

TROMERIC\_REGION, GO\_CHROMOSOME\_CENTROMERIC\_REGION

GO\_SISTER\_CHROMATID\_SEGREGATION, GO\_SISTER\_CHROMATID\_SEGREGATION  
GO\_MITOTIC\_SISTER\_CHROMATID\_SEGREGATION, GO\_MITOTIC\_SISTER\_CHROMATID\_SEGREGATION  
GO\_HISTONE\_BINDING, GO\_HISTONE\_BINDING  
GO\_SPINDLE\_ORGANIZATION, GO\_SPINDLE\_ORGANIZATION  
GO\_SPINDLE\_POLE, GO\_SPINDLE\_POLE  
GO\_MICROTUBULE\_CYTOSKELETON\_ORGANIZATION\_INVOLVED\_IN\_MITOSIS, GO\_MICROTUBULE\_CYTOSKELETON\_ORGANIZATION\_INVOLVED\_IN\_MITOSIS  
GO\_REGULATION\_OF\_SISTER\_CHROMATID\_SEGREGATION, GO\_REGULATION\_OF\_SISTER\_CHROMATID\_SEGREGATION  
GO\_CONDENSED\_CHROMOSOME\_CENTROMERIC\_REGION, GO\_CONDENSED\_CHROMOSOME\_CENTROMERIC\_REGION  
GO\_REGULATION\_OF\_CHROMOSOME\_SEGREGATION, GO\_REGULATION\_OF\_CHROMOSOME\_SEGREGATION  
GO\_KINETOCHORE, GO\_KINETOCHORE  
GO\_MITOTIC\_CELL\_CYCLE\_CHECKPOINT, GO\_MITOTIC\_CELL\_CYCLE\_CHECKPOINT  
GO\_MITOTIC\_SPINDLE\_ORGANIZATION, GO\_MITOTIC\_SPINDLE\_ORGANIZATION  
GO\_PROTEIN\_METHYLATION, GO\_PROTEIN\_METHYLATION  
GO\_ANAPHASE\_PROMOTING\_COMPLEX\_DEPENDENT\_CATABOLIC\_PROCESS, GO\_ANAPHASE\_PROMOTING\_COMPLEX\_DEPENDENT\_CATABOLIC\_PROCESS  
GO\_CHROMOSOME\_SEPARATION, GO\_CHROMOSOME\_SEPARATION  
GO\_REGULATION\_OF\_CHROMOSOME\_SEPARATION, GO\_REGULATION\_OF\_CHROMOSOME\_SEPARATION  
GO\_CHROMATIN\_REMODELING, GO\_CHROMATIN\_REMODELING  
GO\_SPINDLE\_ASSEMBLY, GO\_SPINDLE\_ASSEMBLY  
GO\_CILIARY\_BASAL\_BODY\_PLASMA\_MEMBRANE\_DOCKING, GO\_CILIARY\_BASAL\_BODY\_PLASMA\_MEMBRANE\_DOCKING  
GO\_PEPTIDYL\_LYSINE\_METHYLATION, GO\_PEPTIDYL\_LYSINE\_METHYLATION  
GO\_METAPHASE\_ANAPHASE\_TRANSITION\_OF\_CELL\_CYCLE, GO\_METAPHASE\_ANAPHASE\_TRANSITION\_OF\_CELL\_CYCLE  
GO\_CHROMATIN\_ASSEMBLY\_OR\_DISASSEMBLY, GO\_CHROMATIN\_ASSEMBLY\_OR\_DISASSEMBLY  
GO\_NEGATIVE\_REGULATION\_OF\_CHROMOSOME\_ORGANIZATION, GO\_NEGATIVE\_REGULATION\_OF\_CHROMOSOME\_ORGANIZATION  
GO\_NUCLEOSOME\_ORGANIZATION, GO\_NUCLEOSOME\_ORGANIZATION  
GO\_CHROMOSOME\_TELOMERIC\_REGION, GO\_CHROMOSOME\_TELOMERIC\_REGION  
GO\_NEGATIVE\_REGULATION\_OF\_CHROMOSOME\_SEGREGATION, GO\_NEGATIVE\_REGULATION\_OF\_CHROMOSOME\_SEGREGATION  
GO\_CHROMATIN\_ASSEMBLY, GO\_CHROMATIN\_ASSEMBLY  
