

GOBP\_PROTEIN\_METHYLTRANSFERASE\_COMPLEX, GOBP\_PROTEIN\_METHYLTRANSFERASE\_COMPLEX  
GOCC\_METHYLTRANSFERASE\_COMPLEX, GOCC\_METHYLTRANSFERASE\_COMPLEX  
GOCC\_PROTEIN\_ACETYLTTRANSFERASE\_COMPLEX, GOCC\_PROTEIN\_ACETYLTTRANSFERASE\_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  
GOBP\_REGULATION\_OF\_PROTEASOMAL\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_PROTEASOMAL\_PROTEIN\_CATABOLIC\_PROCESS  
GOBP\_PEPTIDYL\_LYSINE\_METHYLATION, GOBP\_PEPTIDYL\_LYSINE\_METHYLATION  
GOBP\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS  
GOBP\_HISTONE\_METHYLATION, GOBP\_HISTONE\_METHYLATION  
GOCC\_HISTONE\_METHYLTRANSFERASE\_COMPLEX, GOCC\_HISTONE\_METHYLTRANSFERASE\_COMPLEX  
GOCC\_ENDOPEPTIDASE\_COMPLEX, GOCC\_ENDOPEPTIDASE\_COMPLEX  
GOBP\_CELLULAR\_RESPONSE\_TO\_TOPOLOGICALLY\_INCORRECT\_PROTEIN, GOBP\_CELLULAR\_RESPONSE\_TO\_TOPOLOGICALLY\_INCORRECT\_PROTEIN  
GOCC\_PEPTIDASE\_COMPLEX, GOCC\_PEPTIDASE\_COMPLEX  
GOBP\_ERAD\_PATHWAY, GOBP\_ERAD\_PATHWAY  
GOBP\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS  
GOCC\_SWI\_SNF\_SUPERFAMILY\_TYPE\_COMPLEX, GOCC\_SWI\_SNF\_SUPERFAMILY\_TYPE\_COMPLEX  
GOMF\_UBIQUITIN\_BINDING, GOMF\_UBIQUITIN\_BINDING  
GOBP\_VIRAL\_GENOME\_REPLICATION, GOBP\_VIRAL\_GENOME\_REPLICATION  
GOMF\_POLYUBIQUITIN\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING, GOMF\_POLYUBIQUITIN\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING  
HP\_THICK\_EYEBROW, HP\_THICK\_EYEBROW  
GOMF\_PROTEIN\_METHYLTRANSFERASE\_ACTIVITY, GOMF\_PROTEIN\_METHYLTRANSFERASE\_ACTIVITY  
GOMF\_TRANSCRIPTION\_COREPRESSOR\_ACTIVITY, GOMF\_TRANSCRIPTION\_COREPRESSOR\_ACTIVITY  
GOMF\_NUCLEAR\_HORMONE\_RECEPTOR\_BINDING, GOMF\_NUCLEAR\_HORMONE\_RECEPTOR\_BINDING  
GOCC\_H4\_HISTONE\_ACETYLTTRANSFERASE\_COMPLEX, GOCC\_H4\_HISTONE\_ACETYLTTRANSFERASE\_COMPLEX  
GOBP\_PEPTIDYL\_LYSINE\_ACETYLTATION, GOBP\_PEPTIDYL\_LYSINE\_ACETYLTATION  
GOBP\_HISTONE\_H4\_ACETYLTATION, GOBP\_HISTONE\_H4\_ACETYLTATION  
GOMF\_N\_METHYLTRANSFERASE\_ACTIVITY, GOMF\_N\_METHYLTRANSFERASE\_ACTIVITY  
GOMF\_METHYLATED\_HISTONE\_BINDING, GOMF\_METHYLATED\_HISTONE\_BINDING  
GOBP\_UBIQUITIN\_DEPENDENT\_ERAD\_PATHWAY, GOBP\_UBIQUITIN\_DEPENDENT\_ERAD\_PATHWAY  
GOCC\_AUTOPHAGOSOME, GOCC\_AUTOPHAGOSOME  
GOBP\_NEGATIVE\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_NEGATIVE\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS  
GOBP\_POSITIVE\_REGULATION\_OF\_WNT\_SIGNALING\_PATHWAY, GOBP\_POSITIVE\_REGULATION\_OF\_WNT\_SIGNALING\_PATHWAY  
