

GOBP\_DNA\_INTEGRITY\_CHECKPOINT

GOBP\_TELOMERE\_ORGANIZATION, GOBP\_TELOMERE\_ORGANIZATION, GOBP\_CATALYTIC\_ACTIVITY\_ACTING\_ON\_DNA, GOMF\_CATALYTIC\_ACTIVITY\_ACTING\_ON\_DNA, GOBP\_DNA\_DEPENDENT\_DNA\_REPLICATION, GOBP\_DNA\_DEPENDENT\_DNA\_REPLICATION, GOBP\_REGULATION\_OF\_PROTEASOMAL\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_PROTEASOMAL\_PROTEIN\_CATABOLIC\_PROCESS, GOMF\_SINGLE\_STRANDED\_DNA\_BINDING, GOMF\_SINGLE\_STRANDED\_DNA\_BINDING, GOBP\_REGULATION\_OF\_SIGNAL\_TRANSDUCTION\_BY\_P53\_CLASS\_MEDIATOR, GOBP\_REGULATION\_OF\_SIGNAL\_TRANSDUCTION\_BY\_P53\_CLASS\_MEDIATOR, GOBP\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_RESPONSE\_TO\_IONIZING\_RADIATION, GOBP\_RESPONSE\_TO\_IONIZING\_RADIATION, GOBP\_DNA\_GEOMETRIC\_CHANGE, GOBP\_DNA\_GEOMETRIC\_CHANGE, GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR, GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR, GOBP\_RECOMBINATIONAL\_REPAIR, GOBP\_RECOMBINATIONAL\_REPAIR, GOCC\_REPLICATION\_FORK, GOCC\_REPLICATION\_FORK, GOMF\_PROTEIN\_C\_TERMINUS\_BINDING, GOMF\_PROTEIN\_C\_TERMINUS\_BINDING, GOMF\_3\_5\_EXONUCLEASE\_ACTIVITY, GOMF\_3\_5\_EXONUCLEASE\_ACTIVITY, GOBP\_DNA\_BIOSYNTHETIC\_PROCESS, GOBP\_DNA\_BIOSYNTHETIC\_PROCESS, GOBP\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE, GOBP\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_MODIFICATION\_BY\_SMALL\_PROTEIN\_CONJUGATION\_OR\_REMOVAL, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_MODIFICATION\_BY\_SMALL\_PROTEIN\_CONJUGATION\_OR\_REMOVAL, GOBP\_NON\_RECOMBINATIONAL\_REPAIR, GOBP\_NON\_RECOMBINATIONAL\_REPAIR, GOBP\_CELL\_CYCLE\_DNA\_REPLICATION, GOBP\_CELL\_CYCLE\_DNA\_REPLICATION, GOMF\_EXONUCLEASE\_ACTIVITY, GOMF\_EXONUCLEASE\_ACTIVITY, GOBP\_SIGNAL\_TRANSDUCTION\_IN\_RESPONSE\_TO\_DNA\_DAMAGE, GOBP\_SIGNAL\_TRANSDUCTION\_IN\_RESPONSE\_TO\_DNA\_DAMAGE, GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_CYCLE\_G2\_M\_PHASE\_TRANSITION, GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_CYCLE\_G2\_M\_PHASE\_TRANSITION, GOCC\_CULLIN\_RING\_UBIQUITIN\_LIGASE\_COMPLEX, GOCC\_CULLIN\_RING\_UBIQUITIN\_LIGASE\_COMPLEX, HP\_NEOPLASM\_OF\_THE\_NERVOUS\_SYSTEM, HP\_NEOPLASM\_OF\_THE\_NERVOUS\_SYSTEM, GOBP\_MITOTIC\_DNA\_INTEGRITY\_CHECKPOINT, GOBP\_MITOTIC\_DNA\_INTEGRITY\_CHECKPOINT, GOBP\_DNA\_DAMAGE\_RESPONSE\_