

# Reuben J. Peters

## List of Publications by Year in descending order

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155  
papers

12,687  
citations

27035

58  
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31191

106  
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169  
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169  
docs citations

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times ranked

9259  
citing authors

#	ARTICLE	IF	CITATIONS
1	Dissecting the labdane-related diterpenoid biosynthetic gene clusters in rice reveals directional cross-cluster phytotoxicity. <i>New Phytologist</i> , 2022, 233, 878-889.	3.5	17
2	Diterpene synthases from <i>Leonurus japonicus</i> elucidate epoxy-bridge formation of spiro-labdane diterpenoids. <i>Plant Physiology</i> , 2022, 189, 99-111.	2.3	5
3	Tanshinones: Leading the way into Lamiaceae labdane-related diterpenoid biosynthesis. <i>Current Opinion in Plant Biology</i> , 2022, 66, 102189.	3.5	20
4	Deceptive Complexity in Formation of Cleistantha-8,12-diene. <i>Organic Letters</i> , 2022, 24, 2646-2649.	2.4	2
5	Origin and early evolution of the plant terpene synthase family. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2100361119.	3.3	48
6	Production of the plant hormone gibberellin by rhizobia increases host legume nodule size. <i>ISME Journal</i> , 2022, 16, 1809-1817.	4.4	13
7	Rice diterpenoid phytoalexins are involved in defence against parasitic nematodes and shape rhizosphere nematode communities. <i>New Phytologist</i> , 2022, 235, 1231-1245.	3.5	12
8	Investigation of Acid-Base Catalysis in Halimadienyl Diphosphate Synthase Involved in <i>Mycobacterium tuberculosis</i> Virulence. <i>ACS Bio &amp; Med Chem Au</i> , 2022, 2, 490-498.	1.7	4
9	A (conditional) role for labdane-related diterpenoid natural products in rice stomatal closure. <i>New Phytologist</i> , 2021, 230, 698-709.	3.5	18
10	A pair of threonines mark ent-kaurane synthases for phytohormone biosynthesis. <i>Phytochemistry</i> , 2021, 184, 112672.	1.4	7
11	Rice contains a biosynthetic gene cluster associated with production of the casbane-type diterpenoid phytoalexin ent-10-oxodepressin. <i>New Phytologist</i> , 2021, 231, 85-93.	3.5	21
12	Expansion within the CYP71D subfamily drives the heterocyclization of tanshinones synthesis in <i>Salvia miltiorrhiza</i> . <i>Nature Communications</i> , 2021, 12, 685.	5.8	94
13	Interdependent evolution of biosynthetic gene clusters for momilactone production in rice. <i>Plant Cell</i> , 2021, 33, 290-305.	3.1	34
14	Mining of the <i>Catharanthus roseus</i> Genome Leads to Identification of a Biosynthetic Gene Cluster for Fungicidal Sesquiterpenes. <i>Journal of Natural Products</i> , 2021, 84, 2709-2716.	1.5	5
15	Magnesium-specific ring expansion/contraction catalysed by the class II diterpene cyclase from pleuromutilin biosynthesis. <i>Organic and Biomolecular Chemistry</i> , 2020, 18, 5586-5588.	1.5	2
16	Why are momilactones always associated with biosynthetic gene clusters in plants?. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 13867-13869.	3.3	21
17	Doing the gene shuffle to close synteny: dynamic assembly of biosynthetic gene clusters. <i>New Phytologist</i> , 2020, 227, 992-994.	3.5	13
18	Oil Body Formation in <i>Marchantia polymorpha</i> Is Controlled by MpC1HDZ and Serves as a Defense against Arthropod Herbivores. <i>Current Biology</i> , 2020, 30, 2815-2828.e8.	1.8	48