GOBP\_HETEROCHROMATIN\_ORGANIZATION, GOBP\_HETEROCHROMATIN\_ORGANIZATION  
GOBP\_AUTOPHAGOSOME\_MATURATION, GOBP\_AUTOPHAGOSOME\_MATURATION  
GOBP\_HISTONE\_H3\_K4\_METHYLATION, GOBP\_HISTONE\_H3\_K4\_METHYLATION  
GOBP\_PROTEIN\_ACETYLTATION, GOBP\_PROTEIN\_ACETYLTATION  
GOBP\_REGULATION\_OF\_HISTONE\_METHYLATION, GOBP\_REGULATION\_OF\_HISTONE\_METHYLATION  
GOBP\_POSITIVE\_REGULATION\_OF\_PROTEOLYSIS\_INVOLVED\_IN\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEOLYSIS\_INVOLVED\_IN\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS  
GOBP\_HEMATOPOIETIC\_PROGENITOR\_CELL\_DIFFERENTIATION, GOBP\_HEMATOPOIETIC\_PROGENITOR\_CELL\_DIFFERENTIATION  
GOBP\_POSITIVE\_REGULATION\_OF\_I\_KAPPAB\_KINASE\_NF\_KAPPAB\_SIGNALING, GOBP\_POSITIVE\_REGULATION\_OF\_I\_KAPPAB\_KINASE\_NF\_KAPPAB\_SIGNALING  
HP\_FRONTOTEMPORAL\_DEMENTIA, HP\_FRONTOTEMPORAL\_DEMENTIA  
GOBP\_INTERLEUKIN\_1\_MEDIATED\_SIGNALING\_PATHWAY, GOBP\_INTERLEUKIN\_1\_MEDIATED\_SIGNALING\_PATHWAY  
GOBP\_POSITIVE\_REGULATION\_OF\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS  
GOCC\_MLL1\_2\_COMPLEX, GOCC\_MLL1\_2\_COMPLEX  
GOMF\_LYSINE\_N\_METHYLTRANSFERASE\_ACTIVITY, GOMF\_LYSINE\_N\_METHYLTRANSFERASE\_ACTIVITY  
GOBP\_NEGATIVE\_REGULATION\_OF\_PROTEOLYSIS\_INVOLVED\_IN\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_NEGATIVE\_REGULATION\_OF\_PROTEOLYSIS\_INVOLVED\_IN\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS  
GOBP\_REGULATION\_OF\_HEMATOPOIETIC\_PROGENITOR\_CELL\_DIFFERENTIATION, GOBP\_REGULATION\_OF\_HEMATOPOIETIC\_PROGENITOR\_CELL\_DIFFERENTIATION  
GOCC\_HISTONE\_DEACETYLASE\_COMPLEX, GOCC\_HISTONE\_DEACETYLASE\_COMPLEX  
GOBP\_DNA\_MODIFICATION, GOBP\_DNA\_MODIFICATION  
GOBP\_NEGATIVE\_REGULATION\_OF\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_NEGATIVE\_REGULATION\_OF\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS  
GOBP\_POSITIVE\_REGULATION\_OF\_RESPONSE\_TO\_ENDOPLASMIC\_RETICULUM\_STRESS, GOBP\_POSITIVE\_REGULATION\_OF\_RESPONSE\_TO\_ENDOPLASMIC\_RETICULUM\_STRESS  
GOBP\_PROTEIN\_QUALITY\_CONTROL\_FOR\_MISFOLDED\_OR\_INCOMPLETELY\_SYNTHESIZED\_PROTEINS, GOBP\_PROTEIN\_QUALITY\_CONTROL\_FOR\_MISFOLDED\_OR\_INCOMPLETELY\_SYNTHESIZED\_PROTEINS  
HP\_TELECANTHUS, HP\_TELECANTHUS  
GOMF\_CORE\_PROMOTER\_SEQUENCE\_SPECIFIC\_DNA\_BINDING, GOMF\_CORE\_PROMOTER\_SEQUENCE\_SPECIFIC\_DNA\_BINDING  
GOBP\_ACTIVATION\_OF\_INNATE\_IMMUNE\_RESPONSE, GOBP\_ACTIVATION\_OF\_INNATE\_IMMUNE\_RESPONSE  
GOBP\_CYTOPLASMIC\_PATTERN\_RECOGNITION\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_CYTOPLASMIC\_PATTERN\_RECOGNITION\_RECEPTOR\_SIGNALING\_PATHWAY  
GOBP\_G0\_TO\_G1\_TRANSITION, GOBP\_G0\_TO\_G1\_TRANSITION  
GOBP\_PROTEIN\_K63\_LINKED\_DEUBIQUITINATION, GOBP\_PROTEIN\_K63\_LINKED\_DEUBIQUITINATION  
GOBP\_TRANSCRIPTION\_BY\_RNA\_POLYMERASE\_I, GOBP\_TRANSCRIPTION\_BY\_RNA\_POLYMERASE\_I  
HP\_EMOTIONAL\_LABILITY, HP\_EMOTIONAL\_LABILITY  
GOMF\_TRANSCRIPTION\_COREGULATOR\_BINDING, GOMF\_TRANSCRIPTION\_COREGULATOR\_BINDING  
HP\_HIP\_DYSPLASIA, HP\_HIP\_DYSPLASIA  
GOBP\_HISTONE\_DEUBIQUITINATION, GOBP\_HISTONE\_DEUBIQUITINATION  
GOBP\_HISTONE\_H4\_K5\_ACETYLTATION, GOBP\_HISTONE\_H4\_K5\_ACETYLTATION  
GOMF\_NUCLEOSOME\_BINDING, GOMF\_NUCLEOSOME\_BINDING  
HP\_SHORT\_PHILTRUM, HP\_SHORT\_PHILTRUM  
GOMF\_P53\_BINDING, GOMF\_P53\_BINDING  
HP\_PARAGANGLIOMA, HP\_PARAGANGLIOMA  
GOBP\_POSITIVE\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS  
GOBP\_POSITIVE\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS  
GOBP\_RESPONSE\_TO\_MISFOLDED\_PROTEIN, GOBP\_RESPONSE\_TO\_MISFOLDED\_PROTEIN  
HP\_BROAD\_EYEBROW, HP\_BROAD\_EYEBROW  
GOBP\_HEMATOPOIETIC\_STEM\_CELL\_DIFFERENTIATION, GOBP\_HEMATOPOIETIC\_STEM\_CELL\_DIFFERENTIATION  
GOBP\_HISTONE\_H3\_ACETYLTATION, GOBP\_HISTONE\_H3\_ACETYLTATION  
GOBP\_INCLUSION\_BODY\_ASSEMBLY, GOBP\_INCLUSION\_BODY\_ASSEMBLY  
GOMF\_GENERAL\_TRANSCRIPTION\_INITIATION\_FACTOR\_BINDING, GOMF\_GENERAL\_TRANSCRIPTION\_INITIATION\_FACTOR\_BINDING  
GOBP\_HISTONE\_UBIQUITINATION, GOBP\_HISTONE\_UBIQUITINATION  
GOMF\_PHOSPHOPROTEIN\_BINDING, GOMF\_PHOSPHOPROTEIN\_BINDING  
GOMF\_2\_OXOGLUTARATE\_DEPENDENT\_DIOXYGENASE\_ACTIVITY, GOMF\_2\_OXOGLUTARATE\_DEPENDENT\_DIOXYGENASE\_ACTIVITY  
GOBP\_HISTONE\_H2A\_ACETYLTATION, GOBP\_HISTONE\_H2A\_ACETYLTATION  
GOBP\_MACROMOLECULE\_DEACYLTATION, GOBP\_MACROMOLECULE\_DEACYLTATION  
HP\_BROAD\_NASAL\_TIP, HP\_BROAD\_NASAL\_TIP  
GOBP\_HISTONE\_H4\_K16\_ACETYLTATION, GOBP\_HISTONE\_H4\_K16\_ACETYLTATION  
GOCC\_H4\_H2A\_HISTONE\_ACETYLTTRANSFERASE\_COMPLEX, GOCC\_H4\_H2A\_HISTONE\_ACETYLTTRANSFERASE\_COMPLEX  
GOBP\_NEGATIVE\_REGULATION\_OF\_TYPE\_I\_INTERFERON\_PRODUCTION, GOBP\_NEGATIVE\_REGULATION\_OF\_TYPE\_I\_INTERFERON\_PRODUCTION  
GOBP\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_BY\_P53\_CLASS\_MEDIATOR, GOBP\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_BY\_P53\_CLASS\_MEDIATOR  
