Hong-Ye Li

List of Publications by Year in descending order

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31902 60497 9,600 298 53 81 citations h-index g-index papers 304 304 304 9243 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Adsorption characteristics of acrylonitrile, p-toluenesulfonic acid, 1-naphthalenesulfonic acid and methyl blue on graphene in aqueous solutions. Chemical Engineering Journal, 2011, 173, 144-149.	6.6	322
2	Molecular and cellular mechanisms of neutral lipid accumulation in diatom following nitrogen deprivation. Biotechnology for Biofuels, 2013, 6, 67.	6.2	296
3	Genetic improvement of the microalga Phaeodactylum tricornutum for boosting neutral lipid accumulation. Metabolic Engineering, 2015, 27, 1-9.	3.6	250
4	Improvement of Neutral Lipid and Polyunsaturated Fatty Acid Biosynthesis by Overexpressing a Type 2 Diacylglycerol Acyltransferase in Marine Diatom Phaeodactylum tricornutum. Marine Drugs, 2013, 11, 4558-4569.	2.2	229
5	Genetic Bottlenecks Reduce Population Variation in an Experimental RNA Virus Population. Journal of Virology, 2004, 78, 10582-10587.	1.5	186
6	Effect of chitosan on incidence of brown rot, quality and physiological attributes of postharvest peach fruit. Journal of the Science of Food and Agriculture, 2001, 81, 269-274.	1.7	179
7	Arabidopsis Acyl-CoA-Binding Protein ACBP2 Interacts With an Ethylene-Responsive Element-Binding Protein, AtEBP, via its Ankyrin Repeats. Plant Molecular Biology, 2004, 54, 233-243.	2.0	167
8	Glucose-6-phosphate dehydrogenase as a target for highly efficient fatty acid biosynthesis in microalgae by enhancing NADPH supply. Metabolic Engineering, 2017, 41, 212-221.	3.6	156
9	Analysis of Genetic Bottlenecks during Horizontal Transmission of Cucumber Mosaic Virus. Journal of Virology, 2006, 80, 8345-8350.	1.5	146
10	<i>Arabidopsis thaliana</i> acylâ€CoAâ€binding protein ACBP2 interacts with heavyâ€metalâ€binding farnesylated protein AtFP6. New Phytologist, 2009, 181, 89-102.	3.5	141
11	Genetic Structure and Population Variability of Tomato Yellow Leaf Curl China Virus. Journal of Virology, 2007, 81, 5902-5907.	1.5	120
12	Colletotrichum species associated with cultivated citrus in China. Fungal Diversity, 2013, 61, 61-74.	4.7	120
13	Occurrence of plastidial triacylglycerol synthesis and the potential regulatory role of AGPAT in the model diatom Phaeodactylum tricornutum. Biotechnology for Biofuels, 2017, 10, 97.	6.2	115
14	Acyl-CoA-binding protein 2 binds lysophospholipase 2 and lysoPC to promote tolerance to cadmium-induced oxidative stress in transgenic Arabidopsis. Plant Journal, 2010, 62, no-no.	2.8	114
15	A type 2 diacylglycerol acyltransferase accelerates the triacylglycerol biosynthesis in heterokont oleaginous microalga Nannochloropsis oceanica. Journal of Biotechnology, 2016, 229, 65-71.	1.9	113
16	Proteomics to reveal metabolic network shifts towards lipid accumulation following nitrogen deprivation in the diatom Phaeodactylum tricornutum. Journal of Applied Phycology, 2014, 26, 73-82.	1.5	112
17	Ethylene- and pathogen-inducible Arabidopsis acyl-CoA-binding protein 4 interacts with an ethylene-responsive element binding protein. Journal of Experimental Botany, 2008, 59, 3997-4006.	2.4	105
18	Arabidopsis ACBP3 is an extracellularly targeted acyl-CoA-binding protein. Planta, 2006, 223, 871-881.	1.6	101

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19	Molecular characterization of a glycerol-3-phosphate acyltransferase reveals key features essential for triacylglycerol production in Phaeodactylum tricornutum. Biotechnology for Biofuels, 2016, 9, 60.	6.2	101
