

CTIV_PATHWAY, PID_MYC_ACTIV_PATHWAY

BIOCARTA_VDR_PATHWAY, BIOCARTA_VDR_PATHWAY
PID_AURORA_B_PATHWAY, PID_AURORA_B_PATHWAY
WP_TRANSLATION_FACTORS, WP_TRANSLATION_FACTORS
REACTOME_RUNX1_INTERACTS_WITH_CO_FACTORS_WHOSE_PRECISE_EFFECT_ON_RUNX1_TARGETS_IS_NOT_KNOWN, REACTOME_RUNX1_INTERACTS_WITH_CO_FACTORS_WHOSE_PRECISE_EFFECT_ON_RUNX1_TARGETS_IS_NOT_KNOWN
REACTOME_DEADENYLATION_OF_MRNA, REACTOME_DEADENYLATION_OF_MRNA
REACTOME_DISEASES_OF_MITOTIC_CELL_CYCLE, REACTOME_DISEASES_OF_MITOTIC_CELL_CYCLE
PID_TAP63_PATHWAY, PID_TAP63_PATHWAY
PID_MYC_PATHWAY, PID_MYC_PATHWAY
REACTOME_PYROPTOSIS, REACTOME_PYROPTOSIS
REACTOME_MTORC1_MEDIATED_SIGNALLING, REACTOME_MTORC1_MEDIATED_SIGNALLING
REACTOME_REGULATION_OF_PTEN_GENE_TRANSCRIPTION, REACTOME_REGULATION_OF_PTEN_GENE_TRANSCRIPTION
REACTOME_SUMOYLATION_OF_TRANSCRIPTION_COFACTORS, REACTOME_SUMOYLATION_OF_TRANSCRIPTION_COFACTORS
WP_UNFOLDED_PROTEIN_RESPONSE, WP_UNFOLDED_PROTEIN_RESPONSE
REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION, REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION
PID_SMAD2_3NUCLEAR_PATHWAY, PID_SMAD2_3NUCLEAR_PATHWAY
REACTOME_TRANSCRIPTION_OF_E2F_TARGETS_UNDER_NEGATIVE_CONTROL_BY_DREAM_COMPLEX, REACTOME_TRANSCRIPTION_OF_E2F_TARGETS_UNDER_NEGATIVE_CONTROL_BY_DREAM_COMPLEX
KEGG_BASAL_TRANSCRIPTION_FACTORS, KEGG_BASAL_TRANSCRIPTION_FACTORS
BIOCARTA_EIF_PATHWAY, BIOCARTA_EIF_PATHWAY
REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_ADDITIONAL_CELL_CYCLE_GENES_WHOSE_EXACT_ROLE_IN_THE_P53_PATHWAY_REMAIN_UNCERTAIN, REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_ADDITIONAL_CELL_CYCLE_GENES_WHOSE_EXACT_ROLE_IN_THE_P53_PATHWAY
PID_MYC_REPRESS_PATHWAY, PID_MYC_REPRESS_PATHWAY
PID_RB_1PATHWAY, PID_RB_1PATHWAY