

TOPOLOGICALLY\_INCORRECT\_PROTEIN, GOBP\_CELLULAR\_RESPONSE\_TO\_TOPOLOGICALLY\_INCORRECT\_PROTEIN

- GOBP\_ERAD\_PATHWAY, GOBP\_ERAD\_PATHWAY
- GOBP\_UBIQUITIN\_DEPENDENT\_ERAD\_PATHWAY, GOBP\_UBIQUITIN\_DEPENDENT\_ERAD\_PATHWAY
- GOBP\_MYELOID\_CELL\_HOMEOSTASIS, GOBP\_MYELOID\_CELL\_HOMEOSTASIS
- GOBP\_POSITIVE\_REGULATION\_OF\_CYSSTEINE\_TYPE\_ENDOPEPTIDASE\_ACTIVITY, GOBP\_POSITIVE\_REGULATION\_OF\_CYSSTEINE\_TYPE\_ENDOPEPTIDASE\_ACTIVITY
- GOBP\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE, GOBP\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE
- GOBP\_VASCULAR\_ENDOTHELIAL\_GROWTH\_FACTOR\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_VASCULAR\_ENDOTHELIAL\_GROWTH\_FACTOR\_RECEPTOR\_SIGNALING\_PATHWAY
- GOBP\_NEGATIVE\_REGULATION\_OF\_RESPONSE\_TO\_ENDOPLASMIC\_RETICULUM\_STRESS, GOBP\_NEGATIVE\_REGULATION\_OF\_RESPONSE\_TO\_ENDOPLASMIC\_RETICULUM\_STRESS
- GOBP\_PROTEIN\_DEGLYCOSYLATION, GOBP\_PROTEIN\_DEGLYCOSYLATION
- GOBP\_PROTEIN\_K48\_LINKED\_UBIQUITINATION, GOBP\_PROTEIN\_K48\_LINKED\_UBIQUITINATION
- GOBP\_PROTEIN\_DENEDDYLATION, GOBP\_PROTEIN\_DENEDDYLATION
- GOBP\_ACTIVATION\_OF\_CYSSTEINE\_TYPE\_ENDOPEPTIDASE\_ACTIVITY\_INVOLVED\_IN\_APOPTOTIC\_PROCESS, GOBP\_ACTIVATION\_OF\_CYSSTEINE\_TYPE\_ENDOPEPTIDASE\_ACTIVITY\_INVOLVED\_IN\_APOPTOTIC\_PROCESS
- GOBP\_MITOCHONDRIAL\_FUSION, GOBP\_MITOCHONDRIAL\_FUSION
- GOBP\_RESPONSE\_TO\_MISFOLDED\_PROTEIN, GOBP\_RESPONSE\_TO\_MISFOLDED\_PROTEIN
- GOBP\_REGULATION\_OF\_ENDOPLASMIC\_RETICULUM\_STRESS\_INDUCED\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY, GOBP\_REGULATION\_OF\_ENDOPLASMIC\_RETICULUM\_STRESS\_INDUCED\_INTRINSIC\_APOPTOTIC\_SIGNALING\_P
- GOBP\_CELLULAR\_RESPONSE\_TO\_STEROL\_DEPLETION, GOBP\_CELLULAR\_RESPONSE\_TO\_STEROL\_DEPLETION
- GOBP\_ENDOPLASMIC\_RETICULUM\_TO\_CYTOSOL\_TRANSPORT, GOBP\_ENDOPLASMIC\_RETICULUM\_TO\_CYTOSOL\_TRANSPORT
- GOBP\_MEMBRANE\_PROTEIN\_INTRACELLULAR\_DOMAIN\_PROTEOLYSIS, GOBP\_MEMBRANE\_PROTEIN\_INTRACELLULAR\_DOMAIN\_PROTEOLYSIS
- GOBP\_NUCLEOTIDE\_SUGAR\_METABOLIC\_PROCESS, GOBP\_NUCLEOTIDE\_SUGAR\_METABOLIC\_PROCESS
- GOBP\_PROTEIN\_QUALITY\_CONTROL\_FOR\_MISFOLDED\_OR\_INCOMPLETELY\_SYNTHESIZED\_PROTEINS, GOBP\_PROTEIN\_QUALITY\_CONTROL\_FOR\_MISFOLDED\_OR\_INCOMPLETELY\_SYNTHESIZED\_PROTEINS
- GOBP\_ENDOCRINE\_PANCREAS\_DEVELOPMENT, GOBP\_ENDOCRINE\_PANCREAS\_DEVELOPMENT
- GOBP\_RESPONSE\_TO\_STEROL\_DEPLETION, GOBP\_RESPONSE\_TO\_STEROL\_DEPLETION
