David P. Fewer

List of Publications by Year in descending order

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70961 62479 6,924 91 41 80 citations h-index g-index papers 97 97 97 6942 docs citations times ranked citing authors all docs

| # | Article | IF | CITATIONS |
|----|--|-------------|-----------|
| 1 | The Natural Products Atlas 2.0: a database of microbially-derived natural products. Nucleic Acids Research, 2022, 50, D1317-D1323. | 6.5 | 112 |
| 2 | Discovery of varlaxins, new aeruginosin-type inhibitors of human trypsins. Organic and Biomolecular Chemistry, 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. ACS Omega, 2022, 7, 11818-11828. | 1.6 | 2 |
| 4 | Single cell mutant selection for metabolic engineering of actinomycetes. Metabolic Engineering, 2022, 73, 124-133. | 3.6 | 7 |
| 5 | Semi-synthetic puwainaphycin/minutissamide cyclic lipopeptides with improved antifungal activity and limited cytotoxicity. RSC Advances, 2021, 11, 30873-30886. | 1.7 | 7 |
| 6 | A community resource for paired genomic and metabolomic data mining. Nature Chemical Biology, 2021, 17, 363-368. | 3.9 | 81 |
| 7 | CyanoMetDB, a comprehensive public database of secondary metabolites from cyanobacteria. Water Research, 2021, 196, 117017. | 5. 3 | 142 |
| 8 | Genome Reduction and Secondary Metabolism of the Marine Sponge-Associated Cyanobacterium Leptothoe. Marine Drugs, 2021, 19, 298. | 2.2 | 4 |
| 9 | Doing synthetic biology with photosynthetic microorganisms. Physiologia Plantarum, 2021, 173, 624-638. | 2.6 | 20 |
| 10 | Chemical diversity and cellular effects of antifungal cyclic lipopeptides from cyanobacteria. Physiologia Plantarum, 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. Organic and Biomolecular Chemistry, 2021, 19, 5577-5588. | 1.5 | 5 |
| 12 | Potent Inhibitor of Human Trypsins from the Aeruginosin Family of Natural Products. ACS Chemical Biology, 2021, 16, 2537-2546. | 1.6 | 11 |
| 13 | Mining of Cyanobacterial Genomes Indicates Natural Product Biosynthetic Gene Clusters Located in Conjugative Plasmids. Frontiers in Microbiology, 2021, 12, 684565. | 1.5 | 12 |
| 14 | A pharmaceutical model for the molecular evolution of microbial natural products. FEBS Journal, 2020, 287, 1429-1449. | 2.2 | 22 |
| 15 | Dereplication of Natural Products with Antimicrobial and Anticancer Activity from Brazilian Cyanobacteria. Toxins, 2020, 12, 12. | 1.5 | 27 |
| 16 | Shared PKS Module in Biosynthesis of Synergistic Laxaphycins. Frontiers in Microbiology, 2020, 11, 578878. | 1.5 | 14 |
| 17 | Phylogenomic Analysis of Secondary Metabolism in the Toxic Cyanobacterial Genera Anabaena, Dolichospermum and Aphanizomenon. Toxins, 2020, 12, 248. | 1.5 | 34 |
| 18 | Biosynthesis of the Bis-Prenylated Alkaloids Muscoride A and B. ACS Chemical Biology, 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. Marine Drugs, 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> . Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 26909-26917. | 3.3 | 39 |
| 21 | Alternative Biosynthetic Starter Units Enhance the Structural Diversity of Cyanobacterial Lipopeptides. Applied and Environmental Microbiology, 2019, 85, . | 1.4 | 24 |
| 22 | Strains of the toxic and bloom-forming <i>Nodularia spumigena</i> (cyanobacteria) can degrade methylphosphonate and release methane. ISME Journal, 2018, 12, 1619-1630. | 4.4 | 75 |
| 23 | Discovery of a Pederin Family Compound in a Nonsymbiotic Bloom-Forming Cyanobacterium. ACS Chemical Biology, 2018, 13, 1123-1129. | 1.6 | 27 |
| 24 | The Swinholide Biosynthesis Gene Cluster from a Terrestrial Cyanobacterium, Nostoc sp. Strain UHCC 0450. Applied and Environmental Microbiology, 2018, 84, . | 1.4 | 21 |
| 25 | Sphaerocyclamide, a prenylated cyanobactin from the cyanobacterium Sphaerospermopsis sp. LEGE 00249. Scientific Reports, 2018, 8, 14537. | 1.6 | 27 |
| 26 | N-Prenylation of Tryptophan by an Aromatic Prenyltransferase from the Cyanobactin Biosynthetic Pathway. Biochemistry, 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. ACS Chemical Biology, 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. ACS Chemical Biology, 2017, 12, 1538-1546. | 1.6 | 45 |
| 29 | Rearranged Biosynthetic Gene Cluster and Synthesis of Hassallidin E in <i>Planktothrix serta</i> PCC 8927. ACS Chemical Biology, 2017, 12, 1796-1804. | 1.6 | 25 |
| 30 | Cyclic peptide production using a macrocyclase with enhanced substrate promiscuity and relaxed recognition determinants. Chemical Communications, 2017, 53, 10656-10659. | 2.2 | 19 |
| 31 | Simultaneous Production of Anabaenopeptins and Namalides by the Cyanobacterium <i>Nostoc</i> sp. CENA543. ACS Chemical Biology, 2017, 12, 2746-2755. | 1.6 | 35 |
| 32 | Production of High Amounts of Hepatotoxin Nodularin and New Protease Inhibitors Pseudospumigins by the Brazilian Benthic Nostoc sp. CENA543. Frontiers in Microbiology, 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> '). Environmental Microbiology, 2016, 18, 3728-3741. | 1.8 | 15 |
| 34 | A Unique Tryptophan Câ€Prenyltransferase from the Kawaguchipeptin Biosynthetic Pathway. Angewandte Chemie - International Edition, 2016, 55, 3596-3599. | 7.2 | 49 |
| 35 | A Unique Tryptophan Câ€Prenyltransferase from the Kawaguchipeptin Biosynthetic Pathway. Angewandte Chemie, 2016, 128, 3660-3663. | 1.6 | 6 |
| 36 | A liquid chromatography–mass spectrometric method for the detection of cyclic β-amino fatty acid lipopeptides. Journal of Chromatography A, 2016, 1438, 76-83. | 1.8 | 13 |

