Clay C C Wang

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63
papers

3,398
citations

4,153
ext. papers

28
h-index

58
g-index

5.23
L-index

#	Paper	IF	Citations
63	Chromatin-level regulation of biosynthetic gene clusters. <i>Nature Chemical Biology</i> , 2009 , 5, 462-4	11.7	292
62	A gene cluster containing two fungal polyketide synthases encodes the biosynthetic pathway for a polyketide, asperfuranone, in Aspergillus nidulans. <i>Journal of the American Chemical Society</i> , 2009 , 131, 2965-70	16.4	226
61	Advances in Aspergillus secondary metabolite research in the post-genomic era. <i>Natural Product Reports</i> , 2012 , 29, 351-71	15.1	205
60	Two separate gene clusters encode the biosynthetic pathway for the meroterpenoids austinol and dehydroaustinol in Aspergillus nidulans. <i>Journal of the American Chemical Society</i> , 2012 , 134, 4709-20	16.4	188
59	Plumbagin induces cell cycle arrest and apoptosis through reactive oxygen species/c-Jun N-terminal kinase pathways in human melanoma A375.S2 cells. <i>Cancer Letters</i> , 2008 , 259, 82-98	9.9	174
58	Molecular genetic mining of the Aspergillus secondary metabolome: discovery of the emericellamide biosynthetic pathway. <i>Chemistry and Biology</i> , 2008 , 15, 527-32		161
57	Recent advances in awakening silent biosynthetic gene clusters and linking orphan clusters to natural products in microorganisms. <i>Current Opinion in Chemical Biology</i> , 2011 , 15, 137-43	9.7	160
56	An efficient system for heterologous expression of secondary metabolite genes in Aspergillus nidulans. <i>Journal of the American Chemical Society</i> , 2013 , 135, 7720-31	16.4	146
55	Genome-based deletion analysis reveals the prenyl xanthone biosynthesis pathway in Aspergillus nidulans. <i>Journal of the American Chemical Society</i> , 2011 , 133, 4010-7	16.4	134
54	Illuminating the diversity of aromatic polyketide synthases in Aspergillus nidulans. <i>Journal of the American Chemical Society</i> , 2012 , 134, 8212-21	16.4	131
53	Identification and characterization of the asperthecin gene cluster of Aspergillus nidulans. <i>Applied and Environmental Microbiology</i> , 2008 , 74, 7607-12	4.8	126
52	Characterization of the Aspergillus nidulans monodictyphenone gene cluster. <i>Applied and Environmental Microbiology</i> , 2010 , 76, 2067-74	4.8	124
51	Development of Genetic Dereplication Strains in Aspergillus nidulans Results in the Discovery of Aspercryptin. <i>Angewandte Chemie - International Edition</i> , 2016 , 55, 1662-5	16.4	87
50	Toward awakening cryptic secondary metabolite gene clusters in filamentous fungi. <i>Methods in Enzymology</i> , 2012 , 517, 303-24	1.7	86
49	Recent advances in genome mining of secondary metabolite biosynthetic gene clusters and the development of heterologous expression systems in Aspergillus nidulans. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2014 , 41, 433-42	4.2	84
48	Overexpression of the Aspergillus nidulans histone 4 acetyltransferase EsaA increases activation of secondary metabolite production. <i>Molecular Microbiology</i> , 2012 , 86, 314-30	4.1	82
47	Resistance Gene-Guided Genome Mining: Serial Promoter Exchanges in Aspergillus nidulans Reveal the Biosynthetic Pathway for Fellutamide B, a Proteasome Inhibitor. <i>ACS Chemical Biology</i> , 2016 , 11, 2275-84	4.9	<i>75</i>