GOBP\_RRNA\_TRANSCRIPTION, GOBP\_RRNA\_TRANSCRIPTION  
GOMF\_HISTONE\_METHYLTRANSFERASE\_ACTIVITY, GOMF\_HISTONE\_METHYLTRANSFERASE\_ACTIVITY  
HP\_ABNORMAL\_NEURON\_MORPHOLOGY, HP\_ABNORMAL\_NEURON\_MORPHOLOGY  
GOCC\_DNA\_REPAIR\_COMPLEX, GOCC\_DNA\_REPAIR\_COMPLEX  
GOBP\_NEGATIVE\_REGULATION\_OF\_B\_CELL\_ACTIVATION, GOBP\_NEGATIVE\_REGULATION\_OF\_B\_CELL\_ACTIVATION  
GOBP\_REGULATION\_OF\_TYPE\_I\_INTERFERON\_MEDIATED\_SIGNALING\_PATHWAY, GOBP\_REGULATION\_OF\_TYPE\_I\_INTERFERON\_MEDIATED\_SIGNALING\_PATHWAY  
GOBP\_MODULATION\_OF\_PROCESS\_OF\_OTHER\_ORGANISM\_INVOLVED\_IN\_SYMBIOTIC\_INTERACTION, GOBP\_MODULATION\_OF\_PROCESS\_OF\_OTHER\_ORGANISM\_INVOLVED\_IN\_SYMBIOTIC\_INTERACTION  
GOBP\_TUMOR\_NECROSIS\_FACTOR\_MEDIATED\_SIGNALING\_PATHWAY, GOBP\_TUMOR\_NECROSIS\_FACTOR\_MEDIATED\_SIGNALING\_PATHWAY  
HP\_WIDE\_NASAL\_RIDGE, HP\_WIDE\_NASAL\_RIDGE  
GOBP\_SELECTIVE\_AUTOPHAGY, GOBP\_SELECTIVE\_AUTOPHAGY  
GOBP\_INNATE\_IMMUNE\_RESPONSE\_ACTIVATING\_SIGNAL\_TRANSDUCTION, GOBP\_INNATE\_IMMUNE\_RESPONSE\_ACTIVATING\_SIGNAL\_TRANSDUCTION  
GOBP\_PEPTIDYL\_LYSINE\_TRIMETHYLATION, GOBP\_PEPTIDYL\_LYSINE\_TRIMETHYLATION  
GOBP\_TRANSCRIPTION\_INITIATION\_FROM\_RNA\_POLYMERASE\_I\_PROMOTER, GOBP\_TRANSCRIPTION\_INITIATION\_FROM\_RNA\_POLYMERASE\_I\_PROMOTER  
GOBP\_TOLERANCE\_INDUCION, GOBP\_TOLERANCE\_INDUCION  
GOMF\_HISTONE\_LYSINE\_N\_METHYLTRANSFERASE\_ACTIVITY, GOMF\_HISTONE\_LYSINE\_N\_METHYLTRANSFERASE\_ACTIVITY  
HP\_RESTLESSNESS, HP\_RESTLESSNESS  
HP\_ABNORMAL\_LARGE\_INTESTINE\_PHYSIOLOGY, HP\_ABNORMAL\_LARGE\_INTESTINE\_PHYSIOLOGY  
GOBP\_NEGATIVE\_REGULATION\_OF\_CHROMATIN\_ORGANIZATION, GOBP\_NEGATIVE\_REGULATION\_OF\_CHROMATIN\_ORGANIZATION  
GOMF\_LYSINE\_ACETYLTATED\_HISTONE\_BINDING, GOMF\_LYSINE\_ACETYLTATED\_HISTONE\_BINDING  
GOBP\_REGULATION\_OF\_HISTONE\_H3\_K4\_METHYLATION, GOBP\_REGULATION\_OF\_HISTONE\_H3\_K4\_METHYLATION  
GOBP\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE\_BY\_P53\_CLASS\_MEDIATOR, GOBP\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE\_BY\_P53\_CLASS\_MEDIATOR  
HP\_ABNORMAL\_PULMONARY\_VALVE\_MORPHOLOGY, HP\_ABNORMAL\_PULMONARY\_VALVE\_MORPHOLOGY  
GOBP\_REGULATION\_OF\_CHROMATIN\_ASSEMBLY\_OR\_DISASSEMBLY, GOBP\_REGULATION\_OF\_CHROMATIN\_ASSEMBLY\_OR\_DISASSEMBLY  
GOMF\_UBIQUITIN\_SPECIFIC\_PROTEASE\_BINDING, GOMF\_UBIQUITIN\_SPECIFIC\_PROTEASE\_BINDING  
GOBP\_POSITIVE\_REGULATION\_OF\_ATPASE\_ACTIVITY, GOBP\_POSITIVE\_REGULATION\_OF\_ATPASE\_ACTIVITY  