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19	Probing Enzymatic Structure and Function in the Dihydroxylating Sesquiterpene Synthase ZmEDS. <i>Biochemistry</i> , 2020, 59, 2660-2666.	1.2	5
20	The honeysuckle genome provides insight into the molecular mechanism of carotenoid metabolism underlying dynamic flower coloration. <i>New Phytologist</i> , 2020, 227, 930-943.	3.5	68
21	Unraveling a Tangled Skein: Evolutionary Analysis of the Bacterial Gibberellin Biosynthetic Operon. <i>MSphere</i> , 2020, 5, .	1.3	7
22	Genome of <i>Tripterygium wilfordii</i> and identification of cytochrome P450 involved in triptolide biosynthesis. <i>Nature Communications</i> , 2020, 11, 971.	5.8	103
23	Introducing selective agrochemical manipulation of gibberellin metabolism into a cereal crop. <i>Nature Plants</i> , 2020, 6, 67-72.	4.7	17
24	Identification of RoCYP01 (CYP716A15) enables construction of engineered yeast for high-yield production of betulinic acid. <i>Applied Microbiology and Biotechnology</i> , 2019, 103, 7029-7039.	1.7	28
25	Switching on a Nontraditional Enzymatic Baseâ€”Deprotonation by Serine in the <i>ent</i>-Kaurene Synthase from <i>Bradyrhizobium japonicum</i>. <i>ACS Catalysis</i> , 2019, 9, 8867-8871.	5.5	18
26	Combinatorial biosynthesis and the basis for substrate promiscuity in class I diterpene synthases. <i>Metabolic Engineering</i> , 2019, 55, 44-58.	3.6	24
27	CYP72A enzymes catalyse 13-hydrolyzation of gibberellins. <i>Nature Plants</i> , 2019, 5, 1057-1065.	4.7	53
28	The genome of the medicinal plant <i>Andrographis paniculata</i> provides insight into the biosynthesis of the bioactive diterpenoid neoandrographolide. <i>Plant Journal</i> , 2019, 97, 841-857.	2.8	75
29	Isoprenyl diphosphate synthases: the chain length determining step in terpene biosynthesis. <i>Planta</i> , 2019, 249, 9-20.	1.6	29
30	Conserved bases for the initial cyclase in gibberellin biosynthesis: from bacteria to plants. <i>Biochemical Journal</i> , 2019, 476, 2607-2621.	1.7	22
31	Changing Face: A Key Residue for the Addition of Water by Sclareol Synthase. <i>ACS Catalysis</i> , 2018, 8, 3133-3137.	5.5	14
32	Diverging Mechanisms: Cytochromeâ€”P450â€”Catalyzed Demethylation and Î³â€”Lactone Formation in Bacterial Gibberellin Biosynthesis. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 6082-6085.	7.2	18
33	Inferring Roles in Defense from Metabolic Allocation of Rice Diterpenoids. <i>Plant Cell</i> , 2018, 30, 1119-1131.	3.1	55
34	Direct production of dihydroxylated sesquiterpenoids by a maize terpene synthase. <i>Plant Journal</i> , 2018, 94, 847-856.	2.8	27
35	Premutilin Synthase: Ring Rearrangement by a Class II Diterpene Cyclase. <i>Organic Letters</i> , 2018, 20, 1200-1202.	2.4	21
36	Diverging Mechanisms: Cytochromeâ€”P450â€”Catalyzed Demethylation and Î³â€”Lactone Formation in Bacterial Gibberellin Biosynthesis. <i>Angewandte Chemie</i> , 2018, 130, 6190-6193.	1.6	3

