

James M Bradeen

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8105346/publications.pdf>

Version: 2024-02-01

52
papers

2,315
citations

361413

20
h-index

243625

44
g-index

55
all docs

55
docs citations

55
times ranked

3092
citing authors

#	ARTICLE	IF	CITATIONS
1	Gene <i>RB</i> cloned from <i>Solanum bulbocastanum</i> confers broad spectrum resistance to potato late blight. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 9128-9133.	7.1	532
2	Feeding the future. Nature, 2013, 499, 23-24.	27.8	464
3	Plant community richness and microbial interactions structure bacterial communities in soil. Ecology, 2015, 96, 134-142.	3.2	196
4	The <i>Solanum commersonii</i> Genome Sequence Provides Insights into Adaptation to Stress Conditions and Genome Evolution of Wild Potato Relatives. Plant Cell, 2015, 27, 954-968.	6.6	149
5	Insights into organ-specific pathogen defense responses in plants: RNA-seq analysis of potato tuber- <i>Phytophthora infestans</i> interactions. BMC Genomics, 2013, 14, 340.	2.8	101
6	Higher Copy Numbers of the Potato <i>RB</i> Transgene Correspond to Enhanced Transcript and Late Blight Resistance Levels. Molecular Plant-Microbe Interactions, 2009, 22, 437-446.	2.6	92
7	Plant monocultures produce more antagonistic soil <i>Streptomyces</i> communities than high-diversity plant communities. Soil Biology and Biochemistry, 2013, 65, 304-312.	8.8	61
8	Molecular Diversity Analysis of Cultivated Carrot (<i>Daucus carota</i> L.) and Wild <i>Daucus</i> Populations Reveals a Genetically Nonstructured Composition. Journal of the American Society for Horticultural Science, 2002, 127, 383-391.	1.0	49
9	Randomly Amplified Polymorphic DNA in Bulb Onion and Its Use to Assess Inbred Integrity. Journal of the American Society for Horticultural Science, 1995, 120, 752-758.	1.0	47
10	Rdr3, a novel locus conferring black spot disease resistance in tetraploid rose: genetic analysis, LRR profiling, and SCAR marker development. Theoretical and Applied Genetics, 2010, 120, 573-585.	3.6	45
11	Blocking primers reduce co-amplification of plant DNA when studying bacterial endophyte communities. Journal of Microbiological Methods, 2015, 117, 1-3.	1.6	43
12	Effects of plant host species and plant community richness on streptomycete community structure. FEMS Microbiology Ecology, 2013, 83, 596-606.	2.7	39
13	The Fractionated Orthology of Bs2 and Rx/Gpa2 Supports Shared Synteny of Disease Resistance in the Solanaceae. Genetics, 2009, 182, 1351-1364.	2.9	38
14	Common Scab Trials of Potato Varieties and Advanced Selections at Three U.S. Locations. American Journal of Potato Research, 2010, 87, 261-276.	0.9	36
15	Evolutionary Meta-Analysis of Solanaceous Resistance Gene and <i>Solanum</i> Resistance Gene Analog Sequences and a Practical Framework for Cross-Species Comparisons. Molecular Plant-Microbe Interactions, 2012, 25, 603-612.	2.6	33
16	Contrasting Potato Foliage and Tuber Defense Mechanisms against the Late Blight Pathogen <i>Phytophthora infestans</i> . PLoS ONE, 2016, 11, e0159969.	2.5	29
17	A consensus "Honeycrisp" apple (<i>Malus domestica</i>) genetic linkage map from three full-sib progeny populations. Tree Genetics and Genomes, 2014, 10, 627-639.	1.6	27
18	Changes in Disease Resistance Phenotypes Associated With Plant Physiological Age Are Not Caused by Variation in <i>R</i> Gene Transcript Abundance. Molecular Plant-Microbe Interactions, 2009, 22, 362-368.	2.6	26

