

MRNA_SPLICING, REACTOME_MRNA_SPLICING

- PUJANA_BRCA_CENTERED_NETWORK, PUJANA_BRCA_CENTERED_NETWORK
- YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_13, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_13
- REACTOME_NUCLEOTIDE_EXCISION_REPAIR, REACTOME_NUCLEOTIDE_EXCISION_REPAIR
- REACTOME_CHROMOSOME_MAINTENANCE, REACTOME_CHROMOSOME_MAINTENANCE
- WP_MRNA_PROCESSING, WP_MRNA_PROCESSING
- REACTOME_RNA_POLYMERASE_II_TRANSCRIBES_SNRNA_GENES, REACTOME_RNA_POLYMERASE_II_TRANSCRIBES_SNRNA_GENES
- REACTOME_TRANSCRIPTION_COUPLED_NUCLEOTIDE_EXCISION_REPAIR_TC_NER, REACTOME_TRANSCRIPTION_COUPLED_NUCLEOTIDE_EXCISION_REPAIR_TC_NER
- REACTOME_TELOMERE_MAINTENANCE, REACTOME_TELOMERE_MAINTENANCE
- REACTOME_B_WICH_COMPLEX_POSITIVELY_REGULATES_RRNA_EXPRESSION, REACTOME_B_WICH_COMPLEX_POSITIVELY_REGULATES_RRNA_EXPRESSION
- REACTOME_MRNA_SPLICING_MINOR_PATHWAY, REACTOME_MRNA_SPLICING_MINOR_PATHWAY
- REACTOME_POSITIVE_EPIGENETIC_REGULATION_OF_RRNA_EXPRESSION, REACTOME_POSITIVE_EPIGENETIC_REGULATION_OF_RRNA_EXPRESSION
- REACTOME_INHIBITION_OF_DNA_RECOMBINATION_AT_TELOMERE, REACTOME_INHIBITION_OF_DNA_RECOMBINATION_AT_TELOMERE
- MISSIAGLIA_REGULATED_BY_METHYLATION_DN, MISSIAGLIA_REGULATED_BY_METHYLATION_DN
- KEGG_RNA_DEGRADATION, KEGG_RNA_DEGRADATION
- REACTOME_TRANSCRIPTION_OF_THE_HIV_GENOME, REACTOME_TRANSCRIPTION_OF_THE_HIV_GENOME
- REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION, REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION
- GROSS_HYPOXIA_VIA_ELK3_AND_HIF1A_DN, GROSS_HYPOXIA_VIA_ELK3_AND_HIF1A_DN
- REACTOME_DUAL_INCISION_IN_TC_NER, REACTOME_DUAL_INCISION_IN_TC_NER
- MORI_EMU_MYC_LYMPHOMA_BY_ONSET_TIME_UP, MORI_EMU_MYC_LYMPHOMA_BY_ONSET_TIME_UP
- REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_DNA_REPAIR_GENES, REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_DNA_REPAIR_GENES
- REACTOME_DEADENYLATION_DEPENDENT_MRNA_DECAY, REACTOME_DEADENYLATION_DEPENDENT_MRNA_DECAY
- HAHTOLA_MYCOSIS_FUNGOIDES_SKIN_UP, HAHTOLA_MYCOSIS_FUNGOIDES_SKIN_UP
- REACTOME_HIV_TRANSCRIPTION_INITIATION, REACTOME_HIV_TRANSCRIPTION_INITIATION
- KEGG_PYRIMIDINE_METABOLISM, KEGG_PYRIMIDINE_METABOLISM
- ZHONG_RESPONSE_TO_AZACITIDINE_AND_TSA_DN, ZHONG_RESPONSE_TO_AZACITIDINE_AND_TSA_DN
- REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION_INITIATION_FROM_TYPE_1_PROMOTER, REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION_INITIATION_FROM_TYPE_1_PROMOTER
- PURBEY_TARGETS_OF_CTBP1_AND_SATB1_DN, PURBEY_TARGETS_OF_CTBP1_AND_SATB1_DN
- REACTOME_HIV_ELONGATION_ARREST_AND_RECOVERY, REACTOME_HIV_ELONGATION_ARREST_AND_RECOVERY
- REACTOME_FORMATION_OF_RNA_POL_II_ELONGATION_COMPLEX, REACTOME_FORMATION_OF_RNA_POL_II_ELONGATION_COMPLEX
- SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX4_DN, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX4_DN
- YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_10, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_10