GO\_NUCLEAR\_CHROMOSOME\_TELOMERIC\_REGION, GO\_NUCLEAR\_CHROMOSOME\_TELOMERIC\_REGION  
GO\_REGULATION\_OF\_GENE\_SILENCING, GO\_REGULATION\_OF\_GENE\_SILENCING  
GO\_HISTONE\_METHYLATION, GO\_HISTONE\_METHYLATION  
GO\_REGULATION\_OF\_CELLULAR\_RESPONSE\_TO\_HEAT, GO\_REGULATION\_OF\_CELLULAR\_RESPONSE\_TO\_HEAT  
GO\_NEGATIVE\_REGULATION\_OF\_METAPHASE\_ANAPHASE\_TRANSITION\_OF\_CELL\_CYCLE, GO\_NEGATIVE\_REGULATION\_OF\_METAPHASE\_ANAPHASE\_TRANSITION\_OF\_CELL\_CYCLE  
GO\_MICROTUBULE\_ORGANIZING\_CENTER\_ORGANIZATION, GO\_MICROTUBULE\_ORGANIZING\_CENTER\_ORGANIZATION  
GO\_NUCLEAR\_PORE, GO\_NUCLEAR\_PORE  
GO\_CENTROMERE\_COMPLEX\_ASSEMBLY, GO\_CENTROMERE\_COMPLEX\_ASSEMBLY  
GO\_PROTEIN\_LOCALIZATION\_TO\_CHROMOSOME, GO\_PROTEIN\_LOCALIZATION\_TO\_CHROMOSOME  
GO\_REGULATION\_OF\_POSTTRANSCRIPTIONAL\_GENE\_SILENCING, GO\_REGULATION\_OF\_POSTTRANSCRIPTIONAL\_GENE\_SILENCING  
GO\_MIDBODY, GO\_MIDBODY  
GO\_DNA\_REPLICATION\_INDEPENDENT\_NUCLEOSOME\_ORGANIZATION, GO\_DNA\_REPLICATION\_INDEPENDENT\_NUCLEOSOME\_ORGANIZATION  
GO\_MITOTIC\_SPINDLE\_ASSEMBLY, GO\_MITOTIC\_SPINDLE\_ASSEMBLY  
GO\_MEMBRANE\_DOCKING, GO\_MEMBRANE\_DOCKING  
GO\_HISTONE\_LYSINE\_N\_METHYLTRANSFERASE\_ACTIVITY, GO\_HISTONE\_LYSINE\_N\_METHYLTRANSFERASE\_ACTIVITY  
GO\_MITOTIC\_SPINDLE, GO\_MITOTIC\_SPINDLE  
GO\_ATP\_DEPENDENT\_CHROMATIN\_REMODELING, GO\_ATP\_DEPENDENT\_CHROMATIN\_REMODELING  
GO\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING, GO\_MODIFICATION\_DEPENDENT\_PROTEIN\_BINDING  
GO\_NEGATIVE\_REGULATION\_OF\_CELL\_CYCLE\_G2\_M\_PHASE\_TRANSITION, GO\_NEGATIVE\_REGULATION\_OF\_CELL\_CYCLE\_G2\_M\_PHASE\_TRANSITION  
GO\_SITE\_OF\_DNA\_DAMAGE, GO\_SITE\_OF\_DNA\_DAMAGE  
GO\_REGULATION\_OF\_NUCLEAR\_DIVISION, GO\_REGULATION\_OF\_NUCLEAR\_DIVISION  
GO\_CHROMATIN\_REMODELING\_AT\_CENTROMERE, GO\_CHROMATIN\_REMODELING\_AT\_CENTROMERE  
GO\_CONDENSED\_NUCLEAR\_CHROMOSOME, GO\_CONDENSED\_NUCLEAR\_CHROMOSOME  
GO\_REGULATION\_OF\_DNA\_REPLICATION, GO\_REGULATION\_OF\_DNA\_REPLICATION  
GO\_NCRNA\_EXPORT\_FROM\_NUCLEUS, GO\_NCRNA\_EXPORT\_FROM\_NUCLEUS  
