

# David P. Fewer

## List of Publications by Year in descending order

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91  
papers

6,924  
citations

71061

41  
h-index

62565

80  
g-index

97  
all docs

97  
docs citations

97  
times ranked

6942  
citing authors

#	ARTICLE	IF	CITATIONS
1	The Natural Products Atlas 2.0: a database of microbially-derived natural products. <i>Nucleic Acids Research</i> , 2022, 50, D1317-D1323.	6.5	112
2	Discovery of varlaxins, new aeruginosin-type inhibitors of human trypsins. <i>Organic and Biomolecular Chemistry</i> , 2022, 20, 2681-2692.	1.5	8
3	Fatty Acid Substitutions Modulate the Cytotoxicity of Puwainaphycins/Minutissamides Isolated from the Baltic Sea Cyanobacterium <i>Nodularia harveyana</i> UHCC-0300. <i>ACS Omega</i> , 2022, 7, 11818-11828.	1.6	2
4	Single cell mutant selection for metabolic engineering of actinomycetes. <i>Metabolic Engineering</i> , 2022, 73, 124-133.	3.6	7
5	Semi-synthetic puwainaphycin/minutesamide cyclic lipopeptides with improved antifungal activity and limited cytotoxicity. <i>RSC Advances</i> , 2021, 11, 30873-30886.	1.7	7
6	A community resource for paired genomic and metabolomic data mining. <i>Nature Chemical Biology</i> , 2021, 17, 363-368.	3.9	81
7	CyanoMetDB, a comprehensive public database of secondary metabolites from cyanobacteria. <i>Water Research</i> , 2021, 196, 117017.	5.3	142
8	Genome Reduction and Secondary Metabolism of the Marine Sponge-Associated Cyanobacterium <i>Leptothoe</i> . <i>Marine Drugs</i> , 2021, 19, 298.	2.2	4
9	Doing synthetic biology with photosynthetic microorganisms. <i>Physiologia Plantarum</i> , 2021, 173, 624-638.	2.6	20
10	Chemical diversity and cellular effects of antifungal cyclic lipopeptides from cyanobacteria. <i>Physiologia Plantarum</i> , 2021, 173, 639-650.	2.6	16
11	The structure and biosynthesis of heinamides A1-A3 and B1-B5, antifungal members of the laxaphycin lipopeptide family. <i>Organic and Biomolecular Chemistry</i> , 2021, 19, 5577-5588.	1.5	5
12	Potent Inhibitor of Human Trypsins from the Aeruginosin Family of Natural Products. <i>ACS Chemical Biology</i> , 2021, 16, 2537-2546.	1.6	11
13	Mining of Cyanobacterial Genomes Indicates Natural Product Biosynthetic Gene Clusters Located in Conjugative Plasmids. <i>Frontiers in Microbiology</i> , 2021, 12, 684565.	1.5	12
14	A pharmaceutical model for the molecular evolution of microbial natural products. <i>FEBS Journal</i> , 2020, 287, 1429-1449.	2.2	22
15	Dereplication of Natural Products with Antimicrobial and Anticancer Activity from Brazilian Cyanobacteria. <i>Toxins</i> , 2020, 12, 12.	1.5	27
16	Shared PKS Module in Biosynthesis of Synergistic Laxaphycins. <i>Frontiers in Microbiology</i> , 2020, 11, 578878.	1.5	14
17	Phylogenomic Analysis of Secondary Metabolism in the Toxic Cyanobacterial Genera <i>Anabaena</i> , <i>Dolichospermum</i> and <i>Aphanizomenon</i> . <i>Toxins</i> , 2020, 12, 248.	1.5	34
18	Biosynthesis of the Bis-Prenylated Alkaloids Muscoride A and B. <i>ACS Chemical Biology</i> , 2019, 14, 2683-2690.	1.6	32

