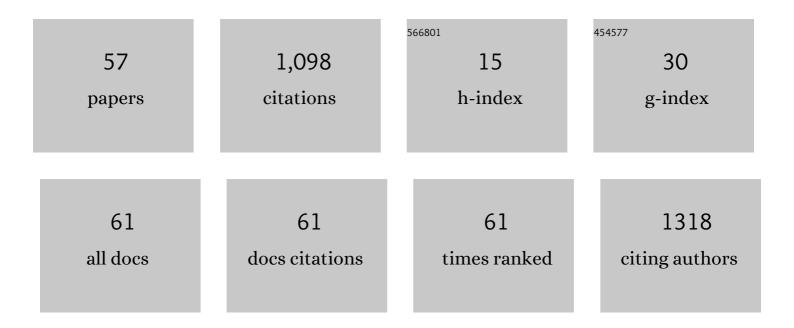
Karen R Harris-Shultz

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Pearl millet genome sequence provides a resource to improve agronomic traits in arid environments. Nature Biotechnology, 2017, 35, 969-976.	9.4	356
2	The Dynamic Changes of the Plasma Membrane Proteins and the Protective Roles of Nitric Oxide in Rice Subjected to Heavy Metal Cadmium Stress. Frontiers in Plant Science, 2016, 7, 190.	1.7	66
3	Exogenous melatonin improves cotton (Cossypium hirsutum L.) pollen fertility under drought by regulating carbohydrate metabolism in male tissues. Plant Physiology and Biochemistry, 2020, 151, 579-588.	2.8	54
4	A major QTL associated with Fusarium oxysporum race 1 resistance identified in genetic populations derived from closely related watermelon lines using selective genotyping and genotyping-by-sequencing for SNP discovery. Theoretical and Applied Genetics, 2014, 127, 2105-2115.	1.8	53
5	Microsatellite Markers Reveal a Predominant Sugarcane Aphid (Homoptera: Aphididae) Clone is Found on Sorghum in Seven States and One Territory of the USA. Crop Science, 2017, 57, 2064-2072.	0.8	41
6	Surveying the genome and constructing a high-density genetic map of napiergrass (Cenchrus) Tj ETQq0 0 0 rg	BT /Qverloc	k 19 Tf 50 54
7	Molecular evolution of the plant ECERIFERUM1 and ECERIFERUM3 genes involved in aliphatic hydrocarbon production. Computational Biology and Chemistry, 2019, 80, 1-9.	1.1	26
8	Development, Linkage Mapping, and Use of Microsatellites in Bermudagrass. Journal of the American Society for Horticultural Science, 2010, 135, 511-520.	0.5	26
9	Genetic Relationships in <i>Zoysia</i> Species and the Identification of Putative Interspecific Hybrids Using Simple Sequence Repeat Markers and Inflorescence Traits. Crop Science, 2013, 53, 285-295.	0.8	24
10	Carbon Monoxide Interacts with Auxin and Nitric Oxide to Cope with Iron Deficiency in Arabidopsis. Frontiers in Plant Science, 2016, 7, 112.	1.7	23
11	Assessment of Genetic Diversity in Napier Grass (Pennisetum purpureum Schum.) using Microsatellite, Single-Nucleotide Polymorphism and Insertion-Deletion Markers from Pearl Millet (Pennisetum) Tj ETQq1 1 0.7	843 1.4 0rgB ⁻	「/Ozværlock 10
12	Phylogenetic analysis reveals multiple introductions of Cynodon species in Australia. Molecular Phylogenetics and Evolution, 2012, 65, 390-396.	1.2	21
13	DNA Polymorphisms at Bermudagrass Microsatellite Loci and Their Use in Genotype Fingerprinting. Crop Science, 2011, 51, 1122-1131.	0.8	20
14	Analysis of genetic diversity and population structure of peanut cultivars and breeding lines from China, India and the US using simple sequence repeat markers. Journal of Integrative Plant Biology, 2016, 58, 452-465.	4.1	18
15	Genetic and phenotypic diversity of Fusarium oxysporum f. sp. niveum populations from watermelon in the southeastern United States. PLoS ONE, 2019, 14, e0219821.	1.1	18
16	Identification of Simple Sequence Repeat Markers that Differentiate Bermudagrass Cultivars Derived from â€Tifgreen'. Journal of the American Society for Horticultural Science, 2011, 136, 211-218.	0.5	18
17	Chemical Analysis of Fermentable Sugars and Secondary Products in 23 Sweet Sorghum Cultivars. Journal of Agricultural and Food Chemistry, 2017, 65, 7629-7637.	2.4	15
18	Mapping QTLs and Identification of Genes Associated with Drought Resistance in Sorghum. Methods in	0.4	15