20	Membrane localization of Arabidopsis acyl-CoA binding protein ACBP2. Plant Molecular Biology, 2003, 51, 483-492.	2.0	95
21	Single amino acid substitutions at the acyl-CoA-binding domain interrupt 14[C]palmitoyl-CoA binding of ACBP2, an Arabidopsis acyl-CoA-binding protein with ankyrin repeats. Plant Molecular Biology, 2000, 44, 711-721.	2.0	94
22	Antibacterial activity and mechanism of action of chitosan solutions against apricot fruit rot pathogen Burkholderia seminalis. Carbohydrate Research, 2011, 346, 1294-1301.	1.1	94
23	Delta 5 Fatty Acid Desaturase Upregulates the Synthesis of Polyunsaturated Fatty Acids in the Marine Diatom <i>Phaeodactylum tricornutum</i> . Journal of Agricultural and Food Chemistry, 2014, 62, 8773-8776.	2.4	91
24	Endophytic Diaporthe associated with Citrus: A phylogenetic reassessment with seven new species from China. Fungal Biology, 2015, 119, 331-347.	1.1	91
25	PdCYP51B, a new putative sterol 14α-demethylase gene of Penicillium digitatum involved in resistance to imazalil and other fungicides inhibiting ergosterol synthesis. Applied Microbiology and Biotechnology, 2011, 91, 1107-1119.	1.7	89
26	The pH signaling transcription factor PacC is required for full virulence in Penicillium digitatum. Applied Microbiology and Biotechnology, 2013, 97, 9087-9098.	1.7	88
27	Glycerol and neutral lipid production in the oleaginous marine diatom Phaeodactylum tricornutum promoted by overexpression of glycerol-3-phosphate dehydrogenase. Biotechnology for Biofuels, 2014, 7, .	6.2	87
28	ACBP4 and ACBP5, novel Arabidopsis acyl-CoA-binding proteins with kelch motifs that bind oleoyl-CoA. Plant Molecular Biology, 2004, 55, 297-309.	2.0	86
29	Antibacterial activity of two chitosan solutions and their effect on rice bacterial leaf blight and leaf streak. Pest Management Science, 2013, 69, 312-320.	1.7	86
30	Synergistic effect of chitosan and Cryptococcus laurentii on inhibition of Penicillium expansum infections. International Journal of Food Microbiology, 2007, 114, 261-266.	2.1	83
31	Phyllosticta species associated with citrus diseases in China. Fungal Diversity, 2012, 52, 209-224.	4.7	80
32	Systemsâ€level analysis of the metabolic responses of the diatom <i><scp>P</scp>haeodactylum tricornutum</i> to phosphorus stress. Environmental Microbiology, 2014, 16, 1793-1807.	1.8	78
33	Molecular characterization of field azoxystrobin-resistant isolates of Botrytis cinerea. Pesticide Biochemistry and Physiology, 2009, 93, 72-76.	1.6	77
34	Chlorophyll Fluorescence Imaging Uncovers Photosynthetic Fingerprint of Citrus Huanglongbing. Frontiers in Plant Science, 2017, 8, 1509.	1.7	77
35	Transformation of diatom <i>Phaeodactylum tricornutum < i> by electroporation and establishment of inducible selection marker. BioTechniques, 2012, 52, 1-3.</i>	0.8	75
36	Diaporthe species occurring on citrus in China. Fungal Diversity, 2013, 61, 237-250.	4.7	73

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37	Identification of a malonyl CoAâ€acyl carrier protein transacylase and its regulatory role in fatty acid biosynthesis in oleaginous microalga <i>Nannochloropsis oceanica</i> . Biotechnology and Applied Biochemistry, 2017, 64, 620-626.	1.4	73
38	Arabidopsis acyl-CoA-binding proteins ACBP4 and ACBP5 are subcellularly localized to the cytosol and ACBP4 depletion affects membrane lipid composition. Plant Molecular Biology, 2008, 68, 571-583.	2.0	71
39	The vacuolar transport of aleurainâ€GFP and 2S albuminâ€GFP fusions is mediated by the same preâ€vacuolar compartments in tobacco BYâ€2 and Arabidopsis suspension cultured cells. Plant Journal, 2008, 56, 824-839.	2.8	69