HP\_NEURODEGENERATION, HP\_NEURODEGENERATION  
GOBP\_NEGATIVE\_REGULATION\_OF\_VIRAL\_GENOME\_REPLICATION, GOBP\_NEGATIVE\_REGULATION\_OF\_VIRAL\_GENOME\_REPLICATION  
GOMF\_ACETYLTTRANSFERASE\_ACTIVITY, GOMF\_ACETYLTTRANSFERASE\_ACTIVITY  
GOBP\_REGULATION\_OF\_PROTEIN\_DEUBIQUITINATION, GOBP\_REGULATION\_OF\_PROTEIN\_DEUBIQUITINATION  
GOCC\_PCG\_PROTEIN\_COMPLEX, GOCC\_PCG\_PROTEIN\_COMPLEX  
HP\_APLASIA\_HYPOPLASIA\_OF\_TOE, HP\_APLASIA\_HYPOPLASIA\_OF\_TOE  
GOBP\_RDNA\_HETEROCHROMATIN\_ASSEMBLY, GOBP\_RDNA\_HETEROCHROMATIN\_ASSEMBLY  
GOMF\_PEPTIDE\_N\_ACETYLTTRANSFERASE\_ACTIVITY, GOMF\_PEPTIDE\_N\_ACETYLTTRANSFERASE\_ACTIVITY  
GOMF\_K63\_LINKED\_POLYUBIQUITIN\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING, GOMF\_K63\_LINKED\_POLYUBIQUITIN\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING  
GOCC\_TRANSCRIPTION\_FACTOR\_TFIID\_COMPLEX, GOCC\_TRANSCRIPTION\_FACTOR\_TFIID\_COMPLEX  
GOCC\_CHROMATIN\_SILENCING\_COMPLEX, GOCC\_CHROMATIN\_SILENCING\_COMPLEX  
GOBP\_ER\_ASSOCIATED\_MISFOLDED\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_ER\_ASSOCIATED\_MISFOLDED\_PROTEIN\_CATABOLIC\_PROCESS  
HP\_DEATH\_IN\_EARLY\_ADULTHOOD, HP\_DEATH\_IN\_EARLY\_ADULTHOOD  
GOBP\_POSITIVE\_REGULATION\_OF\_HISTONE\_METHYLATION, GOBP\_POSITIVE\_REGULATION\_OF\_HISTONE\_METHYLATION  
GOMF\_CYSSTEINE\_TYPE\_PEPTIDASE\_ACTIVITY, GOMF\_CYSSTEINE\_TYPE\_PEPTIDASE\_ACTIVITY  
GOBP\_REGULATION\_OF\_CHROMATIN\_ASSEMBLY, GOBP\_REGULATION\_OF\_CHROMATIN\_ASSEMBLY  
GOMF\_TRANSCRIPTION\_COREPRESSOR\_BINDING, GOMF\_TRANSCRIPTION\_COREPRESSOR\_BINDING  
HP\_PANCREATIC\_ADENOCARCINOMA, HP\_PANCREATIC\_ADENOCARCINOMA  
GOBP\_MYD88\_INDEPENDENT\_TOLL\_LIKE\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_MYD88\_INDEPENDENT\_TOLL\_LIKE\_RECEPTOR\_SIGNALING\_PATHWAY  
GOBP\_INTRACELLULAR\_STEROID\_HORMONE\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_INTRACELLULAR\_STEROID\_HORMONE\_RECEPTOR\_SIGNALING\_PATHWAY  
GOBP\_REGULATION\_OF\_HISTONE\_H3\_K27\_METHYLATION, GOBP\_REGULATION\_OF\_HISTONE\_H3\_K27\_METHYLATION  
HP\_ABNORMAL\_BRONCHUS\_PHYSIOLOGY, HP\_ABNORMAL\_BRONCHUS\_PHYSIOLOGY  
GOCC\_PRC1\_COMPLEX, GOCC\_PRC1\_COMPLEX  
GOMF\_UBIQUITINATION\_LIKE\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING, GOMF\_UBIQUITINATION\_LIKE\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING  
GOBP\_HISTONE\_DEACETYLTATION, GOBP\_HISTONE\_DEACETYLTATION