

MRNA_SPLICING, REACTOME_MRNA_SPLICING

GOCC.SPPLICOSOMAL_COMPLEX, GOCC.SPPLICOSOMAL_COMPLEX
GOBP.VIRAL_GENE_EXPRESSION, GOBP.VIRAL_GENE_EXPRESSION
GSE9006.HEALTHY_VS_TYPE_1_DIABETES_PBMIC_1MONTH_POST_DX_UP, GSE9006.HEALTHY_VS_TYPE_1_DIABETES_PBMIC_1MONTH_POST_DX_UP
GSE3082.DN_VS_CD4.SP_THYMOCYTE_UP, GSE3082.DN_VS_CD4.SP_THYMOCYTE_UP
GOBP.NUCLEAR_EXPORT, GOBP.NUCLEAR_EXPORT
GOBP.ESTABLISHMENT_OF_RNA_LOCALIZATION, GOBP.ESTABLISHMENT_OF_RNA_LOCALIZATION
GSE9006.TYPE_1_VS_TYPE_2_DIABETES_PBMIC_AT_DX_UP, GSE9006.TYPE_1_VS_TYPE_2_DIABETES_PBMIC_AT_DX_UP
MORF.CDC10, MORF.CDC10
MORF.RPA2, MORF.RPA2
CNF2.FBL, CNF2.FBL
CNF2.FBL, CNF2.FBL
MORF.EIF4A2, MORF.EIF4A2
KEGG.SPPLICOSOME, KEGG.SPPLICOSOME
GOBP.RNA_EXPORT_FROM_NUCLEUS, GOBP.RNA_EXPORT_FROM_NUCLEUS
REACTOME.INFLUENZA_INFECTION, REACTOME.INFLUENZA_INFECTION
GSE7568_IL4_TGFB, DEXAMETHASONE_VS_IL4_TGFB_TREATED_MACROPHAGE_UP, GSE7568_IL4_TGFB, DEXAMETHASONE_VS_IL4_TGFB_TREATED_MACROPHAGE_UP
GSE41867_1CMV_ARMSTRONG_VS_CLONIE31_DAY6_EFFECTOR_CD8_TCELL_DN, GSE41867_1CMV_ARMSTRONG_VS_CLONIE31_DAY6_EFFECTOR_CD8_TCELL_DN
GSE24210_RESTING_TREG_VS_TCONV_DN, GSE24210_RESTING_TREG_VS_TCONV_DN
GCM.ACTG1, GCM.ACTG1
GOBP.MRNA_TRANSPORT, GOBP.MRNA_TRANSPORT
GSE9006.TYPE_1_DIABETES_AT_DX_VS_1MONTH_POST_DX_PBMIC_UP, GSE9006.TYPE_1_DIABETES_AT_DX_VS_1MONTH_POST_DX_PBMIC_UP
CNF2.DAP3, CNF2.DAP3
GCM.NPM1, GCM.NPM1
GOCC.U2.TYPE.SPPLICOSOMAL_COMPLEX, GOCC.U2.TYPE.SPPLICOSOMAL_COMPLEX
GOBP.MRNA_EXPORT_FROM_NUCLEUS, GOBP.MRNA_EXPORT_FROM_NUCLEUS
GSE2405_HEAT_KILLED_VS_LIVE_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_9H_UP, GSE2405_HEAT_KILLED_VS_LIVE_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_9H_UP
GOBP.NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_NONSENSE_MEDIATED_DECAY, GOBP.NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_NONSENSE_MEDIATED_DECAY
MORF.RAF1, MORF.RAF1
GSE17721_0.5H_VS_24H_GARDQUIMOD_BMDC_UP, GSE17721_0.5H_VS_24H_GARDQUIMOD_BMDC_UP
MORF.TERF2IP, MORF.TERF2IP
CNF2.HDAC1, CNF2.HDAC1
GSE25146_UNSTIM_VS_HELIOBACTER_PYLORI_LIP5_STIM_AGS_CELL_DN, GSE25146_UNSTIM_VS_HELIOBACTER_PYLORI_LIP5_STIM_AGS_CELL_DN
GOBP.RIBONUCLEOPROTEIN_COMPLEX_BINDING, GOBP.RIBONUCLEOPROTEIN_COMPLEX_BINDING
GCM.CSNK2B, GCM.CSNK2B
REACTOME.NONSENSE_MEDIATED_DECAY_NMD, REACTOME.NONSENSE_MEDIATED_DECAY_NMD
JSON.SICKLE_CELL_DISEASE_DN, JSON.SICKLE_CELL_DISEASE_DN
GOBP.RNA_3_END_PROCESSING, GOBP.RNA_3_END_PROCESSING
GSE1530_WT_VS_IKAROS_KO_LYMPHOID_MULTIPOTENT_PROGENITOR_DN, GSE1530_WT_VS_IKAROS_KO_LYMPHOID_MULTIPOTENT_PROGENITOR_DN
GSE3649_NORMAL_VS_STAPH1_AUREUS_INF_SKIN_UP, GSE3649_NORMAL_VS_STAPH1_AUREUS_INF_SKIN_UP
HAHTOLA_MYCOSIS_FUNGICIDES_SKIN_UP, HAHTOLA_MYCOSIS_FUNGICIDES_SKIN_UP
GSE17721_PAMC5K4_VS_CPG_24H_BMDC_DN, GSE17721_PAMC5K4_VS_CPG_24H_BMDC_DN
GSE1930_STIM_VS_STIM_AND_IL12_24H_CD4_T_CELL_UP, GSE1930_STIM_VS_STIM_AND_IL12_24H_CD4_T_CELL_UP
MORF.PHB, MORF.PHB
GOCC.CATALYTIC_STEP_2.SPPLICOSOME, GOCC.CATALYTIC_STEP_2.SPPLICOSOME
CNF2.RBBP6, CNF2.RBBP6
GSE19401_PLN_VS_PEBERS_PATCH_POLLICULAR_DC_DN, GSE19401_PLN_VS_PEBERS_PATCH_POLLICULAR_DC_DN
MORF.CCNI, MORF.CCNI
GOBP.DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GOBP.DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
GOMF.NUCLEOTIDYLTRANSFERASE_ACTIVITY, GOMF.NUCLEOTIDYLTRANSFERASE_ACTIVITY
GOBP.REGULATION_OF_RNA_SPLICING, GOBP.REGULATION_OF_RNA_SPLICING
