

id	source	term_id	term_name	term_size	intersection_size	p_value
1	GO:BP	GO:0019509	L–methionine salvage from methylthioadenosine	4	2	1.0e–02
2	GO:BP	GO:0071267	L–methionine salvage	4	2	1.0e–02
3	GO:BP	GO:0043102	amino acid salvage	4	2	1.0e–02
4	GO:BP	GO:0071265	L–methionine biosynthetic process	5	2	1.2e–02
5	GO:BP	GO:0009086	methionine biosynthetic process	19	2	1.4e–01
6	GO:BP	GO:0043094	cellular metabolic compound salvage	18	2	1.4e–01
7	GO:BP	GO:0006555	methionine metabolic process	21	2	1.4e–01
8	GO:BP	GO:0090306	meiotic spindle assembly	1	1	1.4e–01
9	GO:BP	GO:0015868	purine ribonucleotide transport	4	1	1.8e–01
10	GO:BP	GO:0015867	ATP transport	4	1	1.8e–01
11	GO:BP	GO:0015866	ADP transport	4	1	1.8e–01
12	GO:BP	GO:0010089	xylem development	4	1	1.8e–01
13	GO:BP	GO:0009067	aspartate family amino acid biosynthetic process	32	2	1.8e–01
14	GO:BP	GO:0009066	aspartate family amino acid metabolic process	36	2	1.8e–01
15	GO:BP	GO:1990544	mitochondrial ATP transmembrane transport	4	1	1.8e–01
16	GO:BP	GO:0140021	mitochondrial ADP transmembrane transport	4	1	1.8e–01
17	GO:BP	GO:0002949	tRNA threonylcarbamoyladenosine modification	2	1	1.8e–01
18	GO:BP	GO:0060548	negative regulation of cell death	3	1	1.8e–01
19	GO:BP	GO:0048759	xylem vessel member cell differentiation	3	1	1.8e–01
20	GO:BP	GO:0000212	meiotic spindle organization	3	1	1.8e–01
21	GO:BP	GO:0070525	tRNA threonylcarbamoyladenosine metabolic process	2	1	1.8e–01
22	GO:BP	GO:0072530	purine–containing compound transmembrane transport	4	1	1.8e–01
23	GO:BP	GO:0000097	sulfur amino acid biosynthetic process	32	2	1.8e–01
24	GO:BP	GO:1905177	tracheary element differentiation	3	1	1.8e–01
25	GO:BP	GO:0000096	sulfur amino acid metabolic process	38	2	1.8e–01
26	GO:BP	GO:1901679	nucleotide transmembrane transport	4	1	1.8e–01
27	GO:BP	GO:0051503	adenine nucleotide transport	5	1	1.9e–01
28	GO:BP	GO:0015969	guanosine tetraphosphate metabolic process	5	1	1.9e–01
29	GO:BP	GO:0034035	purine ribonucleoside bisphosphate metabolic process	5	1	1.9e–01
30	GO:BP	GO:0015865	purine nucleotide transport	5	1	1.9e–01
31	GO:BP	GO:0006520	amino acid metabolic process	252	4	2.1e–01
32	GO:BP	GO:0044272	sulfur compound biosynthetic process	58	2	2.2e–01
33	GO:BP	GO:0010467	gene expression	1650	12	2.6e–01
34	GO:BP	GO:0006595	polyamine metabolic process	9	1	2.6e–01
35	GO:BP	GO:0009116	nucleoside metabolic process	9	1	2.6e–01
36	GO:BP	GO:0051254	positive regulation of RNA metabolic process	81	2	2.6e–01
37	GO:BP	GO:0000184	nuclear–transcribed mRNA catabolic process, nonsense–mediated decay	9	1	2.6e–01
38	GO:BP	GO:0006862	nucleotide transport	11	1	2.6e–01
39	GO:BP	GO:0009891	positive regulation of biosynthetic process	81	2	2.6e–01
40	GO:BP	GO:0009926	auxin polar transport	10	1	2.6e–01
41	GO:BP	GO:0010557	positive regulation of macromolecule biosynthetic process	78	2	2.6e–01
42	GO:BP	GO:1903508	positive regulation of nucleic acid–templated transcription	76	2	2.6e–01
43	GO:BP	GO:0045935	positive regulation of nucleobase–containing compound metabolic process	82	2	2.6e–01
44	GO:BP	GO:0019348	dolichol metabolic process	10	1	2.6e–01
45	GO:BP	GO:0016093	polyprenol metabolic process	10	1	2.6e–01
46	GO:BP	GO:0045893	positive regulation of DNA–templated transcription	76	2	2.6e–01
47	GO:BP	GO:0031328	positive regulation of cellular biosynthetic process	79	2	2.6e–01
48	GO:BP	GO:0051225	spindle assembly	8	1	2.6e–01
49	GO:BP	GO:0010941	regulation of cell death	8	1	2.6e–01
50	GO:BP	GO:1901657	glycosyl compound metabolic process	12	1	2.6e–01
