id	source	term_id	term_name	term_size	intersection_size		p_value
1	GO:MF	GO:0061135	endopeptidase regulator activity	70	12		5.5e−16
2	GO:MF	GO:0061134	peptidase regulator activity	71	12		5.5e−16
3	GO:MF	GO:0030414	peptidase inhibitor activity	69	12		5.5e−16
4	GO:MF	GO:0004866	endopeptidase inhibitor activity	69	12		5.5e−16
5	GO:MF	GO:0004867	serine−type endopeptidase inhibitor activity	38	8		2.1e−11
6	GO:MF	GO:0140678	molecular function inhibitor activity	206	12		1.5e−10
7	GO:MF	GO:0004857	enzyme inhibitor activity	205	12		1.5e−10
8	GO:BP	GO:0009611	response to wounding	48	7		8.8e−08
9	GO:MF	GO:0030234	enzyme regulator activity	418	12		4.2e−07
10	GO:MF	GO:0098772	molecular function regulator activity	452	12		8.9e−07
11	GO:MF	GO:0016835	carbon−oxygen lyase activity	149	6		2.1e−04
12	GO:MF	GO:0000287	magnesium ion binding	153	6		2.2e−04
13	GO:MF	GO:0070006	metalloaminopeptidase activity	15	3		2.2e−04
14	GO:MF	GO:0010333	terpene synthase activity	46	4		2.3e−04
15	GO:MF	GO:0016838	carbon−oxygen lyase activity, acting on phosphates	54	4		3.8e−04
16	GO:MF	GO:0008235	metalloexopeptidase activity	19	3		3.8e−04
17	GO:MF	GO:0004177	aminopeptidase activity	20	3		4.1e−04
18	GO:MF	GO:0030145	manganese ion binding	34	3		2.0e−03
19	GO:MF	GO:0016829	lyase activity	367	7		2.5e−03
20	GO:MF	GO:0004144	diacylglycerol O−acyltransferase activity	10	2		4.1e−03
21	GO:BP	GO:0034214	protein hexamerization	3	2		4.7e−03
22	GO:MF	GO:0016411	acylglycerol O−acyltransferase activity	15	2		8.9e−03
23	GO:BP	GO:0019346	transsulfuration	6	2		1.2e−02
24	GO:BP	GO:0050667	homocysteine metabolic process	6	2		1.2e−02
25	GO:BP	GO:0009092	homoserine metabolic process	7	2		1.3e−02
26	GO:MF	GO:0008238	exopeptidase activity	70	3		1.3e−02
27	GO:MF	GO:0008237	metallopeptidase activity	89	3		2.5e−02
28	GO:MF	GO:0000234	phosphoethanolamine N−methyltransferase activity	2	1		4.3e−02