

SE\_COMPLEX, GO\_ATPASE\_COMPLEX

GO\_H4\_H2A\_HISTONE\_ACETYLTRANSFERASE\_COMPLEX, GO\_H4\_H2A\_HISTONE\_ACETYLTRANSFERASE\_COMPLEX  
GO\_HISTONE\_H2A\_ACETYLTATION, GO\_HISTONE\_H2A\_ACETYLTATION  
GO\_PROTEIN\_ACETYLTRANSFERASE\_COMPLEX, GO\_PROTEIN\_ACETYLTRANSFERASE\_COMPLEX  
GO\_N\_ACYLTRANSFERASE\_ACTIVITY, GO\_N\_ACYLTRANSFERASE\_ACTIVITY  
GO\_PEPTIDASE\_COMPLEX, GO\_PEPTIDASE\_COMPLEX  
GO\_INO80\_TYPE\_COMPLEX, GO\_INO80\_TYPE\_COMPLEX  
GO\_N\_ACETYLTRANSFERASE\_ACTIVITY, GO\_N\_ACETYLTRANSFERASE\_ACTIVITY  
GO\_CHROMATIN\_ASSEMBLY\_OR\_DISASSEMBLY, GO\_CHROMATIN\_ASSEMBLY\_OR\_DISASSEMBLY  
GO\_H4\_HISTONE\_ACETYLTRANSFERASE\_COMPLEX, GO\_H4\_HISTONE\_ACETYLTRANSFERASE\_COMPLEX  
GO\_HISTONE\_H4\_ACETYLTATION, GO\_HISTONE\_H4\_ACETYLTATION  
GO\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GO\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS  
GO\_DNA\_DEPENDENT\_DNA\_REPLICATION, GO\_DNA\_DEPENDENT\_DNA\_REPLICATION  
GO\_LYSINE\_ACETYLATED\_HISTONE\_BINDING, GO\_LYSINE\_ACETYLATED\_HISTONE\_BINDING  
GO\_SIN3\_TYPE\_COMPLEX, GO\_SIN3\_TYPE\_COMPLEX  
GO\_STEROID\_HORMONE\_RECEPTOR\_BINDING, GO\_STEROID\_HORMONE\_RECEPTOR\_BINDING  
GO\_NUCLEAR\_RECEPTOR\_BINDING, GO\_NUCLEAR\_RECEPTOR\_BINDING  
GO\_CHROMATIN\_REMODELING\_AT\_CENTROMERE, GO\_CHROMATIN\_REMODELING\_AT\_CENTROMERE  
GO\_CENTROMERE\_COMPLEX\_ASSEMBLY, GO\_CENTROMERE\_COMPLEX\_ASSEMBLY  
GO\_SWR1\_COMPLEX, GO\_SWR1\_COMPLEX  
GO\_ATP\_DEPENDENT\_CHROMATIN\_REMODELING, GO\_ATP\_DEPENDENT\_CHROMATIN\_REMODELING  
GO\_RESPONSE\_TO\_UV, GO\_RESPONSE\_TO\_UV  
GO\_CHROMATIN\_REMODELING, GO\_CHROMATIN\_REMODELING  
GO\_POSITIVE\_REGULATION\_OF\_TELOMERASE\_RNA\_LOCALIZATION\_TO\_CAJAL\_BODY, GO\_POSITIVE\_REGULATION\_OF\_TELOMERASE\_RNA\_LOCALIZATION\_TO\_CAJAL\_BODY  
GO\_PEPTIDE\_N\_ACETYLTRANSFERASE\_ACTIVITY, GO\_PEPTIDE\_N\_ACETYLTRANSFERASE\_ACTIVITY