SIGNAL\_TRANSDUCTION\_BY\_P53\_CLASS\_MEDIATOR, GOBP\_DNA\_DAMAGE\_RESPONSE\_SIGNAL\_TRANSDUCTION\_BY\_P53\_CLASS\_MEDIATOR, GOBP\_REGULATION\_OF\_DNA\_REPAIR, GOBP\_REGULATION\_OF\_DNA\_REPAIR, GOBP\_POSITIVE\_REGULATION\_OF\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSTREPLICATION\_REPAIR, GOBP\_POSTREPLICATION\_REPAIR, GOBP\_POSITIVE\_REGULATION\_OF\_INTRACELLULAR\_PROTEIN\_TRANSPORT, GOBP\_POSITIVE\_REGULATION\_OF\_INTRACELLULAR\_PROTEIN\_TRANSPORT, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEOLYSIS\_INVOLVED\_IN\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEOLYSIS\_INVOLVED\_IN\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_INTERSTRAND\_CROSS\_LINK\_REPAIR, GOBP\_INTERSTRAND\_CROSS\_LINK\_REPAIR, HP\_SOMATIC\_MUTATION, HP\_SOMATIC\_MUTATION, HP\_PREMATURELY\_AGED\_APPEARANCE, HP\_PREMATURELY\_AGED\_APPEARANCE, GOBP\_MISMATCH\_REPAIR, GOBP\_MISMATCH\_REPAIR, HP\_PROMINENT\_NOSE, HP\_PROMINENT\_NOSE, GOBP\_SIGNAL\_TRANSDUCTION\_INVOLVED\_IN\_CELL\_CYCLE\_CHECKPOINT, GOBP\_SIGNAL\_TRANSDUCTION\_INVOLVED\_IN\_CELL\_CYCLE\_CHECKPOINT, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_RESPONSE\_TO\_UV, GOBP\_RESPONSE\_TO\_UV, GOMF\_EXONUCLEASE\_ACTIVITY\_ACTIVE\_WITH\_EITHER\_RIBO\_OR\_DEOXYRIBONUCLEIC\_ACIDS\_AND\_PRODUCING\_5\_PHOSPHOMONOESTERS, GOMF\_EXONUCLEASE\_ACTIVITY\_ACTIVE\_WITH\_EITHER\_RIBO\_OR\_DEOXYRIBONUCLEIC\_ACIDS\_AND\_PRODUCING\_5\_PHOSPHOMONOESTERS, HP\_NARROW\_MOUTH, HP\_NARROW\_MOUTH, HP\_CRANIOSYNOSTOSIS, HP\_CRANIOSYNOSTOSIS, GOBP\_MITOTIC\_G1\_S\_TRANSITION\_CHECKPOINT, GOBP\_MITOTIC\_G1\_S\_TRANSITION\_CHECKPOINT, GOBP\_POSITIVE\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_DNA\_SYNTHESIS\_INVOLVED\_IN\_DNA\_REPAIR, GOBP\_DNA\_SYNTHESIS\_INVOLVED\_IN\_DNA\_REPAIR, GOCC\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_REGULATOR\_COMPLEX, GOCC\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_REGULATOR\_COMPLEX, GOMF\_DAMAGED\_DNA\_BINDING, GOMF\_DAMAGED\_DNA\_BINDING, GOBP\_CELLULAR\_RESPONSE\_TO\_IONIZING\_RADIATION, GOBP\_CELLULAR\_RESPONSE\_TO\_IONIZING\_RADIATION, 