GOBP.REGULATION_OF_MRNA_PROCESSING, GOBP.REGULATION_OF_MRNA_PROCESSING
GOBP.MRNA_3_END_PROCESSING, GOBP.MRNA_3_END_PROCESSING
GOBP.TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_IL_PROMOTER, GOBP.TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_IL_PROMOTER
GSE3701_MULTIPOTENT_PROGENITOR_VS_RAG2_KO_NK_CELL_DN, GSE3701_MULTIPOTENT_PROGENITOR_VS_RAG2_KO_NK_CELL_DN
GOCC.RNA_POLYMERASE_COMPLEX, GOCC.RNA_POLYMERASE_COMPLEX
GOBP.REGULATION_OF_GENE_SILENCING, GOBP.REGULATION_OF_GENE_SILENCING
GOBP.RNA_PHOSPHODIESTER_BOND_HYDROLYSIS, GOBP.RNA_PHOSPHODIESTER_BOND_HYDROLYSIS
GOBP.MRNA_TRANSCRIPTION, GOBP.MRNA_TRANSCRIPTION
SESTO_RESPONSE_TO_UV_C0, SESTO_RESPONSE_TO_UV_C0
GSE3982_CTRL_VS_IGE_STIM_MAST_CELL_UP, GSE3982_CTRL_VS_IGE_STIM_MAST_CELL_UP
GOBP.REGULATION_OF_GENE_SILENCING_BY_RNA, GOBP.REGULATION_OF_GENE_SILENCING_BY_RNA
GOBP.POSITIVE_REGULATION_OF_TRANSLATION, GOBP.POSITIVE_REGULATION_OF_TRANSLATION
GSE4363_TTH_VS_1Y6C_INT_CXC3R5P_EFFECTOR_CD4_TCELL_UP, GSE4363_TTH_VS_1Y6C_INT_CXC3R5P_EFFECTOR_CD4_TCELL_UP
GOCC.METHYLTRANSFERASE_COMPLEX, GOCC.METHYLTRANSFERASE_COMPLEX
MORF.BMI1, MORF.BMI1
GOBP.POSITIVE_REGULATION_OF_VIRAL_PROCESS, GOBP.POSITIVE_REGULATION_OF_VIRAL_PROCESS
CNF2.SF3, CNF2.SF3
REACTOME.GENE_SILENCING_BY_RNA, REACTOME.GENE_SILENCING_BY_RNA
GOCC.PRECATALYTIC.SPPLICOSOME, GOCC.PRECATALYTIC.SPPLICOSOME
GOBP.TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_IL_PROMOTER, GOBP.TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_IL_PROMOTER
CNF2.NPM1, CNF2.NPM1
GOBP.DNA_TEMPLATED_TRANSCRIPTION_TERMINATION, GOBP.DNA_TEMPLATED_TRANSCRIPTION_TERMINATION
KCCGNSWTTT, UNKNOWN, KCCGNSWTTT, UNKNOWN
KYN, WERNER_SYNDROM_AND_NORMAL_AGING_DN, KYN, WERNER_SYNDROM_AND_NORMAL_AGING_DN
CREB12_TARGET_GENES, CREB12_TARGET_GENES
CNF2.DD5, CNF2.DD5
GOCC.SM_LIKE_PROTEIN_FAMILY_COMPLEX, GOCC.SM_LIKE_PROTEIN_FAMILY_COMPLEX
REACTOME.RNA_POLYMERASE_IL_TRANSCRIBES_SNRNA_GENES, REACTOME.RNA_POLYMERASE_IL_TRANSCRIBES_SNRNA_GENES
GOBP.TYPE_1_INTERFERON_PRODUCTION, GOBP.TYPE_1_INTERFERON_PRODUCTION
GSE1460_DP_THYMOCYTE_VS_THYMIC_STROMAL_CELL_UP, GSE1460_DP_THYMOCYTE_VS_THYMIC_STROMAL_CELL_UP
GSE2508_CTRL_VS_ROSGLITAZONE_STIM_MACROPHAGE_UP, GSE2508_CTRL_VS_ROSGLITAZONE_STIM_MACROPHAGE_UP
REACTOME.SNRNP_ASSEMBLY, REACTOME.SNRNP_ASSEMBLY
MORF.SNRP70, MORF.SNRP70
GOBP.SNRNA_TRANSCRIPTION, GOBP.SNRNA_TRANSCRIPTION
REACTOME_MRNA_SPLICING_MINOR_PATHWAY, REACTOME_MRNA_SPLICING_MINOR_PATHWAY
MORF.TERF1, MORF.TERF1
SAKAI_CHRONIC_HEPATITIS_VS_LIVER_CANCER_UP, SAKAI_CHRONIC_HEPATITIS_VS_LIVER_CANCER_UP
GOCC.RNA_POLYMERASE_IL_HOLOENZYME, GOCC.RNA_POLYMERASE_IL_HOLOENZYME
GOCC.SMALL_NUCLEAR_RIBONUCLEOPROTEIN_COMPLEX, GOCC.SMALL_NUCLEAR_RIBONUCLEOPROTEIN_COMPLEX
REACTOME_TRANSCRIPTION_OF_THE_HIV_GENOME, REACTOME_TRANSCRIPTION_OF_THE_HIV_GENOME
GCM.PRN1, GCM.PRN1
ZHONG_RESPONSE_TO_AZACITIDINE_AND_TSA_DN, ZHONG_RESPONSE_TO_AZACITIDINE_AND_TSA_DN
GOBP.REGULATION_OF_MRNA_SPLICING_VIA.SPPLICOSOME, GOBP.REGULATION_OF_MRNA_SPLICING_VIA.SPPLICOSOME
GCM.HBP1, GCM.HBP1
REACTOME_VIRAL_MESSENGER_RNA_SYNTHESIS, REACTOME_VIRAL_MESSENGER_RNA_SYNTHESIS
GSE1685_IPNG_TNF_VS_IPNG_KO_LYMPHOID_MULTIPOTENT_PROGENITOR_UP, GSE1685_IPNG_TNF_VS_IPNG_KO_LYMPHOID_MULTIPOTENT_PROGENITOR_UP
GOBP.TERMINATION_OF_RNA_POLYMERASE_IL_TRANSCRIPTION, GOBP.TERMINATION_OF_RNA_POLYMERASE_IL_TRANSCRIPTION
REACTOME.DEADENYLATION_DEPENDENT_MRNA_DECAY, REACTOME.DEADENYLATION_DEPENDENT_MRNA_DECAY