40	Transcriptional regulation of microalgae for concurrent lipid overproduction and secretion. Science Advances, 2019, 5, eaau3795.	4.7	68
41	Construction of Novel Chloroplast Expression Vector and Development of an Efficient Transformation System for the Diatom Phaeodactylum tricornutum. Marine Biotechnology, 2014, 16, 538-546.	1.1	65
42	The mitogen-activated protein kinase kinase BOS5 is involved in regulating vegetative differentiation and virulence in Botrytis cinerea. Fungal Genetics and Biology, 2010, 47, 753-760.	0.9	64
43	Dual expression of plastidial GPAT1 and LPAT1 regulates triacylglycerol production and the fatty acid profile in Phaeodactylum tricornutum. Biotechnology for Biofuels, 2018, 11, 318.	6.2	64
44	The calcineurin-responsive transcription factor Crz1 is required for conidation, full virulence and DMI resistance in Penicillium digitatum. Microbiological Research, 2013, 168, 211-222.	2.5	63
45	CD40 ligation stimulates MCP-1 and IL-8 production, TRAF6 recruitment, and MAPK activation in proximal tubule cells. American Journal of Physiology - Renal Physiology, 2002, 282, F1020-F1033.	1.3	62
46	Biochemical and Genetic Engineering of Diatoms for Polyunsaturated Fatty Acid Biosynthesis. Marine Drugs, 2014, 12, 153-166.	2.2	62
47	Genomic and transcriptomic analyses of the tangerine pathotype of Alternaria alternata in response to oxidative stress. Scientific Reports, 2016, 6, 32437.	1.6	62
48	Engineering Variants of the I-Scel Homing Endonuclease with Strand-specific and Site-specific DNA-nicking Activity. Journal of Molecular Biology, 2008, 382, 188-202.	2.0	61
49	Antifungal effect and mechanism of chitosan against the rice sheath blight pathogen, Rhizoctonia solani. Biotechnology Letters, 2012, 34, 2291-2298.	1.1	61
50	A new inducible expression system in a transformed green alga, Chlorella vulgaris. Genetics and Molecular Research, 2011, 10, 3427-3434.	0.3	59
51	Accumulation of Recombinant SARS-CoV Spike Protein in Plant Cytosol and Chloroplasts Indicate Potential for Development of Plant-Derived Oral Vaccines. Experimental Biology and Medicine, 2006, 231, 1346-1352.	1.1	58
52	\hat{l}^2 -elemene inhibits tumor-promoting effect of M2 macrophages in lung cancer. Biochemical and Biophysical Research Communications, 2017, 490, 514-520.	1.0	58
53	Cell-Wall-Degrading Enzymes Required for Virulence in the Host Selective Toxin-Producing Necrotroph Alternaria alternata of Citrus. Frontiers in Microbiology, 2019, 10, 2514.	1.5	58
54	The pivotal role of malic enzyme in enhancing oil accumulation in green microalga Chlorella pyrenoidosa. Microbial Cell Factories, 2016, 15, 120.	1.9	57

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55	PdbrlA, PdabaA and PdwetA control distinct stages of conidiogenesis in Penicillium digitatum. Research in Microbiology, 2015, 166, 56-65.	1.0	56
56	Adaptive evolution of microalgal strains empowered by fulvic acid for enhanced polyunsaturated fatty acid production. Bioresource Technology, 2019, 277, 204-210.	4.8	55
57	Agrobacterium tumefaciens-mediated genetic transformation of the phytopathogenic fungus Penicillium digitatum. Journal of Zhejiang University: Science B, 2008, 9, 823-828.	1.3	54
58	The phyllosphere microbiome shifts toward combating melanose pathogen. Microbiome, 2022, 10, 56.	4.9	54
59	Examination of metabolic responses to phosphorus limitation via proteomic analyses in the marine diatom Phaeodactylum tricornutum. Scientific Reports, 2015, 5, 10373.	1.6	53
