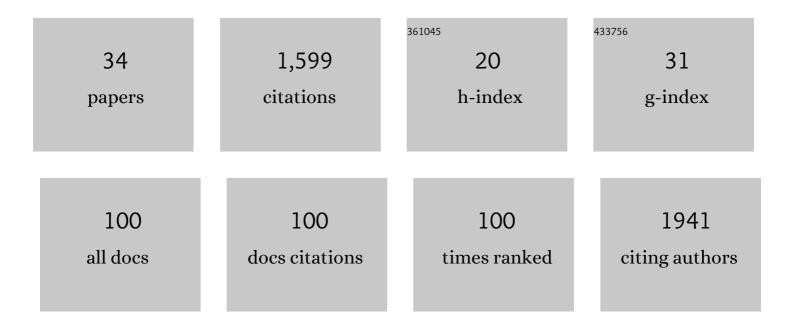
## Eric Lam

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1659240/publications.pdf Version: 2024-02-01



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#	Article	IF	CITATIONS
1	Integrative analysis of chromatin states in <scp>A</scp> rabidopsis identified potential regulatory mechanisms for natural antisense transcript production. Plant Journal, 2013, 73, 77-90.	2.8	147
2	Arabidopsis metacaspase 2d is a positive mediator of cell death induced during biotic and abiotic stresses. Plant Journal, 2011, 66, 969-982.	2.8	144
3	Comprehensive definition of genome features in <i>Spirodela polyrhiza</i> by highâ€depth physical mapping and shortâ€read <scp>DNA</scp> sequencing strategies. Plant Journal, 2017, 89, 617-635.	2.8	115
4	Return of the Lemnaceae: duckweed as a model plant system in the genomics and postgenomics era. Plant Cell, 2021, 33, 3207-3234.	3.1	111
5	Roles of Plant-Associated Microbiota in Traditional Herbal Medicine. Trends in Plant Science, 2018, 23, 559-562.	4.3	91
6	Calcium-dependent Activation and Autolysis of Arabidopsis Metacaspase 2d. Journal of Biological Chemistry, 2011, 286, 10027-10040.	1.6	84
7	Bacterial Production of Indole Related Compounds Reveals Their Role in Association Between Duckweeds and Endophytes. Frontiers in Chemistry, 2018, 6, 265.	1.8	75
8	Regulating the reapers: activating metacaspases for programmed cell death. Trends in Plant Science, 2012, 17, 487-494.	4.3	73
9	Classification and Nomenclature of Metacaspases and Paracaspases: No More Confusion with Caspases. Molecular Cell, 2020, 77, 927-929.	4.5	71
10	Generating a highâ€confidence reference genome map of the Greater Duckweed by integration of cytogenomic, optical mapping, and Oxford Nanopore technologies. Plant Journal, 2018, 96, 670-684.	2.8	64
11	Natural variance in salt tolerance and induction of starch accumulation in duckweeds. Planta, 2015, 241, 1395-1404.	1.6	61
12	Expression of <i>Arabidopsis Bax Inhibitorâ€1</i> in transgenic sugarcane confers drought tolerance. Plant Biotechnology Journal, 2016, 14, 1826-1837.	4.1	59
13	Genome and time-of-day transcriptome of <i>Wolffia australiana</i> link morphological minimization with gene loss and less growth control. Genome Research, 2021, 31, 225-238.	2.4	56
14	Duckweed hosts a taxonomically similar bacterial assemblage as the terrestrial leaf microbiome. PLoS ONE, 2020, 15, e0228560.	1.1	51
15	RNA interference as a gene silencing tool to control <i><i>Tuta absoluta</i></i> in tomato (Solanum) Tj ETQq1 I	0.784314	4 rgBT /Overic
16	Structural basis for Ca2+-dependent activation of a plant metacaspase. Nature Communications, 2020, 11, 2249.	5.8	38
17	Characterization of a photosynthetic mutant of Lemna lacking the cytochrome b6-f complex. Biochimica Et Biophysica Acta - Bioenergetics, 1985, 810, 106-109.	0.5	25
18	Correlation of Apiose Levels and Growth Rates in Duckweeds. Frontiers in Chemistry, 2018, 6, 291.	1.8	25

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19	Linkage structure of cell-wall polysaccharides from three duckweed species. Carbohydrate Polymers, 2019, 223, 115119.	5.1	23
20	Flavonoids from duckweeds: potential applications in the human diet. RSC Advances, 2020, 10, 44981-44988.	1.7	21
21	Domain swap between two typeâ€∢scp>II metacaspases defines key elements for their biochemical properties. Plant Journal, 2018, 96, 921-936.	2.8	17
22	Host-specific and tissue-dependent orchestration of microbiome community structure in traditional rice paddy ecosystems. Plant and Soil, 2020, 452, 379-395.	1.8	14
23	Auxin-Producing Bacteria from Duckweeds Have Different Colonization Patterns and Effects on Plant Morphology. Plants, 2022, 11, 721.	1.6	14
24	Wolffia, a minimalist plant and synthetic biology chassis. Trends in Plant Science, 2022, 27, 430-439.	4.3	13
25	Defining the Functional Network of Epigenetic Regulators in Arabidopsis thaliana. Molecular Plant, 2009, 2, 661-674.	3.9	12
26	High Saccharification, Low Lignin, and High Sustainability Potential Make Duckweeds Adequate as Bioenergy Feedstocks. Bioenergy Research, 2021, 14, 1082-1092.	2.2	12
27	Sequence-guided approach to genotyping plant clones and species using polymorphic NB-ARC-related genes. Plant Molecular Biology, 2018, 98, 219-231.	2.0	8
28	Decoding the role of chromatin architecture in development: coming closer to the end of the tunnel. Frontiers in Plant Science, 2014, 5, 374.	1.7	3
29	Mixing genomes alters nuclear architecture. Nature Plants, 2018, 4, 65-66.	4.7	2
30	NDP-Sugar Pathways Overview of Spirodela polyrhiza and Their Relevance for Bioenergy and Biorefinery. Bioenergy Research, 0, , 1.	2.2	1
31	Duckweed hosts a taxonomically similar bacterial assemblage as the terrestrial leaf microbiome. , 2020, 15, e0228560.		0
32	Duckweed hosts a taxonomically similar bacterial assemblage as the terrestrial leaf microbiome. , 2020, 15, e0228560.		0
33	Duckweed hosts a taxonomically similar bacterial assemblage as the terrestrial leaf microbiome. , 2020, 15, e0228560.		0
34	Duckweed hosts a taxonomically similar bacterial assemblage as the terrestrial leaf microbiome. , 2020, 15, e0228560.		0