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19	The Biosynthesis of Rare Homo-Amino Acid Containing Variants of Microcystin by a Benthic Cyanobacterium. <i>Marine Drugs</i> , 2019, 17, 271.	2.2	20
20	Antitumor astins originate from the fungal endophyte <i>Cyanodermella asteris</i> living within the medicinal plant <i>Aster tataricus</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 26909-26917.	3.3	39
21	Alternative Biosynthetic Starter Units Enhance the Structural Diversity of Cyanobacterial Lipopeptides. <i>Applied and Environmental Microbiology</i> , 2019, 85, .	1.4	24
22	Strains of the toxic and bloom-forming <i>Nodularia spumigena</i> (cyanobacteria) can degrade methylphosphonate and release methane. <i>ISME Journal</i> , 2018, 12, 1619-1630.	4.4	75
23	Discovery of a Pederin Family Compound in a Nonsymbiotic Bloom-Forming Cyanobacterium. <i>ACS Chemical Biology</i> , 2018, 13, 1123-1129.	1.6	27
24	The Swinholide Biosynthesis Gene Cluster from a Terrestrial Cyanobacterium, <i>Nostoc</i> sp. Strain UHCC 0450. <i>Applied and Environmental Microbiology</i> , 2018, 84, .	1.4	21
25	Sphaerocyclamide, a prenylated cyanobactin from the cyanobacterium <i>Sphaerospermopsis</i> sp. LEGE 00249. <i>Scientific Reports</i> , 2018, 8, 14537.	1.6	27
26	N-Prenylation of Tryptophan by an Aromatic Prenyltransferase from the Cyanobactin Biosynthetic Pathway. <i>Biochemistry</i> , 2018, 57, 6860-6867.	1.2	26
27	Genetic Organization of Anabaenopeptin and Spumigin Biosynthetic Gene Clusters in the Cyanobacterium <i>Sphaerospermopsis torques-reginae</i> ITEP-024. <i>ACS Chemical Biology</i> , 2017, 12, 769-778.	1.6	25
28	Phylogenomic Analysis of the Microviridin Biosynthetic Pathway Coupled with Targeted Chemo-Enzymatic Synthesis Yields Potent Protease Inhibitors. <i>ACS Chemical Biology</i> , 2017, 12, 1538-1546.	1.6	45
29	Rearranged Biosynthetic Gene Cluster and Synthesis of Hassallidin E in <i>Planktothrix sarta</i> PCC 8927. <i>ACS Chemical Biology</i> , 2017, 12, 1796-1804.	1.6	25
30	Cyclic peptide production using a macrocyclase with enhanced substrate promiscuity and relaxed recognition determinants. <i>Chemical Communications</i> , 2017, 53, 10656-10659.	2.2	19
31	Simultaneous Production of Anabaenopeptins and Namalides by the Cyanobacterium <i>Nostoc</i> sp. CENA543. <i>ACS Chemical Biology</i> , 2017, 12, 2746-2755.	1.6	35
32	Production of High Amounts of Hepatotoxin Nodularin and New Protease Inhibitors Pseudospumigins by the Brazilian Benthic <i>Nostoc</i> sp. CENA543. <i>Frontiers in Microbiology</i> , 2017, 8, 1963.	1.5	35
33	The cyclochlorotine mycotoxin is produced by the nonribosomal peptide synthetase CctN in <i>Talaromyces islandicus</i> (â€” <i>Penicillium islandicum</i> â€™). <i>Environmental Microbiology</i> , 2016, 18, 3728-3741.	1.8	15
34	A Unique Tryptophan Prenyltransferase from the Kawaguchipeptin Biosynthetic Pathway. <i>Angewandte Chemie - International Edition</i> , 2016, 55, 3596-3599.	7.2	49
35	A Unique Tryptophan Prenyltransferase from the Kawaguchipeptin Biosynthetic Pathway. <i>Angewandte Chemie</i> , 2016, 128, 3660-3663.	1.6	6
36	A liquid chromatographyâ€”mass spectrometric method for the detection of cyclic Î²-amino fatty acid lipopeptides. <i>Journal of Chromatography A</i> , 2016, 1438, 76-83.	1.8	13