60	High-efficiency promoter-driven coordinated regulation of multiple metabolic nodes elevates lipid accumulation in the model microalga Phaeodactylum tricornutum. Microbial Cell Factories, 2018, 17, 54.	1.9	53
61	Thioredoxin and Glutaredoxin Systems Required for Oxidative Stress Resistance, Fungicide Sensitivity, and Virulence of Alternaria alternata. Applied and Environmental Microbiology, 2018, 84, .	1.4	53
62	PdSNF1, a sucrose non-fermenting protein kinase gene, is required for Penicillium digitatum conidiation and virulence. Applied Microbiology and Biotechnology, 2013, 97, 5433-5445.	1.7	52
63	Evolution of Gene Regulation during Transcription and Translation. Genome Biology and Evolution, 2015, 7, 1155-1167.	1.1	52
64	QUASIMODO 3 (QUA3) is a putative homogalacturonan methyltransferase regulating cell wall biosynthesis in Arabidopsis suspension-cultured cells. Journal of Experimental Botany, 2011, 62, 5063-5078.	2.4	50
65	In vitro inhibition of postharvest pathogens of fruit and control of gray mold of strawberry and green mold of citrus by aureobasidin A. International Journal of Food Microbiology, 2007, 119, 223-229.	2.1	48
66	Proteomic profile in Perna viridis after exposed to Prorocentrum lima, a dinoflagellate producing DSP toxins. Environmental Pollution, 2015, 196, 350-357.	3.7	48
67	Effect of chitosan solution on the inhibition of <i>Acidovorax citrulli</i> causing bacterial fruit blotch of watermelon. Journal of the Science of Food and Agriculture, 2013, 93, 1010-1015.	1.7	47
68	Identification of a novel phylogenetic lineage of Alternaria alternata causing citrus brown spot in China. Fungal Biology, 2015, 119, 320-330.	1.1	46
69	Hyperspectral reflectance imaging combined with carbohydrate metabolism analysis for diagnosis of citrus Huanglongbing in different seasons and cultivars. Sensors and Actuators B: Chemical, 2018, 275, 50-60.	4.0	46
70	Emerging waste valorisation techniques to moderate the hazardous impacts, and their path towards sustainability. Journal of Hazardous Materials, 2022, 423, 127023.	6.5	46
71	The clinical data from 19 critically ill patients with coronavirus disease 2019: a single-centered, retrospective, observational study. Zeitschrift Fur Gesundheitswissenschaften, 2022, 30, 361-364.	0.8	45
72	Occurrence of imazalil-resistant biotype of Penicillium digitatum in China and the resistant molecular mechanism. Journal of Zhejiang University: Science A, 2006, 7, 362-365.	1.3	42

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73	Glucosylceramides are required for mycelial growth and full virulence in Penicillium digitatum. Biochemical and Biophysical Research Communications, 2014, 455, 165-171.	1.0	42
74	Genomewide investigation into DNA elements and ABC transporters involved in imazalil resistance in <i>Penicillium digitatum</i> . FEMS Microbiology Letters, 2013, 348, 11-18.	0.7	41
75	Os2 MAP kinase-mediated osmostress tolerance in Penicillium digitatum is associated with its positive regulation on glycerol synthesis and negative regulation on ergosterol synthesis. Microbiological Research, 2014, 169, 511-521.	2.5	41
76	High Performance Liquid Chromatography and Metabolomics Analysis of Tannase Metabolism of Gallic Acid and Gallates in Tea Leaves. Journal of Agricultural and Food Chemistry, 2020, 68, 4946-4954.	2.4	41
77	Enhanced polyunsaturated fatty acid production using food wastes and biofuels byproducts by an evolved strain of Phaeodactylum tricornutum. Bioresource Technology, 2020, 296, 122351.	4.8	40
78	A diet high in sugar and fat influences neurotransmitter metabolism and then affects brain function by altering the gut microbiota. Translational Psychiatry, 2021, 11, 328.	2.4	40