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GOBP\_TYPE\_I\_INTERFERON\_PRODUCTION, GOBP\_TYPE\_I\_INTERFERON\_PRODUCTION, GOBP\_POSITIVE\_REGULATION\_OF\_TYPE\_I\_INTERFERON\_PRODUCTION, GOBP\_POSITIVE\_REGULATION\_OF\_TYPE\_I\_INTERFERON\_PRODUCTION, GOMF\_PROTEIN\_SERINE\_THREONINE\_KINASE\_ACTIVATOR\_ACTIVITY, GOBP\_POSITIVE\_REGULATION\_OF\_CYSINE\_TYPE\_ENDOPEPTIDASE\_ACTIVITY, GOBP\_POSITIVE\_REGULATION\_OF\_CYSINE\_TYPE\_ENDOPEPTIDASE\_ACTIVITY, HP\_MACULE, HP\_MACULE, GOMF\_EXORIBONUCLEASE\_ACTIVITY, GOMF\_EXORIBONUCLEASE\_ACTIVITY, GOBP\_NEGATIVE\_REGULATION\_OF\_RESPONSE\_TO\_DNA\_DAMAGE\_STIMULUS, GOBP\_NEGATIVE\_REGULATION\_OF\_RESPONSE\_TO\_DNA\_DAMAGE\_STIMULUS, GOBP\_PPTIDYL\_THREONINE\_MODIFICATION, GOBP\_PPTIDYL\_THREONINE\_MODIFICATION, GOMF\_DEOXYRIBONUCLEASE\_ACTIVITY, GOMF\_DEOXYRIBONUCLEASE\_ACTIVITY, GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_PRECISION\_COMPLEX\_ASSEMBLY, GOBP\_NUCLEOTIDE\_EXCISION\_REPAIR\_PRECISION\_COMPLEX\_ASSEMBLY, GOBP\_RESPONSE\_TO\_GAMMA\_RADIATION, GOBP\_RESPONSE\_TO\_GAMMA\_RADIATION, HP\_LARGE\_BASAL\_GANGLIA, HP\_LARGE\_BASAL\_GANGLIA, GOCC\_PROTEIN\_KINASE\_COMPLEX, GOCC\_PROTEIN\_KINASE\_COMPLEX, HP\_ABNORMALITY\_OF\_CHROMOSOME\_STABILITY, HP\_ABNORMALITY\_OF\_CHROMOSOME\_STABILITY, GOBP\_REGULATION\_OF\_DNA\_DAMAGE\_CHECKPOINT, GOBP\_REGULATION\_OF\_DNA\_DAMAGE\_CHECKPOINT, GOMF\_NUCLEAR\_RECEPTOR\_COACTIVATOR\_ACTIVITY, GOMF\_NUCLEAR\_RECEPTOR\_COACTIVATOR\_ACTIVITY, GOCC\_CYCLIN\_DEPENDENT\_PROTEIN\_KINASE\_HOLOENZYME\_COMPLEX, GOCC\_CYCLIN\_DEPENDENT\_PROTEIN\_KINASE\_HOLOENZYME\_COMPLEX, GOMF\_DNA\_REPLICATION\_ORIGIN\_BINDING, GOMF\_DNA\_REPLICATION\_ORIGIN\_BINDING, HP\_ABNORMAL\_FEMORAL\_NECK\_HEAD\_MORPHOLOGY, HP\_ABNORMAL\_FEMORAL\_NECK\_HEAD\_MORPHOLOGY, HP\_SUBMUCOUS\_CLEFT\_OF\_SOFT\_AND\_HARD\_PALATE, HP\_SUBMUCOUS\_CLEFT\_OF\_SOFT\_AND\_HARD\_PALATE, HP\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_BY\_P53\_CLASS\_MEDIATOR, HP\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_BY\_P53\_CLASS\_MEDIATOR, GOBP\_POSITIVE\_REGULATION\_OF\_TELOMERE\_MAINTENANCE, GOBP\_POSITIVE\_REGULATION\_OF\_TELOMERE\_MAINTENANCE, GOBP\_POSITIVE\_REGULATION\_OF\_ORGANELLE\_ASSEMBLY, GOBP\_POSITIVE\_REGULATION\_OF\_ORGANELLE\_ASSEMBLY, GOBP\_REGULATION\_OF\_INTRACELLULAR\_STEROID\_HORMONE\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_REGULATION\_OF\_INTRACELLULAR\_STEROID\_HORMONE\_RECEPTOR\_SIGNALING\_PATHWAY, HP\_ABNORMALITY\_OF\_THE\_UVULA, HP\_ABNORMALITY\_OF\_THE\_UVULA, GOBP\_NEGATIVE\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