

Leslie R Goertzen

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	CYTOKININ RESPONSE FACTOR 2 is involved in modulating the salt stress response. <i>Plant Journal</i> , 2022, 110, 1097-1110.	2.8	10
2	Detection of subgenome bias using an anchored syntenic approach in <i>Eleusine coracana</i> (finger) Tj ETQq0 0 0 rgBT, /Overlock, 10 Tf 50 7	1.2	2
3	The Vascular Flora of Chewacla State Park, Lee County, Alabama. <i>Castanea</i> , 2020, 85, 169.	0.2	0
4	Identification and functional characterization of the <i>Marshallia</i> (Asteraceae) Clade III Cytokinin Response Factor (CRF). <i>Plant Signaling and Behavior</i> , 2019, 14, e1633886.	1.2	5
5	Lineage specific conservation of cis-regulatory elements in Cytokinin Response Factors. <i>Scientific Reports</i> , 2019, 9, 13387.	1.6	11
6	Transcriptome Analysis Reveals Unique Relationships Among <i>Eleusine</i> Species and Heritage of <i>Eleusine coracana</i> . <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 2029-2036.	0.8	18
7	Development of a goosegrass (<i>Eleusine indica</i>) draft genome and application to weed science research. <i>Pest Management Science</i> , 2019, 75, 2776-2784.	1.7	29
8	The Mitochondrial Genome of <i>Eleusine indica</i> and Characterization of Gene Content Within Poaceae. <i>Genome Biology and Evolution</i> , 2019, 12, 3684-3697.	1.1	3
9	Cytokinin Response Factor 5 has transcriptional activity governed by its C-terminal domain. <i>Plant Signaling and Behavior</i> , 2017, 12, e1276684.	1.2	22
10	Complete plastid genome sequence of goosegrass (<i>Eleusine indica</i>) and comparison with other Poaceae. <i>Gene</i> , 2017, 600, 36-43.	1.0	15
11	Vascular Expression and C-Terminal Sequence Divergence of Cytokinin Response Factors in Flowering Plants. <i>Plant and Cell Physiology</i> , 2012, 53, 1683-1695.	1.5	39
12	The CRF domain defines Cytokinin Response Factor proteins in plants. <i>BMC Plant Biology</i> , 2010, 10, 74.	1.6	86
13	Horticulture, hybrid cultivars and exotic plant invasion: a case study of <i>Wisteria</i> (Fabaceae). <i>Botanical Journal of the Linnean Society</i> , 2008, 158, 593-601.	0.8	16
14	Bi-Parental Cytoplasmic DNA Inheritance in <i>Wisteria</i> (Fabaceae): Evidence from a Natural Experiment. <i>Plant and Cell Physiology</i> , 2007, 48, 662-665.	1.5	17
15	Invasive <i>Wisteria</i> in the Southeastern United States: genetic diversity, hybridization and the role of urban centers. <i>Urban Ecosystems</i> , 2007, 10, 379-395.	1.1	16
16	Massive horizontal transfer of mitochondrial genes from diverse land plant donors to the basal angiosperm <i>Amborella</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 17747-17752.	3.3	240
17	ITS secondary structure derived from comparative analysis: implications for sequence alignment and phylogeny of the Asteraceae. <i>Molecular Phylogenetics and Evolution</i> , 2003, 29, 216-234.	1.2	141
18	EFFECT OF TAXON SAMPLING, CHARACTER WEIGHTING, AND COMBINED DATA ON THE INTERPRETATION OF RELATIONSHIPS AMONG THE HETEROKONT ALGAE. <i>Journal of Phycology</i> , 2003, 39, 423-443.	1.0	35

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19	The Complete External Transcribed Spacer of 18S-26S rDNA: Amplification and Phylogenetic Utility at Low Taxonomic Levels in Asteraceae and Closely Allied Families. <i>Molecular Phylogenetics and Evolution</i> , 2000, 14, 285-303.	1.2	147
20	Molecular Systematics of the Asteriscus Alliance (Asteraceae: Inuleae) I: Evidence from the Internal Transcribed Spacers of Nuclear Ribosomal DNA. <i>Systematic Botany</i> , 1999, 24, 249.	0.2	33
21	The defensive role of trichomes in black medick (<i>Medicago lupulina</i> , Fabaceae). <i>Plant Systematics and Evolution</i> , 1993, 184, 101-111.	0.3	21