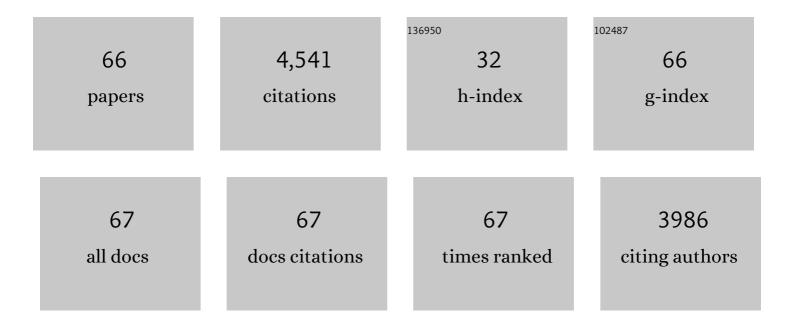
Clay C C Wang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Chromatin-level regulation of biosynthetic gene clusters. Nature Chemical Biology, 2009, 5, 462-464.	8.0	358
2	A Gene Cluster Containing Two Fungal Polyketide Synthases Encodes the Biosynthetic Pathway for a Polyketide, Asperfuranone, in <i>Aspergillus nidulans</i> . Journal of the American Chemical Society, 2009, 131, 2965-2970.	13.7	292
3	Advances in Aspergillus secondary metabolite research in the post-genomic era. Natural Product Reports, 2012, 29, 351.	10.3	233
4	Two Separate Gene Clusters Encode the Biosynthetic Pathway for the Meroterpenoids Austinol and Dehydroaustinol in <i>Aspergillus nidulans</i> . Journal of the American Chemical Society, 2012, 134, 4709-4720.	13.7	223
5	Molecular Genetic Mining of the Aspergillus Secondary Metabolome: Discovery of the Emericellamide Biosynthetic Pathway. Chemistry and Biology, 2008, 15, 527-532.	6.0	193
6	Plumbagin induces cell cycle arrest and apoptosis through reactive oxygen species/c-Jun N-terminal kinase pathways in human melanoma A375.S2 cells. Cancer Letters, 2008, 259, 82-98.	7.2	189
7	Recent advances in awakening silent biosynthetic gene clusters and linking orphan clusters to natural products in microorganisms. Current Opinion in Chemical Biology, 2011, 15, 137-143.	6.1	181
8	An Efficient System for Heterologous Expression of Secondary Metabolite Genes in Aspergillus nidulans. Journal of the American Chemical Society, 2013, 135, 7720-7731.	13.7	180
9	Illuminating the Diversity of Aromatic Polyketide Synthases in <i>Aspergillus nidulans</i> . Journal of the American Chemical Society, 2012, 134, 8212-8221.	13.7	168
10	Characterization of the <i>Aspergillus nidulans</i> Monodictyphenone Gene Cluster. Applied and Environmental Microbiology, 2010, 76, 2067-2074.	3.1	159
11	Genome-Based Deletion Analysis Reveals the Prenyl Xanthone Biosynthesis Pathway in <i>Aspergillus nidulans</i> . Journal of the American Chemical Society, 2011, 133, 4010-4017.	13.7	154
12	Identification and Characterization of the Asperthecin Gene Cluster of <i>Aspergillus nidulans</i> . Applied and Environmental Microbiology, 2008, 74, 7607-7612.	3.1	149
13	Development of Genetic Dereplication Strains in <i>Aspergillus nidulans</i> Results in the Discovery of Aspercryptin. Angewandte Chemie - International Edition, 2016, 55, 1662-1665.	13.8	139
14	Toward Awakening Cryptic Secondary Metabolite Gene Clusters in Filamentous Fungi. Methods in Enzymology, 2012, 517, 303-324.	1.0	116
15	Overexpression of the <i><scp>A</scp>spergillus nidulans</i> histone 4 acetyltransferase <scp>EsaA</scp> increases activation of secondary metabolite production. Molecular Microbiology, 2012, 86, 314-330.	2.5	116
16	Recent advances in genome mining of secondary metabolite biosynthetic gene clusters and the development of heterologous expression systems in <i>Aspergillus nidulans</i> . Journal of Industrial Microbiology and Biotechnology, 2014, 41, 433-442.	3.0	115
17	Characterization of Aspergillus fumigatus Isolates from Air and Surfaces of the International Space Station. MSphere, 2016, 1, .	2.9	108
18	Resistance Gene-Guided Genome Mining: Serial Promoter Exchanges in <i>Aspergillus nidulans</i> Reveal the Biosynthetic Pathway for Fellutamide B, a Proteasome Inhibitor. ACS Chemical Biology, 2016, 11, 2275-2284.	3.4	105

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19	bZIP transcription factors affecting secondary metabolism, sexual development and stress responses in Aspergillus nidulans. Microbiology (United Kingdom), 2013, 159, 77-88.	1.8	89
20	Microbial metabolomics in open microscale platforms. Nature Communications, 2016, 7, 10610.	12.8	86
21	Recent advances in the genome mining of <i>Aspergillus</i> secondary metabolites (covering) Tj ETQq1 1 0.784	314 rgBT /	Overlock 10 T