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

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| 55 | Convergent evolution of [D-Leucine1] microcystin-LR in taxonomically disparate cyanobacteria. BMC Evolutionary Biology, 2013, 13, 86. | 3.2 | 29 |
| 56 | Improving the coverage of the cyanobacterial phylum using diversity-driven genome sequencing. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 1053-1058. | 3.3 | 769 |
| 57 | Insights into the Physiology and Ecology of the Brackish-Water-Adapted Cyanobacterium Nodularia spumigena CCY9414 Based on a Genome-Transcriptome Analysis. PLoS ONE, 2013, 8, e60224. | 1.1 | 95 |
| 58 | New Structural Variants of Aeruginosin Produced by the Toxic Bloom Forming Cyanobacterium Nodularia spumigena. PLoS ONE, 2013, 8, e73618. | 1.1 | 65 |
| 59 | Genome-derived insights into the biology of the hepatotoxic bloom-forming cyanobacterium Anabaena sp. strain 90. BMC Genomics, 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 Microcystis. PLoS ONE, 2012, 7, e43002. | 1.1 | 54 |
| 61 | Cyanobacteria produce a high variety of hepatotoxic peptides in lichen symbiosis. Proceedings of the National Academy of Sciences of the United States of America, 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. Polar Biology, 2012, 35, 875-889. | 0.5 | 60 |
| 63 | Cyanobacterial toxins: biosynthetic routes and evolutionary roots. FEMS Microbiology Reviews, 2012, , n/a-n/a. | 3.9 | 2 |
| 64 | Deinobacterium chartae gen. nov., sp. nov., an extremely radiation-resistant, biofilm-forming bacterium isolated from a Finnish paper mill. International Journal of Systematic and Evolutionary Microbiology, 2011, 61, 540-548. | 0.8 | 16 |
| 65 | Speciation in Red Algae: Members of the Ceramiales as Model Organisms. Integrative and Comparative Biology, 2011, 51, 492-504. | 0.9 | 17 |
| 66 | Nonâ€autonomous transposable elements associated with inactivation of microcystin gene clusters in strains of the genus <i>Anabaena</i> isolated from the Baltic Sea. Environmental Microbiology Reports, 2011, 3, 189-194. | 1.0 | 20 |
| 67 | Galega orientalis is more diverse than Galega officinalis in Caucasus-whole-genome AFLP analysis and phylogenetics of symbiosis-related genes. Molecular Ecology, 2011, 20, 4808-4821. | 2.0 | 18 |
| 68 | Nostophycin Biosynthesis Is Directed by a Hybrid Polyketide Synthase-Nonribosomal Peptide Synthetase in the Toxic Cyanobacterium Nostoc sp. Strain 152. Applied and Environmental Microbiology, 2011, 77, 8034-8040. | 1.4 | 29 |
| 69 | Genome Mining Demonstrates the Widespread Occurrence of Gene Clusters Encoding Bacteriocins in Cyanobacteria. PLoS ONE, 2011, 6, e22384. | 1.1 | 78 |
| 70 | Cyanobactinsâ€"ribosomal cyclic peptides produced by cyanobacteria. Applied Microbiology and Biotechnology, 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 Trichophilus welckeri(Chlorophyta, Ulvophyceae). BMC Evolutionary Biology, 2010, 10, 86. | 3.2 | 58 |
| 72 | Screening for biohydrogen production by cyanobacteria isolated from the Baltic Sea and Finnish lakes. International Journal of Hydrogen Energy, 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 Anabaena (Cyanobacteria). Chemistry and Biology, 2010, 17, 265-273. | 6.2 | 100 |