GSE25147_UNSTIM_VS_HELIOBACTER_PYLORI_LIP5_STIM_MKN45_CELL_UP, GSE25147_UNSTIM_VS_HELIOBACTER_PYLORI_LIP5_STIM_MKN45_CELL_UP
GSE360_DC_VS_MAC_UP, GSE360_DC_VS_MAC_UP
GOBP.REGULATION_OF_BIOLOGICAL_PROCESS_INVOLVED_IN_SYMBIOTIC_INTERACTION, GOBP.REGULATION_OF_BIOLOGICAL_PROCESS_INVOLVED_IN_SYMBIOTIC_INTERACTION
KEGG.RNA_DEGRADATION, KEGG.RNA_DEGRADATION
REACTOME_POSITIVE_EPIGENETIC_REGULATION_OF_RRNA_EXPRESSION, REACTOME_POSITIVE_EPIGENETIC_REGULATION_OF_RRNA_EXPRESSION
GOBP.POSITIVE_REGULATION_OF_TYPE_1_INTERFERON_PRODUCTION, GOBP.POSITIVE_REGULATION_OF_TYPE_1_INTERFERON_PRODUCTION
WOOD_EBV_FBNA1_TARGETS_UP, WOOD_EBV_FBNA1_TARGETS_UP
REACTOME_CYTOSOLIC_SENSORS_OF_PATHOGEN_ASSOCIATED_DNA, REACTOME_CYTOSOLIC_SENSORS_OF_PATHOGEN_ASSOCIATED_DNA
REACTOME_NEGATIVE_EPIGENETIC_REGULATION_OF_RRNA_EXPRESSION, REACTOME_NEGATIVE_EPIGENETIC_REGULATION_OF_RRNA_EXPRESSION
HP.TYPE_1_DIABETES_MELLITUS, HP.TYPE_1_DIABETES_MELLITUS
MIR2038_3P, MIR2038_3P
MODULE_350, MODULE_350
GOMF.RNA_HELICASE_ACTIVITY, GOMF.RNA_HELICASE_ACTIVITY
MIR5848AT_3P, MIR5848AT_3P
REACTOME_FORMATION_OF_TC_NER_PRE_INCISION_COMPLEX, REACTOME_FORMATION_OF_TC_NER_PRE_INCISION_COMPLEX
GOMF_SINGLE_STRANDED_RNA_BINDING, GOMF_SINGLE_STRANDED_RNA_BINDING
GOBP.SPPLICOSOMAL_COMPLEX_ASSEMBLY, GOBP.SPPLICOSOMAL_COMPLEX_ASSEMBLY
MIR5849V_3P, MIR5849V_3P
MIR5848B_3P, MIR5848B_3P
GCM.MSN, GCM.MSN
GCM.HDAC1, GCM.HDAC1
CNF2.LIBZ1, CNF2.LIBZ1
GOBP.REGULATION_OF_TELOMERASE_MAINTENANCE_VIA_TELOMERASE_LENGTHENING, GOBP.REGULATION_OF_TELOMERASE_MAINTENANCE_VIA_TELOMERASE_LENGTHENING
GSE1930_STIM_VS_STIM_AND_IFNAB_24H_CD8_T_CELL_UP, GSE1930_STIM_VS_STIM_AND_IFNAB_24H_CD8_T_CELL_UP
GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_DN, GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_DN
GCM.TTR, GCM.TTR
ATCGGGA, UNKNOWN, ATCGGGA, UNKNOWN
GOCC.U2.TYPE.CATALYTIC_STEP_2.SPPLICOSOME, GOCC.U2.TYPE.CATALYTIC_STEP_2.SPPLICOSOME
REACTOME_FORMATION_OF_RNA_POL_IL_ELONGATION_COMPLEX, REACTOME_FORMATION_OF_RNA_POL_IL_ELONGATION_COMPLEX
REACTOME_HIV_TRANSCRIPTION_INITIATION, REACTOME_HIV_TRANSCRIPTION_INITIATION
GOBP_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GOBP_TRANSCRIPTION_BY_RNA_POLYMERASE_I
GOMF.RNA_POLYMERASE_ACTIVITY, GOMF.RNA_POLYMERASE_ACTIVITY
GOCC.CYTOPLASMIC_STRESS_GRANULE, GOCC.CYTOPLASMIC_STRESS_GRANULE
CNF2.PPP6C, CNF2.PPP6C
GOBP.ALTERNATIVE_MRNA_SPLICING_VIA.SPPLICOSOME, GOBP.ALTERNATIVE_MRNA_SPLICING_VIA.SPPLICOSOME
HP.ABNORMAL_ELECTRORETINOGRAM, HP.ABNORMAL_ELECTRORETINOGRAM
MORF.CIRC1, MORF.CIRC1
REACTOME_B_WICH_COMPLEX_POSITIVELY_REGULATES_RRNA_EXPRESSION, REACTOME_B_WICH_COMPLEX_POSITIVELY_REGULATES_RRNA_EXPRESSION
GOMF.DOUBLE_STRANDED_RNA_BINDING, GOMF.DOUBLE_STRANDED_RNA_BINDING
ACAWNRNSRCG, UNKNOWN, ACAWNRNSRCG, UNKNOWN
GOCC.SPPLICOSOMAL_TRISNRNP_COMPLEX, GOCC.SPPLICOSOMAL_TRISNRNP_COMPLEX
GSE12963_ENV_NEF_VS_ENV_NEF_AND_VPR_DEFICIENT_HIV1_INF_CD4_TCELL_UP, GSE12963_ENV_NEF_VS_ENV_NEF_AND_VPR_DEFICIENT_HIV1_INF_CD4_TCELL_UP
GCM.RAF1, GCM.RAF1
GOBP.RNA_POLYADENYLATION, GOBP.RNA_POLYADENYLATION
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_7, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_7
GCM.CSNK1A, GCM.CSNK1A
REACTOME_RNA_POLYMERASE_IL_TRANSCRIPTION, REACTOME_RNA_POLYMERASE_IL_TRANSCRIPTION
MORF.HEAB, MORF.HEAB
