTICOSPINAL_TRACTS, HP_DEGENERATION_OF_THE_LATERAL_CORTICOSPINAL_TRACTS
	CAO_BLOOD_FLUMIST_AGE_05_14YO_IDY_UP, CAO_BLOOD_FLUMIST_AGE_05_14YO_IDY_UP
	chr12q22, chr12q22
	GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H4_K20_SPECIFIC, GOMF_HISTONE_METHYLTRANSFERASE_ACTIVITY_H4_K20_SPECIFIC
	LET_7D_3P, LET_7D_3P
	GOBP_REGULATION_OF_DNA_CATABOLIC_PROCESS, GOBP_REGULATION_OF_DNA_CATABOLIC_PROCESS
	GOMF_GUANYLYLTRANSFERASE_ACTIVITY, GOMF_GUANYLYLTRANSFERASE_ACTIVITY
	GOCC_PRC1_COMPLEX, GOCC_PRC1_COMPLEX
	MIR4445_3P, MIR4445_3P
	HP_MORNING_MYOCLONIC_JERKS, HP_MORNING_MYOCLONIC_JERKS
	GOBP_NEGATIVE_REGULATION_OF_CALCINEURIN_MEDIATED_SIGNALING, GOBP_NEGATIVE_REGULATION_OF_CALCINEURIN_MEDIATED_SIGNALING
	HP_DUPLICATION_OF_THE_DISTAL_PHALANX_OF_HAND, HP_DUPLICATION_OF_THE_DISTAL_PHALANX_OF_HAND
	GOBP_EMBRYONIC_CAMERA_TYPE_EYE_FORMATION, GOBP_EMBRYONIC_CAMERA_TYPE_EYE_FORMATION
	GOBP_DIADENOSINE_POLYPHOSPHATE_CATABOLIC_PROCESS, GOBP_DIADENOSINE_POLYPHOSPHATE_CATABOLIC_PROCESS
	GOBP_POSITIVE_REGULATION_OF_HEMATOPOIETIC_STEM_CELL_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_HEMATOPOIETIC_STEM_CELL_PROLIFERATION
	HP_POLYPHAGIA, HP_POLYPHAGIA
	GOBP_REGULATION_OF_RIG_1_SIGNALING_PATHWAY, GOBP_REGULATION_OF_RIG_1_SIGNALING_PATHWAY
	MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_2, MYLLYKANGAS_AMPLIFICATION_HOT_SPOT_2
	GOBP_ACYL_CARNITINE_TRANSPORT, GOBP_ACYL_CARNITINE_TRANSPORT
	GOCC_PERINUCLEAR_ENDOPLASMIC_RETICULUM, GOCC_PERINUCLEAR_ENDOPLASMIC_RETICULUM
	GOBP_POSITIVE_REGULATION_OF_PHOSPHOLIPID_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_PHOSPHOLIPID_BIOSYNTHETIC_PROCESS
	GOBP_OPTIC_CUP_MORPHOGENESIS_INVOLVED_IN_CAMERA_TYPE_EYE_DEVELOPMENT, GOBP_OPTIC_CUP_MORPHOGENESIS_INVOLVED_IN_CAMERA_TYPE_EYE_DEVELOPMENT
	HP_RETROCEBELLAR_CYST, HP_RETROCEBELLAR_CYST
	HP_ABNORMALITY_OF_CARTILAGE_OF_EXTERNAL_EAR, HP_ABNORMALITY_OF_CARTILAGE_OF_EXTERNAL_EAR
	ZNF530_TARGET_GENES, ZNF530_TARGET_GENES
	GOBP_QUATERNARY_AMMONIUM_GROUP_TRANSPORT, GOBP_QUATERNARY_AMMONIUM_GROUP_TRANSPORT
	HP_DILATATION_OF_THE_BLADDER, HP_DILATATION_OF_THE_BLADDER
