-	source	term_id	term_name	term_size	intersection_size	p_value
1 2	GO:BP	GO:0019509 GO:0071267	L-methionine salvage from methylthioadenosine L-methionine salvage	4	2	1.0e-02 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 6	GO:BP GO:BP	GO:0009086 GO:0043094	methionine biosynthetic process cellular metabolic compound salvage	19 18	2	1.4e-01 1.4e-01
7	GO:BP	GO:0006555	methionine metabolic process	21	2	1.4e-01
8 9	GO:BP	GO:0090306 GO:0015868	meiotic spindle assembly purine ribonucleotide transport	4	1	1.4e-01 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 13	GO:BP GO:BP	GO:0010089 GO:0009067	xylem development aspartate family amino acid biosynthetic process	32	2	1.8e-01 1.8e-01
14	GO:BP	GO:0009066	aspartate family amino acid metabolic process	36	2	1.8e-01
15 16	GO:BP	GO:1990544 GO:0140021	mitochondrial ATP transmembrane transport mitochondrial ADP transmembrane transport	4	1	1.8e-01 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 20	GO:BP	GO:0048759 GO:0000212	xylem vessel member cell differentiation meiotic spindle organization	3	1	1.8e-01 1.8e-01
21	GO:BP	GO:0070525	tRNA threonylcarbamoyladenosine metabolic process	2	1	1.8e-01
22 23	GO:BP	GO:0072530 GO:0000097	purine–containing compound transmembrane transport sulfur amino acid biosynthetic process	32	2	1.8e-01 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 27	GO:BP GO:BP	GO:1901679 GO:0051503	nucleotide transmembrane transport adenine nucleotide transport	5	1	1.8e-01 1.9e-01
28	GO:BP	GO:0015969	guanosine tetraphosphate metabolic process	5	1	1.9e-01
29 30	GO:BP	GO:0034035 GO:0015865	purine ribonucleoside bisphosphate metabolic process purine nucleotide transport	5	1	1.9e-01 1.9e-01
31	GO:BP	GO:0006520	amino acid metabolic process	252	4	2.1e-01
32 33	GO:BP	GO:0044272	sulfur compound biosynthetic process	58 1650	2	2.2e-01
33 34	GO:BP GO:BP	GO:0010467 GO:0006595	gene expression polyamine metabolic process	9	12	2.6e-01 2.6e-01
35	GO:BP	GO:0009116	nucleoside metabolic process	9	1	2.6e-01
36 37	GO:BP GO:BP	GO:0051254 GO:0000184	positive regulation of RNA metabolic process nuclear–transcribed mRNA catabolic process, nonsense–mediated decay	81 9	1	2.6e-01 2.6e-01
38	GO:BP	GO:0006862	nucleotide transport	11	1	2.6e-01
39 40	GO:BP	GO:0009891 GO:0009926	positive regulation of biosynthetic process	81	2	2.6e-01 2.6e-01
40 41	GO:BP	GO:0009926 GO:0010557	auxin polar transport positive regulation of macromolecule biosynthetic process	78	2	2.6e-01 2.6e-01
42	GO:BP	GO:1903508	positive regulation of nucleic acid–templated transcription	76	2	2.6e-01
43 44	GO:BP GO:BP	GO:0045935 GO:0019348	positive regulation of nucleobase–containing compound metabolic process dolichol metabolic process	10	1	2.6e-01 2.6e-01
45	GO:BP	GO:0016093	polyprenol metabolic process	10	1	2.6e-01
46 47	GO:BP	GO:0045893 GO:0031328	positive regulation of DNA-templated transcription positive regulation of cellular biosynthetic process	76 79	2	2.6e-01 2.6e-01
4 <i>7</i> 48	GO:BP	GO:0031328 GO:0051225	spindle assembly	8	1	2.6e-01 2.6e-01
49	GO:BP	GO:0010941	regulation of cell death	8	1	2.6e-01
50 51	GO:BP	GO:1901657 GO:1902680	glycosyl compound metabolic process positive regulation of RNA biosynthetic process	12 76	2	2.6e-01 2.6e-01
52	GO:BP	GO:0045132	meiotic chromosome segregation	13	1	2.8e-01
53 54	GO:BP	GO:0031325 GO:1901607	positive regulation of cellular metabolic process alpha–amino acid biosynthetic process	92	2	3.0e-01 3.0e-01
55	GO:BP	GO:1901607	RNA metabolic process			U_U_U_U I
56 57		GO.0016070	TATA MOLUDONO PIOCESS	1340	10	3.0e-01
	GO:BP	GO:0007051	spindle organization	15	10	3.0e-01 3.0e-01
58	GO:BP GO:BP				10	3.0e-01
58 59	GO:BP GO:BP GO:BP	GO:0007051 GO:0051173 GO:0010604 GO:0044281	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process	15 101 104 677	10 1 2 2 6	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01
58 59 60	GO:BP	GO:0007051 GO:0051173 GO:0010604	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process	15 101 104	10 1 2 2	3.0e-01 3.0e-01 3.1e-01 3.1e-01
58 59 60 61 62	GO:BP GO:BP GO:BP GO:BP GO:BP	GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process	15 101 104 677 102 17	10 1 2 2 6 2 1	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01
