

source	term name	term ID	n. of term genes	corrected p-value	visualization icons
BP	response to organic substance	GO:0010033	206	1.08e-02	visualization icons
BP	response to pheromone	GO:0019236	107	7.44e-05	visualization icons
BP	response to stimulus	GO:0050896	1148	1.15e-04	visualization icons
BP	response to stress	GO:0006950	730	1.14e-04	visualization icons
BP	single-organism process	GO:0044699	3755	4.43e-13	visualization icons
BP	cellular process	GO:0009987	4884	4.35e-07	visualization icons
BP	cell wall organization or biogenesis	GO:0071554	251	2.89e-05	visualization icons
BP	single-organism cellular process	GO:0044763	3397	7.87e-17	visualization icons
BP	cell division	GO:0051301	355	6.69e-17	visualization icons
BP	cell cycle	GO:0007049	741	2.29e-28	visualization icons
BP	cell cycle process	GO:0022402	629	1.61e-23	visualization icons
BP	cell cycle phase transition	GO:0044770	140	8.95e-03	visualization icons
BP	cell cycle G1/S phase transition	GO:0044843	66	3.17e-11	visualization icons
BP	spindle organization	GO:0007051	40	3.44e-03	visualization icons
BP	cytokinesis	GO:0000910	83	4.47e-02	visualization icons
BP	cytoskeleton-dependent cytokinesis	GO:0061640	72	1.40e-02	visualization icons
BP	cytokinetic process	GO:0032506	67	7.67e-03	visualization icons
BP	spindle pole body separation	GO:0000073	13	1.95e-04	visualization icons
BP	mitotic cell cycle	GO:0000278	379	8.03e-22	visualization icons
BP	mitotic cell cycle process	GO:1903047	367	1.16e-20	visualization icons
BP	mitotic cytokinesis	GO:0000281	65	4.80e-02	visualization icons
BP	mitotic cell cycle phase transition	GO:0044772	137	5.56e-10	visualization icons
BP	G1/S transition of mitotic cell cycle	GO:0000082	66	3.17e-11	visualization icons
BP	regulation of transcription involved in G1/S transition of ...	GO:0000083	27	2.52e-11	visualization icons
BP	negative regulation of transcription involved in G1/S tr ...	GO:0071930	7	1.45e-02	visualization icons
BP	mitotic cytokinetic process	GO:1902410	55	1.38e-02	visualization icons
BP	mitotic spindle organization	GO:0007052	94	1.17e-02	visualization icons
BP	cellular response to stimulus	GO:0051716	383	4.98e-05	visualization icons
BP	cellular response to stress	GO:0033554	631	2.35e-05	visualization icons
BP	cellular response to DNA damage stimulus	GO:0006974	331	2.48e-08	visualization icons
BP	cell separation after cytokinesis	GO:0000920	20	2.24e-04	visualization icons
BP	multi-organism process	GO:0051704	305	6.83e-04	visualization icons
BP	multi-organism cellular process	GO:0044764	186	3.03e-06	visualization icons
BP	conjugation	GO:0000746	127	2.44e-08	visualization icons
BP	regulation of gene expression, epigenetic	GO:0040029	151	2.23e-04	visualization icons
BP	sex determination	GO:0007530	35	2.38e-03	visualization icons
BP	reproduction	GO:0000003	486	4.56e-14	visualization icons
BP	reproduction of a single-celled organism	GO:0032505	203	1.20e-03	visualization icons
BP	meiotic cell cycle	GO:0051321	303	3.46e-03	visualization icons
BP	meiotic cell cycle process	GO:1903046	236	2.01e-02	visualization icons
BP	biological regulation	GO:0065007	1791	7.41e-06	visualization icons
BP	regulation of molecular function	GO:0065009	247	1.00e-02	visualization icons
BP	regulation of biological process	GO:0050789	158	8.11e-05	visualization icons
BP	regulation of cellular process	GO:0050794	1506	1.24e-05	visualization icons
BP	regulation of cell cycle	GO:0051726	214	1.55e-07	visualization icons
BP	regulation of cell cycle process	GO:0010564	154	1.05e-02	visualization icons