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37	Antifungal Compounds from Cyanobacteria. <i>Marine Drugs</i> , 2015, 13, 2124-2140.	2.2	83
38	Natural Product Biosynthetic Diversity and Comparative Genomics of the Cyanobacteria. <i>Trends in Microbiology</i> , 2015, 23, 642-652.	3.5	266
39	Antifungal activity improved by coproduction of cyclodextrins and anabaenolysins in Cyanobacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 13669-13674.	3.3	27
40	Minimum Information about a Biosynthetic Gene cluster. <i>Nature Chemical Biology</i> , 2015, 11, 625-631.	3.9	715
41	Draft genome sequence of <i>Talaromyces islandicus</i> (â€œ <i>Penicillium islandicum</i> â€) WF-38-12, a neglected mold with significant biotechnological potential. <i>Journal of Biotechnology</i> , 2015, 211, 101-102.	1.9	17
42	Genomic insights into the distribution, genetic diversity and evolution of polyketide synthases and nonribosomal peptide synthetases. <i>Current Opinion in Genetics and Development</i> , 2015, 35, 79-85.	1.5	33
43	Pseudoaeruginosins, Nonribosomal Peptides in <i>Nodularia spumigena</i> . <i>ACS Chemical Biology</i> , 2015, 10, 725-733.	1.6	22
44	Identification of geosmin and 2-methylisoborneol in cyanobacteria and molecular detection methods for the producers of these compounds. <i>Water Research</i> , 2015, 68, 56-66.	5.3	114
45	Phylum-wide comparative genomics unravel the diversity of secondary metabolism in Cyanobacteria. <i>BMC Genomics</i> , 2014, 15, 977.	1.2	175
46	4-Methylproline Guided Natural Product Discovery: Co-Occurrence of 4-Hydroxy- and 4-Methylprolines in Nostoweipeptins and Nostopeptolides. <i>ACS Chemical Biology</i> , 2014, 9, 2646-2655.	1.6	28
47	Reply to Sasso et al.: Distribution and phylogeny of nonribosomal peptide and polyketide biosynthetic pathways in eukaryotes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E3947-E3947.	3.3	2
48	Hassallidins, antifungal glycolipopeptides, are widespread among cyanobacteria and are the end-product of a nonribosomal pathway. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E1909-17.	3.3	102
49	Nostosins, Trypsin Inhibitors Isolated from the Terrestrial Cyanobacterium <i>Nostoc</i> sp. Strain FSN. <i>Journal of Natural Products</i> , 2014, 77, 1784-1790.	1.5	41
50	Atlas of nonribosomal peptide and polyketide biosynthetic pathways reveals common occurrence of nonmodular enzymes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 9259-9264.	3.3	310
51	The Genetic Basis for O-Acetylation of the Microcystin Toxin in Cyanobacteria. <i>Chemistry and Biology</i> , 2013, 20, 861-869.	6.2	20
52	Genome Mining Expands the Chemical Diversity of the Cyanobactin Family to Include Highly Modified Linear Peptides. <i>Chemistry and Biology</i> , 2013, 20, 1033-1043.	6.2	90
53	Cyanobacterial toxins: biosynthetic routes and evolutionary roots. <i>FEMS Microbiology Reviews</i> , 2013, 37, 23-43.	3.9	282
54	Lichen species identity and diversity of cyanobacterial toxins in symbiosis. <i>New Phytologist</i> , 2013, 198, 647-651.	3.5	22

