

METRIC\_CHANGE, GOBP\_DNA\_GEOMETRIC\_CHANGE

- GOMF\_CATALYTIC\_ACTIVITY\_ACTING\_ON\_DNA, GOMF\_CATALYTIC\_ACTIVITY\_ACTING\_ON\_DNA
- GOBP\_TELOMERE\_ORGANIZATION, GOBP\_TELOMERE\_ORGANIZATION
- GOBP\_RECOMBINATIONAL\_REPAIR, GOBP\_RECOMBINATIONAL\_REPAIR
- GOMF\_DNA\_HELICASE\_ACTIVITY, GOMF\_DNA\_HELICASE\_ACTIVITY
- GOBP\_DNA\_INTEGRITY\_CHECKPOINT, GOBP\_DNA\_INTEGRITY\_CHECKPOINT
- GOBP\_REGULATION\_OF\_TRANSCRIPTION\_FROM\_RNA\_POLYMERASE\_II\_PROMOTER\_IN\_RESPONSE\_TO\_HYPOXIA, GOBP\_REGULATION\_OF\_TRANSCRIPTION\_FROM\_RNA\_POLYMERASE\_II\_PROMOTER\_IN\_RESPONSE\_TO\_HYPOXIA
- GOBP\_VIRION\_ASSEMBLY, GOBP\_VIRION\_ASSEMBLY
- GOBP\_TELOMERE\_MAINTENANCE\_VIA\_TELOMERE\_LENGTHENING, GOBP\_TELOMERE\_MAINTENANCE\_VIA\_TELOMERE\_LENGTHENING
- GOBP\_CELL\_CYCLE\_DNA\_REPLICATION, GOBP\_CELL\_CYCLE\_DNA\_REPLICATION
- GOBP\_REGULATION\_OF\_DNA\_TEMPLATED\_TRANSCRIPTION\_IN\_RESPONSE\_TO\_STRESS, GOBP\_REGULATION\_OF\_DNA\_TEMPLATED\_TRANSCRIPTION\_IN\_RESPONSE\_TO\_STRESS
- GOBP\_PEROXISOMAL\_TRANSPORT, GOBP\_PEROXISOMAL\_TRANSPORT
- GOMF\_SINGLE\_STRANDED\_DNA\_HELICASE\_ACTIVITY, GOMF\_SINGLE\_STRANDED\_DNA\_HELICASE\_ACTIVITY
- GOBP\_DNA\_STRAND\_ELONGATION, GOBP\_DNA\_STRAND\_ELONGATION
- GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_DNA\_INCISION, GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_DNA\_INCISION
- GOMF\_3\_5\_EXONUCLEASE\_ACTIVITY, GOMF\_3\_5\_EXONUCLEASE\_ACTIVITY
- GOBP\_INTERFERON\_BETA\_PRODUCTION, GOBP\_INTERFERON\_BETA\_PRODUCTION
- GOBP\_INTERSTRAND\_CROSS\_LINK\_REPAIR, GOBP\_INTERSTRAND\_CROSS\_LINK\_REPAIR
- GOBP\_TELOMERE\_MAINTENANCE\_VIA\_SEMI\_CONSERVATIVE\_REPLICATION, GOBP\_TELOMERE\_MAINTENANCE\_VIA\_SEMI\_CONSERVATIVE\_REPLICATION
- GOCC\_SWI\_SNF\_SUPERFAMILY\_TYPE\_COMPLEX, GOCC\_SWI\_SNF\_SUPERFAMILY\_TYPE\_COMPLEX
- GOCC\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_REGULATOR\_COMPLEX, GOCC\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_REGULATOR\_COMPLEX
- GOBP\_POSTREPLICATION\_REPAIR, GOBP\_POSTREPLICATION\_REPAIR
- GOBP\_BASE\_EXCISION\_REPAIR, GOBP\_BASE\_EXCISION\_REPAIR
- GOBP\_MITOTIC\_DNA\_INTEGRITY\_CHECKPOINT, GOBP\_MITOTIC\_DNA\_INTEGRITY\_CHECKPOINT
- GOMF\_KINASE\_ACTIVATOR\_ACTIVITY, GOMF\_KINASE\_ACTIVATOR\_ACTIVITY
- GOBP\_POSITIVE\_REGULATION\_OF\_TELOMERE\_MAINTENANCE, GOBP\_POSITIVE\_REGULATION\_OF\_TELOMERE\_MAINTENANCE
- GOBP\_DNA\_STRAND\_ELONGATION\_INVOLVED\_IN\_DNA\_REPLICATION, GOBP\_DNA\_STRAND\_ELONGATION\_INVOLVED\_IN\_DNA\_REPLICATION
- GOBP\_TELOMERASE\_RNA\_LOCALIZATION, GOBP\_TELOMERASE\_RNA\_LOCALIZATION
