

...TARGET_GENES, TOX4_TARGET_GENES

- TMTCGCGANR_UNKNOWN, TMTCGCGANR_UNKNOWN
- REACTOME_CELLULAR_SENESCENCE, REACTOME_CELLULAR_SENESCENCE
- ZBTB5_TARGET_GENES, ZBTB5_TARGET_GENES
- CREB3L2_TARGET_GENES, CREB3L2_TARGET_GENES
- REACTOME_HATS_ACETYLATE_HISTONES, REACTOME_HATS_ACETYLATE_HISTONES
- MXD1_TARGET_GENES, MXD1_TARGET_GENES
- TSHZ1_TARGET_GENES, TSHZ1_TARGET_GENES
- REACTOME_B_WICH_COMPLEX_POSITIVELY_REGULATES_RRNA_EXPRESSION, REACTOME_B_WICH_COMPLEX_POSITIVELY_REGULATES_RRNA_EXPRESSION
- REACTOME_POSITIVE_EPIGENETIC_REGULATION_OF_RRNA_EXPRESSION, REACTOME_POSITIVE_EPIGENETIC_REGULATION_OF_RRNA_EXPRESSION
- REACTOME_SENESCENCE_ASSOCIATED_SECRETORY_PHENOTYPE_SASP, REACTOME_SENESCENCE_ASSOCIATED_SECRETORY_PHENOTYPE_SASP
- REACTOME_EPIGENETIC_REGULATION_OF_GENE_EXPRESSION, REACTOME_EPIGENETIC_REGULATION_OF_GENE_EXPRESSION
- REACTOME_RNA_POLYMERASE_I_PROMOTER_ESCAPE, REACTOME_RNA_POLYMERASE_I_PROMOTER_ESCAPE
- REACTOME_DNA_DAMAGE_TELOMERE_STRESS_INDUCED_SENESCENCE, REACTOME_DNA_DAMAGE_TELOMERE_STRESS_INDUCED_SENESCENCE
- REACTOME_HDACS_DEACETYLATE_HISTONES, REACTOME_HDACS_DEACETYLATE_HISTONES
- HOXC13_TARGET_GENES, HOXC13_TARGET_GENES
- GOBP_VESICLE_MEDIATED_TRANSPORT_TO_THE_PLASMA_MEMBRANE, GOBP_VESICLE_MEDIATED_TRANSPORT_TO_THE_PLASMA_MEMBRANE
- REACTOME_FORMATION_OF_THE_BETA_CATENIN_TCF_TRANSACTIVATING_COMPLEX, REACTOME_FORMATION_OF_THE_BETA_CATENIN_TCF_TRANSACTIVATING_COMPLEX
- ZNF584_TARGET_GENES, ZNF584_TARGET_GENES
- REACTOME_MEIOSIS, REACTOME_MEIOSIS
- REACTOME_SIRT1_NEGATIVELY_REGULATES_RRNA_EXPRESSION, REACTOME_SIRT1_NEGATIVELY_REGULATES_RRNA_EXPRESSION
- TERT_TARGET_GENES, TERT_TARGET_GENES
- REACTOME_NEGATIVE_EPIGENETIC_REGULATION_OF_RRNA_EXPRESSION, REACTOME_NEGATIVE_EPIGENETIC_REGULATION_OF_RRNA_EXPRESSION
- REACTOME_INHIBITION_OF_DNA_RECOMBINATION_AT_TELOMERE, REACTOME_INHIBITION_OF_DNA_RECOMBINATION_AT_TELOMERE
- REACTOME_ERCC6_CSB_AND_EHMT2_G9A_POSITIVELY_REGULATE_RRNA_EXPRESSION, REACTOME_ERCC6_CSB_AND_EHMT2_G9A_POSITIVELY_REGULATE_RRNA_EXPRESSION
- REACTOME_DISEASES_OF_PROGRAMMED_CELL_DEATH, REACTOME_DISEASES_OF_PROGRAMMED_CELL_DEATH
- REACTOME_REPRODUCTION, REACTOME_REPRODUCTION
- REACTOME_RMTS_METHYLATE_HISTONE_ARGININES, REACTOME_RMTS_METHYLATE_HISTONE_ARGININES
