

PENDENT\_PROTEIN\_BINDING, GOMF\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING

- GOBP\_PROTEIN\_METHYLATION, GOBP\_PROTEIN\_METHYLATION
- GOCC\_PROTEIN\_ACETYLTRANSFERASE\_COMPLEX, GOCC\_PROTEIN\_ACETYLTRANSFERASE\_COMPLEX
- GOBP\_NEGATIVE\_REGULATION\_OF\_GENE\_EXPRESSION\_EPIGENETIC, GOBP\_NEGATIVE\_REGULATION\_OF\_GENE\_EXPRESSION\_EPIGENETIC
- GOMF\_UBIQUITIN\_LIKE\_PROTEIN\_BINDING, GOMF\_UBIQUITIN\_LIKE\_PROTEIN\_BINDING
- GOCC\_ENDOPEPTIDASE\_COMPLEX, GOCC\_ENDOPEPTIDASE\_COMPLEX
- GOBP\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS
- GOBP\_REGULATION\_OF\_PROTEASOMAL\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_PROTEASOMAL\_PROTEIN\_CATABOLIC\_PROCESS
- GOCC\_HISTONE\_METHYLTRANSFERASE\_COMPLEX, GOCC\_HISTONE\_METHYLTRANSFERASE\_COMPLEX
- GOCC\_PEPTIDASE\_COMPLEX, GOCC\_PEPTIDASE\_COMPLEX
- GOBP\_CELLULAR\_RESPONSE\_TO\_TOPOLOGICALLY\_INCORRECT\_PROTEIN, GOBP\_CELLULAR\_RESPONSE\_TO\_TOPOLOGICALLY\_INCORRECT\_PROTEIN
- GOBP\_ERAD\_PATHWAY, GOBP\_ERAD\_PATHWAY
- GOBP\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PRO
- GOCC\_SWI\_SNF\_SUPERFAMILY\_TYPE\_COMPLEX, GOCC\_SWI\_SNF\_SUPERFAMILY\_TYPE\_COMPLEX
- GOMF\_UBIQUITIN\_BINDING, GOMF\_UBIQUITIN\_BINDING
- GOMF\_TRANSCRIPTION\_COREPRESSOR\_ACTIVITY, GOMF\_TRANSCRIPTION\_COREPRESSOR\_ACTIVITY
- GOMF\_POLYUBIQUITIN\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING, GOMF\_POLYUBIQUITIN\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING
- GOCC\_H4\_HISTONE\_ACETYLTRANSFERASE\_COMPLEX, GOCC\_H4\_HISTONE\_ACETYLTRANSFERASE\_COMPLEX
- GOMF\_NUCLEAR\_HORMONE\_RECEPTOR\_BINDING, GOMF\_NUCLEAR\_HORMONE\_RECEPTOR\_BINDING
- GOBP\_HISTONE\_H4\_ACETYLATION, GOBP\_HISTONE\_H4\_ACETYLATION
- GOMF\_N\_METHYLTRANSFERASE\_ACTIVITY, GOMF\_N\_METHYLTRANSFERASE\_ACTIVITY
- GOMF\_PROTEIN\_METHYLTRANSFERASE\_ACTIVITY, GOMF\_PROTEIN\_METHYLTRANSFERASE\_ACTIVITY
- GOBP\_POSITIVE\_REGULATION\_OF\_WNT\_SIGNALING\_PATHWAY, GOBP\_POSITIVE\_REGULATION\_OF\_WNT\_SIGNALING\_PATHWAY
- GOBP\_UBIQUITIN\_DEPENDENT\_ERAD\_PATHWAY, GOBP\_UBIQUITIN\_DEPENDENT\_ERAD\_PATHWAY
- GOBP\_POSITIVE\_REGULATION\_OF\_CHROMATIN\_ORGANIZATION, GOBP\_POSITIVE\_REGULATION\_OF\_CHROMATIN\_ORGANIZATION
- GOBP\_AUTOPHAGOSOME\_MATURATION, GOBP\_AUTOPHAGOSOME\_MATURATION
- GOBP\_NEGATIVE\_REGULATION\_OF\_PROTEOLYSIS\_INVOLVED\_IN\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_NEGATIVE\_REGULATION\_OF\_PROTEOLYSIS\_INVOLVED\_IN\_CELLULAR\_PROT
- GOBP\_HISTONE\_H3\_K4\_METHYLATION, GOBP\_HISTONE\_H3\_K4\_METHYLATION
- GOCC\_NUCLEAR\_UBIQUITIN\_LIGASE\_COMPLEX, GOCC\_NUCLEAR\_UBIQUITIN\_LIGASE\_COMPLEX
- GOCC\_AUTOPHAGOSOME, GOCC\_AUTOPHAGOSOME
- GOBP\_HEMATOPOIETIC\_PROGENITOR\_CELL\_DIFFERENTIATION, GOBP\_HEMATOPOIETIC\_PROGENITOR\_CELL\_DIFFERENTIATION
- GOBP\_REGULATION\_OF\_HEMATOPOIETIC\_PROGENITOR\_CELL\_DIFFERENTIATION, GOBP\_REGULATION\_OF\_HEMATOPOIETIC\_PROGENITOR\_CELL\_DIFFERENTIATION
- GOCC\_MLL1\_2\_COMPLEX, GOCC\_MLL1\_2\_COMPLEX
- GOBP\_INTERLEUKIN\_1\_MEDIATED\_SIGNALING\_PATHWAY, GOBP\_INTERLEUKIN\_1\_MEDIATED\_SIGNALING\_PATHWAY
- GOBP\_DNA\_MODIFICATION, GOBP\_DNA\_MODIFICATION