- GOMF\_EXONUCLEASE\_ACTIVITY, GOMF\_EXONUCLEASE\_ACTIVITY
- GOBP\_PROTEIN\_MONOUBIQUITINATION, GOBP\_PROTEIN\_MONOUBIQUITINATION
- GOBP\_POSITIVE\_REGULATION\_OF\_TELOMERE\_MAINTENANCE\_VIA\_TELOMERE\_LENGTHENING, GOBP\_POSITIVE\_REGULATION\_OF\_TELOMERE\_MAINTENANCE\_VIA\_TELOMERE\_LENGTHENING
- GOBP\_ERROR\_PRONE\_TRANSLESION\_SYNTHESIS, GOBP\_ERROR\_PRONE\_TRANSLESION\_SYNTHESIS
- GOMF\_CULLIN\_FAMILY\_PROTEIN\_BINDING, GOMF\_CULLIN\_FAMILY\_PROTEIN\_BINDING
- GOCC\_DNA\_HELICASE\_COMPLEX, GOCC\_DNA\_HELICASE\_COMPLEX
- GOBP\_TRANSLESION\_SYNTHESIS, GOBP\_TRANSLESION\_SYNTHESIS
- GOCC\_INO80\_TYPE\_COMPLEX, GOCC\_INO80\_TYPE\_COMPLEX
- GOCC\_SCF\_UBIQUITIN\_LIGASE\_COMPLEX, GOCC\_SCF\_UBIQUITIN\_LIGASE\_COMPLEX
- GOBP\_POSITIVE\_REGULATION\_OF\_DNA\_BIOSYNTHETIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_DNA\_BIOSYNTHETIC\_PROCESS
- GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_DNA\_DAMAGE\_RECOGNITION, GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_DNA\_DAMAGE\_RECOGNITION
- GOBP\_HISTONE\_UBIQUITINATION, GOBP\_HISTONE\_UBIQUITINATION
- GOCC\_ATPASE\_COMPLEX, GOCC\_ATPASE\_COMPLEX
- GOBP\_ERROR\_FREE\_TRANSLESION\_SYNTHESIS, GOBP\_ERROR\_FREE\_TRANSLESION\_SYNTHESIS
- GOCC\_DNA\_REPLICATION\_PREINITIATION\_COMPLEX, GOCC\_DNA\_REPLICATION\_PREINITIATION\_COMPLEX
- GOBP\_INTERLEUKIN\_1\_MEDIATED\_SIGNALING\_PATHWAY, GOBP\_INTERLEUKIN\_1\_MEDIATED\_SIGNALING\_PATHWAY
- GOBP\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_CHROMOSOME, GOBP\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_CHROMOSOME
- GOCC\_CMG\_COMPLEX, GOCC\_CMG\_COMPLEX
- GOMF\_DAMAGED\_DNA\_BINDING, GOMF\_DAMAGED\_DNA\_BINDING
- GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_DNA\_GAP\_FILLING, GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_DNA\_GAP\_FILLING
- GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_PREINCISION\_COMPLEX\_STABILIZATION, GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_PREINCISION\_COMPLEX\_STABILIZATION
- GOBP\_MITOCHONDRION\_MORPHOGENESIS, GOBP\_MITOCHONDRION\_MORPHOGENESIS
- GOBP\_POSITIVE\_REGULATION\_OF\_CYTOPLASMIC\_TRANSLATION, GOBP\_POSITIVE\_REGULATION\_OF\_CYTOPLASMIC\_TRANSLATION
- GOCC\_CUL4\_RING\_E3\_UBIQUITIN\_LIGASE\_COMPLEX, GOCC\_CUL4\_RING\_E3\_UBIQUITIN\_LIGASE\_COMPLEX
- GOBP\_CELLULAR\_RESPONSE\_TO\_GAMMA\_RADIATION, GOBP\_CELLULAR\_RESPONSE\_TO\_GAMMA\_RADIATION
- GOBP\_NEGATIVE\_REGULATION\_OF\_DNA\_DEPENDENT\_DNA\_REPLICATION, GOBP\_NEGATIVE\_REGULATION\_OF\_DNA\_DEPENDENT\_DNA\_REPLICATION
- GOBP\_REGULATION\_OF\_DNA\_RECOMBINATION, GOBP\_REGULATION\_OF\_DNA\_RECOMBINATION