22	Discovery of McrA, a master regulator of <i>Aspergillus</i> secondary metabolism. Molecular Microbiology, 2017, 103, 347-365.	2.5	73
23	Biosynthetic Pathway of the Reduced Polyketide Product Citreoviridin in <i>Aspergillus terreus</i> var. <i>aureus</i> Revealed by Heterologous Expression in <i>Aspergillus nidulans</i> . Organic Letters, 2016, 18, 1366-1369.	4.6	57
24	Recent advances in genome mining of secondary metabolites in Aspergillus terreus. Frontiers in Microbiology, 2014, 5, 717.	3.5	51
25	Identification and molecular genetic analysis of the cichorine gene cluster in Aspergillus nidulans. MedChemComm, 2012, 3, 997.	3.4	48
26	Methylobacterium ajmalii sp. nov., Isolated From the International Space Station. Frontiers in Microbiology, 2021, 12, 639396.	3.5	46
27	The fungal natural product azaphilone-9 binds to HuR and inhibits HuR-RNA interaction in vitro. PLoS ONE, 2017, 12, e0175471.	2.5	45
28	Azaphilones Inhibit Tau Aggregation and Dissolve Tau Aggregates <i>in Vitro</i> . ACS Chemical Neuroscience, 2015, 6, 751-760.	3.5	42
29	Characterization of Aspergillus niger Isolated from the International Space Station. MSystems, 2018, 3,	3.8	42
30	Advances in space microbiology. IScience, 2021, 24, 102395.	4.1	42
31	Reengineering an Azaphilone Biosynthesis Pathway in <i>Aspergillus nidulans</i> To Create Lipoxygenase Inhibitors. Organic Letters, 2012, 14, 972-975.	4.6	38
32	Inhibition of Tau Aggregation by Three Aspergillus nidulans Secondary Metabolites: 2,ï‰-Dihydroxyemodin, Asperthecin, and Asperbenzaldehyde. Planta Medica, 2014, 80, 77-85.	1.3	38
33	Proteomic and Metabolomic Characteristics of Extremophilic Fungi Under Simulated Mars Conditions. Frontiers in Microbiology, 2019, 10, 1013.	3.5	36
34	Hybrid Transcription Factor Engineering Activates the Silent Secondary Metabolite Gene Cluster for (+)-Asperlin in <i>Aspergillus nidulans</i> . ACS Chemical Biology, 2018, 13, 3193-3205.	3.4	35
35	Genome mining and molecular characterization of the biosynthetic gene cluster of a diterpenic meroterpenoid, 15-deoxyoxalicine B, in Penicillium canescens. Chemical Science, 2015, 6, 6537-6544.	7.4	33
36	International Space Station conditions alter genomics, proteomics, and metabolomics in Aspergillus nidulans. Applied Microbiology and Biotechnology, 2019, 103, 1363-1377.	3.6	32

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37	Contributions of Spore Secondary Metabolites to UV-C Protection and Virulence Vary in Different Aspergillus fumigatus Strains. MBio, 2020, 11, .	4.1	32
38	Spatial regulation of a common precursor from two distinct genes generates metabolite diversity. Chemical Science, 2015, 6, 5913-5921.	7.4	31
39	Telomere position effect is regulated by heterochromatin-associated proteins and NkuA in Aspergillus nidulans. Microbiology (United Kingdom), 2010, 156, 3522-3531.	1.8	29
40	Proteomic characterization of Aspergillus fumigatus isolated from air and surfaces of the International Space Station. Fungal Genetics and Biology, 2019, 124, 39-46.	2.1	28
41	Engineering Fungal Nonribosomal Peptide Synthetase-like Enzymes by Heterologous Expression and Domain Swapping. Organic Letters, 2016, 18, 6236-6239.	4.6	27
42	Characterization of the product of a nonribosomal peptide synthetase-like (NRPS-like) gene using the doxycycline dependent Tet-on system in Aspergillus terreus. Fungal Genetics and Biology, 2016, 89, 84-88.	2.1	24
43	Asperfuranone from <i>Aspergillus nidulans</i> Inhibits Proliferation of Human Non‣mall Cell Lung Cancer A549 Cells via Blocking Cell Cycle Progression and Inducing Apoptosis. Basic and Clinical Pharmacology and Toxicology, 2010, 107, 583-589.	2.5	22
44	Overexpression of a three-gene conidial pigment biosynthetic pathway in Aspergillus nidulans reveals the first NRPS known to acetylate tryptophan. Fungal Genetics and Biology, 2017, 101, 1-6.	2.1	21