(2015-2013)

bZIP transcription factors affecting secondary metabolism, sexual development and stress responses in Aspergillus nidulans. <i>Microbiology (United Kingdom)</i> , 2013 , 159, 77-88	2.9	74
Microbial metabolomics in open microscale platforms. <i>Nature Communications</i> , 2016 , 7, 10610	17.4	67
Characterization of Isolates from Air and Surfaces of the International Space Station. <i>MSphere</i> , 2016 , 1,	5	61
Discovery of McrA, a master regulator of Aspergillus secondary metabolism. <i>Molecular Microbiology</i> , 2017 , 103, 347-365	4.1	45
Recent advances in the genome mining of secondary metabolites (covering 2012-2018). MedChemComm, 2019, 10, 840-866	5	40
Biosynthetic Pathway of the Reduced Polyketide Product Citreoviridin in Aspergillus terreus var. aureus Revealed by Heterologous Expression in Aspergillus nidulans. <i>Organic Letters</i> , 2016 , 18, 1366-9	6.2	39
Recent advances in genome mining of secondary metabolites in Aspergillus terreus. <i>Frontiers in Microbiology</i> , 2014 , 5, 717	5.7	37
Azaphilones inhibit tau aggregation and dissolve tau aggregates in vitro. <i>ACS Chemical Neuroscience</i> , 2015 , 6, 751-60	5.7	35
Reengineering an azaphilone biosynthesis pathway in Aspergillus nidulans to create lipoxygenase inhibitors. <i>Organic Letters</i> , 2012 , 14, 972-5	6.2	32
The fungal natural product azaphilone-9 binds to HuR and inhibits HuR-RNA interaction in vitro. <i>PLoS ONE</i> , 2017 , 12, e0175471	3.7	28
Inhibition of Tau aggregation by three Aspergillus nidulans secondary metabolites: 2,Edihydroxyemodin, asperthecin, and asperbenzaldehyde. <i>Planta Medica</i> , 2014 , 80, 77-85	3.1	28
Characterization of Aspergillus niger Isolated from the International Space Station. <i>MSystems</i> , 2018 , 3,	7.6	27
Telomere position effect is regulated by heterochromatin-associated proteins and NkuA in Aspergillus nidulans. <i>Microbiology (United Kingdom)</i> , 2010 , 156, 3522-3531	2.9	26
Spatial regulation of a common precursor from two distinct genes generates metabolite diversity. <i>Chemical Science</i> , 2015 , 6, 5913-5921	9.4	23
Identification and molecular genetic analysis of the cichorine gene cluster in. <i>MedChemComm</i> , 2012 , 3,	5	23
Hybrid Transcription Factor Engineering Activates the Silent Secondary Metabolite Gene Cluster for (+)-Asperlin in Aspergillus nidulans. <i>ACS Chemical Biology</i> , 2018 , 13, 3193-3205	4.9	22
Characterization of the product of a nonribosomal peptide synthetase-like (NRPS-like) gene using the doxycycline dependent Tet-on system in Aspergillus terreus. <i>Fungal Genetics and Biology</i> , 2016 , 89, 84-88	3.9	21
Genome mining and molecular characterization of the biosynthetic gene cluster of a diterpenic meroterpenoid, 15-deoxyoxalicine B, in. <i>Chemical Science</i> , 2015 , 6, 6537-6544	9.4	19
	responses in Aspergillus nidulans. Microbiology (United Kingdom), 2013, 159, 77-88 Microbial metabolomics in open microscale platforms. Nature Communications, 2016, 7, 10610 Characterization of Isolates from Air and Surfaces of the International Space Station. MSphere, 2016, 1, Discovery of McrA, a master regulator of Aspergillus secondary metabolism. Molecular Microbiology, 2017, 103, 347-365 Recent advances in the genome mining of secondary metabolites (covering 2012-2018), MedChemComm, 2019, 10, 840-866 Biosynthetic Pathway of the Reduced Polyketide Product Citreoviridin in Aspergillus terreus var. aureus Revealed by Heterologous Expression in Aspergillus nidulans. Organic Letters, 2016, 18, 1366-9 Recent advances in genome mining of secondary metabolites in Aspergillus terreus. Frontiers in Microbiology, 2014, 5, 717 Azaphilones inhibit tau aggregation and dissolve tau aggregates in vitro. ACS Chemical Neuroscience, 2015, 6, 751-60 Reengineering an azaphilone biosynthesis pathway in Aspergillus nidulans to create lipoxygenase inhibitors. Organic Letters, 2012, 14, 972-5 The fungal natural product azaphilone-9 binds to HuR and inhibits HuR-RNA interaction in vitro. PLoS ONE, 2017, 12, e0175-471 Inhibition of Tau aggregation by three Aspergillus nidulans secondary metabolites: 2,Bilhydroxyemodin, asperthecin, and asperbenzaldehyde. Planta Medica, 2014, 80, 77-85 Characterization of Aspergillus niger Isolated from the International Space Station. MSystems, 2018, 3, 3, 1913-35921 Identification and molecular genetic analysis of the cichorine gene cluster in. MedChemComm, 2012, 3, 1914 (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1914) (1	responses in Aspergillus nidulans. Microbiology (United Kingdom), 2013, 159, 77-88 Microbial metabolomics in open microscale platforms. Nature Communications, 2016, 7, 10610 174 Characterization of Isolates from Air and Surfaces of the International Space Station. MSphere, 2016, 1, Discovery of McrA, a master regulator of Aspergillus secondary metabolism. Molecular Microbiology, 2017, 103, 347-365 Recent advances in the genome mining of secondary metabolites (covering 2012-2018). MedChemComm, 2019, 10, 840-866 Biosynthetic Pathway of the Reduced Polyketide Product Citreoviridin in Aspergillus terreus var. aureus Revealed by Heterologous Expression in Aspergillus nidulans. Organic Letters, 2016, 18, 1366-9 Recent advances in genome mining of secondary metabolites in Aspergillus terreus. Frontiers in Microbiology, 2014, 5, 717 Azaphilones inhibit tau aggregation and dissolve tau aggregates in vitro. ACS Chemical Neuroscience, 2015, 6, 751-60 Reengineering an azaphilone biosynthesis pathway in Aspergillus nidulans to create lipoxygenase inhibitors. Organic Letters, 2012, 14, 972-5 Reengineering an azaphilone biosynthesis pathway in Aspergillus nidulans to create lipoxygenase inhibitors. Organic Letters, 2012, 14, 972-5 Inhibition of Tau aggregation by three Aspergillus nidulans secondary metabolites: 2, Elihydroxyemodin, asperthecin, and asperbenzaldehyde. Planta Medica, 2014, 80, 77-85 Characterization of Aspergillus niger Isolated from the International Space Station. MSystems, 2018, 13. Telomere position effect is regulated by heterochromatin-associated proteins and NkuA in Aspergillus nidulans. Microbiology (United Kingdom), 2010, 156, 3522-3531 Spatial regulation of a common precursor from two distinct genes generates metabolite diversity. Chemical Science, 2015, 6, 5913-5921 Identification and molecular genetic analysis of the cichorine gene cluster in. MedChemComm, 2012, 3. Characterization of the product of a nonribosomal peptide synthetase-like (NRPS-like) gene using the doxycycline depend