- GOBP\_POSITIVE\_REGULATION\_OF\_TELOMERASE\_RNA\_LOCALIZATION\_TO\_CAJAL\_BODY, GOBP\_POSITIVE\_REGULATION\_OF\_TELOMERASE\_RNA\_LOCALIZATION\_TO\_CAJAL\_BODY
- GOBP\_GLOBAL\_GENOME\_NUCLEOTIDE\_EXCISION\_REPAIR, GOBP\_GLOBAL\_GENOME\_NUCLEOTIDE\_EXCISION\_REPAIR
- GOBP\_DNA\_SYNTHESIS\_INVOLVED\_IN\_DNA\_REPAIR, GOBP\_DNA\_SYNTHESIS\_INVOLVED\_IN\_DNA\_REPAIR
- GOBP\_REGULATION\_OF\_DOUBLE\_STRAND\_BREAK\_REPAIR\_VIA\_HOMOLOGOUS\_RECOMBINATION, GOBP\_REGULATION\_OF\_DOUBLE\_STRAND\_BREAK\_REPAIR\_VIA\_HOMOLOGOUS\_RECOMBINATION
- GOCC\_HISTONE\_DEACETYLASE\_COMPLEX, GOCC\_HISTONE\_DEACETYLASE\_COMPLEX
- GOBP\_MITOCHONDRIAL\_GENOME\_MAINTENANCE, GOBP\_MITOCHONDRIAL\_GENOME\_MAINTENANCE
- GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_DNA\_DUPLEX\_UNWINDING, GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_DNA\_DUPLEX\_UNWINDING
- GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_PREINCISION\_COMPLEX\_ASSEMBLY, GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_PREINCISION\_COMPLEX\_ASSEMBLY
- GOCC\_MLL1\_2\_COMPLEX, GOCC\_MLL1\_2\_COMPLEX
- GOBP\_POSITIVE\_REGULATION\_OF\_RESPONSE\_TO\_DNA\_DAMAGE\_STIMULUS, GOBP\_POSITIVE\_REGULATION\_OF\_RESPONSE\_TO\_DNA\_DAMAGE\_STIMULUS
- GOBP\_MITOTIC\_DNA\_REPLICATION, GOBP\_MITOTIC\_DNA\_REPLICATION
- GOBP\_MITOCHONDRIAL\_DNA\_REPLICATION, GOBP\_MITOCHONDRIAL\_DNA\_REPLICATION
- GOBP\_DOUBLE\_STRAND\_BREAK\_REPAIR\_VIA\_BREAK\_INDUCED\_REPLICATION, GOBP\_DOUBLE\_STRAND\_BREAK\_REPAIR\_VIA\_BREAK\_INDUCED\_REPLICATION
- GOBP\_MISMATCH\_REPAIR, GOBP\_MISMATCH\_REPAIR
- GOCC\_NURD\_COMPLEX, GOCC\_NURD\_COMPLEX
- GOMF\_3\_5\_DNA\_HELICASE\_ACTIVITY, GOMF\_3\_5\_DNA\_HELICASE\_ACTIVITY
- GOBP\_G2\_DNA\_DAMAGE\_CHECKPOINT, GOBP\_G2\_DNA\_DAMAGE\_CHECKPOINT
- GOBP\_INTERFERON\_ALPHA\_PRODUCTION, GOBP\_INTERFERON\_ALPHA\_PRODUCTION
- GOBP\_HEMATOPOIETIC\_PROGENITOR\_CELL\_DIFFERENTIATION, GOBP\_HEMATOPOIETIC\_PROGENITOR\_CELL\_DIFFERENTIATION
- GOBP\_POSITIVE\_REGULATION\_OF\_TOLL\_LIKE\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_POSITIVE\_REGULATION\_OF\_TOLL\_LIKE\_RECEPTOR\_SIGNALING\_PATHWAY
- GOBP\_POSITIVE\_REGULATION\_OF\_INTERFERON\_BETA\_PRODUCTION, GOBP\_POSITIVE\_REGULATION\_OF\_INTERFERON\_BETA\_PRODUCTION
- GOBP\_NEUTROPHIL\_HOMEOSTASIS, GOBP\_NEUTROPHIL\_HOMEOSTASIS
- GOMF\_DNA\_POLYMERASE\_BINDING, GOMF\_DNA\_POLYMERASE\_BINDING
- GOBP\_NEGATIVE\_REGULATION\_OF\_TRANSFORMING\_GROWTH\_FACTOR\_BETA\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_NEGATIVE\_REGULATION\_OF\_TRANSFORMING\_GROWTH\_FACTOR\_BETA\_RECEPTOR\_SIGNALING\_PATHWAY