58 59 60 61 62 63	GO:BP GO:BP GO:BP GO:BP GO:BP GO:BP	GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport	15 101 104 677 102 17 102 16	10 1 2 2 2 6 2	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01
58 59 60 61 62 63 64	GO:BP GO:BP GO:BP GO:BP GO:BP	GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process	15 101 104 677 102 17	10 1 2 2 6 2 1 2	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01
58 59 60 61 62 63 64 65 66	GO:BP GO:BP GO:BP GO:BP GO:BP GO:BP GO:BP GO:BP GO:BP	GO:0007051 GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0060918 GO:0048366 GO:0010468	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression	15 101 104 677 102 17 102 16 16 18 840	10 1 2 2 6 2 1 2 1 1 1	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01
58 59 60 61	GO:BP GO:BP GO:BP GO:BP GO:BP GO:BP GO:BP GO:BP	GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0060918 GO:0048366	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development	15 101 104 677 102 17 102 16 16 16	10 1 2 2 6 2 1 2 1 1 1	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01
58 59 60 61 62 63 64 65 66 67 68	GO:BP	GO:0007051 GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0060918 GO:0010468 GO:0010468 GO:0009893 GO:0006807 GO:1901264	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport	15 101 104 677 102 17 102 16 16 18 840 109 4126 20	10 1 2 2 6 2 1 1 2 1 7 2 23 1	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01
58 59 60 61 62 63 64 65 66 67 68 69 70	GO:BP	GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0060918 GO:0048366 GO:0010468 GO:0009893 GO:0006807	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport tRNA metabolic process	15 101 104 677 102 17 102 16 16 18 840 109 4126	10 1 2 2 6 2 1 1 1 1 7 2 23	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01
58 59 60 61 62 63 64 65 66 67 68	GO:BP	GO:0007051 GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0060918 GO:0010468 GO:0010468 GO:0009893 GO:0006807 GO:1901264 GO:0006399	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport	15 101 104 677 102 17 102 16 16 18 840 109 4126 20 109	10 1 2 2 6 2 1 1 2 1 7 2 23 1 2	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01 3.2e-01
58 59 60 61 62 63 64 65 66 67 68 69 70 71 72	GO:BP	GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0048366 GO:0010468 GO:0006807 GO:1901264 GO:0006399 GO:0048522 GO:0070925 GO:0015748	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport tRNA metabolic process positive regulation of cellular process organelle assembly organophosphate ester transport	15 101 104 677 102 17 102 16 16 18 840 109 4126 20 109 112 25 23	10 1 2 2 6 2 1 1 2 1 1 7 2 23 1 2 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01 3.2e-01 3.2e-01 3.3e-01 3.3e-01 3.3e-01
58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73	GO:BP	GO:0007051 GO:0051173 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0060918 GO:0010468 GO:0010468 GO:0006807 GO:1901264 GO:0006399 GO:0048522 GO:0070925	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport tRNA metabolic process positive regulation of cellular process organelle assembly	15 101 104 677 102 17 102 16 16 18 840 109 4126 20 109 112 25	10 1 2 2 6 2 1 1 2 1 1 7 2 23 1 2 2 1	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01 3.2e-01 3.2e-01 3.3e-01 3.3e-01
58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76	GO:BP	GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0060918 GO:0010468 GO:0010468 GO:0006807 GO:1901264 GO:0006399 GO:0048522 GO:0070925 GO:0015748 GO:0010087 GO:0008219 GO:00033875	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport tRNA metabolic process positive regulation of cellular process organelle assembly organophosphate ester transport phloem or xylem histogenesis cell death ribonucleoside bisphosphate metabolic process	15 101 104 677 102 17 102 16 16 18 840 109 4126 20 109 112 25 23 25 26 27	10 1 2 2 6 2 1 1 2 1 1 1 1 7 2 2 2 3 1 2 2 1 1 1 1 1 1 1 1 1 1	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01 3.2e-01 3.2e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01