51	GO:BP	GO:1902680	positive regulation of RNA biosynthetic process	76	2	2.6e–01
52	GO:BP	GO:0045132	meiotic chromosome segregation	13	1	2.8e–01
53	GO:BP	GO:0031325	positive regulation of cellular metabolic process	92	2	3.0e–01
54	GO:BP	GO:1901607	alpha–amino acid biosynthetic process	91	2	3.0e–01
55	GO:BP	GO:0016070	RNA metabolic process	1340	10	3.0e–01
56	GO:BP	GO:0007051	spindle organization	15	1	3.0e–01
57	GO:BP	GO:0051173	positive regulation of nitrogen compound metabolic process	101	2	3.1e–01
58	GO:BP	GO:0010604	positive regulation of macromolecule metabolic process	104	2	3.1e–01
59	GO:BP	GO:0044281	small molecule metabolic process	677	6	3.1e–01
60	GO:BP	GO:0006790	sulfur compound metabolic process	102	2	3.1e–01
61	GO:BP	GO:0140694	non–membrane–bounded organelle assembly	17	1	3.1e–01
62	GO:BP	GO:0008652	amino acid biosynthetic process	102	2	3.1e–01
63	GO:BP	GO:0009914	hormone transport	16	1	3.1e–01
64	GO:BP	GO:0060918	auxin transport	16	1	3.1e–01
65	GO:BP	GO:0048366	leaf development	18	1	3.1e–01
66	GO:BP	GO:0010468	regulation of gene expression	840	7	3.1e–01
67	GO:BP	GO:0009893	positive regulation of metabolic process	109	2	3.2e–01
68	GO:BP	GO:0006807	nitrogen compound metabolic process	4126	23	3.2e–01
69	GO:BP	GO:1901264	carbohydrate derivative transport	20	1	3.2e–01
70	GO:BP	GO:0006399	tRNA metabolic process	109	2	3.2e–01
71	GO:BP	GO:0048522	positive regulation of cellular process	112	2	3.2e–01
72	GO:BP	GO:0070925	organelle assembly	25	1	3.3e–01
73	GO:BP	GO:0015748	organophosphate ester transport	23	1	3.3e–01
74	GO:BP	GO:0010087	phloem or xylem histogenesis	25	1	3.3e–01
75	GO:BP	GO:0008219	cell death	26	1	3.3e–01
76	GO:BP	GO:0033875	ribonucleoside bisphosphate metabolic process	27	1	3.3e–01
77	GO:BP	GO:0000956	nuclear–transcribed mRNA catabolic process	22	1	3.3e–01
78	GO:BP	GO:0044238	primary metabolic process	4847	26	3.3e–01
79	GO:BP	GO:0031347	regulation of defense response	27	1	3.3e–01
80	GO:BP	GO:0006576	biogenic amine metabolic process	25	1	3.3e–01
81	GO:BP	GO:0034641	cellular nitrogen compound metabolic process	2297	14	3.3e–01
82	GO:BP	GO:0033865	nucleoside bisphosphate metabolic process	27	1	3.3e–01
83	GO:BP	GO:0006355	regulation of DNA–templated transcription	739	6	3.3e–01
84	GO:BP	GO:0006139	nucleobase–containing compound metabolic process	1874	12	3.3e–01
85	GO:BP	GO:0034032	purine nucleoside bisphosphate metabolic process	27	1	3.3e–01
86	GO:BP	GO:0051252	regulation of RNA metabolic process	752	6	3.3e–01
87	GO:BP	GO:0060255	regulation of macromolecule metabolic process	915	7	3.3e–01
88	GO:BP	GO:2001141	regulation of RNA biosynthetic process	739	6	3.3e–01
89	GO:BP	GO:0140013	meiotic nuclear division	23	1	3.3e–01
90	GO:BP	GO:0098813	nuclear chromosome segregation	28	1	3.3e–01
91	GO:BP	GO:1901605	alpha–amino acid metabolic process	132	2	3.3e–01
92	GO:BP	GO:1903046	meiotic cell cycle process	28	1	3.3e–01
93	GO:BP	GO:1903506	regulation of nucleic acid–templated transcription	739	6	3.3e–01
94	GO:BP	GO:0019222	regulation of metabolic process	951	7	3.3e–01
95	GO:BP	GO:0019752	carboxylic acid metabolic process	432	4	3.3e–01
96	GO:BP	GO:0010556	regulation of macromolecule biosynthetic process	774	6	3.3e–01
97	GO:BP	GO:0031326	regulation of cellular biosynthetic process	776	6	3.3e–01
98	GO:BP	GO:0019219	regulation of nucleobase–containing compound metabolic process	774	6	3.3e–01
99	GO:BP	GO:0071704	organic substance metabolic process	5200	27	3.3e–01
100	GO:BP	GO:0006082	organic acid metabolic process	437	4	3.3e–01