source	term_id	term_name	term_size	intersection_size		p_value
GO:BP	GO:0005975	carbohydrate metabolic process	963	16		4.9e-04
GO:CC	GO:0048046	apoplast	146	5		6.8e-03
GO:MF	GO:0030145	manganese ion binding	54	4		1.4e-02
GO:MF	GO:0003824	catalytic activity	8918	52		2.0e-02
GO:MF	GO:0045735	nutrient reservoir activity	75	4		2.0e-02
GO:MF	GO:0016835	carbon-oxygen lyase activity	152	5		2.0e-02
GO:MF	GO:0004096	catalase activity	9	2		2.1e-02
GO:CC	GO:0005576	extracellular region	487	7		2.9e-02
GO:MF	GO:0016491	oxidoreductase activity	1821	17		3.1e-02
GO:MF	GO:0004869	cysteine-type endopeptidase inhibitor activity	12	2		3.1e-02
GO:MF	GO:0010333	terpene synthase activity	54	3		3.3e-02
GO:MF	GO:0016838	carbon-oxygen lyase activity, acting on phosphates	61	3		4.2e-02
GO:BP	GO:0016102	diterpenoid biosynthetic process	33	3		4.3e-02
GO:BP	GO:0016101	diterpenoid metabolic process	34	3		4.3e-02
GO:MF	GO:0016829	lyase activity	336	6		4.7e-02
GO:MF	GO:0008453	alanine-glyoxylate transaminase activity	1	1		4.7e-02
GO:MF	GO:0043169	cation binding	2798	21		4.7e-02
GO:MF	GO:0004857	enzyme inhibitor activity	151	4		4.7e-02
GO:MF	GO:0046872	metal ion binding	2785	21		4.7e-02
GO:MF	GO:0016798	hydrolase activity, acting on glycosyl bonds	496	7		4.7e-02
GO:MF	GO:0004553	hydrolase activity, hydrolyzing O-glycosyl compounds	457	7		4.7e-02
GO:MF	GO:0016174	NAD(P)H oxidase H2O2–forming activity	1	1		4.7e-02
GO:MF	GO:0033946	xyloglucan-specific endo-beta-1,4-glucanase activity	1	1		4.7e-02
	GO:BP GO:CC GO:MF	GO:BP GO:0005975 GO:CC GO:0048046 GO:MF GO:0030145 GO:MF GO:0003824 GO:MF GO:0045735 GO:MF GO:0004096 GO:CC GO:0005576 GO:MF GO:0016491 GO:MF GO:0016491 GO:MF GO:0016838 GO:MF GO:0016838 GO:BP GO:0016102 GO:BP GO:0016101 GO:MF GO:0016829 GO:MF GO:0004857 GO:MF GO:004857 GO:MF GO:0046872 GO:MF GO:0016798 GO:MF GO:0004553 GO:MF GO:0004553 GO:MF GO:0004553 GO:MF GO:0004553 GO:MF GO:0004553 GO:MF GO:0004553	GO:BP GO:0005975 carbohydrate metabolic process GO:CC GO:0048046 apoplast GO:MF GO:0030145 manganese ion binding GO:MF GO:0003824 catalytic activity GO:MF GO:0045735 nutrient reservoir activity GO:MF GO:0016835 carbon-oxygen lyase activity GO:MF GO:0005576 extracellular region GO:MF GO:0016491 oxidoreductase activity GO:MF GO:0016491 oxidoreductase activity GO:MF GO:0016838 carbon-oxygen lyase activity GO:MF GO:0016839 cysteine-type endopeptidase inhibitor activity GO:MF GO:0016838 carbon-oxygen lyase activity, acting on phosphates GO:MF GO:0016838 carbon-oxygen lyase activity, acting on phosphates GO:BP GO:0016102 diterpenoid biosynthetic process GO:BP GO:0016101 diterpenoid metabolic process GO:MF GO:0016829 lyase activity GO:MF GO:0008453 alanine-glyoxylate transaminase activity GO:MF GO:0043169 cation binding GO:MF GO:004857 enzyme inhibitor activity GO:MF GO:0046872 metal ion binding GO:MF GO:0016798 hydrolase activity, acting on glycosyl bonds GO:MF GO:0016174 NAD(P)H oxidase H2O2-forming activity	GO:BP         GO:0005975         carbohydrate metabolic process         963           GO:CC         GO:0048046         apoplast         146           GO:MF         GO:0030145         manganese ion binding         54           GO:MF         GO:0003824         catalytic activity         8918           GO:MF         GO:0045735         nutrient reservoir activity         75           GO:MF         