

**ACETYLTRANSFERASE\_COMPLEX, GOCC\_PROTEIN\_ACETYLTRANSFERASE\_COMPLEX**

- GOMF\_NUCLEAR\_RECEPTOR\_COACTIVATOR\_ACTIVITY, GOMF\_NUCLEAR\_RECEPTOR\_COACTIVATOR\_ACTIVITY
- GOCC\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_REGULATOR\_COMPLEX, GOCC\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_REGULATOR\_COMPLEX
- GOCC\_HISTONE\_METHYLTRANSFERASE\_COMPLEX, GOCC\_HISTONE\_METHYLTRANSFERASE\_COMPLEX
- GOCC\_TRANSCRIPTION\_FACTOR\_TFIID\_COMPLEX, GOCC\_TRANSCRIPTION\_FACTOR\_TFIID\_COMPLEX
- GOCC\_SWI\_SNF\_SUPERFAMILY\_TYPE\_COMPLEX, GOCC\_SWI\_SNF\_SUPERFAMILY\_TYPE\_COMPLEX
- GOBP\_NEGATIVE\_REGULATION\_OF\_GENE\_EXPRESSION\_EPIGENETIC, GOBP\_NEGATIVE\_REGULATION\_OF\_GENE\_EXPRESSION\_EPIGENETIC
- GOCC\_INO80\_TYPE\_COMPLEX, GOCC\_INO80\_TYPE\_COMPLEX
- GOCC\_MLL1\_2\_COMPLEX, GOCC\_MLL1\_2\_COMPLEX
- GOBP\_HISTONE\_DEUBIQUITINATION, GOBP\_HISTONE\_DEUBIQUITINATION
- GOCC\_HISTONE\_DEACETYLASE\_COMPLEX, GOCC\_HISTONE\_DEACETYLASE\_COMPLEX
- GOBP\_HISTONE\_H2A\_ACETYLATION, GOBP\_HISTONE\_H2A\_ACETYLATION
- GOCC\_SAGA\_TYPE\_COMPLEX, GOCC\_SAGA\_TYPE\_COMPLEX
- GOBP\_POSITIVE\_REGULATION\_OF\_HISTONE\_METHYLATION, GOBP\_POSITIVE\_REGULATION\_OF\_HISTONE\_METHYLATION
- GOBP\_CHROMATIN\_SILENCING, GOBP\_CHROMATIN\_SILENCING
- GOBP\_POSITIVE\_REGULATION\_OF\_CELLULAR\_RESPONSE\_TO\_TRANSFORMING\_GROWTH\_FACTOR\_BETA\_STIMULUS, GOBP\_POSITIVE\_REGULATION\_OF\_CELLULAR\_RESPONSE\_TO\_TRANSFORMING\_GROWTH\_FACTOR\_BETA\_STIMULUS
- GOCC\_SWR1\_COMPLEX, GOCC\_SWR1\_COMPLEX
- GOMF\_ADP\_BINDING, GOMF\_ADP\_BINDING
- GOCC\_DNA\_HELICASE\_COMPLEX, GOCC\_DNA\_HELICASE\_COMPLEX
- GOBP\_REGULATION\_OF\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE\_BY\_P53\_CLASS\_MEDIATOR, GOBP\_REGULATION\_OF\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE\_BY\_P53\_CLASS\_MEDIATOR
- GOCC\_H4\_H2A\_HISTONE\_ACETYLTRANSFERASE\_COMPLEX, GOCC\_H4\_H2A\_HISTONE\_ACETYLTRANSFERASE\_COMPLEX