GSE5503_LIVER_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_UP, GSE5503_LIVER_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_UP
WP.EUKARYOTIC_TRANSCRIPTION_INITIATION, WP.EUKARYOTIC_TRANSCRIPTION_INITIATION
GCM.MAX, GCM.MAX
GOBP.POSITIVE_REGULATION_OF_MRNA_METABOLIC_PROCESS, GOBP.POSITIVE_REGULATION_OF_MRNA_METABOLIC_PROCESS
SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX4_DN, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX4_DN
REACTOME_RNA_POLYMERASE_L_PROMOTER_ESCAPE, REACTOME_RNA_POLYMERASE_L_PROMOTER_ESCAPE
HP.SCARING, HP.SCARING
CCAGGTT, MIR499, CCAGGTT, MIR499
REACTOME_RNA_POLYMERASE_IL_TRANSCRIPTION_INITIATION_FROM_TYPE_1_PROMOTER, REACTOME_RNA_POLYMERASE_IL_TRANSCRIPTION_INITIATION_FROM_TYPE_1_PROMOTER
GOBP_POSITIVE_REGULATION_OF_RNA_SPLICING, GOBP_POSITIVE_REGULATION_OF_RNA_SPLICING
GOBP_MAINTENANCE_OF_CELL_NUMBER, GOBP_MAINTENANCE_OF_CELL_NUMBER
GOBP_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_EXONUCLEOLYTIC, GOBP_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_EXONUCLEOLYTIC
GOCC.CA1AL_BODY, GOCC.CA1AL_BODY
REACTOME_HIV_TRANSCRIPTION_ELONGATION, REACTOME_HIV_TRANSCRIPTION_ELONGATION
REACTOME_INHIBITION_OF_DNA_RECOMBINATION_AT_TELOMERE, REACTOME_INHIBITION_OF_DNA_RECOMBINATION_AT_TELOMERE
GOMF_RNA_POLYMERASE_BINDING, GOMF_RNA_POLYMERASE_BINDING
CNF2.CBRF, CNF2.CBRF
GOBP_RNA_CAPPING, GOBP_RNA_CAPPING
GSE41978_ID2_KO_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN, GSE41978_ID2_KO_VS_BIM_KO_KLRG1_LOW_EFFECTOR_CD8_TCELL_DN
GOCC_P_BODY, GOCC_P_BODY
GOMF_SNRNA_BINDING, GOMF_SNRNA_BINDING
GOBP_HISTONE_MRNA_METABOLIC_PROCESS, GOBP_HISTONE_MRNA_METABOLIC_PROCESS
REACTOME_RNA_POLYMERASE_IL_TRANSCRIPTION_INITIATION_FROM_TYPE_3_PROMOTER, REACTOME_RNA_POLYMERASE_IL_TRANSCRIPTION_INITIATION_FROM_TYPE_3_PROMOTER
GOMF_MRNA_3_UTR_BINDING, GOMF_MRNA_3_UTR_BINDING
GOBP_NEGATIVE_REGULATION_OF_MRNA_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_MRNA_METABOLIC_PROCESS
GSE2615_CTRL_VS_DAY3_LAYV_IFLU_VACCINE_PBMIC_DN, GSE2615_CTRL_VS_DAY3_LAYV_IFLU_VACCINE_PBMIC_DN
LIN_APC_TARGETS_LIN_APC_TARGETS
HP_ABNORMAL_INSULIN_LEVEL, HP_ABNORMAL_INSULIN_LEVEL
MIR495_5P, MIR495_5P
HP_APLASIA_HYPOPLASIA_INVOLVING_THE_NOSE, HP_APLASIA_HYPOPLASIA_INVOLVING_THE_NOSE
GOMF_BASAL_TRANSCRIPTION_MACHINERY_BINDING, GOMF_BASAL_TRANSCRIPTION_MACHINERY_BINDING
GOBP_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE, GOBP_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE
REACTOME_RHOBTB_GTPASE_CYCLE, REACTOME_RHOBTB_GTPASE_CYCLE
GOBP.SPPLICOSOMAL_SNRNP_ASSEMBLY, GOBP.SPPLICOSOMAL_SNRNP_ASSEMBLY
MODULE_486, MODULE_486
GCM.SMARCC1, GCM.SMARCC1
HP_HORSESHOE_KIDNEY, HP_HORSESHOE_KIDNEY
CHIARADONNA_NEOPLASTIC_TRANSFORMATION_CD25_UP, CHIARADONNA_NEOPLASTIC_TRANSFORMATION_CD25_UP
REACTOME_GENE_AND_PROTEIN_EXPRESSION_BY_JAK_STAT_SIGNALING_AFTER_INTERLEUKIN_12, REACTOME_GENE_AND_PROTEIN_EXPRESSION_BY_JAK_STAT_SIGNALING_AFTER_INTERLEUKIN_12
GOBP_POSITIVE_REGULATION_OF_MRNA_PROCESSING, GOBP_POSITIVE_REGULATION_OF_MRNA_PROCESSING
GOBP_REGULATION_OF_ALTERNATIVE_MRNA_SPLICING_VIA.SPPLICOSOME, GOBP_REGULATION_OF_ALTERNATIVE_MRNA_SPLICING_VIA.SPPLICOSOME
REACTOME_FORMATION_OF_THE_EARLY_ELONGATION_COMPLEX, REACTOME_FORMATION_OF_THE_EARLY_ELONGATION_COMPLEX
MORF_MYST2, MORF_MYST2
REACTOME_RNA_POLYMERASE_L_TRANSCRIPTION_INITIATION, REACTOME_RNA_POLYMERASE_L_TRANSCRIPTION_INITIATION
KEGG_RNA_POLYMERASE, KEGG_RNA_POLYMERASE
