

Robert Henry

List of Publications by Year in descending order

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Version: 2024-02-01

58
papers

2,217
citations

304743

22
h-index

233421

45
g-index

60
all docs

60
docs citations

60
times ranked

2931
citing authors

#	ARTICLE	IF	CITATIONS
1	Cyanogenesis in the Sorghum Genus: From Genotype to Phenotype. <i>Genes</i> , 2022, 13, 140.	2.4	7
2	New Hybrid Spikelet Sterility Gene Found in Interspecific Cross between <i>Oryza sativa</i> and <i>O. meridionalis</i> . <i>Plants</i> , 2022, 11, 378.	3.5	2
3	Supporting in situ conservation of the genetic diversity of crop wild relatives using genomic technologies. <i>Molecular Ecology</i> , 2022, 31, 2207-2222.	3.9	20
4	Reticulate Evolution in AA-Genome Wild Rice in Australia. <i>Frontiers in Plant Science</i> , 2022, 13, 767635.	3.6	2
5	Potential of Genome Editing to Capture Diversity From Australian Wild Rice Relatives. <i>Frontiers in Genome Editing</i> , 2022, 4, 875243.	5.2	3
6	Genetics and Genomics of African Rice (<i>Oryza glaberrima</i> Steud) Domestication. <i>Rice</i> , 2021, 14, 6.	4.0	13
7	Wheat Grain Transcriptome. , 2021, , 501-512.		0
8	Improving rice salt tolerance by precision breeding in a new era. <i>Current Opinion in Plant Biology</i> , 2021, 60, 101996.	7.1	61
9	Phylogenetic relationships in the <i>Sorghum</i> genus based on sequencing of the chloroplast and nuclear genes. <i>Plant Genome</i> , 2021, 14, e20123.	2.8	13
10	Identification of genes associated with chapatti quality using transcriptome analysis. <i>Journal of Cereal Science</i> , 2021, 101, 103276.	3.7	0
11	Transcriptome profiling of wheat genotypes under heat stress during grain-filling. <i>Journal of Cereal Science</i> , 2020, 91, 102895.	3.7	32
12	Genome-wide association studies for yield component traits in a macadamia breeding population. <i>BMC Genomics</i> , 2020, 21, 199.	2.8	25
13	Advances in Molecular Genetics and Genomics of African Rice (<i>Oryza glaberrima</i> Steud). <i>Plants</i> , 2019, 8, 376.	3.5	10
14	Segregation Distortion Observed in the Progeny of Crosses Between <i>Oryza sativa</i> and <i>O. meridionalis</i> Caused by Abortion During Seed Development. <i>Plants</i> , 2019, 8, 398.	3.5	8
15	Relationships between Iraqi Rice Varieties at the Nuclear and Plastid Genome Levels. <i>Plants</i> , 2019, 8, 481.	3.5	5
16	DIFFERENTIAL RESPONSE OF WHEAT GENOTYPES TO HEAT STRESS DURING GRAIN FILLING. <i>Experimental Agriculture</i> , 2019, 55, 818-827.	0.9	6
17	Genomes of 13 domesticated and wild rice relatives highlight genetic conservation, turnover and innovation across the genus <i>Oryza</i> . <i>Nature Genetics</i> , 2018, 50, 285-296.	21.4	413
18	Sequencing of bulks of segregants allows dissection of genetic control of amylose content in rice. <i>Plant Biotechnology Journal</i> , 2018, 16, 100-110.	8.3	52

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19	A mosaic monoploid reference sequence for the highly complex genome of sugarcane. <i>Nature Communications</i> , 2018, 9, 2638.	12.8	299
20	Grain physical characteristic of the Australian wild rices. <i>Plant Genetic Resources: Characterisation and Utilisation</i> , 2017, 15, 409-420.	0.8	10
21	Comparison of Chapatti and Breadmaking Quality of Wheat Genotypes. <i>Cereal Chemistry</i> , 2017, 94, 409-416.	2.2	5
22	Thirty-three years of 2-acetyl-1-pyrroline, a principal basmati aroma compound in scented rice (<i>Oryza sativa</i> L.): a status review. <i>Journal of the Science of Food and Agriculture</i> , 2017, 97, 384-395.	3.5	123
23	<i>Plant Genetic Resources.</i> , 2017, , 15-29.		1
24	Global agricultural intensification during climate change: a role for genomics. <i>Plant Biotechnology Journal</i> , 2016, 14, 1095-1098.	8.3	221
25	Genomics Strategies for Germplasm Characterization and the Development of Climate Resilient Crops. , 2016, , 3-10.		2
26	Analysis of adaptive ribosomal gene diversity in wild plant populations from contrasting climatic environments. <i>Plant Signaling and Behavior</i> , 2012, 7, 602-604.	2.4	15
27	Structural and Chemical Characterization of Hardwood from Tree Species with Applications as Bioenergy Feedstocks. <i>PLoS ONE</i> , 2012, 7, e52820.	2.5	32
28	Genomics for Bioenergy Production. , 2012, , 21-29.		2
29	The Sugarcane Genome Challenge: Strategies for Sequencing a Highly Complex Genome. <i>Tropical Plant Biology</i> , 2011, 4, 145-156.	1.9	91
30	Eucalypts as a biofuel feedstock. <i>Biofuels</i> , 2011, 2, 639-657.	2.4	40
31	The role of plant biotechnology in bio-energy production. <i>Plant Biotechnology Journal</i> , 2010, 8, 243-243.	8.3	5
32	An Overview of Advances in Plant Genomics in the New Millennium. , 2010, , 1-23.		0
33	New tools for single nucleotide polymorphism (SNP) discovery and analysis accelerating plant biotechnology. <i>Plant Biotechnology Journal</i> , 2009, 7, 311-311.	8.3	22
34	Is Malting Barley Better Feed for Cattle than Feed Barley?. <i>Journal of the Institute of Brewing</i> , 2009, 115, 95-104.	2.3	15
35	Mapping species differences for adventitious rooting in a <i>Corymbia torelliana</i> – <i>Corymbia citriodora</i> subspecies <i>variegata</i> hybrid. <i>Tree Genetics and Genomes</i> , 2008, 4, 715-725.	1.6	28
36	Assessing for genetic and environmental effects on ruminant feed quality in barley (<i>Hordeum</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62 T	1.2	15