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37	Identification and functional characterization of diterpene synthases for triptolide biosynthesis from <i>Tripterygium wilfordii</i> . <i>Plant Journal</i> , 2018, 93, 50-65.	2.8	52
38	A Third Class: Functional Gibberellin Biosynthetic Operon in Beta-Proteobacteria. <i>Frontiers in Microbiology</i> , 2018, 9, 2916.	1.5	23
39	Arginine in the FARM and SARM: A Role in Chain-Length Determination for Arginine in the Aspartate-Rich Motifs of Isoprenyl Diphosphate Synthases from <i>Mycobacterium tuberculosis</i> . <i>Molecules</i> , 2018, 23, 2546.	1.7	6
40	Terpenoid Secondary Metabolites in Bryophytes: Chemical Diversity, Biosynthesis and Biological Functions. <i>Critical Reviews in Plant Sciences</i> , 2018, 37, 210-231.	2.7	57
41	Catalytic Bases and Stereocontrol in Lamiaceae Class II Diterpene Cyclases. <i>Biochemistry</i> , 2018, 57, 3473-3479.	1.2	20
42	Probing the specificity of CYP112 in bacterial gibberellin biosynthesis. <i>Biochemical Journal</i> , 2018, 475, 2167-2177.	1.7	7
43	Probing Labdane-Related Diterpenoid Biosynthesis in the Fungal Genus <i>Aspergillus</i> . <i>Journal of Natural Products</i> , 2017, 80, 328-333.	1.5	19
44	Characterization of CYP115 As a Gibberellin 3-Oxidase Indicates That Certain Rhizobia Can Produce Bioactive Gibberellin A <sub>4</sub> . <i>ACS Chemical Biology</i> , 2017, 12, 912-917.	1.6	28
45	An operon for production of bioactive gibberellin A <sub>4</sub> phytohormone with wide distribution in the bacterial rice leaf streak pathogen <i>Xanthomonas oryzae</i> pv. <i>oryzicola</i> . <i>New Phytologist</i> , 2017, 214, 1260-1266.	3.5	26
46	A Pair of Residues That Interactively Affect Diterpene Synthase Product Outcome. <i>ACS Chemical Biology</i> , 2017, 12, 862-867.	1.6	34
47	<i>cis</i> or <i>trans</i> with class II diterpene cyclases. <i>Organic and Biomolecular Chemistry</i> , 2017, 15, 3158-3160.	1.5	14
48	Investigating the Phylogenetic Range of Gibberellin Biosynthesis in Bacteria. <i>Molecular Plant-Microbe Interactions</i> , 2017, 30, 343-349.	1.4	25
49	Insights into Land Plant Evolution Garnered from the <i>Marchantia polymorpha</i> Genome. <i>Cell</i> , 2017, 171, 287-304.e15.	13.5	973
50	18O <sub>2</sub> labeling experiments illuminate the oxidation of ent-kaurene in bacterial gibberellin biosynthesis. <i>Organic and Biomolecular Chemistry</i> , 2017, 15, 7566-7571.	1.5	11
51	Biosynthesis of Diterpenoids in <i>Tripterygium</i> Adventitious Root Cultures. <i>Plant Physiology</i> , 2017, 175, 92-103.	2.3	27
52	Elucidation of gibberellin biosynthesis in bacteria reveals convergent evolution. <i>Nature Chemical Biology</i> , 2017, 13, 69-74.	3.9	103
53	Identification of a Dolabellane Type Diterpene Synthase and other Root-Expressed Diterpene Synthases in <i>Arabidopsis</i> . <i>Frontiers in Plant Science</i> , 2016, 7, 1761.	1.7	24
54	Extending a Single Residue Switch for Abbreviating Catalysis in Plant ent-Kaurene Synthases. <i>Frontiers in Plant Science</i> , 2016, 7, 1765.	1.7	22