Molecular Biology, 2019, 1931, 11-40.

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19	The Genes <i>Bm2</i> and <i>Blmc</i> that Affect Epicuticular Wax Deposition in Sorghum are Allelic. Crop Science, 2017, 57, 1552-1556.	0.8	13
20	Use of sequence-related amplified polymorphism (SRAP) markers for comparing levels of genetic diversity in centipedegrass (Eremochloa ophiuroides (Munro) Hack.) germplasm. Genetic Resources and Crop Evolution, 2012, 59, 1517-1526.	0.8	12
21	Electrochemical Evaluation of Sweet Sorghum Fermentable Sugar Bioenergy Feedstock. ACS Sustainable Chemistry and Engineering, 2017, 5, 7352-7364.	3.2	12
22	Development of Simple Sequence Repeat Markers and the Analysis of Genetic Diversity and Ploidy Level in a Centipedegrass Collection. Crop Science, 2012, 52, 383-392.	0.8	11
23	Population Structure and Genetic Diversity of Phytophthora nicotianae from Tobacco in Georgia. Plant Disease, 2017, 101, 1113-1118.	0.7	11
24	A Sugarcane Aphid "Superâ€Clone―Predominates on Sorghum and Johnsongrass from Four US States. Crop Science, 2018, 58, 2533-2541.	0.8	11
25	Transferability of SSR and RGA Markers Developed in Cynodon spp. to Zoysia spp Plant Molecular Biology Reporter, 2012, 30, 1264-1269.	1.0	10
26	Stand Maintenance and Genetic Diversity of Bermudagrass Pastures under Different Grazing Management Strategies during a 38‥ear Period. Crop Science, 2011, 51, 2886-2894.	0.8	9
27	Inheritance and Identification of a Major Quantitative Trait Locus (QTL) that Confers Resistance to <i>Meloidogyne incognita</i> and a Novel QTL for Plant Height in Sweet Sorghum. Phytopathology, 2015, 105, 1522-1528.	1.1	9
28	ldentification of fungal pathogens and analysis of genetic diversity of Fusarium tricinctum causing root rots of alfalfa in northâ€east China. Plant Pathology, 2021, 70, 804-814.	1.2	9
29	Development and Characterization of Microsatellite Markers for a Little Bluestem Collection. Journal of the American Society for Horticultural Science, 2015, 140, 78-87.	0.5	9
30	Creation of Hexaploid and Octaploid Zoysiagrass Using Colchicine and Breeding. Crop Science, 2013, 53, 2218-2224.	0.8	8
31	Use of Benzimidazole Agar Plates to Assess Fall Armyworm (Lepidoptera: Noctuidae) Feeding on Excised Maize and Sorghum Leaves. Florida Entomologist, 2015, 98, 394-397.	0.2	8
32	Evidence of Pollinators Foraging on Centipedegrass Inflorescences. Insects, 2020, 11, 795.	1.0	8
33	Development and Characterization of Seashore Paspalum SSR Markers. Crop Science, 2013, 53, 2679-2685.	0.8	7
34	Effects of Genotype and Isolate on Expression of Dollar Spot in Seashore Paspalum. Hortscience: A Publication of the American Society for Hortcultural Science, 2016, 51, 67-73.	0.5	7
35	Genome-wide association mapping of resistance to the sorghum aphid in Sorghum bicolor. Genomics, 2022, 114, 110408.	1.3	7
36	The Environment Strongly Affects Estimates of Heterosis in Hybrid Sweet Sorghum. Sugar Tech, 2018, 20, 261-274.	0.9	6