| 74 | Highly Diverse Cyanobactins in Strains of the Genus <i>Anabaena</i> . Applied and Environmental Microbiology, 2010, 76, 701-709. | 1.4 | 73 |
| 75 | Widespread Occurrence and Lateral Transfer of the Cyanobactin Biosynthesis Gene Cluster in Cyanobacteria. Applied and Environmental Microbiology, 2009, 75, 853-857. | 1.4 | 57 |
| 76 | The nonâ€ribosomal assembly and frequent occurrence of the protease inhibitors spumigins in the bloomâ€forming cyanobacterium <i>Nodularia spumigena</i> . Molecular Microbiology, 2009, 73, 924-937. | 1.2 | 63 |
| 77 | Horizontal gene transfer and recombination shape mesorhizobial populations in the gene center of the host plantsAstragalus luteolusandAstragalus ernestiiin Sichuan, China. FEMS Microbiology Ecology, 2009, 70, 227-235. | 1.3 | 18 |
| 78 | Cultureâ€independent evidence for the persistent presence and genetic diversity of microcystinâ€producing <i>Anabaena</i> (<i>Cyanobacteria</i>) in the Gulf of Finland. Environmental Microbiology, 2009, 11, 855-866. | 1.8 | 64 |
| 79 | Microcystin Production in the Tripartite Cyanolichen <i>Peltigera leucophlebia</i> Plant-Microbe Interactions, 2009, 22, 695-702. | 1.4 | 43 |
| 80 | Genetic diversity in strains of the genus Anabaena isolated from planktonic and benthic habitats of the Gulf of Finland (Baltic Sea). FEMS Microbiology Ecology, 2008, 64, 199-208. | 1.3 | 38 |
| 81 | Evidence for positive selection acting on microcystin synthetase adenylation domains in three cyanobacterial genera. BMC Evolutionary Biology, 2008, 8, 256. | 3.2 | 46 |
| 82 | Natural occurrence of microcystin synthetase deletion mutants capable of producing microcystins in strains of the genus Anabaena (Cyanobacteria). Microbiology (United Kingdom), 2008, 154, 1007-1014. | 0.7 | 36 |
| 83 | The Diversity and Evolution of Rhizobia. Microbiology Monographs, 2007, , 3-41. | 0.3 | 16 |
| 84 | Direct Evidence for Production of Microcystins by <i>Anabaena</i> Strains from the Baltic Sea. Applied and Environmental Microbiology, 2007, 73, 6543-6550. | 1.4 | 86 |
| 85 | Strains of the cyanobacterial genera Calothrix and Rivularia isolated from the Baltic Sea display cryptic diversity and are distantly related to Gloeotrichia and Tolypothrix. FEMS Microbiology Ecology, 2007, 61, 74-84. | 1.3 | 60 |
| 86 | Recurrent adenylation domain replacement in the microcystin synthetase gene cluster. BMC Evolutionary Biology, 2007, 7, 183. | 3.2 | 97 |
| 87 | Discovery of Rare and Highly Toxic Microcystins from Lichen-Associated Cyanobacterium Nostoc sp. Strain IO-102-I. Applied and Environmental Microbiology, 2004, 70, 5756-5763. | 1.4 | 131 |
| 88 | Phylogenetic evidence for the early evolution of microcystin synthesis. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 568-573. | 3.3 | 432 |
| 89 | Phylogeny and Self-Splicing Ability of the Plastid tRNA-Leu Group I Intron. Journal of Molecular Evolution, 2003, 57, 710-720. | 0.8 | 48 |
| 90 | Novel morphology in <i>Enteromorpha </i> (Ulvophyceae) forming green tides. American Journal of Botany, 2002, 89, 1756-1763. | 0.8 | 167 |

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| 91 | Chroococcidiopsis and Heterocyst-Differentiating Cyanobacteria Are Each Other's Closest Living Relatives. Molecular Phylogenetics and Evolution, 2002, 23, 82-90. | 1.2 | 100 |