BP	regulation of spindle pole body separation	GO:0010695	9	7.01e-04	visualization icons
BP	positive regulation of spindle pole body separation	GO:0010696	7	9.12e-03	visualization icons
BP	regulation of mitotic cell cycle	GO:0007346	121	5.18e-03	visualization icons
BP	regulation of metabolic process	GO:0019222	1194	9.23e-04	visualization icons
BP	regulation of catalytic activity	GO:0050790	228	1.28e-02	visualization icons
BP	regulation of transferase activity	GO:0051338	104	4.28e-04	visualization icons
BP	regulation of nitrogen compound metabolic process	GO:0051171	796	3.12e-03	visualization icons
BP	cellular component organization or biogenesis	GO:0071840	2397	2.18e-04	visualization icons
BP	cellular component organization	GO:0016043	2045	2.33e-07	visualization icons
BP	external encapsulating structure organization	GO:0045229	212	4.73e-05	visualization icons
BP	cell wall organization	GO:0071555	212	4.73e-05	visualization icons
BP	organelle fission	GO:0048285	1491	1.87e-04	visualization icons
BP	nuclear division	GO:0000280	341	1.12e-08	visualization icons
BP	meiotic nuclear division	GO:0007126	170	4.85e-04	visualization icons
BP	single-organism organelle organization	GO:1902589	1007	1.13e-02	visualization icons
BP	mitotic nuclear division	GO:0007067	215	2.76e-08	visualization icons
BP	meiosis I	GO:0007127	98	4.56e-04	visualization icons
BP	chromosome organization	GO:0051276	523	3.36e-06	visualization icons
BP	DNA conformation change	GO:0071103	90	8.11e-03	visualization icons
BP	DNA geometric change	GO:0032392	32	5.55e-04	visualization icons
BP	DNA duplex unwinding	GO:0032508	31	4.24e-04	visualization icons
BP	sister chromatid cohesion	GO:0007062	49	3.33e-02	visualization icons
BP	mitotic sister chromatid cohesion	GO:0007064	38	4.62e-03	visualization icons
BP	regulation of cellular component organization	GO:0051128	393	5.87e-03	visualization icons
BP	regulation of organelle organization	GO:0033043	290	6.59e-03	visualization icons
BP	reproductive process	GO:0022414	273	2.35e-08	visualization icons
BP	reproductive process in single-celled organism	GO:0022413	151	2.66e-02	visualization icons
BP	multi-organism reproductive process	GO:0044703	238	4.14e-05	visualization icons
BP	sexual reproduction	GO:0019953	238	4.14e-05	visualization icons
BP	conjugation with cellular fusion	GO:0000747	126	6.11e-03	visualization icons
BP	cytogamy	GO:0000755	13	2.10e-08	visualization icons
BP	cellular response to organic substance	GO:0071310	169	4.10e-03	visualization icons
BP	cellular response to pheromone	GO:0071444	97	1.69e-04	visualization icons
BP	response to pheromone involved in conjugation with cellular fusion	GO:0000749	83	2.49e-05	visualization icons
BP	regulation of macromolecule metabolic process	GO:0060255	109	1.42e-04	visualization icons
BP	regulation of primary metabolic process	GO:0080090	1066	1.75e-04	visualization icons
BP	negative regulation of biological process	GO:0048519	554	3.66e-08	visualization icons
BP	negative regulation of cellular process	GO:0048523	527	7.71e-09	visualization icons
BP	negative regulation of cell cycle	GO:0045786	97	1.68e-02	visualization icons
BP	negative regulation of macromolecule metabolic process	GO:0009892	444	7.83e-06	visualization icons
BP	negative regulation of gene expression	GO:0010605	399	1.08e-05	visualization icons
BP	negative regulation of gene expression, epigenetic	GO:0045814	150	2.04e-04	visualization icons
BP	gene silencing	GO:0016458	154	2.94e-04	visualization icons