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55	Convergent evolution of [D-Leucine1] microcystin-LR in taxonomically disparate cyanobacteria. <i>BMC Evolutionary Biology</i> , 2013, 13, 86.	3.2	29
56	Improving the coverage of the cyanobacterial phylum using diversity-driven genome sequencing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 1053-1058.	3.3	769
57	Insights into the Physiology and Ecology of the Brackish-Water-Adapted Cyanobacterium <i>Nodularia spumigena</i> CCY9414 Based on a Genome-Transcriptome Analysis. <i>PLoS ONE</i> , 2013, 8, e60224.	1.1	95
58	New Structural Variants of Aeruginosin Produced by the Toxic Bloom Forming Cyanobacterium <i>Nodularia spumigena</i> . <i>PLoS ONE</i> , 2013, 8, e73618.	1.1	65
59	Genome-derived insights into the biology of the hepatotoxic bloom-forming cyanobacterium <i>Anabaena</i> sp. strain 90. <i>BMC Genomics</i> , 2012, 13, 613.	1.2	52
60	Analysis of an Inactive Cyanobactin Biosynthetic Gene Cluster Leads to Discovery of New Natural Products from Strains of the Genus <i>Microcystis</i> . <i>PLoS ONE</i> , 2012, 7, e43002.	1.1	54
61	Cyanobacteria produce a high variety of hepatotoxic peptides in lichen symbiosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 5886-5891.	3.3	138
62	Comparison of wintertime eukaryotic community from sea ice and open water in the Baltic Sea, based on sequencing of the 18S rRNA gene. <i>Polar Biology</i> , 2012, 35, 875-889.	0.5	60
63	Cyanobacterial toxins: biosynthetic routes and evolutionary roots. <i>FEMS Microbiology Reviews</i> , 2012, , n/a-n/a.	3.9	2
64	<i>Deinobacterium chartae</i> gen. nov., sp. nov., an extremely radiation-resistant, biofilm-forming bacterium isolated from a Finnish paper mill. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 540-548.	0.8	16
65	Speciation in Red Algae: Members of the Ceramiales as Model Organisms. <i>Integrative and Comparative Biology</i> , 2011, 51, 492-504.	0.9	17
66	Non-autonomous transposable elements associated with inactivation of microcystin gene clusters in strains of the genus <i>Anabaena</i> isolated from the Baltic Sea. <i>Environmental Microbiology Reports</i> , 2011, 3, 189-194.	1.0	20
67	<i>Galega orientalis</i> is more diverse than <i>Galega officinalis</i> in Caucasus-whole-genome AFLP analysis and phylogenetics of symbiosis-related genes. <i>Molecular Ecology</i> , 2011, 20, 4808-4821.	2.0	18
68	Nostophycin Biosynthesis Is Directed by a Hybrid Polyketide Synthase-Nonribosomal Peptide Synthetase in the Toxic Cyanobacterium <i>Nostoc</i> sp. Strain 152. <i>Applied and Environmental Microbiology</i> , 2011, 77, 8034-8040.	1.4	29
69	Genome Mining Demonstrates the Widespread Occurrence of Gene Clusters Encoding Bacteriocins in Cyanobacteria. <i>PLoS ONE</i> , 2011, 6, e22384.	1.1	78
70	Cyanobactins—ribosomal cyclic peptides produced by cyanobacteria. <i>Applied Microbiology and Biotechnology</i> , 2010, 86, 1213-1225.	1.7	258
71	Molecular evidence for a diverse green algal community growing in the hair of sloths and a specific association with <i>Trichophilus welckeri</i> (Chlorophyta, Ulvophyceae). <i>BMC Evolutionary Biology</i> , 2010, 10, 86.	3.2	58
72	Screening for biohydrogen production by cyanobacteria isolated from the Baltic Sea and Finnish lakes. <i>International Journal of Hydrogen Energy</i> , 2010, 35, 1117-1127.	3.8	45