- GOBP\_POSITIVE\_REGULATION\_OF\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS
- GOBP\_POSITIVE\_REGULATION\_OF\_PROTEOLYSIS\_INVOLVED\_IN\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEOLYSIS\_INVOLVED\_IN\_CELLULAR\_PROTEIN
- GOCC\_HISTONE\_DEACETYLASE\_COMPLEX, GOCC\_HISTONE\_DEACETYLASE\_COMPLEX
- GOBP\_POSITIVE\_REGULATION\_OF\_RESPONSE\_TO\_ENDOPLASMIC\_RETICULUM\_STRESS, GOBP\_POSITIVE\_REGULATION\_OF\_RESPONSE\_TO\_ENDOPLASMIC\_RETICULUM\_STRESS
- GOBP\_DNA\_METHYLATION\_OR\_DEMETHYLATION, GOBP\_DNA\_METHYLATION\_OR\_DEMETHYLATION
- GOBP\_PROTEIN\_QUALITY\_CONTROL\_FOR\_MISFOLDED\_OR\_INCOMPLETELY\_SYNTHESIZED\_PROTEINS, GOBP\_PROTEIN\_QUALITY\_CONTROL\_FOR\_MISFOLDED\_OR\_INCOMPLETELY\_SYNTHESIZED\_
- GOBP\_CYTOPLASMIC\_PATTERN\_RECOGNITION\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_CYTOPLASMIC\_PATTERN\_RECOGNITION\_RECEPTOR\_SIGNALING\_PATHWAY
- GOBP\_G0\_TO\_G1\_TRANSITION, GOBP\_G0\_TO\_G1\_TRANSITION
- GOMF\_2\_OXOGLUTARATE\_DEPENDENT\_DIOXYGENASE\_ACTIVITY, GOMF\_2\_OXOGLUTARATE\_DEPENDENT\_DIOXYGENASE\_ACTIVITY
- GOBP\_PROTEIN\_K63\_LINKED\_DEUBIQUITINATION, GOBP\_PROTEIN\_K63\_LINKED\_DEUBIQUITINATION
- GOMF\_TRANSCRIPTION\_COREGULATOR\_BINDING, GOMF\_TRANSCRIPTION\_COREGULATOR\_BINDING
- GOBP\_HISTONE\_DEUBIQUITINATION, GOBP\_HISTONE\_DEUBIQUITINATION
- GOBP\_POSITIVE\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN
- GOBP\_RESPONSE\_TO\_MISFOLDED\_PROTEIN, GOBP\_RESPONSE\_TO\_MISFOLDED\_PROTEIN
- GOMF\_NUCLEOSOME\_BINDING, GOMF\_NUCLEOSOME\_BINDING
- GOBP\_POSITIVE\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS
- GOBP\_TUMOR\_NECROSIS\_FACTOR\_MEDIATED\_SIGNALING\_PATHWAY, GOBP\_TUMOR\_NECROSIS\_FACTOR\_MEDIATED\_SIGNALING\_PATHWAY
- GOMF\_PHOSPHOPROTEIN\_BINDING, GOMF\_PHOSPHOPROTEIN\_BINDING
- GOBP\_HISTONE\_H2A\_ACETYLATION, GOBP\_HISTONE\_H2A\_ACETYLATION
- GOBP\_HISTONE\_H4\_K16\_ACETYLATION, GOBP\_HISTONE\_H4\_K16\_ACETYLATION
- GOCC\_H4\_H2A\_HISTONE\_ACETYLTRANSFERASE\_COMPLEX, GOCC\_H4\_H2A\_HISTONE\_ACETYLTRANSFERASE\_COMPLEX
- GOBP\_MACROMOLECULE\_DEACYLATION, GOBP\_MACROMOLECULE\_DEACYLATION
- GOBP\_RRNA\_TRANSCRIPTION, GOBP\_RRNA\_TRANSCRIPTION
- GOBP\_ACTIVATION\_OF\_INNATE\_IMMUNE\_RESPONSE, GOBP\_ACTIVATION\_OF\_INNATE\_IMMUNE\_RESPONSE
- GOCC\_PCG\_PROTEIN\_COMPLEX, GOCC\_PCG\_PROTEIN\_COMPLEX
- GOBP\_NEGATIVE\_REGULATION\_OF\_B\_CELL\_ACTIVATION, GOBP\_NEGATIVE\_REGULATION\_OF\_B\_CELL\_ACTIVATION
- GOBP\_SELECTIVE\_AUTOPHAGY, GOBP\_SELECTIVE\_AUTOPHAGY
- GOBP\_INCLUSION\_BODY\_ASSEMBLY, GOBP\_INCLUSION\_BODY\_ASSEMBLY
- GOMF\_CYSTEINE\_TYPE\_PEPTIDASE\_ACTIVITY, GOMF\_CYSTEINE\_TYPE\_PEPTIDASE\_ACTIVITY
- GOMF\_LYSINE\_ACETYLATED\_HISTONE\_BINDING, GOMF\_LYSINE\_ACETYLATED\_HISTONE\_BINDING
- GOBP\_HISTONE\_UBIQUITINATION, GOBP\_HISTONE\_UBIQUITINATION
- GOBP\_REGULATION\_OF\_HISTONE\_H3\_K4\_METHYLATION, GOBP\_REGULATION\_OF\_HISTONE\_H3\_K4\_METHYLATION
- GOBP\_REGULATION\_OF\_CHROMATIN\_ASSEMBLY\_OR\_DISASSEMBLY, GOBP\_REGULATION\_OF\_CHROMATIN\_ASSEMBLY\_OR\_DISASSEMBLY
- GOMF\_UBIQUITIN\_SPECIFIC\_PROTEASE\_BINDING, GOMF\_UBIQUITIN\_SPECIFIC\_PROTEASE\_BINDING