BP	negative regulation of nitrogen compound metabolic process	GO:0051172	309	7.20e-05	visualization icons
BP	regulation of protein modification process	GO:0031399	137	2.12e-03	visualization icons
BP	cell fate commitment	GO:0045165	35	2.38e-03	visualization icons
BP	regulation of biological process with cellular fusion	GO:0032005	33	6.41e-03	visualization icons
BP	negative regulation of biosynthetic process	GO:0009890	323	2.61e-03	visualization icons
BP	regulation of cellular metabolic process	GO:0031323	1086	1.15e-04	visualization icons
BP	regulation of phosphate metabolic process	GO:0051174	141	1.58e-02	visualization icons
BP	regulation of nucleobase-containing compound metabolic process	GO:0019220	143	2.08e-03	visualization icons
BP	regulation of RNA metabolic process	GO:0051252	722	4.31e-02	visualization icons
BP	negative regulation of cellular metabolic process	GO:0031324	430	3.60e-06	visualization icons
BP	negative regulation of nucleobase-containing compound metabolic process	GO:0045934	304	5.26e-05	visualization icons
BP	negative regulation of RNA metabolic process	GO:0051253	270	5.24e-04	visualization icons
BP	sulfate reduction	GO:0019419	3	1.25e-02	visualization icons
BP	sulfate assimilation, phosphoadenylyl sulfate reduction by phosphoaden ...	GO:0019379	3	2.52e-02	visualization icons
BP	mating type determination	GO:0007531	35	2.38e-03	visualization icons
BP	regulation of macromolecule biosynthetic process	GO:0010556	860	4.86e-02	visualization icons
BP	regulation of cellular macromolecule biosynthetic process	GO:2000112	854	4.16e-02	visualization icons
BP	negative regulation of macromolecule biosynthetic process	GO:0010558	302	8.37e-04	visualization icons
BP	negative regulation of cellular biosynthetic process	GO:0031327	323	2.61e-03	visualization icons
BP	negative regulation of cellular macromolecule biosynthetic process	GO:2000113	301	7.91e-04	visualization icons
BP	nucleobase-containing compound biosynthetic process	GO:0034654	966	8.95e-03	visualization icons
BP	RNA biosynthetic process	GO:0032774	769	2.54e-02	visualization icons
BP	regulation of RNA biosynthetic process	GO:2001141	705	2.63e-02	visualization icons
BP	regulation of nucleic acid-templated transcription	GO:1903506	705	2.63e-02	visualization icons
BP	regulation of transcription, DNA-templated	GO:0006355	704	2.56e-02	visualization icons
BP	negative regulation of RNA biosynthetic process	GO:1902679	268	4.62e-04	visualization icons
BP	negative regulation of nucleic acid-templated transcription	GO:1903507	268	4.62e-04	visualization icons
BP	negative regulation of transcription, DNA-templated	GO:0045892	267	4.34e-04	visualization icons
BP	chromatin silencing	GO:0006342	150	2.04e-04	visualization icons
BP	chromatin silencing at silent mating-type cassette	GO:0030466	53	3.05e-02	visualization icons
BP	chromatin silencing at telomere	GO:0006348	84	1.76e-02	visualization icons
BP	phosphorylation	GO:0016310	338	1.51e-03	visualization icons
BP	regulation of phosphorylation	GO:0042325	89	5.79e-05	visualization icons
BP	regulation of kinase activity	GO:0043549	69	1.86e-05	visualization icons
BP	G-protein coupled receptor signaling pathway	GO:0007186	44	4.71e-02	visualization icons
BP	pheromone-dependent signal transduction involved in conjugation with c ...	GO:0000750	33	6.41e-03	visualization icons
BP	DNA metabolic process	GO:0006259	542	1.86e-08	visualization icons
BP	DNA strand elongation	GO:0022616	37	8.85e-06	visualization icons
BP	heteroduplex formation	GO:0030491	9	3.24e-02	visualization icons