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73	Two Alternative Starter Modules for the Non-Ribosomal Biosynthesis of Specific Anabaenopeptin Variants in <i>Anabaena</i> (Cyanobacteria). <i>Chemistry and Biology</i> , 2010, 17, 265-273.	6.2	100
74	Highly Diverse Cyanobactins in Strains of the Genus <i>Anabaena</i> . <i>Applied and Environmental Microbiology</i> , 2010, 76, 701-709.	1.4	73
75	Widespread Occurrence and Lateral Transfer of the Cyanobactin Biosynthesis Gene Cluster in Cyanobacteria. <i>Applied and Environmental Microbiology</i> , 2009, 75, 853-857.	1.4	57
76	The non-ribosomal assembly and frequent occurrence of the protease inhibitors spumigins in the bloom-forming cyanobacterium <i>Nodularia spumigena</i> . <i>Molecular Microbiology</i> , 2009, 73, 924-937.	1.2	63
77	Horizontal gene transfer and recombination shape mesorhizobial populations in the gene center of the host plants <i>Astragalus luteolus</i> and <i>Astragalus ernestii</i> in Sichuan, China. <i>FEMS Microbiology Ecology</i> , 2009, 70, 227-235.	1.3	18
78	Culture-independent evidence for the persistent presence and genetic diversity of microcystin-producing <i>Anabaena</i> (Cyanobacteria) in the Gulf of Finland. <i>Environmental Microbiology</i> , 2009, 11, 855-866.	1.8	64
79	Microcystin Production in the Tripartite Cyanolichen <i>Peltigera leucophlebia</i> . <i>Molecular Plant-Microbe Interactions</i> , 2009, 22, 695-702.	1.4	43
80	Genetic diversity in strains of the genus <i>Anabaena</i> isolated from planktonic and benthic habitats of the Gulf of Finland (Baltic Sea). <i>FEMS Microbiology Ecology</i> , 2008, 64, 199-208.	1.3	38
81	Evidence for positive selection acting on microcystin synthetase adenylation domains in three cyanobacterial genera. <i>BMC Evolutionary Biology</i> , 2008, 8, 256.	3.2	46
82	Natural occurrence of microcystin synthetase deletion mutants capable of producing microcystins in strains of the genus <i>Anabaena</i> (Cyanobacteria). <i>Microbiology (United Kingdom)</i> , 2008, 154, 1007-1014.	0.7	36
83	The Diversity and Evolution of Rhizobia. <i>Microbiology Monographs</i> , 2007, , 3-41.	0.3	16
84	Direct Evidence for Production of Microcystins by <i>Anabaena</i> Strains from the Baltic Sea. <i>Applied and Environmental Microbiology</i> , 2007, 73, 6543-6550.	1.4	86
85	Strains of the cyanobacterial genera <i>Calothrix</i> and <i>Rivularia</i> isolated from the Baltic Sea display cryptic diversity and are distantly related to <i>Gloeotrichia</i> and <i>Tolypothrix</i> . <i>FEMS Microbiology Ecology</i> , 2007, 61, 74-84.	1.3	60
86	Recurrent adenylation domain replacement in the microcystin synthetase gene cluster. <i>BMC Evolutionary Biology</i> , 2007, 7, 183.	3.2	97
87	Discovery of Rare and Highly Toxic Microcystins from Lichen-Associated Cyanobacterium <i>Nostoc</i> sp. Strain IO-102-I. <i>Applied and Environmental Microbiology</i> , 2004, 70, 5756-5763.	1.4	131
88	Phylogenetic evidence for the early evolution of microcystin synthesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 568-573.	3.3	432
89	Phylogeny and Self-Splicing Ability of the Plastid tRNA-Leu Group I Intron. <i>Journal of Molecular Evolution</i> , 2003, 57, 710-720.	0.8	48
90	Novel morphology in <i>Enteromorpha</i> (Ulvophyceae) forming green tides. <i>American Journal of Botany</i> , 2002, 89, 1756-1763.	0.8	167

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91	Chroococciopsis and Heterocyst-Differentiating Cyanobacteria Are Each Other's Closest Living Relatives. <i>Molecular Phylogenetics and Evolution</i> , 2002, 23, 82-90.	1.2	100