GO\_RETROGRADE\_VESICLE\_MEDIATED\_TRANSPORT\_GOLGI\_TO\_ENDOPLASMIC\_RETICULUM, GO\_RETROGRADE\_VESICLE\_MEDIATED\_TRANSPORT\_GOLGI\_TO\_ENDOPLASMIC\_RETICULUM  
GO\_HISTONE\_DEMETHYLASE\_ACTIVITY, GO\_HISTONE\_DEMETHYLASE\_ACTIVITY  
GO\_POSITIVE\_REGULATION\_OF\_CHROMOSOME\_ORGANIZATION, GO\_POSITIVE\_REGULATION\_OF\_CHROMOSOME\_ORGANIZATION  
GO\_REGULATION\_OF\_ORGANELLE\_ASSEMBLY, GO\_REGULATION\_OF\_ORGANELLE\_ASSEMBLY  
GO\_ATTACHMENT\_OF\_SPINDLE\_MICROTUBULES\_TO\_KINETOCHORE, GO\_ATTACHMENT\_OF\_SPINDLE\_MICROTUBULES\_TO\_KINETOCHORE  
GO\_MICROTUBULE\_ASSOCIATED\_COMPLEX, GO\_MICROTUBULE\_ASSOCIATED\_COMPLEX  
GO\_SISTER\_CHROMATID\_COHESION, GO\_SISTER\_CHROMATID\_COHESION  
GO\_NUCLEUS\_ORGANIZATION, GO\_NUCLEUS\_ORGANIZATION  
GO\_HISTONE\_EXCHANGE, GO\_HISTONE\_EXCHANGE  
GO\_MITOTIC\_SPINDLE\_POLE, GO\_MITOTIC\_SPINDLE\_POLE  
GO\_METAPHASE\_PLATE\_CONGRESSION, GO\_METAPHASE\_PLATE\_CONGRESSION  
GO\_HETEROCHROMATIN, GO\_HETEROCHROMATIN  
GO\_ANTIGEN\_PROCESSING\_AND\_PRESENTATION\_OF\_PEPTIDE\_ANTIGEN, GO\_ANTIGEN\_PROCESSING\_AND\_PRESENTATION\_OF\_PEPTIDE\_ANTIGEN  
GO\_ESTABLISHMENT\_OF\_MITOTIC\_SPINDLE\_LOCALIZATION, GO\_ESTABLISHMENT\_OF\_MITOTIC\_SPINDLE\_LOCALIZATION  
GO\_POSITIVE\_REGULATION\_OF\_CHROMOSOME\_SEGREGATION, GO\_POSITIVE\_REGULATION\_OF\_CHROMOSOME\_SEGREGATION  
GO\_REGULATION\_OF\_SISTER\_CHROMATID\_COHESION, GO\_REGULATION\_OF\_SISTER\_CHROMATID\_COHESION  
GO\_CONDENSED\_NUCLEAR\_CHROMOSOME\_CENTROMERIC\_REGION, GO\_CONDENSED\_NUCLEAR\_CHROMOSOME\_CENTROMERIC\_REGION  
GO\_DNA\_DEPENDENT\_DNA\_REPLICATION\_MAINTENANCE\_OF\_FIDELITY, GO\_DNA\_DEPENDENT\_DNA\_REPLICATION\_MAINTENANCE\_OF\_FIDELITY  
GO\_NEGATIVE\_REGULATION\_OF\_NUCLEAR\_DIVISION, GO\_NEGATIVE\_REGULATION\_OF\_NUCLEAR\_DIVISION  
GO\_CHROMATIN\_ORGANIZATION\_INVOLVED\_IN\_REGULATION\_OF\_TRANSCRIPTION, GO\_CHROMATIN\_ORGANIZATION\_INVOLVED\_IN\_REGULATION\_OF\_TRANSCRIPTION  
GO\_PROTEIN\_LOCALIZATION\_TO\_CHROMOSOME\_CENTROMERIC\_REGION, GO\_PROTEIN\_LOCALIZATION\_TO\_CHROMOSOME\_CENTROMERIC\_REGION  
GO\_REGULATION\_OF\_CELL\_CYCLE\_CHECKPOINT, GO\_REGULATION\_OF\_CELL\_CYCLE\_CHECKPOINT  
GO\_CHROMOSOME\_LOCALIZATION, GO\_CHROMOSOME\_LOCALIZATION  
GO\_METHYLATED\_HISTONE\_BINDING, GO\_METHYLATED\_HISTONE\_BINDING  