GO:0016835         carbon-oxygen lyase activity         152           GO:MF         GO:0004096         catalase activity         9           GO:MF         GO:0005576         extracellular region         487           GO:MF         GO:0016491         oxidoreductase activity         1821           GO:MF         GO:0016491         oxidoreductase activity         12           GO:MF         GO:0016839         cysteine-type endopeptidase inhibitor activity         12           GO:MF         GO:0016838         carbon-oxygen lyase activity         54           GO:MF         GO:0016838         carbon-oxygen lyase activity, acting on phosphates         61           GO:BP         GO:0016101         diterpenoid metabolic process         33           GO:BP         GO:0016829         lyase activity         336           GO:MF	GO:BP GO:0005975 carbohydrate metabolic process 963 16 GO:CC GO:0048046 apoplast 146 5 GO:MF GO:0030145 manganese ion binding 54 4 GO:MF GO:0003824 catalytic activity 8918 52 GO:MF GO:0045735 nutrient reservoir activity 75 4 GO:MF GO:0016835 carbon-oxygen lyase activity 9 2 GO:MF GO:0004096 catalase activity 9 2 GO:CC GO:0005576 extracellular region 487 7 GO:MF GO:0016491 oxidoreductase activity 1821 17 GO:MF GO:0004869 cysteine-type endopeptidase inhibitor activity 12 2 GO:MF GO:0016838 carbon-oxygen lyase activity 54 3 GO:MF GO:0016838 carbon-oxygen lyase activity 54 3 GO:MF GO:0016838 carbon-oxygen lyase activity 54 3 GO:MF GO:0016838 carbon-oxygen lyase activity, acting on phosphates 61 3 GO:BP GO:0016102 diterpenoid biosynthetic process 33 3 GO:BP GO:0016101 diterpenoid metabolic process 34 3 GO:MF GO:0016829 lyase activity 336 6 GO:MF GO:0004857 alanine-glyoxylate transaminase activity 1 1 1 GO:MF GO:0004857 enzyme inhibitor activity 151 4 GO:MF GO:0004857 metal ion binding 2798 21 GO:MF GO:0048672 metal ion binding 2785 21 GO:MF GO:004553 hydrolase activity, acting on glycosyl bonds 496 7 GO:MF GO:0016798 hydrolase activity, acting on glycosyl compounds 457 7 GO:MF GO:0016798 hydrolase activity, hydrolyzing O-glycosyl compounds 457 7 GO:MF GO:001674 NAD(P)H oxidase H2O2-forming activity 1 1 1	GO:BP GO:0005975 carbohydrate metabolic process 963 16 GO:CC GO:0048046 apoplast 146 5 GO:MF GO:0030145 manganese ion binding 54 4 GO:MF GO:0003824 catalytic activity 8918 52 GO:MF GO:0048735 nutrient reservoir activity 75 4 GO:MF GO:0048735 nutrient reservoir activity 75 4 GO:MF GO:0016835 carbon-oxygen lyase activity 9 2 GO:MF GO:0004096 catalase activity 9 2 GO:CC GO:0005576 extracellular region 487 7 GO:MF GO:0016491 oxidoreductase activity 1821 17 GO:MF GO:0016838 carbon-oxygen lyase activity 12 2 GO:MF GO:0016838 carbon-oxygen lyase activity, acting on phosphates 61 3 GO:MF GO:0016838 carbon-oxygen lyase activity, acting on phosphates 61 3 GO:BP GO:0016101 diterpenoid biosynthetic process 33 3 GO:MF GO:0016829 lyase activity 336 6 GO:MF GO:0016829 lyase activity 336 6 GO:MF GO:0004857 enzyme inhibitor activity 151 4 GO:MF GO:004867 enzyme inhibitor activity 151 4 GO:MF GO:0046872 metal ion binding 2798 21 GO:MF GO:0046873 hydrolase activity, acting on glycosyl bonds 496 7 GO:MF GO:0016798 hydrolase activity, acting on glycosyl compounds 457 7 GO:MF GO:001674 NAD(P)H oxidase H2O2-forming activity 1 1 GO:MF GO:0033946 xyloglucan-specific endo-beta-1,4-glucanase activity 1 1