- REACTOME_ABORTIVE_ELONGATION_OF_HIV_1_TRANSCRIPT_IN_THE_ABSENCE_OF_TAT, REACTOME_ABORTIVE_ELONGATION_OF_HIV_1_TRANSCRIPT_IN_THE_ABSENCE_OF_TAT
- REACTOME_CYTOSOLIC_SENSORS_OF_PATHOGEN_ASSOCIATED_DNA, REACTOME_CYTOSOLIC_SENSORS_OF_PATHOGEN_ASSOCIATED_DNA
- REACTOME_FORMATION_OF_THE_EARLY_ELONGATION_COMPLEX, REACTOME_FORMATION_OF_THE_EARLY_ELONGATION_COMPLEX
- REACTOME_ESTROGEN_DEPENDENT_GENE_EXPRESSION, REACTOME_ESTROGEN_DEPENDENT_GENE_EXPRESSION
- SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5, SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5
- REACTOME_FGFR2_ALTERNATIVE_SPLICING, REACTOME_FGFR2_ALTERNATIVE_SPLICING
- REACTOME_EPIGENETIC_REGULATION_OF_GENE_EXPRESSION, REACTOME_EPIGENETIC_REGULATION_OF_GENE_EXPRESSION
- ASTON_MAJOR_DEPRESSIVE_DISORDER_DN, ASTON_MAJOR_DEPRESSIVE_DISORDER_DN
- VANHARANTA_UTERINE_FIBROID_WITH_7Q_DELETION_UP, VANHARANTA_UTERINE_FIBROID_WITH_7Q_DELETION_UP
- WOOD_EBV_EBNA1_TARGETS_UP, WOOD_EBV_EBNA1_TARGETS_UP
- REACTOME_HIV_TRANSCRIPTION_ELONGATION, REACTOME_HIV_TRANSCRIPTION_ELONGATION
- WP_PYRIMIDINE_METABOLISM, WP_PYRIMIDINE_METABOLISM
- REACTOME_ACTIVATION_OF_ANTERIOR_HOX_GENES_IN_HINDBRAIN_DEVELOPMENT_DURING_EARLY_EMBRYOGENESIS, REACTOME_ACTIVATION_OF_ANTERIOR_HOX_GENES_IN_HINDBRAIN_DEVELOPMENT_DURING
- REACTOME_RNA_POLYMERASE_I_TRANSCRIPTION_INITIATION, REACTOME_RNA_POLYMERASE_I_TRANSCRIPTION_INITIATION
- WP_EUKARYOTIC_TRANSCRIPTION_INITIATION, WP_EUKARYOTIC_TRANSCRIPTION_INITIATION
- MELLMAN_TUT1_TARGETS_UP, MELLMAN_TUT1_TARGETS_UP
- REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION_INITIATION_FROM_TYPE_3_PROMOTER, REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION_INITIATION_FROM_TYPE_3_PROMOTER
- REACTOME_PROCESSING_OF_INTRONLESS_PRE_MRNAS, REACTOME_PROCESSING_OF_INTRONLESS_PRE_MRNAS
- JAZAG_TGFB1_SIGNALING_UP, JAZAG_TGFB1_SIGNALING_UP
- REACTOME_MRNA_CAPPING, REACTOME_MRNA_CAPPING
- STEARMAN_LUNG_CANCER_EARLY_VS_LATE_UP, STEARMAN_LUNG_CANCER_EARLY_VS_LATE_UP
- REACTOME_PROCESSING_OF_CAPPED_INTRONLESS_PRE_MRNA, REACTOME_PROCESSING_OF_CAPPED_INTRONLESS_PRE_MRNA
- LIN_APC_TARGETS, LIN_APC_TARGETS
- CHIARADONNA_NEOPLASTIC_TRANSFORMATION_CDC25_UP, CHIARADONNA_NEOPLASTIC_TRANSFORMATION_CDC25_UP
- REACTOME_PIWI_INTERACTING_RNA_PIRNA_BIOGENESIS, REACTOME_PIWI_INTERACTING_RNA_PIRNA_BIOGENESIS
- KEGG_RNA_POLYMERASE, KEGG_RNA_POLYMERASE
- KIM_WT1_TARGETS_8HR_DN, KIM_WT1_TARGETS_8HR_DN
- REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION_TERMINATION, REACTOME_RNA_POLYMERASE_III_TRANSCRIPTION_TERMINATION
- XU_AKT1_TARGETS_6HR, XU_AKT1_TARGETS_6HR
- REACTOME_RHOBTB_GTPASE_CYCLE, REACTOME_RHOBTB_GTPASE_CYCLE
- HEDENFALK_BREAST_CANCER_BRACX_DN, HEDENFALK_BREAST_CANCER_BRACX_DN
- DAIRKEE_CANCER_PRONE_RESPONSE_BPA, DAIRKEE_CANCER_PRONE_RESPONSE_BPA
- HOLLEMAN_PREDNISOLONE_RESISTANCE_B_ALL_UP, HOLLEMAN_PREDNISOLONE_RESISTANCE_B_ALL_UP
- REACTOME_SIGNALING_BY_FGFR2_IIIA_TM, REACTOME_SIGNALING_BY_FGFR2_IIIA_TM
- LEE_AGING_MUSCLE_UP, LEE_AGING_MUSCLE_UP