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55	Investigating inducible short-chain alcohol dehydrogenases/reductases clarifies rice oryzalexin biosynthesis. <i>Plant Journal</i> , 2016, 88, 271-279.	2.8	30
56	Blocking Deprotonation with Retention of Aromaticity in a Plant <i>ent</i> -Copalyl Diphosphate Synthase Leads to Product Rearrangement. <i>Angewandte Chemie - International Edition</i> , 2016, 55, 634-638.	7.2	61
57	Cytochrome P450 promiscuity leads to a bifurcating biosynthetic pathway for tanshinones. <i>New Phytologist</i> , 2016, 210, 525-534.	3.5	183
58	Extreme promiscuity of a bacterial and a plant diterpene synthase enables combinatorial biosynthesis. <i>Metabolic Engineering</i> , 2016, 37, 24-34.	3.6	63
59	Microbial-type terpene synthase genes occur widely in nonseed land plants, but not in seed plants. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 12328-12333.	3.3	70
60	Molecular Diversity of Terpene Synthases in the Liverwort <i>Marchantia polymorpha</i> . <i>Plant Cell</i> , 2016, 28, tpc.00062.2016.	3.1	48
61	Labeling Studies Clarify the Committed Step in Bacterial Gibberellin Biosynthesis. <i>Organic Letters</i> , 2016, 18, 5974-5977.	2.4	17
62	Analysis of the Genome Sequence of the Medicinal Plant <i>Salvia miltiorrhiza</i> . <i>Molecular Plant</i> , 2016, 9, 949-952.	3.9	255
63	Probing the promiscuity of <i>ent</i> -kaurene oxidases via combinatorial biosynthesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 2526-2531.	3.3	53
64	Product Rearrangement from Altering a Single Residue in the Rice <i>syn</i> -Copalyl Diphosphate Synthase. <i>Organic Letters</i> , 2016, 18, 1060-1063.	2.4	40
65	A Tandem Array of <i>ent</i> -kaurene Synthases in Maize with Roles in Gibberellin and More Specialized Metabolism. <i>Plant Physiology</i> , 2016, 170, 742-751.	2.3	81
66	Characterization of CYP71Z18 indicates a role in maize zealexin biosynthesis. <i>Phytochemistry</i> , 2016, 121, 4-10.	1.4	43
67	Biosynthesis of the Diterpenoid Lycosantalanol via Nerylneryl Diphosphate in <i>Solanum lycopersicum</i> . <i>PLoS ONE</i> , 2015, 10, e0119302.	1.1	42
68	Investigation of the Chemical Interface in the Soybean-Aphid and Rice-Bacteria Interactions Using MALDI-Mass Spectrometry Imaging. <i>Analytical Chemistry</i> , 2015, 87, 5294-5301.	3.2	61
69	The Application of Synthetic Biology to Elucidation of Plant Mono-, Sesqui-, and Diterpenoid Metabolism. <i>Molecular Plant</i> , 2015, 8, 6-16.	3.9	75
70	Investigation of terpene diversification across multiple sequenced plant genomes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E81-8.	3.3	226
71	Optimization of recombinant expression enables discovery of novel cytochrome P450 activity in rice diterpenoid biosynthesis. <i>Applied Microbiology and Biotechnology</i> , 2015, 99, 7549-7558.	1.7	35
72	Efficient heterocyclisation by (di)terpene synthases. <i>Chemical Communications</i> , 2015, 51, 13485-13487.	2.2	33