HP_UNDEVELOPED_NASAL_ALAE, HP_UNDEVELOPED_NASAL_ALAE
GOMF_RNA_POLYMERASE_CORE_ENZYME_BINDING, GOMF_RNA_POLYMERASE_CORE_ENZYME_BINDING
REACTOME_HIV_ELONGATION_ARREST_AND_RECOVERY, REACTOME_HIV_ELONGATION_ARREST_AND_RECOVERY
GOBP_RESPONSE_TO_INTERLEUKIN_12, GOBP_RESPONSE_TO_INTERLEUKIN_12
MIR6936_3P, MIR6936_3P
GOBP_REGULATION_OF_VIRAL_TRANSCRIPTION, GOBP_REGULATION_OF_VIRAL_TRANSCRIPTION
REACTOME_SIGNALING_BY_FGFR, REACTOME_SIGNALING_BY_FGFR
GOCC_U5_SNRNP, GOCC_U5_SNRNP
CTR9_TARGET_GENES, CTR9_TARGET_GENES
GOBP_PROTEIN_PEPTIDYL_ISOMERIZATION, GOBP_PROTEIN_PEPTIDYL_PROLYL_ISOMERIZATION
REACTOME_SIGNALING_BY_FGFR, REACTOME_SIGNALING_BY_FGFR
GOBP_PTYDYL_PROLINE_MODIFICATION, GOBP_PTYDYL_PROLINE_MODIFICATION
GOBP_RESPONSE_TO_FIBROBLAST_GROWTH_FACTOR, GOBP_RESPONSE_TO_FIBROBLAST_GROWTH_FACTOR
HP_OVERLAPPING_TOE, HP_OVERLAPPING_TOE
GOBP_NEGATIVE_REGULATION_OF_RNA_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_RNA_CATABOLIC_PROCESS
HP_PERIAURICULAR_SKIN_TAG_HP, HP_PERIAURICULAR_SKIN_TAG_HP
HP_PROGRESSIVE_NIGHT_BLINDNESS, HP_PROGRESSIVE_NIGHT_BLINDNESS
HP_ABNORMALITY_OF_PERIAURICULAR_REGION, HP_ABNORMALITY_OF_PERIAURICULAR_REGION
REACTOME_MRNA_CAPPING, REACTOME_MRNA_CAPPING
GTACAGG, MIR486, GTACAGG, MIR486
MIR449_3P, MIR449_3P
HP_CHRONIC_INFECTION, HP_CHRONIC_INFECTION
REACTOME_ABORTIVE_ELONGATION_OF_HIV_1_TRANSCRIPT_IN_THE_ABSENCE_OF_TAT, REACTOME_ABORTIVE_ELONGATION_OF_HIV_1_TRANSCRIPT_IN_THE_ABSENCE_OF_TAT
SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5, SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5
GOCC_EXON_EXON_JUNCTION_COMPLEX, GOCC_EXON_EXON_JUNCTION_COMPLEX
GOBP_REGULATION_OF_RNA_EXPORT_FROM_NUCLEUS, GOBP_REGULATION_OF_RNA_EXPORT_FROM_NUCLEUS
GOBP_POSITIVE_REGULATION_OF_VIRAL_TRANSCRIPTION, GOBP_POSITIVE_REGULATION_OF_VIRAL_TRANSCRIPTION
REACTOME_MICRORNA_BIOGENESIS, REACTOME_MICRORNA_BIOGENESIS
GOMF_CIS_TRANS_ISOMERASE_ACTIVITY, GOMF_CIS_TRANS_ISOMERASE_ACTIVITY
HP_ABNORMALITY_OF_CONJUNCTIVAL_THICKNESS, HP_ABNORMALITY_OF_CONJUNCTIVAL_THICKNESS
REACTOME_INTERLEUKIN_12_SIGNALING, REACTOME_INTERLEUKIN_12_SIGNALING
HP_SPARSE_EYELASHES, HP_SPARSE_EYELASHES
MIR8081, MIR8081
REACTOME_RHOBTB1_GTPASE_CYCLE, REACTOME_RHOBTB1_GTPASE_CYCLE
HP_SKIN_TAGS, HP_SKIN_TAGS
REACTOME_PROCESSING_OF_INTRONLESS_PRE_MRNAs, REACTOME_PROCESSING_OF_INTRONLESS_PRE_MRNAs
GOBP_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
GOCC_U12.TYPE.SPPLICOSOMAL_COMPLEX, GOCC_U12.TYPE.SPPLICOSOMAL_COMPLEX
MCRS1_TARGET_GENES, MCRS1_TARGET_GENES
GOBP_REGULATION_OF_NUCLEASE_CONTAINING_COMPOUND_TRANSPORT, GOBP_REGULATION_OF_NUCLEASE_CONTAINING_COMPOUND_TRANSPORT
GOCC_TELOMERASE_HOLOENZYME_COMPLEX, GOCC_TELOMERASE_HOLOENZYME_COMPLEX
REACTOME_FGFR2_ALTERNATIVE_SPLICING, REACTOME_FGFR2_ALTERNATIVE_SPLICING
MIDORIKAWA_AMPLIFIED_IN_LIVER_CANCER, MIDORIKAWA_AMPLIFIED_IN_LIVER_CANCER
GOBP_LRR_DOMAIN_BINDING, GOMF_LRR_DOMAIN_BINDING
MIR493_3P, MIR493_3P
GOBP_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_L_PROMOTER, GOBP_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_L_PROMOTER
MODULE_388, MODULE_388
AGTCACG, MIR345, AGTCACG, MIR345
GOCC_RNA_POLYMERASE_IL_COMPLEX, GOCC_RNA_POLYMERASE_IL_COMPLEX
REACTOME_RNA_POLYMERASE_IL_CHAIN_ELONGATION, REACTOME_RNA_POLYMERASE_IL_CHAIN_ELONGATION
GOMF_PRE_MRNA_BINDING, GOMF_PRE_MRNA_BINDING
KDM1B_TARGET_GENES, KDM1B_TARGET_GENES