	GOBP_HINDBRAIN_RADIAL_GLIA_GUIDED_CELL_MIGRATION, GOBP_HINDBRAIN_RADIAL_GLIA_GUIDED_CELL_MIGRATION
	HP_APLASIA_HYPOPLASIA_OF_THE_EPIGLOTTIS, HP_APLASIA_HYPOPLASIA_OF_THE_EPIGLOTTIS
	DIERICK_SEROTONIN_FUNCTION_GENES, DIERICK_SEROTONIN_FUNCTION_GENES
	MIR1914_5P, MIR1914_5P
	GOBP_BLOOD_VESSEL_MATURATION, GOBP_BLOOD_VESSEL_MATURATION
	GOBP_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_DIFFERENTIATION
	GOBP_DEFINITIVE_ERYTHROCYTE_DIFFERENTIATION, GOBP_DEFINITIVE_ERYTHROCYTE_DIFFERENTIATION
	GOBP_POSITIVE_REGULATION_OF_3_UTR_MEDIATED_MRNA_STABILIZATION, GOBP_POSITIVE_REGULATION_OF_3_UTR_MEDIATED_MRNA_STABILIZATION
	GOCC_ROD_PHOTORECEPTOR_OUTER_SEGMENT, GOCC_ROD_PHOTORECEPTOR_OUTER_SEGMENT
	HP_HYPOPLASIA_OF_THE_EPIGLOTTIS, HP_HYPOPLASIA_OF_THE_EPIGLOTTIS
	GOBP_DIPHOSPHOINOSITOL_POLYPHOSPHATE_METABOLIC_PROCESS, GOBP_DIPHOSPHOINOSITOL_POLYPHOSPHATE_METABOLIC_PROCESS
	GOBP_REGULATION_OF_PHOSPHATIDYLCHOLINE_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_PHOSPHATIDYLCHOLINE_BIOSYNTHETIC_PROCESS
	GOMF_ENDOPOLYPHOSPHATASE_ACTIVITY, GOMF_ENDOPOLYPHOSPHATASE_ACTIVITY
	WP_CARDIAC_PROGENITOR_DIFFERENTIATION, WP_CARDIAC_PROGENITOR_DIFFERENTIATION
	CHESLER_BRAIN_D6MIT150_QTL_CIS, CHESLER_BRAIN_D6MIT150_QTL_CIS
	GOBP_REGULATION_OF_3_UTR_MEDIATED_MRNA_STABILIZATION, GOBP_REGULATION_OF_3_UTR_MEDIATED_MRNA_STABILIZATION
	GOMF_G_QUADRUPLUX_RNA_BINDING, GOMF_G_QUADRUPLUX_RNA_BINDING
	GOBP_SMOOTH_MUSCLE_TISSUE_DEVELOPMENT, GOBP_SMOOTH_MUSCLE_TISSUE_DEVELOPMENT
	HP_SUBCORTICAL_CEREBRAL_ATROPHY, HP_SUBCORTICAL_CEREBRAL_ATROPHY
	GOBP_DNA_CATABOLIC_PROCESS_ENDONUCLEOLYTIC, GOBP_DNA_CATABOLIC_PROCESS_ENDONUCLEOLYTIC
	HP_SKIN_PICKING, HP_SKIN_PICKING
	MIR5092, MIR5092
	GOBP_POSITIVE_REGULATION_OF_CARDIOBLAST_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_CARDIOBLAST_DIFFERENTIATION
	GOBP_REGULATION_OF_DENDRITE_EXTENSION, GOBP_REGULATION_OF_DENDRITE_EXTENSION
	HP_GENERALIZED_OSTEOSCLEROSIS, HP_GENERALIZED_OSTEOSCLEROSIS
	MIR8088, MIR8088
	GOBP_REGULATION_OF_CARDIOBLAST_DIFFERENTIATION, GOBP_REGULATION_OF_CARDIOBLAST_DIFFERENTIATION