45	Overexpression of an LaeA-like Methyltransferase Upregulates Secondary Metabolite Production in <i>Aspergillus nidulans</i> . ACS Chemical Biology, 2019, 14, 1643-1651.	3.4	21
46	Norsolorinic Acid from <i>Aspergillus nidulans</i> Inhibits the Proliferation of Human Breast Adenocarcinoma MCFâ€7 Cells via Fasâ€Mediated Pathway. Basic and Clinical Pharmacology and Toxicology, 2008, 102, 491-497.	2.5	18
47	Draft Genome Sequences of Several Fungal Strains Selected for Exposure to Microgravity at the International Space Station. Genome Announcements, 2017, 5, .	0.8	17
48	Identification and Validation of an Aspergillus nidulans Secondary Metabolite Derivative as an Inhibitor of the Musashi-RNA Interaction. Cancers, 2020, 12, 2221.	3.7	17
49	Draft Genome Sequences of Two Aspergillus fumigatus Strains, Isolated from the International Space Station. Genome Announcements, 2016, 4, .	0.8	16
50	Metabolomic Analysis of Aspergillus niger Isolated From the International Space Station Reveals Enhanced Production Levels of the Antioxidant Pyranonigrin A. Frontiers in Microbiology, 2020, 11, 931.	3.5	16
51	Matingâ€ŧype factorâ€specific regulation of the fumagillin/pseurotin secondary metabolite supercluster in <i>Aspergillus fumigatus</i> . Molecular Microbiology, 2018, 110, 1045-1065.	2.5	15
52	Natural products development under epigenetic modulation in fungi. Phytochemistry Reviews, 2020, 19, 1323-1340.	6.5	14
53	An <i>Aspergillus nidulans</i> Platform for the Complete Cluster Refactoring and Total Biosynthesis of Fungal Natural Products. ACS Synthetic Biology, 2021, 10, 173-182.	3.8	14
54	NORSOLORINIC ACID INHIBITS PROLIFERATION OF T24 HUMAN BLADDER CANCER CELLS BY ARRESTING THE CELL CYCLE AT THE G ₀ /G ₁ PHASE AND INDUCING A FAS/MEMBRANEâ€BOUND FAS LIGANDâ€MEDIATED APOPTOTIC PATHWAY. Clinical and Experimental Pharmacology and Physiology, 2008, 35, 1301-1308.	1.9	13

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55	Identification of the pigment and its role in UV resistance in Paecilomyces variotii, a Chernobyl isolate, using genetic manipulation strategies. Fungal Genetics and Biology, 2021, 152, 103567.	2.1	13
56	Discovery and Elucidation of the Biosynthesis of Aspernidgulenes: Novel Polyenes from <i>Aspergillus Nidulans</i> by Using Serial Promoter Replacement. ChemBioChem, 2019, 20, 329-334.	2.6	12
57	Development of Genetic Dereplication Strains in <i>Aspergillus nidulans</i> Results in the Discovery of Aspercryptin. Angewandte Chemie, 2016, 128, 1694-1697.	2.0	8

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59	Expanding the Chemical Space of Nonribosomal Peptide Synthetase-like Enzymes by Domain and Tailoring Enzyme Recombination. Organic Letters, 2018, 20, 5082-5085.	4.6	7
60	The International Space Station Environment Triggers Molecular Responses in Aspergillus niger. Frontiers in Microbiology, 0, 13, .	3.5	7
61	Epigenetic Manipulation Induces the Production of Coumarinâ€Type Secondary Metabolite from <i>Arthrobotrys foliicola</i> . Israel Journal of Chemistry, 2019, 59, 432-438.	2.3	6
62	Genome-based deletion analysis in Aspergillus terreus reveals the acetylaranotin bis-thiomethyltransferase gene. Fungal Genetics and Biology, 2018, 119, 1-6.	2.1	5
63	Genomic Characterization of the Titan-like Cell Producing Naganishia tulchinskyi, the First Novel Eukaryote Isolated from the International Space Station. Journal of Fungi (Basel, Switzerland), 2022, 8, 165.	3.5	5
64	Prevention of chronic HBV infection induced hepatocellular carcinoma development by using antiplatelet drugs. Hepatobiliary Surgery and Nutrition, 2012, 1, 57-8.	1.5	3
65	The sexual spore pigment asperthecin is required for normal ascospore production and protection from UV light in <i>Aspergillus nidulans</i> . Journal of Industrial Microbiology and Biotechnology, 2021, 48, .	3.0	2
66	Characterization of a silent azaphilone biosynthesis gene cluster in Aspergillus terreus NIH 2624.		