#	Article	IF	CITATIONS
37	Evaluation of strains of <i>Beauveria bassiana</i> and <i>Isaria fumosorosea</i> to control sugarcane aphids on grain sorghum. , 2020, 3, e20047.		6
38	Assessing spatio-temporal patterns of sugarcane aphid (Hemiptera: Aphididae) infestations on silage sorghum yield using unmanned aerial systems (UAS). Crop Protection, 2021, 146, 105681.	1.0	6
39	Incidence and Abundance of Bees and Wasps (Hymenoptera) in Centipedegrass Lawns in Georgia. Journal of Entomological Science, 2020, 55, 547-559.	0.2	6
40	USVL-370, a Zucchini yellow mosaic virus–resistant Watermelon Breeding Line. Hortscience: A Publication of the American Society for Hortcultural Science, 2016, 51, 107-109.	0.5	6
41	Detection of DNA and Ploidy Variation within Vegetatively Propagated Zoysiagrass Cultivars. Journal of the American Society for Horticultural Science, 2014, 139, 547-552.	0.5	6
42	Melanaphis sorghi (Hemiptera: Aphididae) Clonal Diversity in the United States and Brazil. Insects, 2022, 13, 416.	1.0	6
43	Seed Sourcing for Longleaf Pine Herbaceous Understory Restoration: Little Bluestem (Schizachyrium) Tj ETQq1 1 380.	0.784314 0.2	rgBT /Overlo 5
44	A Sugarcane Aphid (Hemiptera: Aphididae) "Super-Clone―Remains on U.S. Sorghum and Johnsongrass and Feeds on Giant Miscanthus. Journal of Entomological Science, 2021, 56, 43-52.	0.2	5
45	Transcript responses to drought in Kentucky bluegrass (Poa pratensis L.) germplasm varying in their tolerance to drought stress. Environmental and Experimental Botany, 2021, 190, 104571.	2.0	5
46	Discovery and Characterization of a Turf-type Triploid Seashore Paspalum. Hortscience: A Publication of the American Society for Hortcultural Science, 2013, 48, 1424-1427.	0.5	5
47	Method for DNA Isolation From Sweetpotato Weevil (Coleoptera: Curculionidae) Collected in Pheromone-Baited Traps. Journal of Economic Entomology, 2019, 112, 1001-1003.	0.8	4
48	Juice chemical properties of 24 sorghum cultivars under varying levels of sugarcane aphid (Melanaphis sacchari) infestation. Arthropod-Plant Interactions, 2021, 15, 707-719.	0.5	4
49	The sorghum epicuticular wax locus Bloomless2 reduces plant damage in P898012 caused by the sugarcane aphid. , 2020, 3, e20008.		3
50	Evaluation of Seashore Paspalum Germplasm for Resistance to Dollar Spot. Itsrj, 2017, 13, 175-184.	0.1	2
51	Colonial bentgrass transcriptâ€expression differences compared with creeping bentgrass in response to waterâ€deficit stress. Crop Science, 2021, 61, 2135-2147.	0.8	2
52	Evaluation of Whorl Damage by Fall Armyworm (Lepidoptera: Noctuidae) on Field- and Greenhouse-Grown Sweet Sorghum Plants. Journal of Entomological Science, 2015, 50, 14-27.	0.2	1
53	A Novel QTL for Root-Knot Nematode Resistance is Identified from a South African Sweet Sorghum Line. Phytopathology, 2019, 109, 1011-1017.	1.1	1

54 Registration of â€~MSBâ€264' and â€~MSBâ€285' bermudagrasses. Journal of Plant Registrations, 2022, 16,485-197.

#	Article	IF	CITATIONS
55	Development and characterization of microsatellites from the sweetpotato weevil, Cylas formicarius elegantulus. Journal of Applied Entomology, 2020, 144, 335-340.	0.8	0
56	Sugarcane aphid resistance in sorghum and its potential bioenergy grass hosts. , 2016, , .		0
57	Transfer of Meloidogyne incognita Resistance Using Marker-assisted Selection in Sorghum. Journal of Nematology, 2021, 53, 1-10.	0.4	0