28	Asperfuranone from Aspergillus nidulans inhibits proliferation of human non-small cell lung cancer A549 cells via blocking cell cycle progression and inducing apoptosis. <i>Basic and Clinical Pharmacology and Toxicology</i> , 2010 , 107, 583-9	3.1	19
27	Proteomic characterization of Aspergillus fumigatus isolated from air and surfaces of the International Space Station. <i>Fungal Genetics and Biology</i> , 2019 , 124, 39-46	3.9	19
26	Proteomic and Metabolomic Characteristics of Extremophilic Fungi Under Simulated Mars Conditions. <i>Frontiers in Microbiology</i> , 2019 , 10, 1013	5.7	17
25	Contributions of Spore Secondary Metabolites to UV-C Protection and Virulence Vary in Different Aspergillus fumigatus Strains. <i>MBio</i> , 2020 , 11,	7.8	17
24	Norsolorinic acid from Aspergillus nidulans inhibits the proliferation of human breast adenocarcinoma MCF-7 cells via Fas-mediated pathway. <i>Basic and Clinical Pharmacology and Toxicology</i> , 2008 , 102, 491-7	3.1	17
23	Engineering Fungal Nonribosomal Peptide Synthetase-like Enzymes by Heterologous Expression and Domain Swapping. <i>Organic Letters</i> , 2016 , 18, 6236-6239	6.2	17
22	Overexpression of a three-gene conidial pigment biosynthetic pathway in Aspergillus nidulans reveals the first NRPS known to acetylate tryptophan. <i>Fungal Genetics and Biology</i> , 2017 , 101, 1-6	3.9	15
21	International Space Station conditions alter genomics, proteomics, and metabolomics in Aspergillus nidulans. <i>Applied Microbiology and Biotechnology</i> , 2019 , 103, 1363-1377	5.7	15
20	Draft Genome Sequences of Several Fungal Strains Selected for Exposure to Microgravity at the International Space Station. <i>Genome Announcements</i> , 2017 , 5,		13
19	sp. nov., Isolated From the International Space Station. <i>Frontiers in Microbiology</i> , 2021 , 12, 639396	5.7	13
18	Norsolorinic acid inhibits proliferation of T24 human bladder cancer cells by arresting the cell cycle at the G0/G1 phase and inducing a Fas/membrane-bound Fas ligand-mediated apoptotic pathway. <i>Clinical and Experimental Pharmacology and Physiology</i> , 2008 , 35, 1301-8	3	12
17	Draft Genome Sequences of Two Aspergillus fumigatus Strains, Isolated from the International Space Station. <i>Genome Announcements</i> , 2016 , 4,		12
16	Overexpression of an LaeA-like Methyltransferase Upregulates Secondary Metabolite Production in. <i>ACS Chemical Biology</i> , 2019 , 14, 1643-1651	4.9	11
15	Metabolomic Analysis of Isolated From the International Space Station Reveals Enhanced Production Levels of the Antioxidant Pyranonigrin A. <i>Frontiers in Microbiology</i> , 2020 , 11, 931	5.7	8
14	Development of Genetic Dereplication Strains in Aspergillus nidulans Results in the Discovery of Aspercryptin. <i>Angewandte Chemie</i> , 2016 , 128, 1694-1697	3.6	8
13	Mating-type factor-specific regulation of the fumagillin/pseurotin secondary metabolite supercluster in Aspergillus fumigatus. <i>Molecular Microbiology</i> , 2018 , 110, 1045-1065	4.1	8
12	Natural products development under epigenetic modulation in fungi. <i>Phytochemistry Reviews</i> , 2020 , 19, 1323-1340	7.7	7
11	Advances in space microbiology. <i>IScience</i> , 2021 , 24, 102395	6.1	7

