

Noah Fahlgren

List of Publications by Year in descending order

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Version: 2024-02-01

58
papers

14,587
citations

87723

38
h-index

155451

55
g-index

84
all docs

84
docs citations

84
times ranked

14531
citing authors

#	ARTICLE	IF	CITATIONS
1	Heat stress changes mineral nutrient concentrations in <i>Chenopodium quinoa</i> seed. <i>Plant Direct</i> , 2022, 6, e384.	0.8	8
2	A protocol for <i>Chenopodium quinoa</i> pollen germination. <i>Plant Methods</i> , 2022, 18, 65.	1.9	3
3	First Plant Cell Atlas symposium report. <i>Plant Direct</i> , 2022, 6, .	0.8	1
4	Vision, challenges and opportunities for a Plant Cell Atlas. <i>ELife</i> , 2021, 10, .	2.8	31
5	Picturing the future of food. <i>The Plant Phenome Journal</i> , 2021, 4, e20014.	1.0	11
6	What Does TERRA-REF's High Resolution, Multi Sensor Plant Sensing Public Domain Data Offer the Computer Vision Community?. , 2021, , .		1
7	Heating quinoa shoots results in yield loss by inhibiting fruit production and delaying maturity. <i>Plant Journal</i> , 2020, 102, 1058-1073.	2.8	19
8	Genome-wide association study (GWAS) of leaf cuticular wax components in <i>Camelina sativa</i> identifies genetic loci related to intracellular wax transport. <i>BMC Plant Biology</i> , 2019, 19, 187.	1.6	22
9	NAPPN: Who We Are, Where We Are Going, and Why You Should Join Us!. <i>The Plant Phenome Journal</i> , 2019, 2, 1-4.	1.0	4
10	Antiviral ARGONAUTES Against <i>Turnip Crinkle Virus</i> Revealed by Image-Based Trait Analysis. <i>Plant Physiology</i> , 2019, 180, 1418-1435.	2.3	22
11	Genetic Diversity and Population Structure of a <i>Camelina sativa</i> Spring Panel. <i>Frontiers in Plant Science</i> , 2019, 10, 184.	1.7	118
12	Classifying cold stress responses of inbred maize seedlings using <i>RGB</i> imaging. <i>Plant Direct</i> , 2019, 3, e00104.	0.8	34
13	Raspberry Pi-powered imaging for plant phenotyping. <i>Applications in Plant Sciences</i> , 2018, 6, e1031.	0.8	68
14	Early Drought Plant Stress Detection with Bi-Directional Long-Term Memory Networks. <i>Photogrammetric Engineering and Remote Sensing</i> , 2018, 84, 459-468.	0.3	5
15	Components of Water Use Efficiency Have Unique Genetic Signatures in the Model C ₄ Grass <i>Setaria</i> . <i>Plant Physiology</i> , 2018, 178, 699-715.	2.3	47
16	TERRA-REF Data Processing Infrastructure. , 2018, , .		22
17	An automated, high-throughput method for standardizing image color profiles to improve image-based plant phenotyping. <i>PeerJ</i> , 2018, 6, e5727.	0.9	31
18	High-throughput phenotyping. <i>American Journal of Botany</i> , 2017, 104, 505-508.	0.8	44

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19	The Next Generation of Training for Arabidopsis Researchers: Bioinformatics and Quantitative Biology. <i>Plant Physiology</i> , 2017, 175, 1499-1509.	2.3	11
20	PlantCV v2: Image analysis software for high-throughput plant phenotyping. <i>PeerJ</i> , 2017, 5, e4088.	0.9	211
21	P-SAMS: a web site for plant artificial microRNA and synthetic <i>trans</i> -acting small interfering RNA design. <i>Bioinformatics</i> , 2016, 32, 157-158.	1.8	67
22	Naïve Bayes pixel-level plant segmentation. , 2016, , .		8
23	Plant scientists: GM technology is safe. <i>Science</i> , 2016, 351, 824-824.	6.0	5
24	Lights, camera, action: high-throughput plant phenotyping is ready for a close-up. <i>Current Opinion in Plant Biology</i> , 2015, 24, 93-99.	3.5	567
25	A Versatile Phenotyping System and Analytics Platform Reveals Diverse Temporal Responses to Water Availability in <i>Setaria</i> . <i>Molecular Plant</i> , 2015, 8, 1520-1535.	3.9	202
26	Roles and Programming of Arabidopsis ARGONAUTE Proteins during Turnip Mosaic Virus Infection. <i>PLoS Pathogens</i> , 2015, 11, e1004755.	2.1	175
27	Highly specific gene silencing in a monocot species by artificial microRNAs derived from chimeric <i>miRNA</i> precursors. <i>Plant Journal</i> , 2015, 82, 1061-1075.	2.8	45
28	CG gene body DNA methylation changes and evolution of duplicated genes in cassava. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 13729-13734.	3.3	129
29	New Generation of Artificial MicroRNA and Synthetic Trans-Acting Small Interfering RNA Vectors for Efficient Gene Silencing in Arabidopsis. <i>Plant Physiology</i> , 2014, 165, 15-29.	2.3	119
30	Preparation of Multiplexed Small RNA Libraries from Plants. <i>Bio-protocol</i> , 2014, 4, .	0.2	7
31	Parallel analysis of RNA ends enhances global investigation of microRNAs and target RNAs of <i>Brachypodium distachyon</i> . <i>Genome Biology</i> , 2013, 14, R145.	13.9	67
32	Phytophthora Have Distinct Endogenous Small RNA Populations That Include Short Interfering and microRNAs. <i>PLoS ONE</i> , 2013, 8, e77181.	1.1	88
33	Functional Analysis of Three Arabidopsis ARGONAUTES Using Slicer-Defective Mutants. <i>Plant Cell</i> , 2012, 24, 3613-3629.	3.1	249
34	The <i>Caenorhabditis elegans</i> RDE-10/RDE-11 Complex Regulates RNAi by Promoting Secondary siRNA Amplification. <i>Current Biology</i> , 2012, 22, 881-890.	1.8	49
35	The Arabidopsis lyrata genome sequence and the basis of rapid genome size change. <i>Nature Genetics</i> , 2011, 43, 476-481.	9.4	814
36	Evolution and Functional Diversification of <i>MIRNA</i> Genes. <i>Plant Cell</i> , 2011, 23, 431-442.	3.1	645