_NEGATIVE\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, HP\_SHORT\_LONG\_BONE, HP\_SHORT\_LONG\_BONE, GOBP\_PROTEIN\_STABILIZATION, GOBP\_PROTEIN\_STABILIZATION, GOBP\_DNA\_STRAND\_ELONGATION, GOBP\_DNA\_STRAND\_ELONGATION, HP\_SEVERE\_SHORT\_STATURE, HP\_SEVERE\_SHORT\_STATURE, HP\_ABNORMAL\_CAROTID\_ARTERY\_MORPHOLOGY, HP\_ABNORMAL\_CAROTID\_ARTERY\_MORPHOLOGY, HP\_MENINGIOMA, HP\_MENINGIOMA, GOBP\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_MITOCHONDRIAL\_MEMBRANE, GOBP\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_MITOCHONDRIAL\_MEMBRANE, HP\_ABNORMALITY\_OF\_THE\_CHOANAE, HP\_ABNORMALITY\_OF\_THE\_CHOANAE, GOBP\_CELLULAR\_RESPONSE\_TO\_RADIATION, GOBP\_CELLULAR\_RESPONSE\_TO\_RADIATION, GOBP\_REGULATION\_OF\_TRANSCRIPTION\_INVOLVED\_IN\_G1\_S\_TRANSITION\_OF\_MITOTIC\_CELL\_CYCLE, GOBP\_REGULATION\_OF\_TRANSCRIPTION\_INVOLVED\_IN\_G1\_S\_TRANSITION\_OF\_MITOTIC\_CELL\_CYCLE, GOMF\_PROTEIN\_SERINE\_THREONINE\_KINASE\_ACTIVATOR\_ACTIVITY, GOMF\_PROTEIN\_SERINE\_THREONINE\_KINASE\_ACTIVATOR\_ACTIVITY, HP\_BLEPHAROPHIMOSIS, HP\_BLEPHAROPHIMOSIS, GOBP\_MITOCHONDRIAL\_OUTER\_MEMBRANE\_PERMEABILIZATION, GOBP\_MITOCHONDRIAL\_OUTER\_MEMBRANE\_PERMEABILIZATION, HP\_SPINAL\_DYSRAPHISM, HP\_SPINAL\_DYSRAPHISM, GOBP\_NEGATIVE\_REGULATION\_OF\_PROTEASOMAL\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_NEGATIVE\_REGULATION\_OF\_PROTEASOMAL\_PROTEIN\_CATABOLIC\_PROCESS, 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GOMF\_MITOGEN\_ACTIVATED\_PROTEIN\_KINASE\_BINDING, HP\_PROPORTIONATE\_SHORT\_STATURE, HP\_PROPORTIONATE\_SHORT\_STATURE, GOBP\_REGULATION\_OF\_INTRACELLULAR\_ESTROGEN\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_REGULATION\_OF\_INTRACELLULAR\_ESTROGEN\_RECEPTOR\_SIGNALING\_PATHWAY, HP\_FACIAL\_ASYMMETRY, HP\_FACIAL\_ASYMMETRY, HP\_CONVEX\_NASAL\_RIDGE, HP\_CONVEX\_NASAL\_RIDGE, HP\_PROPORTIONATE\_TALL\_STATURE, HP\_PROPORTIONATE\_TALL\_STATURE, HP\_IRREGULAR\_HYPERPIGMENTATION, HP\_IRREGULAR\_HYPERPIGMENTATION, GOMF\_ENDODEOXYRIBONUCLEASE\_ACTIVITY, GOMF\_ENDODEOXYRIBONUCLEASE\_ACTIVITY, HP\_RETINAL\_HEMORRHAGE, HP\_RETINAL\_HEMORRHAGE, GOBP\_CELLULAR\_RESPONSE\_TO\_STEROID\_HORMONE\_STIMULUS, GOBP\_CELLULAR\_RESPONSE\_TO\_STEROID\_HORMONE\_STIMULUS, HP\_PELVIC\_KIDNEY, HP\_PELVIC\_KIDNEY, HP\_SOFT\_TISSUE\_SARCOMA, HP\_SOFT\_TISSUE\_SARCOMA, GOMF\_P53\_BINDING, GOMF\_P53\_BINDING, GOBP\_CELLULAR\_RESPONSE\_TO\_GAMMA\_RADIATION, GOBP\_CELLULAR\_RESPONSE\_TO\_GAMMA\_RADIATION, HP\_ABNORMALITY\_OF\_DNA\_REPAIR, HP\_ABNORMALITY\_OF\_DNA\_REPAIR, HP\_ANOTIA, HP\_ANOTIA, HP\_ABNORMAL\_CLAVICLE\_MORPHOLOGY, HP\_ABNORMAL\_CLAVICLE\_MORPHOLOGY, GOBP\_DNA\_CATABOLIC\_PROCESS, GOBP\_DNA\_CATABOLIC\_PROCESS, HP\_COLORECTAL\_POLYPOSIS, HP\_COLORECTAL\_POLYPOSIS, GOBP\_POSITIVE\_REGULATION\_OF\_GENE\_EXPRESSION\_EPIGENETIC, GOBP\_POSITIVE\_REGULATION\_OF\_GENE\_EXPRESSION\_EPIGENETIC, GOBP\_ENDOPLASMIC\_RETICULUM\_CALCIIUM\_ION\_HOMEOSTASIS, GOBP\_ENDOPLASMIC\_RETICULUM\_CALCIIUM\_ION\_HOMEOSTASIS, GOBP\_EXECUTION\_PHASE\_OF\_APOPTOSIS, GOBP\_EXECUTION\_PHASE\_OF\_APOPTOSIS, GOBP\_POSITIVE\_REGULATION\_OF\_NUCLEOCYTOPLASMIC\_TRANSPORT, GOBP\_POSITIVE\_REGULATION\_OF\_NUCLEOCYTOPLASMIC\_TRANSPORT, HP\_DECREASED\_CIRCULATING\_ANTIODY\_LEVEL, HP\_DECREASED\_CIRCULATING\_ANTIODY\_LEVEL, HP\_GLIOMA, HP\_GLIOMA, GOBP\_PROTEIN\_INSERTION\_INTO\_MEMBRANE, GOBP\_PROTEIN\_INSERTION\_INTO\_MEMBRANE, GOBP\_DNA\_CATABOLIC\_PROCESS\_ENDONUCLEOLYTIC, GOBP\_DNA\_CATABOLIC\_PROCESS\_ENDONUCLEOLYTIC, HP\_ECTOPIC\_KIDNEY, HP\_ECTOPIC\_KIDNEY, HP\_DEFECTIVE\_DNA\_REPAIR\_AFTER\_ULTRAVIOLET\_RADIATION\_DAMAGE, HP\_DEFECTIVE\_DNA\_REPAIR\_AFTER\_ULTRAVIOLET\_RADIATION\_DAMAGE, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_LOCALIZATION\_TO\_MEMBRANE, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_LOCALIZATION\_TO\_MEMBRANE, HP\_TELANGIECTASIA\_OF\_THE\_SKIN, HP\_TELANGIECTASIA\_OF\_THE\_SKIN, HP\_ABNORMALITY\_OF\_THE\_ANUS, HP\_ABNORMALITY\_OF\_THE\_ANUS, GOMF\_MISMATCHED\_DNA\_BINDING, GOMF\_MISMATCHED\_DNA\_BINDING, HP\_ABNORMAL\_MEGAKARYOCYTE\_MORPHOLOGY, HP\_ABNORMAL\_MEGAKARYOCYTE\_MORPHOLOGY, HP\_HYPERMELANOTIC\_MACULE, HP\_HYPERMELANOTIC\_MACULE, GOBP\_REGULATION\_OF\_DNA\_RECOMBINATION, GOBP\_REGULATION\_OF\_DNA\_RECOMBINATION, GOBP\_DNA\_DOUBLE\_STRAND\_BREAK\_PROCESSING, GOBP\_DNA\_DOUBLE\_STRAND\_BREAK\_PROCESSING, HP\_NEOPLASM\_OF\_THE\_PANCREAS, HP\_NEOPLASM\_OF\_THE\_PANCREAS, HP\_ABNORMALITY\_OF\_THE\_PREPUTIUM, HP\_ABNORMALITY\_OF\_THE\_PREPUTIUM, HP\_ANAL\_ATRESIA, HP\_ANAL\_ATRESIA, GOBP\_RESPONSE\_TO\_X\_RAY, GOBP\_RESPONSE\_TO\_X\_RAY, GOBP\_REGULATION\_OF\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE, GOBP\_REGULATION\_OF\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE, GOBP\_POSITIVE\_REGULATION\_OF\_CELL\_CYCLE\_ARREST, GOBP\_POSITIVE\_REGULATION\_OF\_CELL\_CYCLE\_ARREST, GOBP\_G0\_TO\_G1\_TRANSITION, GOBP\_G0\_TO\_G1\_TRANSITION, HP\_ABNORMAL\_LABIA\_MINORA\_MORPHOLOGY, HP\_ABNORMAL\_LABIA\_MINORA\_MORPHOLOGY, GOBP\_REGULATION\_OF\_TYPE\_I\_INTERFERON\_MEDIATED\_SIGNALING\_PATHWAY, GOBP\_REGULATION\_OF\_TYPE\_I\_INTERFERON\_MEDIATED\_SIGNALING\_PATHWAY, GOBP\_REGULATION\_OF\_PROTEIN\_EXPORT\_FROM\_NUCLEUS, GOBP\_REGULATION\_OF\_PROTEIN\_EXPORT\_FROM\_NUCLEUS, HP\_ABNORMAL\_ESOPHAGUS\_MORPHOLOGY, HP\_ABNORMAL\_ESOPHAGUS\_MORPHOLOGY, GOBP\_NEGATIVE\_REGULATION\_OF\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_NEGATIVE\_REGULATION\_OF\_PROTEIN\_CATABOLIC\_PROCESS, GOBP\_BASE\_EXCISION\_REPAIR, GOBP\_BASE\_EXCISION\_REPAIR, GOBP\_POSITIVE\_REGULATION\_OF\_DNA\_TEMPLATED\_TRANSCRIPTION\_ELONGATION, GOBP\_POSITIVE\_REGULATION\_OF\_DNA\_TEMPLATED\_TRANSCRIPTION\_ELONGATION, GOBP\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_CHROMOSOME, GOBP\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_CHROMOSOME, HP\_LATERAL\_CLAVICLE\_HOOK, HP\_LATERAL\_CLAVICLE\_HOOK, GOBP\_REGULATION\_OF\_MITOCHONDRIAL\_OUTER\_MEMBRANE\_PERMEABILIZATION\_INVOLVED\_IN\_APOPTOTIC\_SIGNALING\_PATHWAY, GOBP\_REGULATION\_OF\_MITOCHONDRIAL\_OUTER\_MEMBRANE\_PERMEABILIZATION\_INVOLVED\_IN\_APOPTOTIC\_SIGNALING\_PATHWAY, HP\_ADRENAL\_OVERACTIVITY, HP\_ADRENAL\_OVERACTIVITY, GOBP\_REGULATION\_OF\_MEIOTIC\_CELL\_CYCLE, GOBP\_REGULATION\_OF\_MEIOTIC\_CELL\_CYCLE, HP\_ABNORMAL\_ASTROCYTE\_MORPHOLOGY, HP\_ABNORMAL\_ASTROCYTE\_MORPHOLOGY, GOBP\_REGULATION\_OF\_CYCLIN\_DEPENDENT\_PROTEIN\_KINASE\_ACTIVITY, GOBP\_REGULATION\_OF\_CYCLIN\_DEPENDENT\_PROTEIN\_KINASE\_ACTIVITY, GOBP\_MITOTIC\_G2\_M\_TRANSITION\_CHECKPOINT, GOBP\_MITOTIC\_G2\_M\_TRANSITION\_CHECKPOINT, GOBP\_REGULATION\_OF\_HISTONE\_H4\_ACETYLATION, GOBP\_REGULATION\_OF\_HISTONE\_H4\_ACETYLATION, GOCC\_POSITIVE\_REGULATION\_OF\_JNK\_CASCADE, GOCC\_POSITIVE\_REGULATION\_OF\_JNK\_CASCADE, GOMF\_POLYUBIQUITIN\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING, GOMF\_POLYUBIQUITIN\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING, HP\_ABNORMAL\_MORPHOLOGY\_OF\_THE\_CONJUNCTIVAL\_VASCULATURE, HP\_ABNORMAL\_MORPHOLOGY\_OF\_THE\_CONJUNCTIVAL\_VASCULATURE, HP\_HYPOPIGMENTED\_MACULE, HP\_HYPOPIGMENTED\_MACULE, GOBP\_LYMPHOID\_PROGENITOR\_CELL\_DIFFERENTIATION, GOBP\_LYMPHOID\_PROGENITOR\_CELL\_DIFFERENTIATION, GOBP\_DNA\_STRAND\_ELONGATION\_INVOLVED\_IN\_DNA\_REPLICATION, GOBP\_DNA\_STRAND\_ELONGATION\_INVOLVED\_IN\_DNA\_REPLICATION, HP\_BREAST\_CARCINOMA, HP\_BREAST\_CARCINOMA, HP\_ABNORMALITY\_OF\_THE\_THYMUS, HP\_ABNORMALITY\_OF\_THE\_THYMUS, HP\_HYPOPLASTIC\_LABIA\_MINORA, HP\_HYPOPLASTIC\_LABIA\_MINORA, GOBP\_REGULATION\_OF\_PPTIDYL\_LYSINE\_ACETYLATION, GOBP\_REGULATION\_OF\_PPTIDYL\_LYSINE\_ACETYLATION, GOBP\_CELLULAR\_RESPONSE\_TO\_UV, GOBP\_CELLULAR\_RESPONSE\_TO\_UV, HP\_NEOPLASM\_OF\_THE\_RESPIRATORY\_SYSTEM, HP\_NEOPLASM\_OF\_THE\_RESPIRATORY\_SYSTEM, GOBP\_POSITIVE\_REGULATION\_OF\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_MITOCHONDRIUM, GOBP\_POSITIVE\_REGULATION\_OF\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_MITOCHONDRIUM, GOBP\_REGULATION\_OF\_CENTROSOME\_CYCLE, GOBP\_REGULATION\_OF\_CENTROSOME\_CYCLE, GOBP\_POSITIVE\_REGULATION\_OF\_STRESS\_ACTIVATED\_PROTEIN\_KINASE\_SIGNALING\_CASCADE, GOBP\_POSITIVE\_REGULATION\_OF\_STRESS\_ACTIVATED\_PROTEIN\_KINASE\_SIGNALING\_CASCADE, HP\_HIRSUTISM, HP\_HIRSUTISM, GOBP\_MITOTIC\_DNA\_REPLICATION, GOBP\_MITOTIC\_DNA\_REPLICATION, GOBP\_REGULATION\_OF\_MAMMARY\_GLAND\_EPITHELIAL\_CELL\_PROLIFERATION, GOBP\_REGULATION\_OF\_MAMMARY\_GLAND\_EPITHELIAL\_CELL\_PROLIFERATION, HP\_MUCOSAL\_TELANGIECTASIAE, HP\_MUCOSAL\_TELANGIECTASIAE, HP\_PREMATURE\_GRAYING\_OF\_HAIR, HP\_PREMATURE\_GRAYING\_OF\_HAIR, HP\_ABNORMALITY\_OF\_THE\_LARYNX, HP\_ABNORMALITY\_OF\_THE\_LARYNX, GOBP\_TELOMERE\_CAPPING, GOBP\_TELOMERE\_CAPPING, GOMF\_KINASE\_INHIBITOR\_ACTIVITY, GOMF\_KINASE\_INHIBITOR\_ACTIVITY, HP\_NEOPLASM\_OF\_THE\_BREAST, HP\_NEOPLASM\_OF\_THE\_BREAST, HP\_RECURRENT\_URINARY\_TRACT\_INFECTIONS, HP\_RECURRENT\_URINARY\_TRACT\_INFECTIONS, GOBP\_NEGATIVE\_REGULATION\_OF\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE, GOBP\_NEGATIVE\_REGULATION\_OF\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE, GOBP\_POSITIVE\_REGULATION\_OF\_TELOMERE\_MAINTENANCE\_VIA\_TELOMERE\_LENGTHENING, GOBP\_POSITIVE\_REGULATION\_OF\_TELOMERE\_MAINTENANCE\_VIA\_TELOMERE\_LENGTHENING, HP\_PRIMITIVE\_NEUROECTODERMAL\_TUMOR, HP\_PRIMITIVE\_NEUROECTODERMAL\_TUMOR, HP\_