- REACTOME_ACTIVATED_PKN1_STIMULATES_TRANSCRIPTION_OF_AR_ANDROGEN_RECEPTOR_REGULATED_GENES_KLK2_AND_KLK3, REACTOME_ACTIVATED_PKN1_STIMULATES_TRANSCRIPTION_OF_AR_ANDROGEN_RECEPTOR_REGULATED_C
- REACTOME_CONDENSATION_OF_PROPHASE_CHROMOSOMES, REACTOME_CONDENSATION_OF_PROPHASE_CHROMOSOMES
- REACTOME_MEIOTIC_SYNAPSIS, REACTOME_MEIOTIC_SYNAPSIS
- KRCTCNNNNMANAGC_UNKNOWN, KRCTCNNNNMANAGC_UNKNOWN
- REACTOME_DNA_METHYLATION, REACTOME_DNA_METHYLATION
- CENPT_TARGET_GENES, CENPT_TARGET_GENES
- DICER1_TARGET_GENES, DICER1_TARGET_GENES
- REACTOME_MEIOTIC_RECOMBINATION, REACTOME_MEIOTIC_RECOMBINATION
- REACTOME_BASE_EXCISION_REPAIR_AP_SITE_FORMATION, REACTOME_BASE_EXCISION_REPAIR_AP_SITE_FORMATION
- REACTOME_PRC2_METHYLATES_HISTONES_AND_DNA, REACTOME_PRC2_METHYLATES_HISTONES_AND_DNA
- REACTOME_RECOGNITION_AND_ASSOCIATION_OF_DNA_GLYCOSYLASE_WITH_SITE_CONTAINING_AN_AFFECTED_PURINE, REACTOME_RECOGNITION_AND_ASSOCIATION_OF_DNA_GLYCOSYLASE_WITH_SITE_CONTAINING_AN_AFFECTED_PU
- PPARA_02, PPARA_02
- REACTOME_AMYLOID_FIBER_FORMATION, REACTOME_AMYLOID_FIBER_FORMATION
- REACTOME_PRE_NOTCH_EXPRESSION_AND_PROCESSING, REACTOME_PRE_NOTCH_EXPRESSION_AND_PROCESSING
- MIR4665_5P, MIR4665_5P
- WP_HISTONE_MODIFICATIONS, WP_HISTONE_MODIFICATIONS
- MIR3675_3P, MIR3675_3P
- REACTOME_METALLOPROTEASE_DUBS, REACTOME_METALLOPROTEASE_DUBS
- ZNF582_TARGET_GENES, ZNF582_TARGET_GENES
- GOBP_RDNA_HETEROCHROMATIN_ASSEMBLY, GOBP_RDNA_HETEROCHROMATIN_ASSEMBLY
- GOBP_MALE_MEIOTIC_NUCLEAR_DIVISION, GOBP_MALE_MEIOTIC_NUCLEAR_DIVISION
- GOBP_NEGATIVE_REGULATION_OF_CILIUM_ASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_CILIUM_ASSEMBLY
- ASGHARZADEH_NEUROBLASTOMA_POOR_SURVIVAL_DN, ASGHARZADEH_NEUROBLASTOMA_POOR_SURVIVAL_DN
- GOMF_PHOSPHATASE_INHIBITOR_ACTIVITY, GOMF_PHOSPHATASE_INHIBITOR_ACTIVITY
- GOBP_NEGATIVE_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION
- GOCC_POLE_PLASM, GOCC_POLE_PLASM
- GOBP_POSITIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GOBP_POSITIVE_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
- HP_DISTURBANCE_OF_FACIAL_EXPRESSION, HP_DISTURBANCE_OF_FACIAL_EXPRESSION
- GOBP_CELLULAR_RESPONSE_TO_X_RAY, GOBP_CELLULAR_RESPONSE_TO_X_RAY
- ZNF23_TARGET_GENES, ZNF23_TARGET_GENES
- GOBP_NEGATIVE_REGULATION_BY_HOST_OF_VIRAL_TRANSCRIPTION, GOBP_NEGATIVE_REGULATION_BY_HOST_OF_VIRAL_TRANSCRIPTION