

TRANSFERASE_COMPLEX, GO_PROTEIN_ACETYLTRANSFERASE_COMPLEX

- GO_ATPASE_COMPLEX, GO_ATPASE_COMPLEX
- GO_SAGA_TYPE_COMPLEX, GO_SAGA_TYPE_COMPLEX
- GO_N_ACYLTRANSFERASE_ACTIVITY, GO_N_ACYLTRANSFERASE_ACTIVITY
- GO_H4_H2A_HISTONE_ACETYLTRANSFERASE_COMPLEX, GO_H4_H2A_HISTONE_ACETYLTRANSFERASE_COMPLEX
- GO_PEPTIDASE_COMPLEX, GO_PEPTIDASE_COMPLEX
- GO_SNRNA_TRANSCRIPTION, GO_SNRNA_TRANSCRIPTION
- GO_POSITIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION, GO_POSITIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION
- GO_N_ACETYLTRANSFERASE_ACTIVITY, GO_N_ACETYLTRANSFERASE_ACTIVITY
- GO_HISTONE_H2A_ACETYLTATION, GO_HISTONE_H2A_ACETYLTATION
- GO_STAGA_COMPLEX, GO_STAGA_COMPLEX
- GO_H4_HISTONE_ACETYLTRANSFERASE_COMPLEX, GO_H4_HISTONE_ACETYLTRANSFERASE_COMPLEX
- GO_INO80_TYPE_COMPLEX, GO_INO80_TYPE_COMPLEX
- GO_SAGA_COMPLEX, GO_SAGA_COMPLEX
- GO_SWR1_COMPLEX, GO_SWR1_COMPLEX
- GO_PEPTIDE_N_ACETYLTRANSFERASE_ACTIVITY, GO_PEPTIDE_N_ACETYLTRANSFERASE_ACTIVITY
- GO_TRANSCRIPTION_FACTOR_TFTC_COMPLEX, GO_TRANSCRIPTION_FACTOR_TFTC_COMPLEX
- GO_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC, GO_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC
- GO_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_BINDING, GO_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_BINDING
- GO_TRANSCRIPTION_PREINITIATION_COMPLEX_ASSEMBLY, GO_TRANSCRIPTION_PREINITIATION_COMPLEX_ASSEMBLY
- GO_RNA_POLYMERASE_II_PREINITIATION_COMPLEX_ASSEMBLY, GO_RNA_POLYMERASE_II_PREINITIATION_COMPLEX_ASSEMBLY
- GO_POSITIVE_REGULATION_OF_CHROMATIN_ORGANIZATION, GO_POSITIVE_REGULATION_OF_CHROMATIN_ORGANIZATION
- GO_NUCLEAR_RECEPTOR_BINDING, GO_NUCLEAR_RECEPTOR_BINDING
- GO_NUCLEAR_HORMONE_RECEPTOR_BINDING, GO_NUCLEAR_HORMONE_RECEPTOR_BINDING