DOLICHOCEPHALY, HP\_DOLICHOCEPHALY, GOBP\_CARDIAC\_MUSCLE\_TISSUE\_REGENERATION, GOBP\_CARDIAC\_MUSCLE\_TISSUE\_REGENERATION, GOBP\_REGULATION\_OF\_PROTEIN\_ACETYLATION, GOBP\_REGULATION\_OF\_PROTEIN\_ACETYLATION, GOBP\_UV\_DAMAGE\_EXCISION\_REPAIR, GOBP\_UV\_DAMAGE\_EXCISION\_REPAIR, HP\_ABNORMAL\_PHARYNX\_MORPHOLOGY, HP\_ABNORMAL\_PHARYNX\_MORPHOLOGY, GOCC\_CULA\_RING\_UBIQUITIN\_LIGASE\_COMPLEX, GOCC\_CULA\_RING\_UBIQUITIN\_LIGASE\_COMPLEX, GOMF\_DNA\_BINDING\_BENDING, GOMF\_DNA\_BINDING\_BENDING, HP\_EPENDYMOMA, HP\_EPENDYMOMA, GOBP\_POSITIVE\_REGULATION\_OF\_PPTIDYL\_LYSINE\_ACETYLATION, GOBP\_POSITIVE\_REGULATION\_OF\_PPTIDYL\_LYSINE\_ACETYLATION, HP\_ABNORMALITY\_OF\_THE\_ANTERIOR\_FONTANELLE, HP\_ABNORMALITY\_OF\_THE\_ANTERIOR\_FONTANELLE, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_ACETYLATION, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_ACETYLATION, GOMF\_TUMOR\_NECROSIS\_FACTOR\_RECEPTOR\_SUPERFAMILY\_BINDING, GOMF\_TUMOR\_NECROSIS\_FACTOR\_RECEPTOR\_SUPERFAMILY\_BINDING, GOBP\_POSITIVE\_REGULATION\_OF\_TRANSCRIPTION\_FROM\_RNA\_POLYMERASE\_II\_PROMOTER\_IN\_RESPONSE\_TO\_STRESS, GOBP\_POSITIVE\_REGULATION\_OF\_TRANSCRIPTION\_FROM\_RNA\_POLYMERASE\_II\_PROMOTER\_IN\_RESPONSE\_TO\_STRESS, GOMF\_CYCLIN\_DEPENDENT\_PROTEIN\_SERINE\_THREONINE\_KINASE\_REGULATOR\_ACTIVITY, GOMF\_CYCLIN\_DEPENDENT\_PROTEIN\_SERINE\_THREONINE\_KINASE\_REGULATOR\_ACTIVITY, HP\_ABNORMAL\_URETER\_MORPHOLOGY, HP\_ABNORMAL\_URETER\_MORPHOLOGY, GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_CYCLE\_ARREST, GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_CYCLE\_ARREST, HP\_ABNORMALITY\_OF\_THE\_CHOROID\_PLEXUS, HP\_ABNORMALITY\_OF\_THE\_CHOROID\_PLEXUS, GOBP\_G2\_DNA\_DAMAGE\_CHECKPOINT, GOBP\_G2\_DNA\_DAMAGE\_CHECKPOINT, GOBP\_INTRACELLULAR\_ESTROGEN\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_INTRACELLULAR\_ESTROGEN\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_CYCLE\_CHECKPOINT, GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_CYCLE\_CHECKPOINT, HP\_ABNORMALITY\_OF\_EARLOBE, HP\_ABNORMALITY\_OF\_EARLOBE, HP\_ABNORMAL\_CIRCULATING\_CREATININE\_CONCENTRATION, HP\_ABNORMAL\_CIRCULATING\_CREATININE\_CONCENTRATION, GOBP\_ACTIVATION\_OF\_MAPKK\_ACTIVITY, GOBP\_ACTIVATION\_OF\_MAPKK\_ACTIVITY, HP\_SHORT\_PALPEBRAL\_FISSURE, HP\_SHORT\_PALPEBRAL\_FISSURE