79	The TB Structural Genomics Consortium: A decade of progress. Tuberculosis, 2011, 91, 155-172.	0.8	39
80	Enrichment of Long-Chain Polyunsaturated Fatty Acids by Coordinated Expression of Multiple Metabolic Nodes in the Oleaginous Microalga <i>Phaeodactylum tricornutum</i> Agricultural and Food Chemistry, 2017, 65, 7713-7720.	2.4	39
81	Differentiation in MALDI-TOF MS and FTIR spectra between two closely related species Acidovorax oryzae and Acidovorax citrulli. BMC Microbiology, 2012, 12, 182.	1.3	38
82	Identification of a putative patatin-like phospholipase domain-containing protein 3 (PNPLA3) ortholog involved in lipid metabolism in microalga Phaeodactylum tricornutum. Algal Research, 2015, 12, 274-279.	2.4	38
83	Antisense knockdown of pyruvate dehydrogenase kinase promotes the neutral lipid accumulation in the diatom Phaeodactylum tricornutum. Microbial Cell Factories, 2014, 13, 100.	1.9	36
84	Constitutive and Chloroplast Targeted Expression of Acetyl-CoA Carboxylase in Oleaginous Microalgae Elevates Fatty Acid Biosynthesis. Marine Biotechnology, 2018, 20, 566-572.	1.1	36
85	A molecular mechanism of azoxystrobin resistance in Penicillium digitatum UV mutants and a PCR-based assay for detection of azoxystrobin-resistant strains in packing- or store-house isolates. International Journal of Food Microbiology, 2009, 131, 157-161.	2.1	35
86	Ethanol induced jasmonate pathway promotes astaxanthin hyperaccumulation in Haematococcus pluvialis. Bioresource Technology, 2019, 289, 121720.	4.8	34
87	Real-time dynamics of soliton collision in a bound-state soliton fiber laser. Nanophotonics, 2020, 9, 1921-1929.	2.9	34
88	<pre><scp>PdMLE1</scp>, a specific and active transposon acts as a promoter and confers <i><scp>P</scp>enicillium digitatum</i> with <scp>DMI</scp> resistance. Environmental Microbiology Reports, 2013, 5, 135-142.</pre>	1.0	33
89	The methionine biosynthesis regulator AaMetR contributes to oxidative stress tolerance and virulence in Alternaria alternata. Microbiological Research, 2019, 219, 94-109.	2.5	33
90	TAG pathway engineering via GPAT2 concurrently potentiates abiotic stress tolerance and oleaginicity in Phaeodactylum tricornutum. Biotechnology for Biofuels, 2020, 13, 160.	6.2	33

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91	A lipid droplet-associated protein involved in lipid droplet biogenesis and triacylglycerol accumulation in the oleaginous microalga Phaeodactylum tricornutum. Algal Research, 2017, 26, 215-224.	2.4	32
92	A decomposition-based chemical reaction optimization for multi-objective vehicle routing problem for simultaneous delivery and pickup with time windows. Memetic Computing, 2018, 10, 103-120.	2.7	32
93	Gray matter atrophy in progressive supranuclear palsy: meta-analysis of voxel-based morphometry studies. Neurological Sciences, 2013, 34, 1049-1055.	0.9	31
94	Sustainable and stepwise waste-based utilisation strategy for the production of biomass and biofuels by engineered microalgae. Environmental Pollution, 2020, 265, 114854.	3.7	31
95	Functional analysis of two sterol regulatory element binding proteins in Penicillium digitatum. PLoS ONE, 2017, 12, e0176485.	1.1	31
96	Removal of Chattonella marina with clay minerals modified with a gemini surfactant. Applied Clay Science, 2010, 50, 604-607.	2.6	30
97	Biotechnological approaches to enhance biofuel producing potential of microalgae. Fuel, 2021, 302, 121169.	3.4	30
98	Allelopathic effects of Alexandrium spp. on Prorocentrum donghaiense. Harmful Algae, 2010, 10, 116-120.	2.2	29