#	ARTICLE	IF	CITATIONS
19	Region-Specific <i>Cis</i> - and <i>Trans</i> -Acting Factors Contribute to Genetic Variability in Meiotic Recombination in Maize. <i>Genetics</i> , 1997, 146, 1101-1113.	2.9	24
20	Restriction fragment length polymorphisms reveal considerable nuclear divergence within a well-supported maternal clade in <i>Allium</i> section <i>Cepa</i> (Alliaceae). <i>American Journal of Botany</i> , 1995, 82, 1455-1462.	1.7	20
21	Mapping a Novel Black Spot Resistance Locus in the Climbing Rose Brite Eyes [®] (RADbrite [™]). <i>Frontiers in Plant Science</i> , 2018, 9, 1730.	3.6	20
22	Variability for Restriction Fragment-length Polymorphisms (RFLPs) and Relationships among Elite Commercial Inbred and Virtual Hybrid Onion Populations. <i>Journal of the American Society for Horticultural Science</i> , 1998, 123, 1034-1037.	1.0	16
23	Distribution of Rose Black Spot (<i>Diplocarpon rosae</i>) Genetic Diversity in Eastern North America Using Amplified Fragment Length Polymorphism and Implications for Resistance Screening. <i>Journal of the American Society for Horticultural Science</i> , 2007, 132, 534-540.	1.0	16
24	Fruit Texture Phenotypes of the RosBREED U.S. Apple Reference Germplasm Set. <i>Hortscience: A Publication of the American Society for Horticultural Science</i> , 2013, 48, 296-303.	1.0	15
25	Genome Microscale Heterogeneity among Wild Potatoes Revealed by Diversity Arrays Technology Marker Sequences. <i>International Journal of Genomics</i> , 2013, 2013, 1-9.	1.6	14
26	Prediction of Genotypic Values for Apple Fruit Texture Traits in a Breeding Population Derived from Honeycrisp [™] . <i>Journal of the American Society for Horticultural Science</i> , 2011, 136, 408-414.	1.0	14
27	Mapping the black spot resistance locus Rdr3 in the shrub rose "George Vancouver" allows for the development of improved diagnostic markers for DNA-informed breeding. <i>Theoretical and Applied Genetics</i> , 2020, 133, 2011-2020.	3.6	12
28	A DArT marker-based linkage map for wild potato <i>Solanum bulbocastanum</i> facilitates structural comparisons between <i>Solanum</i> A and B genomes. <i>BMC Genetics</i> , 2014, 15, 123.	2.7	11
29	Pushing the boundaries of resistance: insights from <i>Brachypodium</i> -rust interactions. <i>Frontiers in Plant Science</i> , 2015, 6, 558.	3.6	11
30	Characterization of black spot resistance in diploid roses with QTL detection, meta-analysis and candidate-gene identification. <i>Theoretical and Applied Genetics</i> , 2020, 133, 3299-3321.	3.6	11
31	Coexpression gene network analysis of cold-tolerant <i>Solanum commersonii</i> reveals new insights in response to low temperatures. <i>Crop Science</i> , 2021, 61, 3538-3550.	1.8	11
32	Towards Efficient Isolation of R Gene Orthologs from Multiple Genotypes: Optimization of Long Range-PCR. <i>Molecular Breeding</i> , 2006, 17, 137-148.	2.1	10
33	Characterization of the defence response to <i>Venturia inaequalis</i> in Honeycrisp [™] apple, its ancestors, and progeny. <i>European Journal of Plant Pathology</i> , 2014, 140, 69-81.	1.7	10
34	Molecular Linkage Maps. , 2011, , 68-89.		10
35	Resistance traits and AFLP characterization of diploid primitive tuber-bearing potatoes. <i>Genetic Resources and Crop Evolution</i> , 2007, 54, 1797-1806.	1.6	9
36	An Evaluation of Two Seedling Phenotyping Protocols to Assess pH Adaptability in Deciduous Azalea (<i>Rhododendron</i> sect. <i>Pentanthera</i> G. Don). <i>Hortscience: A Publication of the American Society for Horticultural Science</i> , 2018, 53, 268-274.	1.0	9

#	ARTICLE	IF	CITATIONS
37	Carrot. , 2007, , 161-184.		9
38	Molecular Markers and Mapping in Bulb Onion, A Forgotten Monocot. Hortscience: A Publication of the American Society for Horticultural Science, 1996, 31, 1116-1118.	1.0	9
39	Disease resistance gene transcription in transgenic potato is unaltered by temperature extremes and plant physiological age. European Journal of Plant Pathology, 2011, 130, 469-476.	1.7	7
40	Potato Tuber Blight Resistance Phenotypes Correlate with RB Transgene Transcript Levels in an Age-Dependent Manner. Phytopathology, 2015, 105, 1131-1136.	2.2	7
41	Introduction to Potato. , 2011, , 1-19.		7
42	A Novel Class of Simple PCR Markers with SNP-Level Sensitivity for Mapping and Haplotype Characterization in Solanum Species. American Journal of Potato Research, 2011, 88, 269-282.	0.9	5
43	Cloning of Late Blight Resistance Genes. , 2011, , 153-183.		5
44	An Updated Host Differential Due to Two Novel Races of Diplocarpon rosae Wolf, the Causal Agent of Rose Black Spot Disease. Hortscience: A Publication of the American Society for Horticultural Science, 2020, 55, 1756-1758.	1.0	3
45	The many-faced Janus of plant breeding. Plants People Planet, 2019, 1, 306-309.	3.3	2
46	Herbicide tolerance in primitive diploid potato species comprising superseriesstellata: Toward establishment of seedling cultivation conditions for field evaluations. American Journal of Potato Research, 2007, 84, 415.	0.9	1
47	On the Value of Wild Solanum Species for Improved Crop Disease Resistance: Resistances to Nematodes and Viruses. Compendium of Plant Genomes, 2021, , 95-118.	0.5	1
48	AFLP-derived, Codominant Markers for Locus-specific Applications. Hortscience: A Publication of the American Society for Horticultural Science, 1998, 33, 514e-514.	1.0	1
49	A Review of Allium Section Allium.. Systematic Botany, 1997, 22, 593.	0.5	0
50	PHYLOGENETIC ASSESSMENT IN THE GENUS ALLIUM USING RESTRICTION FRAGMENT LENGTH POLYMORPHISMS. Hortscience: A Publication of the American Society for Horticultural Science, 1992, 27, 611e-611.	1.0	0
51	Toward Characterization of and Breeder-friendly Molecular Markers for Genes Affecting Carotene accumulation in Carrot (Daucus carota). Hortscience: A Publication of the American Society for Horticultural Science, 1997, 32, 512D-512.	1.0	0
52	511 Toward Mapping and Cloning Late Blight Resistance Derived from the Wild Solanum bulbocastanum using Potato + S. bulbocastanum Somatic Hybrids. Hortscience: A Publication of the American Society for Horticultural Science, 1999, 34, 533E-534.	1.0	0