GOBP_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERASE_LENGTHENING, GOBP_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERASE_LENGTHENING
MIR6514_3P, MIR6514_3P
GOBP_POSITIVE_REGULATION_OF_MRNA_SPLICING_VIA.SPPLICOSOME, GOBP_POSITIVE_REGULATION_OF_MRNA_SPLICING_VIA.SPPLICOSOME
GOCC_ENDORIBONUCLEASE_COMPLEX, GOCC_ENDORIBONUCLEASE_COMPLEX
MIR548V, MIR548V
HP_ABNORMALITY_OF_FIBRIL_MORPHOLOGY, HP_ABNORMALITY_OF_FIBRIL_MORPHOLOGY
MELLMAN_TUT1_TARGETS_UP, MELLMAN_TUT1_TARGETS_UP
MIR5503_3P, MIR5503_3P
WP_CYTOSOLIC_DNASENSING_PATHWAY, WP_CYTOSOLIC_DNASENSING_PATHWAY
TCTGATC, MIR383, TCTGATC, MIR383
GOCC_MRNA_CLEAVAGE_FACTOR_COMPLEX, GOCC_MRNA_CLEAVAGE_FACTOR_COMPLEX
GOMF_DRUG_BINDING, GOMF_DRUG_BINDING
CAFFAREL_RESPONSE_TO_THE_8HR_5_DN, CAFFAREL_RESPONSE_TO_THE_8HR_5_DN
GOCC_CATALYTIC_STEP_1.SPPLICOSOME, GOCC_CATALYTIC_STEP_1.SPPLICOSOME
GAZDA_DIAMOND_BLACKFAN_ANEMIA_ERYTHROID_UP, GAZDA_DIAMOND_BLACKFAN_ANEMIA_ERYTHROID_UP
GOBP_POSITIVE_REGULATION_OF_MRNA_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_MRNA_CATABOLIC_PROCESS
REACTOME_PWW1_INTERACTING_RNA_PIRNA_BIOGENESIS, REACTOME_PWW1_INTERACTING_RNA_PIRNA_BIOGENESIS
NKX22_O1, NKX22_O1
HP_FRONTOTEMPORAL_DEMENTIA, HP_FRONTOTEMPORAL_DEMENTIA
GCM_AIP, GCM_AIP
HP_GENERALIZED_AMYOTROPHY, HP_GENERALIZED_AMYOTROPHY
YBN1_TARGET_GENES, YBN1_TARGET_GENES
ALX4_TARGET_GENES, ALX4_TARGET_GENES
GOBP_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS, GOBP_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS
DORN_ADENOVIRUS_INFECTION_32HR_DN, DORN_ADENOVIRUS_INFECTION_32HR_DN
MIR638_5P, MIR638_5P
HP_AMYOTROPHIC_LATERAL_SCLEROSIS, HP_AMYOTROPHIC_LATERAL_SCLEROSIS
GOBP_TERMINATION_OF_RNA_POLYMERASE_I_TRANSCRIPTION, GOBP_TERMINATION_OF_RNA_POLYMERASE_I_TRANSCRIPTION
GOMF_POLY_PYRIMIDINE_TRACT_BINDING, GOMF_POLY_PYRIMIDINE_TRACT_BINDING
GOBP_VIRAL_TRANSLATION, GOBP_VIRAL_TRANSLATION
HP_RONE_SPICULE_PIGMENTATION_OF_THE_RETINA, HP_RONE_SPICULE_PIGMENTATION_OF_THE_RETINA
HP_LONG_PALPEBRAL_FISSURE, HP_LONG_PALPEBRAL_FISSURE
AGTTCTC, MIR146A, AGTTCTC, MIR146A, MIR146B
HP_NASOGASTRIC_TUBE_FEEDING_IN_INFANCY, HP_NASOGASTRIC_TUBE_FEEDING_IN_INFANCY
REACTOME_MRNA_DECAY_BY_5_TO_3_EXORIBONUCLEASE, REACTOME_MRNA_DECAY_BY_5_TO_3_EXORIBONUCLEASE
REACTOME_INTERLEUKIN_12_FAMILY_SIGNALING, REACTOME_INTERLEUKIN_12_FAMILY_SIGNALING
PER1_TARGET_GENES, PER1_TARGET_GENES
REACTOME_RNA_POLYMERASE_IL_TRANSCRIPTION_TERMINATION, REACTOME_RNA_POLYMERASE_IL_TRANSCRIPTION_TERMINATION
GOMF_POLY_PURINE_TRACT_BINDING, GOMF_POLY_PURINE_TRACT_BINDING
REACTOME_RHOBTB1_GTPASE_CYCLE, REACTOME_RHOBTB1_GTPASE_CYCLE
GOCC_MRNA_CLEAVAGE_AND_POLYADENYLATION_SPECIFICITY_FACTOR_COMPLEX, GOCC_MRNA_CLEAVAGE_AND_POLYADENYLATION_SPECIFICITY_FACTOR_COMPLEX
CHENG_IMPRINTED_BY_ESTRADIOL, CHENG_IMPRINTED_BY_ESTRADIOL
GOBP_RNA_INTERFERENCE, GOBP_RNA_INTERFERENCE
GOBP_TELOMERASE_RNA_BINDING, GOMF_TELOMERASE_RNA_BINDING
GOCC_RNA_POLYMERASE_IL_CORE_COMPLEX, GOCC_RNA_POLYMERASE_IL_CORE_COMPLEX
GOBP_POSITIVE_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING, GOBP_POSITIVE_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING
REACTOME_SIGNALING_BY_FGFR_IN_DISEASE, REACTOME_SIGNALING_BY_FGFR_IN_DISEASE
HOLLEMAN_VINCISTINE_RESISTANCE_ALL_DN, HOLLEMAN_VINCISTINE_RESISTANCE_ALL_DN
GOMF_REGULATORY_RNA_BINDING, GOMF_REGULATORY_RNA_BINDING
XU_AKT1_TARGETS_AHR_XU_AKT1_TARGETS_AHR