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37	Differential LongSAGE tag abundance analysis in a barley seed germination time course and validation with relative real-time RT-PCR. <i>Plant Science</i> , 2008, 175, 858-867.	3.6	8
38	SAGE of the developing wheat caryopsis. <i>Plant Biotechnology Journal</i> , 2007, 5, 69-83.	8.3	49
39	Measurement of genetic and environmental variation in barley (<i>Hordeum vulgare</i>) grain hardness. <i>Journal of Cereal Science</i> , 2007, 46, 82-92.	3.7	44
40	Abundant transcripts of malting barley identified by serial analysis of gene expression (SAGE). <i>Plant Biotechnology Journal</i> , 2006, 4, 289-301.	8.3	37
41	Congruence in QTL for adventitious rooting in <i>Pinus elliottii</i> – <i>Pinus caribaea</i> hybrids resolves between and within-species effects. <i>Molecular Breeding</i> , 2006, 18, 11-28.	2.1	16
42	Selecting for increased barley grain size. <i>Journal of Cereal Science</i> , 2006, 43, 198-208.	3.7	34
43	DNA banks and their role in facilitating the application of genomics to plant germplasm. <i>Plant Genetic Resources: Characterisation and Utilisation</i> , 2006, 4, 64-70.	0.8	11
44	Isolation and partial characterisation of a putative monoterpene synthase from <i>Melaleuca alternifolia</i> . <i>Plant Physiology and Biochemistry</i> , 2004, 42, 875-882.	5.8	16
45	The promoter of the <i>asi</i> gene directs expression in the maternal tissues of the seed in transgenic barley. <i>Plant Molecular Biology</i> , 2003, 52, 787-800.	3.9	26
46	Genes associated with the end of dormancy in grapes. <i>Functional and Integrative Genomics</i> , 2003, 3, 144-152.	3.5	56
47	Single-nucleotide polymorphism detection in plants using a single-stranded pyrosequencing protocol with a universal biotinylated primer. <i>Analytical Biochemistry</i> , 2003, 317, 166-170.	2.4	38
48	Genetics of physical wood properties and early growth in a tropical pine hybrid. <i>Canadian Journal of Forest Research</i> , 2003, 33, 1923-1932.	1.7	6
49	Branch architecture QTL for <i>Pinus elliottii</i> var. <i>elliottii</i> × $\frac{1}{2}$ <i>Pinus caribaea</i> var. <i>hondurensis</i> hybrids. <i>Annals of Forest Science</i> , 2002, 59, 617-625.	2.0	21
50	Isolation of genes involved in secondary metabolism from <i>Melaleuca alternifolia</i> (Cheel) using expressed sequence tags (ESTs). <i>Plant Science</i> , 2002, 162, 9-15.	3.6	21
51	Measurement of green fluorescent protein concentration in single cells by image analysis. <i>Analytical Biochemistry</i> , 2002, 310, 84-92.	2.4	35
52	Analysis of grape ESTs: global gene expression patterns in leaf and berry. <i>Plant Science</i> , 2000, 159, 87-95.	3.6	64
53	Identification of the creatine binding domain of creatine kinase by photoaffinity labeling. <i>BBA - Proteins and Proteomics</i> , 1998, 1387, 80-88.	2.1	9
54	Dichloroaromatic phosphoguanidines are potent inhibitors but very poor substrates for cytosolic creatine kinase. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 1997, 1357, 49-56.	4.1	2

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55	N-Dibenzylphospho-N ^ε -3-(2,6-dichlorophenyl)propyl-guanidine is a bisubstrate-analog for creatine kinase. <i>BBA - Proteins and Proteomics</i> , 1997, 1342, 83-89.	2.1	5
56	Synthesis and Differential Properties of Creatine Analogues as Inhibitors for Human Creatine Kinase Isoenzymes. <i>FEBS Journal</i> , 1996, 238, 446-452.	0.2	8
57	Interaction of Apoprotein from Porcine High-Density Lipoprotein with Dimyristoyl Lecithin. 2. Nature of Lipid-Protein Interaction. <i>FEBS Journal</i> , 1976, 64, 549-563.	0.2	75
58	The Interaction of Apoprotein from Porcine High-Density Lipoprotein with Dimyristoyl Phosphatidylcholine. <i>FEBS Journal</i> , 1974, 48, 583-594.	0.2	38