58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77	GO:BP	GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0060918 GO:0048366 GO:0010468 GO:0006807 GO:1901264 GO:0006399 GO:0048522 GO:0070925 GO:0015748 GO:0010087 GO:0008219 GO:00033875 GO:0000956	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport tRNA metabolic process positive regulation of cellular process organelle assembly organophosphate ester transport phloem or xylem histogenesis cell death ribonucleoside bisphosphate metabolic process nuclear-transcribed mRNA catabolic process	15 101 104 677 102 17 102 16 16 18 840 109 4126 20 109 112 25 23 25 26 27 22	10 1 2 2 6 2 1 1 2 1 1 1 7 2 23 1 2 2 1 1 1 1 1 1 1 1 1 1 1	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01 3.2e-01 3.2e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01
58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74	GO:BP	GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0060918 GO:0010468 GO:0010468 GO:0006807 GO:1901264 GO:0006399 GO:0048522 GO:0070925 GO:0015748 GO:0010087 GO:0008219 GO:00033875	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport tRNA metabolic process positive regulation of cellular process organelle assembly organophosphate ester transport phloem or xylem histogenesis cell death ribonucleoside bisphosphate metabolic process	15 101 104 677 102 17 102 16 16 18 840 109 4126 20 109 112 25 23 25 26 27	10 1 2 2 6 2 1 1 2 1 1 1 1 7 2 2 2 3 1 2 2 1 1 1 1 1 1 1 1 1 1	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01 3.2e-01 3.2e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01
58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80	GO:BP	GO:0007051 GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0060918 GO:0010468 GO:0010468 GO:0006807 GO:1901264 GO:0006399 GO:0006399 GO:0015748 GO:0015748 GO:0010087 GO:0008219 GO:000956 GO:00044238 GO:0006576	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport tRNA metabolic process positive regulation of cellular process organelle assembly organophosphate ester transport phloem or xylem histogenesis cell death ribonucleoside bisphosphate metabolic process primary metabolic process regulation of defense response biogenic amine metabolic process	15 101 104 677 102 17 102 16 16 18 840 109 4126 20 109 112 25 23 25 26 27 22 4847 27 25	10 1 2 2 6 2 1 1 1 1 1 7 2 23 1 2 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1	3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01 3.2e-01 3.2e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01
58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78	GO:BP	GO:0007051 GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0009914 GO:0060918 GO:0048366 GO:0010468 GO:0006807 GO:1901264 GO:0006399 GO:0048522 GO:0070925 GO:0015748 GO:0010087 GO:0008219 GO:0008219 GO:000956 GO:00044238 GO:0031347	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport tRNA metabolic process positive regulation of cellular process organelle assembly organophosphate ester transport phloem or xylem histogenesis cell death ribonucleoside bisphosphate metabolic process primary metabolic process regulation of defense response	15 101 104 677 102 17 102 16 16 18 840 109 4126 20 109 112 25 23 25 26 27 22 4847 27	10 1 2 2 6 2 1 1 1 1 1 1 7 2 23 1 2 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1	3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01 3.2e-01 3.2e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01 3.3e-01
58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 30 31	GO:BP	GO:0007051 GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0048366 GO:0010468 GO:0006807 GO:1901264 GO:0006399 GO:0048522 GO:0070925 GO:0015748 GO:0015748 GO:001087 GO:0008219 GO:0008219 GO:0033875 GO:000956 GO:0034641 GO:0033865 GO:0006355	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport tRNA metabolic process positive regulation of cellular process organelle assembly organophosphate ester transport phloem or xylem histogenesis cell death ribonucleoside bisphosphate metabolic process primary metabolic process regulation of defense response biogenic amine metabolic process cellular nitrogen compound metabolic process regulation of DNA-templated transcription	15 101 104 677 102 17 102 16 16 18 840 109 4126 20 109 112 25 23 25 26 27 22 4847 27 25 2297 27 739	10 1 2 2 6 2 1 1 1 1 1 1 7 2 2 3 1 1 2 2 1 1 1 1 1 1 1 1 1 1 1 1 1	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01 3.2e-01 3.2e-01 3.3e-01