GOBP_NEGATIVE_REGULATION_OF_MRNA_PROCESSING, GOBP_NEGATIVE_REGULATION_OF_MRNA_PROCESSING
SYED_ESTRADIOL_RESPONSE, SYED_ESTRADIOL_RESPONSE
GOBP_REGULATION_OF_MRNA_POLYADENYLATION, GOBP_REGULATION_OF_MRNA_POLYADENYLATION
GOBP_RESPONSE_TO_ARSINIC_CONTAINING_SUBSTANCE, GOBP_RESPONSE_TO_ARSINIC_CONTAINING_SUBSTANCE
REACTOME_RNA_POLYMERASE_I_TRANSCRIPTION_TERMINATION, REACTOME_RNA_POLYMERASE_I_TRANSCRIPTION_TERMINATION
HP_RETINAL_COLOBOMA, HP_RETINAL_COLOBOMA
GOBP_PROTEIN_REFOLDING, GOBP_PROTEIN_REFOLDING
GOBP_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_I_PROMOTER, GOBP_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_I_PROMOTER
HP_EMG_CHRONIC_DENERVATION_SIGNS, HP_EMG_CHRONIC_DENERVATION_SIGNS
GATA2_O1, GATA2_O1
MODULE_322, MODULE_322
HP_MUSCLE_FIBER_ATROPHY, HP_MUSCLE_FIBER_ATROPHY
HASLINGER_B_CLL_WITH_CHROMOSOME_12_TRISOMY, HASLINGER_B_CLL_WITH_CHROMOSOME_12_TRISOMY
MIR314_3P, MIR314_3P
GOMF_RNA_CAP_BINDING, GOMF_RNA_CAP_BINDING
MIR1178_3P, MIR1178_3P
LIU_BREAST_CANCER_LIU_BREAST_CANCER
GOBP_NEGATIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS
GCM.CAMP2, GCM.CAMP2
DORN_ADENOVIRUS_INFECTION_48HR_DN, DORN_ADENOVIRUS_INFECTION_48HR_DN
GOBP_NEGATIVE_REGULATION_OF_RNA_SPLICING, GOBP_NEGATIVE_REGULATION_OF_RNA_SPLICING
HOLLEMAN_PREDNISOLONE_RESISTANCE_B_ALL_UP, HOLLEMAN_PREDNISOLONE_RESISTANCE_B_ALL_UP
GOCC_PP1P_COMPLEX, GOCC_PP1P_COMPLEX
BOYLAN_MULTIPLE_MYELOMA_C_CLUSTER_UP, BOYLAN_MULTIPLE_MYELOMA_C_CLUSTER_UP
HP_APLASIA_HYPOPLASIA_OF_THE_OPTIC_NERVE, HP_APLASIA_HYPOPLASIA_OF_THE_OPTIC_NERVE
GOMF_POLY_A_BINDING, GOMF_POLY_A_BINDING
MIR6794_3P, MIR6794_3P
HP_CAPILLARY_MALFORMATION, HP_CAPILLARY_MALFORMATION
GOBP_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERASE, GOBP_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERASE
HP_RADIAL_DEVIATION_OF_THE_HAND_OR_OF_FINGERS_OF_THE_HAND_HP, HP_RADIAL_DEVIATION_OF_THE_HAND_OR_OF_FINGERS_OF_THE_HAND_HP
GOBP_NEGATIVE_REGULATION_OF_MRNA_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_MRNA_CATABOLIC_PROCESS
CATRACC, UNKNOWN, CATRACC, UNKNOWN
HP_HYPOPLASIA_OF_THE_RADIUS, HP_HYPOPLASIA_OF_THE_RADIUS
GOMF_MIRNA_BINDING, GOMF_MIRNA_BINDING
HP_SYNOSTOSIS_INVOLVING_BONES_OF_THE_HAND, HP_SYNOSTOSIS_INVOLVING_BONES_OF_THE_HAND
GOBP_IRRS_DEPENDENT_VIRAL_TRANSLATIONAL_INITIATION, GOBP_IRRS_DEPENDENT_VIRAL_TRANSLATIONAL_INITIATION
GOBP_MRNA_CLEAVAGE, GOBP_MRNA_CLEAVAGE
GOCC_SMRN_SMRN_PROTEIN_COMPLEX, GOCC_SMRN_SMRN_PROTEIN_COMPLEX
GOBP.SPPLICOSOMAL_TRISNRNP_COMPLEX_ASSEMBLY, GOBP.SPPLICOSOMAL_TRISNRNP_COMPLEX_ASSEMBLY
HP_ABSENT_FOREARM_BONE, HP_ABSENT_FOREARM_BONE
MODULE_229, MODULE_229
KEGG_BASAL_TRANSCRIPTION_FACTORS, KEGG_BASAL_TRANSCRIPTION_FACTORS
LOPEZ_TRANSLATION_VIA_FNT1_SIGNALING, LOPEZ_TRANSLATION_VIA_FNT1_SIGNALING
MIR2909, MIR2909
GOMF_ANDROGEN_RECEPTOR_BINDING, GOMF_ANDROGEN_RECEPTOR_BINDING
SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX1_DN, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX1_DN
GOBP_RNA_SPLICING_VIA_YAUCH_HEDGEHOG_SIGNALING, GOBP_RNA_SPLICING_VIA_YAUCH_HEDGEHOG_SIGNALING
GOBP_RESPONSE_TO_SALT, GOBP_RESPONSE_TO_SALT
GOBP_POSITIVE_REGULATION_OF_CALCIIUM_MEDIATED_SIGNALING, GOBP_POSITIVE_REGULATION_OF_CALCIIUM_MEDIATED_SIGNALING
SIMBULAN_UV_RESPONSE_NORMAL_DN, SIMBULAN_UV_RESPONSE_NORMAL_DN
GOBP_RESPONSE_TO_DSRNA, GOBP_RESPONSE_TO_DSRNA
GOCC_POST_MRNA_RELEASE.SPPLICOSOMAL_COMPLEX, GOCC_POST_MRNA_RELEASE.SPPLICOSOMAL_COMPLEX