LIST OF PUBLICATIONS

10	Discovery and Elucidation of the Biosynthesis of Aspernidgulenes: Novel Polyenes from Aspergillus Nidulans by Using Serial Promoter Replacement. <i>ChemBioChem</i> , 2019 , 20, 329-334	3.8	6	
9	Epigenetic Manipulation Induces the Production of Coumarin-Type Secondary Metabolite from Arthrobotrys foliicola. <i>Israel Journal of Chemistry</i> , 2019 , 59, 432-438	3.4	5	
8	Expanding the Chemical Space of Nonribosomal Peptide Synthetase-like Enzymes by Domain and Tailoring Enzyme Recombination. <i>Organic Letters</i> , 2018 , 20, 5082-5085	6.2	4	
7	Identification of the pigment and its role in UV resistance in Paecilomyces variotii, a Chernobyl isolate, using genetic manipulation strategies. <i>Fungal Genetics and Biology</i> , 2021 , 152, 103567	3.9	4	
6	Genome-based deletion analysis in Aspergillus terreus reveals the acetylaranotin bis-thiomethyltransferase gene. <i>Fungal Genetics and Biology</i> , 2018 , 119, 1-6	3.9	3	
5	Prevention of chronic HBV infection induced hepatocellular carcinoma development by using antiplatelet drugs. <i>Hepatobiliary Surgery and Nutrition</i> , 2012 , 1, 57-8	2.1	3	
4	An Platform for the Complete Cluster Refactoring and Total Biosynthesis of Fungal Natural Products. <i>ACS Synthetic Biology</i> , 2021 , 10, 173-182	5.7	3	
3	Identification and Validation of an Secondary Metabolite Derivative as an Inhibitor of the Musashi-RNA Interaction. <i>Cancers</i> , 2020 , 12,	6.6	3	
2	Characterization of a silent azaphilone biosynthesis gene cluster in Aspergillus terreus NIH 2624 Fungal Genetics and Biology, 2022 , 160, 103694	3.9	2	
1	Looking Ahead to 2030: Survey of Evolving Needs in Pharmacy Education. <i>Pharmacy (Basel, Switzerland)</i> , 2021 , 9,	2	1	