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73	Functional divergence of diterpene syntheses in the medicinal plant <i>Salvia miltiorrhiza</i> Bunge. <i>Plant Physiology</i> , 2015, 169, pp.00695.2015.	2.3	118
74	Full-length transcriptome sequences and splice variants obtained by a combination of sequencing platforms applied to different root tissues of <i>Salvia miltiorrhiza</i> and tanshinone biosynthesis. <i>Plant Journal</i> , 2015, 82, 951-961.	2.8	337
75	An ent-kaurene-derived diterpenoid virulence factor from <i>Xanthomonas oryzae</i> pv. <i>oryzicola</i> . <i>New Phytologist</i> , 2015, 206, 295-302.	3.5	28
76	Biosynthesis of Lycosantalanol, a cis-Prenyl Derived Diterpenoid. <i>Journal of the American Chemical Society</i> , 2014, 136, 16951-16953.	6.6	41
77	Combining metabolomics and transcriptomics to characterize tanshinone biosynthesis in <i>Salvia miltiorrhiza</i> . <i>BMC Genomics</i> , 2014, 15, 73.	1.2	165
78	Biosynthesis, elicitation and roles of monocot terpenoid phytoalexins. <i>Plant Journal</i> , 2014, 79, 659-678.	2.8	233
79	Characterization of an Orphan Diterpenoid Biosynthetic Operon from <i>Salinispora arenicola</i> . <i>Journal of Natural Products</i> , 2014, 77, 2144-2147.	1.5	27
80	Functional Conservation of the Capacity for ent-Kaurene Biosynthesis and an Associated Operon in Certain Rhizobia. <i>Journal of Bacteriology</i> , 2014, 196, 100-106.	1.0	47
81	Novel Product Chemistry from Mechanistic Analysis of ent-Copalyl Diphosphate Synthases from Plant Hormone Biosynthesis. <i>Angewandte Chemie - International Edition</i> , 2014, 53, 7198-7202.	7.2	64
82	To Gibberellins and Beyond! Surveying the Evolution of (Di)Terpenoid Metabolism. <i>Annual Review of Plant Biology</i> , 2014, 65, 259-286.	8.6	228
83	1.55Å-resolution structure of ent-copalyl diphosphate synthase and exploration of general acid function by site-directed mutagenesis. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2014, 1840, 184-190.	1.1	52
84	Biochemical characterization of the castor bean ent-kaurene synthase(-like) family supports quantum chemical view of diterpene cyclization. <i>Phytochemistry</i> , 2014, 103, 13-21.	1.4	24
85	CYP76AH1 catalyzes turnover of miltiradiene in tanshinones biosynthesis and enables heterologous production of ferruginol in yeasts. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 12108-12113.	3.3	326
86	Characterization of CYP76AH4 clarifies phenolic diterpenoid biosynthesis in the Lamiaceae. <i>Organic and Biomolecular Chemistry</i> , 2013, 11, 7650.	1.5	94
87	The Role of Momilactones in Rice Allelopathy. <i>Journal of Chemical Ecology</i> , 2013, 39, 175-185.	0.9	112
88	Picking sides: distinct roles for CYP76M6 and CYP76M8 in rice oryzalexin biosynthesis. <i>Biochemical Journal</i> , 2013, 454, 209-216.	1.7	48
89	CYP701A8: A Rice ent-Kaurene Oxidase Paralog Diverted to More Specialized Diterpenoid Metabolism. <i>Plant Physiology</i> , 2012, 158, 1418-1425.	2.3	109
90	Characterization of CYP76M5 indicates Metabolic Plasticity within a Plant Biosynthetic Gene Cluster. <i>Journal of Biological Chemistry</i> , 2012, 287, 6159-6168.	1.6	116

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91	Insights into Diterpene Cyclization from Structure of Bifunctional Abietadiene Synthase from <i>Abies grandis</i> . <i>Journal of Biological Chemistry</i> , 2012, 287, 6840-6850.	1.6	91
92	A Single Residue Change Leads to a Hydroxylated Product from the Class II Diterpene Cyclization Catalyzed by Abietadiene Synthase. <i>Organic Letters</i> , 2012, 14, 5828-5831.	2.4	52
93	Isotuberculosinol: the unusual case of an immunomodulatory diterpenoid from <i>Mycobacterium tuberculosis</i> . <i>MedChemComm</i> , 2012, 3, 899.	3.5	15
94	Functional characterization of wheat ent-kaurene(-like) synthases indicates continuing evolution of labdane-related diterpenoid metabolism in the cereals. <i>Phytochemistry</i> , 2012, 84, 47-55.	1.4	60
95	Functional characterization of wheat copalyl diphosphate synthases sheds light on the early evolution of labdane-related diterpenoid metabolism in the cereals. <i>Phytochemistry</i> , 2012, 84, 40-46.	1.4	65
96	Effect of Isotopically Sensitive Branching on Product Distribution for Pentalenene Synthase: Support for a Mechanism Predicted by Quantum Chemistry. <i>Journal of the American Chemical Society</i> , 2012, 134, 11369-11371.	6.6	82
97	Terpenoid synthase structures: a so far incomplete view of complex catalysis. <i>Natural Product Reports</i> , 2012, 29, 1153.	5.2	311
98	Functional characterization and evolution of the isotuberculosinol operon in <i>Mycobacterium tuberculosis</i> and related <i>Mycobacteria</i> . <i>Frontiers in Microbiology</i> , 2012, 3, 368.	1.5	23
99	Genetic evidence for natural product-mediated plant-plant allelopathy in rice ( <i>Oryza sativa</i> ). <i>New Phytologist</i> , 2012, 193, 570-575.	3.5	146
100	Gibberellin Phytohormone Metabolism. , 2012, , 233-249.		3
101	Isotuberculosinol: An immunomodulatory diterpenoid from <i>Mycobacterium tuberculosis</i> . <i>FASEB Journal</i> , 2012, 26, 800.2.	0.2	0
102	To Gibberellins and Beyond! Insights into the Evolution of Diterpenoid Metabolism. <i>FASEB Journal</i> , 2012, 26, 576.1.	0.2	0
103	Electrostatic effects on (di)terpene synthase product outcome. <i>Chemical Communications</i> , 2011, 47, 4074.	2.2	47
104	The <i>Selaginella</i> Genome Identifies Genetic Changes Associated with the Evolution of Vascular Plants. <i>Science</i> , 2011, 332, 960-963.	6.0	794
105	Evident and latent plasticity across the rice diterpene synthase family with potential implications for the evolution of diterpenoid metabolism in the cereals. <i>Biochemical Journal</i> , 2011, 435, 589-595.	1.7	46
106	CYP99A3: functional identification of a diterpene oxidase from the momilactone biosynthetic gene cluster in rice. <i>Plant Journal</i> , 2011, 65, 87-95.	2.8	102
107	Domain loss has independently occurred multiple times in plant terpene synthase evolution. <i>Plant Journal</i> , 2011, 68, 1051-1060.	2.8	64
108	Magnesium depletion triggers production of an immune modulating diterpenoid in <i>Mycobacterium tuberculosis</i> . <i>Molecular Microbiology</i> , 2011, 79, 1594-1601.	1.2	16

