



- REACTOME_COOPERATION_OF_PREFOLDIN_AND_TRIC_CCT_IN_ACTIN_AND_TUBULIN_FOLDING
- REACTOME_FORMATION_OF_TUBULIN_FOLDING_INTERMEDIATES_BY_CCT_TRIC
- GO_PROTEASOME_ACCESSORY_COMPLEX
- REACTOME_CROSS_PRESENTATION_OF_SOLUBLE_EXOGENOUS_ANTIGENS_ENDOSOMES
- GO_ENDOPEPTIDASE_COMPLEX
- KEGG_PROTEASOME
- REACTOME_DNA_REPLICATION_PRE_INITIATION
- GO_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS
- REACTOME_METABOLISM_OF_POLYAMINES
- GO_THREONINE_TYPE_PEPTIDASE_ACTIVITY
- GO_PROTEASOMAL_UBIQUITIN_INDEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GO_PROTEASOME_CORE_COMPLEX
- KEGG_PENTOSE_PHOSPHATE_PATHWAY
- KEGG_FRUCTOSE_AND_MANNOSE_METABOLISM
- GO_GLYCOLYTIC_PROCESS_THROUGH_FRUCTOSE_6_PHOSPHATE
- GO_NUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS
- GO_NUCLEOTIDE_SUGAR_BIOSYNTHETIC_PROCESS
- GO_RIBONUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS
- GO_FICOLIN_1_RICH_GRANULE
- REACTOME_DNA_STRAND_ELONGATION
- GO_CELLULAR_METABOLIC_COMPOUND_SALVAGE
- GO_NUCLEOBASE_BIOSYNTHETIC_PROCESS
- GO_PURINE_NUCLEOBASE_METABOLIC_PROCESS
- KEGG_CYSSTEINE_AND_METHIONINE_METABOLISM
- KEGG_GLUTATHIONE_METABOLISM
- KEGG_DNA_REPLICATION
- KEGG_AMINO_SUGAR_AND_NUCLEOTIDE_SUGAR_METABOLISM
- REACTOME_LAGGING_STRAND_SYNTHESIS