99	Molecular phylogeny and PSP toxin profile of the Alexandrium tamarense species complex along the coast of China. Marine Pollution Bulletin, 2014, 89, 209-219.	2.3	29
100	Molecular exploration of algal interaction between the diatom Phaeodactylum tricornutum and the dinoflagellate Alexandrium tamarense. Algal Research, 2016, 17, 132-141.	2.4	29
101	The citrus postharvest pathogen Penicillium digitatum depends on the PdMpkB kinase for developmental and virulence functions. International Journal of Food Microbiology, 2016, 236, 167-176.	2.1	29
102	Influence of Bragg reflection of chirped tilted fiber Bragg grating on Raman suppression in high-power tandem pumping fiber amplifiers. Optics Express, 2020, 28, 19508.	1.7	29
103	Action of Chitosan Against Xanthomonas Pathogenic Bacteria Isolated from Euphorbia pulcherrima. Molecules, 2012, 17, 7028-7041.	1.7	28
104	P-glycoprotein expression in Perna viridis after exposure to Prorocentrum lima, a dinoflagellate producing DSP toxins. Fish and Shellfish Immunology, 2014, 39, 254-262.	1.6	28
105	Identification of secondary metabolite biosynthetic gene clusters associated with the infection of citrus fruit by Penicillium digitatum. Postharvest Biology and Technology, 2017, 134, 17-21.	2.9	28
106	P-glycoprotein and its inducible expression in three bivalve species after exposure to Prorocentrum lima. Aquatic Toxicology, 2015, 169, 123-132.	1.9	27
107	Molecular identification of green algae from the rafts based infrastructure of Porphyra yezoensis. Marine Pollution Bulletin, 2012, 64, 2077-2082.	2.3	26
108	Expression profile of eight glutathione S-transferase genes in Crassostrea ariakensis after exposure to DSP toxins producing dinoflagellate Prorocentrum lima. Toxicon, 2015, 105, 45-55.	0.8	26

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109	A Genomics Based Discovery of Secondary Metabolite Biosynthetic Gene Clusters in Aspergillus ustus. PLoS ONE, 2015, 10, e0116089.	1.1	25
110	Overproduction of Bioactive Algal Chrysolaminarin by the Critical Carbon Flux Regulator Phosphoglucomutase. Biotechnology Journal, 2019, 14, 1800220.	1.8	25
111	Responses of CYP450 in the mussel Perna viridis after short-term exposure to the DSP toxins-producing dinoflagellate Prorocentrum lima. Ecotoxicology and Environmental Safety, 2019, 176, 178-185.	2.9	25
112	De novo transcriptome analysis of the mussel Perna viridis after exposure to the toxic dinoflagellate Prorocentrum lima. Ecotoxicology and Environmental Safety, 2020, 192, 110265.	2.9	25
113	Bacterial brown stripe of rice in soil-less culture system caused by Acidovorax avenae subsp. avenae in China. Journal of General Plant Pathology, 2011, 77, 64-67.	0.6	24
114	Csn5 Is Required for the Conidiogenesis and Pathogenesis of the Alternaria alternata Tangerine Pathotype. Frontiers in Microbiology, 2018, 9, 508.	1.5	24
115	The basal transcription factor II H subunit Tfb5 is required for stress response and pathogenicity in the tangerine pathotype of <i>Alternaria alternata</i> i>Nolecular Plant Pathology, 2020, 21, 1337-1352.	2.0	24
116	Synergistic bioconversion of lipids and carotenoids from food waste by Dunaliella salina with fulvic acid via a two-stage cultivation strategy. Energy Conversion and Management, 2021, 234, 113908.	4.4	24
117	Exogenous Melatonin Promotes the Salt Tolerance by Removing Active Oxygen and Maintaining Ion Balance in Wheat (Triticum aestivum L.). Frontiers in Plant Science, 2021, 12, 787062.	1.7	24
118	IL-8 amplifies CD40/CD154-mediated ICAM-1 production via the CXCR-1 receptor and p38-MAPK pathway in human renal proximal tubule cells. American Journal of Physiology - Renal Physiology, 2009, 296, F438-F445.	1.3	23