GO\_SPINDLE\_MICROTUBULE, GO\_SPINDLE\_MICROTUBULE  
GO\_NEGATIVE\_REGULATION\_OF\_GENE\_EXPRESSION\_EPIGENETIC, GO\_NEGATIVE\_REGULATION\_OF\_GENE\_EXPRESSION\_EPIGENETIC  
GO\_POSITIVE\_REGULATION\_OF\_CELL\_CYCLE\_G2\_M\_PHASE\_TRANSITION, GO\_POSITIVE\_REGULATION\_OF\_CELL\_CYCLE\_G2\_M\_PHASE\_TRANSITION  
GO\_PROTEIN\_N\_TERMINUS\_BINDING, GO\_PROTEIN\_N\_TERMINUS\_BINDING  
GO\_PROTEIN\_ACETYLATION, GO\_PROTEIN\_ACETYLATION  
GO\_CELLULAR\_RESPONSE\_TO\_HEAT, GO\_CELLULAR\_RESPONSE\_TO\_HEAT  
GO\_PROTEIN\_LOCALIZATION\_TO\_KINETOCHORE, GO\_PROTEIN\_LOCALIZATION\_TO\_KINETOCHORE  
GO\_POSITIVE\_REGULATION\_OF\_CHROMATIN\_SILENCING, GO\_POSITIVE\_REGULATION\_OF\_CHROMATIN\_SILENCING  
GO\_MITOTIC\_SISTER\_CHROMATID\_COHESION, GO\_MITOTIC\_SISTER\_CHROMATID\_COHESION  
GO\_CHROMATIN\_DNA\_BINDING, GO\_CHROMATIN\_DNA\_BINDING  
GO\_MITOTIC\_CHROMOSOME\_CONDENSATION, GO\_MITOTIC\_CHROMOSOME\_CONDENSATION  
GO\_NUCLEAR\_ENVELOPE\_ORGANIZATION, GO\_NUCLEAR\_ENVELOPE\_ORGANIZATION  
GO\_REPLICATION\_FORK\_PROCESSING, GO\_REPLICATION\_FORK\_PROCESSING  
GO\_SPINDLE\_MIDZONE, GO\_SPINDLE\_MIDZONE  
GO\_MICROTUBULE\_MOTOR\_ACTIVITY, GO\_MICROTUBULE\_MOTOR\_ACTIVITY  
GO\_KINETOCHORE\_ORGANIZATION, GO\_KINETOCHORE\_ORGANIZATION  
GO\_NEURON\_PROJECTION\_CYTOPLASM, GO\_NEURON\_PROJECTION\_CYTOPLASM  
GO\_PROTEIN\_LOCALIZATION\_TO\_CHROMATIN, GO\_PROTEIN\_LOCALIZATION\_TO\_CHROMATIN  
GO\_CONDENSED\_NUCLEAR\_CHROMOSOME\_KINETOCHORE, GO\_CONDENSED\_NUCLEAR\_CHROMOSOME\_KINETOCHORE  
GO\_MITOTIC\_METAPHASE\_PLATE\_CONGRESSION, GO\_MITOTIC\_METAPHASE\_PLATE\_CONGRESSION  
GO\_SPINDLE\_POLE\_CENTROSOME, GO\_SPINDLE\_POLE\_CENTROSOME  
GO\_POSITIVE\_REGULATION\_OF\_PROTEOLYSIS\_INVOLVED\_IN\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GO\_POSITIVE\_REGULATION\_OF\_PROTEOLYSIS\_INVOLVED\_IN\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS  
GO\_PROTEIN\_LOCALIZATION\_TO\_CHROMOSOME\_TELOMERIC\_REGION, GO\_PROTEIN\_LOCALIZATION\_TO\_CHROMOSOME\_TELOMERIC\_REGION  
GO\_RAN\_GTPASE\_BINDING, GO\_RAN\_GTPASE\_BINDING  
GO\_CONDENSIN\_COMPLEX, GO\_CONDENSIN\_COMPLEX  
GO\_PEPTIDYL\_LYSINE\_TRIMETHYLATION, GO\_PEPTIDYL\_LYSINE\_TRIMETHYLATION  