BP	regulation of DNA metabolic process	GO:0051052	107	1.74e-06	visualization icons
BP	positive regulation of DNA metabolic process	GO:0051054	27	4.18e-03	visualization icons
BP	DNA recombination	GO:0006310	241	3.57e-04	visualization icons
BP	mitotic recombination	GO:0006312	58	4.45e-03	visualization icons
BP	DNA replication	GO:0006260	211	1.21e-06	visualization icons
BP	cell cycle DNA replication	GO:0044786	36	7.02e-05	visualization icons
BP	nuclear DNA replication	GO:0032620	35	5.37e-05	visualization icons
BP	DNA-dependent DNA replication	GO:0006261	122	1.72e-08	visualization icons
BP	pre-replicative complex assembly	GO:0036388	20	3.87e-03	visualization icons
BP	pre-replicative complex assembly involved in cell cycle DNA rep ...	GO:1902299	20	3.87e-03	visualization icons
BP	pre-replicative complex assembly involved in nuclear cell cyc ...	GO:0006267	20	3.87e-03	visualization icons
BP	DNA strand elongation involved in DNA replication	GO:0006271	34	3.54e-06	visualization icons
BP	lagging strand elongation	GO:0006273	18	1.92e-03	visualization icons
BP	DNA unwinding involved in DNA replication	GO:0006268	13	2.42e-04	visualization icons
BP	regulation of DNA replication	GO:0006275	56	1.22e-02	visualization icons
BP	DNA replication initiation	GO:0006270	53	2.54e-06	visualization icons
BP	maintenance of DNA repeat elements	GO:0043570	23	5.03e-04	visualization icons
BP	maintenance of DNA trinucleotide repeats	GO:0035753	8	1.94e-02	visualization icons
BP	DNA repair	GO:0006281	284	5.44e-07	visualization icons
BP	double-strand break repair	GO:0006302	109	3.46e-05	visualization icons
BP	recombinational repair	GO:0000725	67	4.03e-04	visualization icons
BP	double-strand break repair via homologous recombination	GO:0000724	59	1.01e-03	visualization icons
BP	mating type switching	GO:0007533	29	4.97e-04	visualization icons
BP	protein phosphorylation	GO:0006468	213	1.46e-06	visualization icons
BP	regulation of protein phosphorylation	GO:0001932	77	7.37e-05	visualization icons
BP	regulation of protein kinase activity	GO:0045859	65	8.66e-06	visualization icons
BP	regulation of cyclin-dependent protein kinase activity	GO:1904029	32	1.01e-08	visualization icons
BP	regulation of protein serine/threonine kinase activity	GO:0071900	43	5.35e-07	visualization icons
BP	regulation of cyclin-dependent protein serine/threonine kinase ...	GO:0000079	32	1.01e-08	visualization icons
BP	meiotic mismatch repair	GO:0000710	9	1.96e-02	visualization icons
BP	polysaccharide catabolic process	GO:0000272	18	2.83e-02	visualization icons
BP	spermine transport	GO:0000296	4	3.84e-02	visualization icons
BP	morphogenesis checkpoint	GO:0044879	4	2.57e-04	visualization icons
BP	sulfur amino acid metabolic process	GO:0000096	50	1.75e-02	visualization icons
BP	methionine metabolic process	GO:0006555	42	4.04e-03	visualization icons
source	term name	term ID	n. of term genes	corrected p-value	visualization icons
CC	protein-DNA complex	GO:0032993	61	2.78e-09	visualization icons
CC	pre-replicative complex	GO:0036387	17	1.30e-03	visualization icons
CC	DNA replication preinitiation complex	GO:0031261	22	7.17e-03	visualization icons
CC	DNA packaging complex	GO:0044815	17	1.39e-03	visualization icons
CC	DNA bending complex	GO:1990104	12	1.14e-04	visualization icons
CC	non-membrane-bound organelle	GO:0043228	1276	1.17e-02	visualization icons
CC	intracellular non-membrane-bound organelle	GO:0043232	1276	1.17e-02	visualization icons
CC	cytoskeleton	GO:0005856	233	2.33e-03	visualization icons