119	Quercetin potentiates the concurrent hyper-accumulation of cellular biomass and lipids in Chlorella vulgaris. Bioresource Technology, 2018, 269, 434-442.	4.8	23
120	Genomic features and evolution of the conditionally dispensable chromosome in the tangerine pathotype of <i>Alternaria alternata</i> i>Nolecular Plant Pathology, 2019, 20, 1425-1438.	2.0	23
121	Transmissibility of acute haemorrhagic conjunctivitis in small-scale outbreaks in Hunan Province, China. Scientific Reports, 2020, 10, 119.	1.6	23
122	Physiological and molecular responses in halotolerant Dunaliella salina exposed to molybdenum disulfide nanoparticles. Journal of Hazardous Materials, 2021, 404, 124014.	6.5	23
123	Consensus on the application of negative pressure wound therapy of diabetic foot wounds. Burns and Trauma, 2021, 9, tkab018.	2.3	23
124	G-box binding coincides with increased Solanum melongena cysteine proteinase expression in senescent fruits and circadian-regulated leaves. Plant Molecular Biology, 2003, 51, 9-19.	2.0	22
125	Experimental Methodology for Obtaining the Flow Curve of Sheet Materials in a Wide Range of Strains. Steel Research International, 2013, 84, 146-154.	1.0	22
126	A waste upcycling loop: Two-factor adaptive evolution of microalgae to increase polyunsaturated fatty acid production using food waste. Journal of Cleaner Production, 2022, 331, 130018.	4.6	22

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127	Bare-Bones Teaching-Learning-Based Optimization. Scientific World Journal, The, 2014, 2014, 1-17.	0.8	21
128	Rational design and fabrication of a cancer-targeted chitosan nanocarrier to enhance selective cellular uptake and anticancer efficacy of selenocystine. Journal of Materials Chemistry B, 2015, 3, 2497-2504.	2.9	21
129	A combined light regime and carbon supply regulation strategy for microalgae-based sugar industry wastewater treatment and low-carbon biofuel production to realise a circular economy. Chemical Engineering Journal, 2022, 446, 137422.	6.6	21
130	Functional caveolae are a prerequisite for CD40 signaling in human renal proximal tubule cells. American Journal of Physiology - Renal Physiology, 2004, 286, F711-F719.	1.3	20
131	Identification of a putative seipin ortholog involved in lipid accumulation in marine microalga Phaeodactylum tricornutum. Journal of Applied Phycology, 2017, 29, 2821-2829.	1.5	20
132	Up-regulation of Nrf2-dependent antioxidant defenses in Perna viridis after exposed to Prorocentrum lima. Fish and Shellfish Immunology, 2019, 90, 173-179.	1.6	20
133	Enantioselectivity in toxicity and degradation of dichlorprop-methyl in algal cultures. Journal of Environmental Science and Health - Part B Pesticides, Food Contaminants, and Agricultural Wastes, 2008, 43, 288-292.	0.7	19
134	OSBP-related protein 7 interacts with GATE-16 and negatively regulates GS28 protein stability. Experimental Cell Research, 2011, 317, 2353-2363.	1.2	19
135	Gene expression profiles in zebrafish (Danio rerio) liver after acute exposure to okadaic acid. Environmental Toxicology and Pharmacology, 2014, 37, 791-802.	2.0	19
136	A Destructive New Disease of <i>Citrus</i> in China Caused by <i>Cryptosporiopsis citricarpa</i> sp. nov Plant Disease, 2012, 96, 804-812.	0.7	18
137	Heterogeneous expression of human PNPLA3 triggers algal lipid accumulation and lipid droplet enlargement. Algal Research, 2018, 31, 276-281.	2.4	18
138	Metabarcoding reveals differences in fungal communities between unflooded versus tidal flat soil in coastal saline ecosystem. Science of the Total Environment, 2019, 690, 911-922.	3.9	18