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109	Structure and mechanism of the diterpene cyclase ent-copalyl diphosphate synthase. <i>Nature Chemical Biology</i> , 2011, 7, 431-433.	3.9	166
110	Rv0989c encodes a novel ( <i>ent</i> )-geranyl diphosphate synthase facilitating decaprenyl diphosphate biosynthesis in <i>Mycobacterium tuberculosis</i> . <i>FEBS Letters</i> , 2011, 585, 549-554.	1.3	15
111	Parsing a multifunctional biosynthetic gene cluster from rice: Biochemical characterization of CYP71Z6 & 7. <i>FEBS Letters</i> , 2011, 585, 3446-3451.	1.3	70
112	Diterpenoid biopolymers: New directions for renewable materials engineering. <i>Biopolymers</i> , 2011, 95, 71-76.	1.2	12
113	A Novel Labdane-13-ene-15-ol Producing Bifunctional Diterpene Synthase from <i>Selaginella moellendorffii</i> . <i>ChemBioChem</i> , 2011, 12, 1984-1987.	1.3	43
114	Increasing diterpene yield with a modular metabolic engineering system in <i>E. coli</i> : comparison of MEV and MEP isoprenoid precursor pathway engineering. <i>Applied Microbiology and Biotechnology</i> , 2010, 85, 1893-1906.	1.7	183
115	Diterpene cyclases and the nature of the isoprene fold. <i>Proteins: Structure, Function and Bioinformatics</i> , 2010, 78, 2417-2432.	1.5	131
116	Characterization of the kaurene oxidase CYP701A3, a multifunctional cytochrome P450 from gibberellin biosynthesis. <i>Biochemical Journal</i> , 2010, 431, 337-347.	1.7	91
117	A Single Residue Switch for Mg <sup>2+</sup> -dependent Inhibition Characterizes Plant Class II Diterpene Cyclases from Primary and Secondary Metabolism. <i>Journal of Biological Chemistry</i> , 2010, 285, 20558-20563.	1.6	41
118	Synthesis of ( $\pm$ )-Nosyberkol (Isotuberculosinol, Revised Structure of Edaxadiene) and ( $\pm$ )-Tuberculosinol. <i>Organic Letters</i> , 2010, 12, 2626-2629.	2.4	33
119	Two rings in them all: The labdane-related diterpenoids. <i>Natural Product Reports</i> , 2010, 27, 1521.	5.2	354
120	Characterization and Inhibition of a Class II Diterpene Cyclase from <i>Mycobacterium tuberculosis</i> . <i>Journal of Biological Chemistry</i> , 2009, 284, 23574-23579.	1.6	35
121	CYP76M7 Is an <i>ent</i> -Cassadiene C11-Hydroxylase Defining a Second Multifunctional Diterpenoid Biosynthetic Gene Cluster in Rice. <i>Plant Cell</i> , 2009, 21, 3315-3325.	3.1	199
122	Gibberellin biosynthesis in bacteria: Separate <i>ent</i> -copalyl diphosphate and <i>ent</i> -kaurene synthases in <i>Bradyrhizobium japonicum</i> . <i>FEBS Letters</i> , 2009, 583, 475-480.	1.3	152
123	Investigating the conservation pattern of a putative second terpene synthase divalent metal binding motif in plants. <i>Phytochemistry</i> , 2009, 70, 366-369.	1.4	51
124	A Functional Genomics Approach to Tanshinone Biosynthesis Provides Stereochemical Insights. <i>Organic Letters</i> , 2009, 11, 5170-5173.	2.4	250
125	Edaxadiene: A New Bioactive Diterpene from <i>Mycobacterium tuberculosis</i> . <i>Journal of the American Chemical Society</i> , 2009, 131, 17526-17527.	6.6	55
126	Increasing Complexity of a Diterpene Synthase Reaction with a Single Residue Switch. <i>Journal of the American Chemical Society</i> , 2008, 130, 5400-5401.	6.6	69