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37	<i>mut-16</i> and other <i>mutator</i> class genes modulate 22G and 26G siRNA pathways in <i>Caenorhabditis elegans</i> . Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 1201-1208.	3.3	128
38	The ERI-6/7 Helicase Acts at the First Stage of an siRNA Amplification Pathway That Targets Recent Gene Duplications. PLoS Genetics, 2011, 7, e1002369.	1.5	74
39	Unique functionality of 22-nt miRNAs in triggering RDR6-dependent siRNA biogenesis from target transcripts in Arabidopsis. Nature Structural and Molecular Biology, 2010, 17, 997-1003.	3.6	448
40	Genome sequencing and analysis of the model grass <i>Brachypodium distachyon</i> . Nature, 2010, 463, 763-768.	13.7	1,685
41	MicroRNA Gene Evolution in <i>Arabidopsis lyrata</i> and <i>Arabidopsis thaliana</i> . Plant Cell, 2010, 22, 1074-1089.	3.1	234
42	<i>Arabidopsis</i> RNA-Dependent RNA Polymerases and Dicer-Like Proteins in Antiviral Defense and Small Interfering RNA Biogenesis during <i>Turnip Mosaic Virus</i> Infection. Plant Cell, 2010, 22, 481-496.	3.1	454
43	Identification of <i>MIR390a</i> precursor processing-defective mutants in Arabidopsis by direct genome sequencing. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 466-471.	3.3	137
44	miRNA Target Prediction in Plants. Methods in Molecular Biology, 2010, 592, 51-57.	0.4	246
45	Computational and analytical framework for small RNA profiling by high-throughput sequencing. Rna, 2009, 15, 992-1002.	1.6	112
46	Genome-wide profiling of <i>Populus</i> small RNAs. BMC Genomics, 2009, 10, 620.	1.2	90
47	Genome sequence and analysis of the Irish potato famine pathogen <i>Phytophthora infestans</i> . Nature, 2009, 461, 393-398.	13.7	1,405
48	Distinct Argonaute-Mediated 22G-RNA Pathways Direct Genome Surveillance in the <i>C. elegans</i> Germline. Molecular Cell, 2009, 36, 231-244.	4.5	449
49	PRG-1 and 21U-RNAs Interact to Form the piRNA Complex Required for Fertility in <i>C. elegans</i> . Molecular Cell, 2008, 31, 67-78.	4.5	528
50	Specificity of ARGONAUTE7-miR390 Interaction and Dual Functionality in TAS3 Trans-Acting siRNA Formation. Cell, 2008, 133, 128-141.	13.5	712
51	AGO1-miR173 complex initiates phased siRNA formation in plants. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 20055-20062.	3.3	178
52	Genome-Wide Analysis of the RNA-DEPENDENT RNA POLYMERASE6/DICER-LIKE4 Pathway in Arabidopsis Reveals Dependency on miRNA- and tasiRNA-Directed Targeting. Plant Cell, 2007, 19, 926-942.	3.1	381
53	Update of ASRP: the Arabidopsis Small RNA Project database. Nucleic Acids Research, 2007, 36, D982-D985.	6.5	70
54	High-Throughput Sequencing of Arabidopsis microRNAs: Evidence for Frequent Birth and Death of MIRNA Genes. PLoS ONE, 2007, 2, e219.	1.1	1,100

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55	Genome-Wide Profiling and Analysis of Arabidopsis siRNAs. PLoS Biology, 2007, 5, e57.	2.6	473
56	Repression of <i>AUXIN RESPONSE FACTOR10</i> by microRNA160 is critical for seed germination and post-germination stages. Plant Journal, 2007, 52, 133-146.	2.8	548
57	Regulation of <i>AUXIN RESPONSE FACTOR3</i> by TAS3 ta-siRNA Affects Developmental Timing and Patterning in Arabidopsis. Current Biology, 2006, 16, 939-944.	1.8	545
58	Expression of Arabidopsis MIRNA Genes. Plant Physiology, 2005, 138, 2145-2154.	2.3	626