GO\_REGULATION\_OF\_CENTROSOME\_CYCLE, GO\_REGULATION\_OF\_CENTROSOME\_CYCLE  
GO\_HISTONE\_H3\_K9\_METHYLATION, GO\_HISTONE\_H3\_K9\_METHYLATION  
GO\_STRUCTURAL\_CONSTITUENT\_OF\_NUCLEAR\_PORE, GO\_STRUCTURAL\_CONSTITUENT\_OF\_NUCLEAR\_PORE  
GO\_MICROTUBULE\_ORGANIZING\_CENTER\_LOCALIZATION, GO\_MICROTUBULE\_ORGANIZING\_CENTER\_LOCALIZATION  
GO\_PHOSPHATASE\_REGULATOR\_ACTIVITY, GO\_PHOSPHATASE\_REGULATOR\_ACTIVITY  
GO\_ESTABLISHMENT\_OF\_SPINDLE\_ORIENTATION, GO\_ESTABLISHMENT\_OF\_SPINDLE\_ORIENTATION  
GO\_HOMOLOGOUS\_RECOMBINATION, GO\_HOMOLOGOUS\_RECOMBINATION  
GO\_PERICENTRIC\_HETEROCHROMATIN, GO\_PERICENTRIC\_HETEROCHROMATIN  
GO\_CYTOSKELETON\_DEPENDENT\_CYTOKINESIS, GO\_CYTOSKELETON\_DEPENDENT\_CYTOKINESIS  
GO\_MICROTUBULE\_DEPOLYMERIZATION, GO\_MICROTUBULE\_DEPOLYMERIZATION  
GO\_KINESIN\_COMPLEX, GO\_KINESIN\_COMPLEX  
GO\_MEIOTIC\_CHROMOSOME\_CONDENSATION, GO\_MEIOTIC\_CHROMOSOME\_CONDENSATION  
GO\_REGULATION\_OF\_MITOTIC\_SPINDLE\_ASSEMBLY, GO\_REGULATION\_OF\_MITOTIC\_SPINDLE\_ASSEMBLY  
GO\_REGULATION\_OF\_SPINDLE\_CHECKPOINT, GO\_REGULATION\_OF\_SPINDLE\_CHECKPOINT  
GO\_POSITIVE\_REGULATION\_OF\_DNA\_DEPENDENT\_DNA\_REPLICATION, GO\_POSITIVE\_REGULATION\_OF\_DNA\_DEPENDENT\_DNA\_REPLICATION  
GO\_NUCLEAR\_HETEROCHROMATIN, GO\_NUCLEAR\_HETEROCHROMATIN  
GO\_CONDENSED\_CHROMOSOME\_OUTER\_KINETOCHORE, GO\_CONDENSED\_CHROMOSOME\_OUTER\_KINETOCHORE  
GO\_ATTACHMENT\_OF\_MITOTIC\_SPINDLE\_MICROTUBULES\_TO\_KINETOCHORE, GO\_ATTACHMENT\_OF\_MITOTIC\_SPINDLE\_MICROTUBULES\_TO\_KINETOCHORE  
GO\_REGULATION\_OF\_SPINDLE\_ASSEMBLY, GO\_REGULATION\_OF\_SPINDLE\_ASSEMBLY  
GO\_HISTONE\_H4\_K20\_METHYLATION, GO\_HISTONE\_H4\_K20\_METHYLATION  
GO\_MITOTIC\_CYTOKINESIS, GO\_MITOTIC\_CYTOKINESIS  
GO\_HISTONE\_H4\_K5\_ACETYLATION, GO\_HISTONE\_H4\_K5\_ACETYLATION  
GO\_REGULATION\_OF\_ATTACHMENT\_OF\_SPINDLE\_MICROTUBULES\_TO\_KINETOCHORE, GO\_REGULATION\_OF\_ATTACHMENT\_OF\_SPINDLE\_MICROTUBULES\_TO\_KINETOCHORE  
GO\_PML\_BODY, GO\_PML\_BODY  
GO\_PROTEIN\_SUMOYLATION, GO\_PROTEIN\_SUMOYLATION  
GO\_NUCLEAR\_PORE\_ORGANIZATION, GO\_NUCLEAR\_PORE\_ORGANIZATION  