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127	Following evolution's lead to a single residue switch for diterpene synthase product outcome. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 7397-7401.	3.3	150
128	Synergistic Substrate Inhibition of ent-Copalyl Diphosphate Synthase: A Potential Feed-Forward Inhibition Mechanism Limiting Gibberellin Metabolism. Plant Physiology, 2007, 144, 445-454.	2.3	66
129	A Modular Approach for Facile Biosynthesis of Labdane-Related Diterpenes. Journal of the American Chemical Society, 2007, 129, 6684-6685.	6.6	144
130	A Single Residue Switch Converts Abietadiene Synthase into a Pimaradiene Specific Cyclase. Journal of the American Chemical Society, 2007, 129, 15736-15737.	6.6	88
131	16-Aza-ent-beyerane and 16-Aza-ent-trachylobane:Â Potent Mechanism-Based Inhibitors of Recombinantent-Kaurene Synthase from Arabidopsis thaliana. Journal of the American Chemical Society, 2007, 129, 12453-12460.	6.6	83
132	Probing the Role of the DXDD Motif in Class II Diterpene Cyclases. ChemBioChem, 2007, 8, 869-874.	1.3	80
133	Functional characterization of the rice kaurene synthase-like gene family. Phytochemistry, 2007, 68, 312-326.	1.4	124
134	An unexpected diterpene cyclase from rice: Functional identification of a stemodene synthase. Archives of Biochemistry and Biophysics, 2006, 448, 133-140.	1.4	44
135	Uncovering the complex metabolic network underlying diterpenoid phytoalexin biosynthesis in rice and other cereal crop plants. Phytochemistry, 2006, 67, 2307-2317.	1.4	187
136	The Maize An2 Gene is Induced by Fusarium Attack and Encodes an ent-Copalyl Diphosphate Synthase. Plant Molecular Biology, 2005, 59, 881-894.	2.0	123
137	Identification of Syn-Pimara-7,15-Diene Synthase Reveals Functional Clustering of Terpene Synthases Involved in Rice Phytoalexin/Allelochemical Biosynthesis. Plant Physiology, 2004, 135, 2098-2105.	2.3	195
138	Functional identification of ricesyn-copalyl diphosphate synthase and its role in initiating biosynthesis of diterpenoid phytoalexin/allelopathic natural products. Plant Journal, 2004, 39, 309-318.	2.8	152
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