- GOBP\_NEGATIVE\_REGULATION\_OF\_ENDOPLASMIC\_RETICULUM\_UNFOLDED\_PROTEIN\_RESPONSE, GOBP\_NEGATIVE\_REGULATION\_OF\_ENDOPLASMIC\_RETICULUM\_UNFOLDED\_PROTEIN\_RESPONSE
- GOBP\_ENDOPLASMIC\_RETICULUM\_CALCIIUM\_ION\_HOMEOSTASIS, GOBP\_ENDOPLASMIC\_RETICULUM\_CALCIIUM\_ION\_HOMEOSTASIS
- GOBP\_PROTEIN\_FOLDING\_IN\_ENDOPLASMIC\_RETICULUM, GOBP\_PROTEIN\_FOLDING\_IN\_ENDOPLASMIC\_RETICULUM
- GOBP\_ETHANOL\_CATABOLIC\_PROCESS, GOBP\_ETHANOL\_CATABOLIC\_PROCESS
- GOBP\_REGULATION\_OF\_APOPTOTIC\_DNA\_FRAGMENTATION, GOBP\_REGULATION\_OF\_APOPTOTIC\_DNA\_FRAGMENTATION
- GOBP\_REGULATION\_OF\_DNA\_CATABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_DNA\_CATABOLIC\_PROCESS
- GOBP\_PROTEIN\_K63\_LINKED\_UBIQUITINATION, GOBP\_PROTEIN\_K63\_LINKED\_UBIQUITINATION
- GOBP\_SYNAPTONEMAL\_COMPLEX\_ORGANIZATION, GOBP\_SYNAPTONEMAL\_COMPLEX\_ORGANIZATION
- GOBP\_POST\_EMBRYONIC\_ANIMAL\_MORPHOGENESIS, GOBP\_POST\_EMBRYONIC\_ANIMAL\_MORPHOGENESIS
- GOBP\_REGULATION\_OF\_MITOCHONDRIAL\_OUTER\_MEMBRANE\_PERMEABILIZATION\_INVOLVED\_IN\_APOPTOTIC\_SIGNALING\_PATHWAY, GOBP\_REGULATION\_OF\_MITOCHONDRIAL\_OUTER\_MEMBRANE\_PERMEABILIZATION\_INVOLV
- GOBP\_MANNOSYLATION, GOBP\_MANNOSYLATION
- GOBP\_PROTEIN\_DEMANNOSYLATION, GOBP\_PROTEIN\_DEMANNOSYLATION
- GOBP\_POST\_EMBRYONIC\_ANIMAL\_ORGAN\_MORPHOGENESIS, GOBP\_POST\_EMBRYONIC\_ANIMAL\_ORGAN\_MORPHOGENESIS
- GOBP\_PROTEIN\_MANNOSYLATION, GOBP\_PROTEIN\_MANNOSYLATION
- GOBP\_FLAVONOID\_METABOLIC\_PROCESS, GOBP\_FLAVONOID\_METABOLIC\_PROCESS
- GOBP\_NUCLEAR\_MEMBRANE\_ORGANIZATION, GOBP\_NUCLEAR\_MEMBRANE\_ORGANIZATION
- GOBP\_RESPONSE\_TO\_LITHIUM\_ION, GOBP\_RESPONSE\_TO\_LITHIUM\_ION
- GOBP\_HEPARAN\_SULFATE\_PROTEOGLYCAN\_BIOSYNTHETIC\_PROCESS, GOBP\_HEPARAN\_SULFATE\_PROTEOGLYCAN\_BIOSYNTHETIC\_PROCESS
- GOBP\_ER\_NUCLEUS\_SIGNALING\_PATHWAY, GOBP\_ER\_NUCLEUS\_SIGNALING\_PATHWAY
- GOBP\_PROTEIN\_O\_LINKED\_MANNOSYLATION, GOBP\_PROTEIN\_O\_LINKED\_MANNOSYLATION
- GOBP\_NEGATIVE\_REGULATION\_OF\_ENDOPLASMIC\_RETICULUM\_STRESS\_INDUCED\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY, GOBP\_NEGATIVE\_REGULATION\_OF\_ENDOPLASMIC\_RETICULUM\_STRESS\_INDUCED\_INTRINSIC\_APO
- GOBP\_3\_PHOSPHOADENOSINE\_5\_PHOSPHOSULFATE\_METABOLIC\_PROCESS, GOBP\_3\_PHOSPHOADENOSINE\_5\_PHOSPHOSULFATE\_METABOLIC\_PROCESS
- GOBP\_REGULATION\_OF\_TRANSLATIONAL\_INITIATION\_BY\_EIF2\_ALPHA\_PHOSPHORYLATION, GOBP\_REGULATION\_OF\_TRANSLATIONAL\_INITIATION\_BY\_EIF2\_ALPHA\_PHOSPHORYLATION