58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 30 31 32 33 34	GO:BP	GO:0007051 GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0060918 GO:0010468 GO:0010468 GO:0010468 GO:0006807 GO:1901264 GO:0006399 GO:0006399 GO:0015748 GO:0015748 GO:0008219 GO:0008219 GO:000956 GO:0033875 GO:000956 GO:0033865 GO:0033865 GO:0006355 GO:0006355	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport tRNA metabolic process positive regulation of cellular process organelle assembly organophosphate ester transport phloem or xylem histogenesis cell death ribonucleoside bisphosphate metabolic process primary metabolic process regulation of defense response biogenic amine metabolic process cellular nitrogen compound metabolic process regulation of DNA-templated transcription nucleobase-containing compound metabolic process	15 101 104 677 102 17 102 16 16 18 840 109 4126 20 109 112 25 23 25 26 27 22 4847 27 25 2297 27 739 1874	10 1 2 2 6 2 1 1 1 1 1 1 7 2 2 2 3 1 1 2 2 1 1 1 1 1 1 1 1 1 1 1 1	3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01 3.2e-01 3.2e-01 3.3e-01
58 59 60 61 62 63 64 65 66 67 70 71 72 73 74 75 76 77 78 79 30 31 32 33 34 35	GO:BP	GO:0007051 GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0048366 GO:0010468 GO:0006807 GO:1901264 GO:0006399 GO:0048522 GO:0070925 GO:0015748 GO:0015748 GO:001087 GO:0008219 GO:0008219 GO:0033875 GO:000956 GO:0034641 GO:0033865 GO:0006355	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport tRNA metabolic process positive regulation of cellular process organelle assembly organophosphate ester transport phloem or xylem histogenesis cell death ribonucleoside bisphosphate metabolic process primary metabolic process regulation of defense response biogenic amine metabolic process cellular nitrogen compound metabolic process regulation of DNA-templated transcription	15 101 104 677 102 17 102 16 16 18 840 109 4126 20 109 112 25 23 25 26 27 22 4847 27 25 2297 27 739	10 1 2 2 6 2 1 1 1 1 1 1 7 2 2 3 1 1 2 2 1 1 1 1 1 1 1 1 1 1 1 1 1	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01 3.2e-01 3.2e-01 3.3e-01
58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 30 31 32 33 34 35 36 37	GO:BP	GO:0007051 GO:0051173 GO:0010604 GO:0044281 GO:0006790 GO:0140694 GO:0008652 GO:0009914 GO:0060918 GO:0048366 GO:0010468 GO:0006807 GO:1901264 GO:0006399 GO:0048522 GO:0070925 GO:0015748 GO:0015748 GO:001087 GO:0008219 GO:0033875 GO:000956 GO:0034641 GO:0033865 GO:0006355 GO:0006139 GO:0034032 GO:00060255	spindle organization positive regulation of nitrogen compound metabolic process positive regulation of macromolecule metabolic process small molecule metabolic process sulfur compound metabolic process sulfur compound metabolic process non-membrane-bounded organelle assembly amino acid biosynthetic process hormone transport auxin transport leaf development regulation of gene expression positive regulation of metabolic process nitrogen compound metabolic process carbohydrate derivative transport tRNA metabolic process positive regulation of cellular process organelle assembly organophosphate ester transport phloem or xylem histogenesis cell death ribonucleoside bisphosphate metabolic process nuclear-transcribed mRNA catabolic process primary metabolic process regulation of defense response biogenic amine metabolic process cellular nitrogen compound metabolic process nucleoside bisphosphate metabolic process regulation of DNA-templated transcription nucleobase-containing compound metabolic process regulation of RNA metabolic process regulation of RNA metabolic process regulation of RNA metabolic process regulation of macromolecule metabolic process	15 101 104 677 102 17 102 16 16 18 840 109 4126 20 109 112 25 23 25 26 27 22 4847 27 25 2297 27 739 1874 27 752 915	10 1 2 6 2 1 2 1 1 2 23 1 2 2 1 <	3.0e-01 3.0e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.1e-01 3.2e-01 3.2e-01 3.2e-01 3.2e-01 3.2e-01 3.3e-01
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