139	Glucoseâ€6â€Phosphate Dehydrogenase from the Oleaginous Microalga <i>Nannochloropsis</i> Its Potential Role in Promoting Lipogenesis. Biotechnology Journal, 2020, 15, e1900135.	1.8	18
140	Analysis of melatonin regulation of germination and antioxidant metabolism in different wheat cultivars under polyethylene glycol stress. PLoS ONE, 2020, 15, e0237536.	1.1	18
141	Raman suppression in 5ÂkW fiber amplifier using long period fiber grating fabricated by CO2 laser. Optics and Laser Technology, 2022, 145, 107484.	2.2	18
142	Histone Acetyltransferases and Deacetylases Are Required for Virulence, Conidiation, DNA Damage Repair, and Multiple Stresses Resistance of Alternaria alternata. Frontiers in Microbiology, 2021, 12, 783633.	1.5	18
143	The role of diatom glucose-6-phosphate dehydrogenase on lipogenic NADPH supply in green microalgae through plastidial oxidative pentose phosphate pathway. Applied Microbiology and Biotechnology, 2018, 102, 10803-10815.	1.7	17
144	Systems-level analysis of metabolic mechanism following nitrogen limitation in benthic dinoflagellate Prorocentrum lima. Algal Research, 2018, 33, 389-398.	2.4	17

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145	Potentiation of concurrent expression of lipogenic genes by novel strong promoters in the oleaginous microalga <i>Phaeodactylum tricornutum</i> . Biotechnology and Bioengineering, 2019, 116, 3006-3015.	1.7	17
146	Optimized Planting Density Maintains High Wheat Yield Under Limiting Irrigation in North China Plain. International Journal of Plant Production, 2020, 14, 107-117.	1.0	17
147	Antisense knockdown of pyruvate dehydrogenase kinase promotes the neutral lipid accumulation in the diatom. Microbial Cell Factories, 2014, 13, 100.	1.9	17
148	An Injection-Locked Single-Longitudinal-Mode Fiber Ring Laser With Cylindrical Vector Beam Emission. IEEE Photonics Journal, 2017, 9, 1-8.	1.0	16
149	Isolation and characterization of a novel Escherichia coli Kayfunavirus phage DY1. Virus Research, 2021, 293, 198274.	1.1	16
150	CD40/CD154 ligation induces mononuclear cell adhesion to human renal proximal tubule cells via increased ICAM-1 expression. American Journal of Physiology - Renal Physiology, 2005, 289, F145-F153.	1.3	15
151	Protein interactors of acyl-CoA-binding protein ACBP2 mediate cadmium tolerance in Arabidopsis. Plant Signaling and Behavior, 2010, 5, 1025-1027.	1.2	15
152	Complete mitochondrial genome sequence of the phytopathogenic fungus Penicillium digitatum and comparative analysis of closely related species. FEMS Microbiology Letters, 2011, 323, 29-34.	0.7	15
153	Strong seed-specific protein expression from the Vigna radiata storage protein 8SGα promoter in transgenic Arabidopsis seeds. Journal of Biotechnology, 2014, 174, 49-56.	1.9	15
154	Chemical Response of the Toxic Dinoflagellate <i>Karenia mikimotoi</i> Against Grazing by Three Species of Zooplankton. Journal of Eukaryotic Microbiology, 2015, 62, 470-480.	0.8	15
155	Downregulation of miR-223 and miR-19a induces differentiation and promotes recruitment of osteoclast cells in giant-cell tumor of the bone via the Runx2/TWIST-RANK/RANKL pathway. Biochemical and Biophysical Research Communications, 2018, 505, 1003-1009.	1.0	15
156	Genomic Sequencing of Phyllosticta citriasiana Provides Insight Into Its Conservation and Diversification With Two Closely Related Phyllosticta Species Associated With Citrus. Frontiers in Microbiology, 2019, 10, 2979.	1.5	15
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