GO\_REGULATION\_OF\_GENERATION\_OF\_PRECURSOR\_METABOLITES\_AND\_ENERGY, GO\_REGULATION\_OF\_GENERATION\_OF\_PRECURSOR\_METABOLITES\_AND\_ENERGY  
GO\_SPINDLE\_LOCALIZATION, GO\_SPINDLE\_LOCALIZATION  
GO\_CYTOKINESIS, GO\_CYTOKINESIS  
GO\_POSITIVE\_REGULATION\_OF\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS, GO\_POSITIVE\_REGULATION\_OF\_CELLULAR\_PROTEIN\_CATABOLIC\_PROCESS  
GO\_MICROTUBULE\_NUCLEATION, GO\_MICROTUBULE\_NUCLEATION  
GO\_PROTEIN\_LOCALIZATION\_TO\_CYTOSKELETON, GO\_PROTEIN\_LOCALIZATION\_TO\_CYTOSKELETON  
GO\_NEGATIVE\_REGULATION\_OF\_HISTONE\_MODIFICATION, GO\_NEGATIVE\_REGULATION\_OF\_HISTONE\_MODIFICATION  
GO\_SEX\_CHROMOSOME, GO\_SEX\_CHROMOSOME  
GO\_REGULATION\_OF\_ATP\_METABOLIC\_PROCESS, GO\_REGULATION\_OF\_ATP\_METABOLIC\_PROCESS  
GO\_CHROMOSOME\_ORGANIZATION\_INVOLVED\_IN\_MEIOTIC\_CELL\_CYCLE, GO\_CHROMOSOME\_ORGANIZATION\_INVOLVED\_IN\_MEIOTIC\_CELL\_CYCLE  
GO\_REGULATION\_OF\_SPINDLE\_ORGANIZATION, GO\_REGULATION\_OF\_SPINDLE\_ORGANIZATION  
GO\_SPERM\_CHROMATIN\_CONDENSATION, GO\_SPERM\_CHROMATIN\_CONDENSATION  
GO\_MAINTENANCE\_OF\_DNA\_METHYLATION, GO\_MAINTENANCE\_OF\_DNA\_METHYLATION  
GO\_HISTONE\_H3\_K9\_MODIFICATION, GO\_HISTONE\_H3\_K9\_MODIFICATION  
GO\_MOTOR\_ACTIVITY, GO\_MOTOR\_ACTIVITY  
GO\_NEGATIVE\_REGULATION\_OF\_CENTROSOME\_CYCLE, GO\_NEGATIVE\_REGULATION\_OF\_CENTROSOME\_CYCLE  
GO\_NUCLEOTIDE\_PHOSPHORYLATION, GO\_NUCLEOTIDE\_PHOSPHORYLATION  
GO\_HISTONE\_METHYLTRANSFERASE\_ACTIVITY\_H3\_K9\_SPECIFIC, GO\_HISTONE\_METHYLTRANSFERASE\_ACTIVITY\_H3\_K9\_SPECIFIC  
GO\_MITOTIC\_SPINDLE\_MIDZONE, GO\_MITOTIC\_SPINDLE\_MIDZONE  
GO\_REGULATION\_OF\_PROTEIN\_DEPHOSPHORYLATION, GO\_REGULATION\_OF\_PROTEIN\_DEPHOSPHORYLATION  
GO\_BETA\_CATENIN\_DESTRUCTION\_COMPLEX\_DISASSEMBLY, GO\_BETA\_CATENIN\_DESTRUCTION\_COMPLEX\_DISASSEMBLY  
GO\_EXIT\_FROM\_MITOSIS, GO\_EXIT\_FROM\_MITOSIS  
GO\_REGULATION\_OF\_PROTEIN\_EXPORT\_FROM\_NUCLEUS, GO\_REGULATION\_OF\_PROTEIN\_EXPORT\_FROM\_NUCLEUS  
GO\_MICROTUBULE\_POLYMERIZATION\_OR\_DEPOLYMERIZATION, GO\_MICROTUBULE\_POLYMERIZATION\_OR\_DEPOLYMERIZATION  
GO\_POSITIVE\_REGULATION\_OF\_SISTER\_CHROMATID\_COHESION, GO\_POSITIVE\_REGULATION\_OF\_SISTER\_CHROMATID\_COHESION