MIR6857_3P, MIR6857_3P
BONOME_OVARIAN_CANCER_POOR_SURVIVAL_DN, BONOME_OVARIAN_CANCER_POOR_SURVIVAL_DN
GOBP_SOMATIC_STEM_CELL_POPULATION_MAINTENANCE, GOBP_SOMATIC_STEM_CELL_POPULATION_MAINTENANCE
MCT1ATG, MIR368, MCT1ATG, MIR368
MIR3174, MIR3174
HOEK_NK_CELL_2011_2012_TIV_ADULT_3DY_UP, HOEK_NK_CELL_2011_2012_TIV_ADULT_3DY_UP
REACTOME_CYCLOSOPORIN_A_BINDING, REACTOME_CYCLOSOPORIN_A_BINDING
GOCC_SIGNALING_BY_FGFR2_IIIA_TM, REACTOME_SIGNALING_BY_FGFR2_IIIA_TM
ZNP19_TARGET_GENES, ZNP19_TARGET_GENES
MAYBURD_RESPONSE_TO_166536_UP, MAYBURD_RESPONSE_TO_166536_UP
MIR4766_3P, MIR4766_3P
HP_SPARSE_OR_ABSENT_EYELASHES, HP_SPARSE_OR_ABSENT_EYELASHES
YAUCH_HEDGEHOG_SIGNALING, PARACRINE_UP, YAUCH_HEDGEHOG_SIGNALING, PARACRINE_UP
HOLLEMAN_VINCISTINE_RESISTANCE_B_ALL_DN, HOLLEMAN_VINCISTINE_RESISTANCE_B_ALL_DN
HP_FIBULAR_HYPOPLASIA, HP_FIBULAR_HYPOPLASIA
HP_APLASIA_HYPOPLASIA_OF_THE_EYELED, HP_APLASIA_HYPOPLASIA_OF_THE_EYELED
ATGCACG, MIR5178, ATGCACG, MIR5178
GOBP_MRNA_CLEAVAGE_INVOLVED_IN_MRNA_PROCESSING, GOBP_MRNA_CLEAVAGE_INVOLVED_IN_MRNA_PROCESSING
JIANO_TIPS_TARSIER_DN, JIANO_TIPS_TARSIER_DN
GOBP_SMALL_RNA_LOADING_ONTO_RISC, GOBP_SMALL_RNA_LOADING_ONTO_RISC
HP_MICRORETROGNATHIA, HP_MICRORETROGNATHIA
HP_XEROSTOMIA, HP_XEROSTOMIA
GOMF_RNA_7_METHYLGUANOSINE_CAP_BINDING, GOMF_RNA_7_METHYLGUANOSINE_CAP_BINDING
NKX2_4_TARGET_GENES, NKX2_4_TARGET_GENES
MIR4751, MIR4751
GOMF_RNA_STEM_LOOP_BINDING, GOMF_RNA_STEM_LOOP_BINDING
MIR6870_3P, MIR6870_3P
GOMF_RNA_POLYMERASE_IL_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_BINDING, GOMF_RNA_POLYMERASE_IL_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_BINDING
HP_RIMED_VACUOLIS, HP_RIMED_VACUOLIS
AZARE_STAT3_TARGETS, AZARE_STAT3_TARGETS
GOMF_U6_SNRNA_BINDING, GOMF_U6_SNRNA_BINDING
GOCC_METHYLOSOME, GOCC_METHYLOSOME
GOCC_REGION_OF_CYTOSOL, GOCC_REGION_OF_CYTOSOL
RASHI_RESPONSE_TO_IONIZING_RADIATION_3, RASHI_RESPONSE_TO_IONIZING_RADIATION_3
HOLLEMAN_PREDNISOLONE_RESISTANCE_ALL_UP, HOLLEMAN_PREDNISOLONE_RESISTANCE_ALL_UP
REACTOME_SIGNALING_BY_FGFR2_IN_DISEASE, REACTOME_SIGNALING_BY_FGFR2_IN_DISEASE
GOBP_NEGATIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_NONHOMOLOGOUS_END_JOINING, GOBP_NEGATIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_NONHOMOLOGOUS_END_JOINING
GOCC_U4_SNRNP, GOCC_U4_SNRNP
MIR5198_3P, MIR5198_3P
GOBP_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_BY_HOST, GOBP_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_BY_HOST
HP_SUPERNUMERARY_BONES_OF_THE_AXIAL_SKELETON, HP_SUPERNUMERARY_BONES_OF_THE_AXIAL_SKELETON
MARIADASON_RESPONSE_TO_BUTYRATE_CURCUMIN_SULINDAC_TSA_8, MARIADASON_RESPONSE_TO_BUTYRATE_CURCUMIN_SULINDAC_TSA_8
LOPEZ_MISOTHELIOMA_SURVIVAL_1, LOPEZ_MISOTHELIOMA_SURVIVAL_1
RBMX_TARGET_GENES, RBMX_TARGET_GENES
GOBP_PHOSPHORYLATION_OF_RNA_POLYMERASE_IL_C_TERMINAL_DOMAIN, GOBP_PHOSPHORYLATION_OF_RNA_POLYMERASE_IL_C_TERMINAL_DOMAIN
MIR424_3P, MIR424_3P
MIR3914, MIR3914
GOBP_CELLULAR_RESPONSE_TO_DSRNA, GOBP_CELLULAR_RESPONSE_TO_DSRNA
GOBP_REGULATION_OF_TRANSLATION_AT_SYNAPE_MODULATING_SYNAPTIC_TRANSMISSION, GOBP_REGULATION_OF_TRANSLATION_AT_SYNAPE_MODULATING_SYNAPTIC_TRANSMISSION
SCHLINGEMANN_SKIN_CARCIINOGENESIS_TPA_UP, SCHLINGEMANN_SKIN_CARCIINOGENESIS_TPA_UP
MDM2_TARGET_GENES, MDM2_TARGET_GENES
HP_ABNORMAL_FUNDUS_AUTOFLUORESCENCE_IMAGING, HP_ABNORMAL_FUNDUS_AUTOFLUORESCENCE_IMAGING