

Lawrence B Smart

List of Publications by Year in descending order

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Version: 2024-02-01

80
papers

3,279
citations

136950

32
h-index

168389

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92
all docs

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docs citations

92
times ranked

3298
citing authors

#	ARTICLE	IF	CITATIONS
1	Heterosis for Biomass-Related Traits in Interspecific Triploid Hybrids of Willow (<i>Salix</i> spp.). <i>Bioenergy Research</i> , 2022, 15, 1042-1056.	3.9	5
2	Comparative transcriptomics and eQTL mapping of response to <i>Melampsora americana</i> in selected <i>Salix purpurea</i> F2 progeny. <i>BMC Genomics</i> , 2022, 23, 71.	2.8	5
3	Mapping the sex determination region in the <i>Salix</i> F1 hybrid common parent population confirms a ZW system in six diverse species. <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	1.8	9
4	Nonadditive gene expression is correlated with nonadditive phenotypic expression in interspecific triploid hybrids of willow (<i>Salix</i> spp.). <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	1.8	3
5	Effects of Cold Temperature and Acclimation on Cold Tolerance and Cannabinoid Profiles of <i>Cannabis sativa</i> L. (Hemp). <i>Horticulturae</i> , 2022, 8, 531.	2.8	7
6	Genetic mapping of sexually dimorphic volatile and non-volatile floral secondary chemistry of a dioecious willow. <i>Journal of Experimental Botany</i> , 2022, 73, 6352-6366.	4.8	8
7	A General Model to Explain Repeated Turnovers of Sex Determination in the Salicaceae. <i>Molecular Biology and Evolution</i> , 2021, 38, 968-980.	8.9	53
8	Seasonal characterization of high-cannabinoid hemp (<i>Cannabis sativa</i> L.) reveals variation in cannabinoid accumulation, flowering time, and disease resistance. <i>GCB Bioenergy</i> , 2021, 13, 546-561.	5.6	50
9	Yield and biomass quality of shrub willow hybrids in differing rotation lengths and spacing designs. <i>Biomass and Bioenergy</i> , 2021, 146, 105977.	5.7	4
10	Introduction: United States Hemp Research and Education Conference special issue. <i>GCB Bioenergy</i> , 2021, 13, 516-516.	5.6	0
11	A semi-commercial case study of willow biomass production in the northeastern United States. <i>Agronomy Journal</i> , 2021, 113, 1287-1302.	1.8	3
12	Limited effect of environmental stress on cannabinoid profiles in high-cannabidiol hemp (<i>Cannabis</i>) Tj ETQq0 0.0 rgBT /Overlock 10	5.6	32
13	Morphometric relationships and their contribution to biomass and cannabinoid yield in hybrids of hemp (<i>Cannabis sativa</i>). <i>Journal of Experimental Botany</i> , 2021, 72, 7694-7709.	4.8	18
14	Integrative genomics reveals paths to sex dimorphism in <i>Salix purpurea</i> L. <i>Horticulture Research</i> , 2021, 8, 170.	6.3	12
15	Sexual dimorphism in the dioecious willow <i>Salix purpurea</i> . <i>American Journal of Botany</i> , 2021, 108, 1374-1387.	1.7	14
16	The <i>Melampsora americana</i> population on <i>Salix purpurea</i> in the Great Lakes region is highly diverse with a contributory influence of clonality. <i>Phytopathology</i> , 2021, , .	2.2	2
17	Sex determination through X-Y heterogamety in <i>Salix nigra</i> . <i>Heredity</i> , 2021, 126, 630-639.	2.6	26
18	Microbiome of Field Grown Hemp Reveals Potential Microbial Interactions With Root and Rhizosphere Soil. <i>Frontiers in Microbiology</i> , 2021, 12, 741597.	3.5	9

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19	Phylogenomics of the genus <i>Populus</i> reveals extensive interspecific gene flow and balancing selection. <i>New Phytologist</i> , 2020, 225, 1370-1382.	7.3	93
20	Evaluating the Microbiome of Hemp. <i>Phytobiomes Journal</i> , 2020, 4, 351-363.	2.7	12
21	Transcriptome analysis of contrasting resistance to herbivory by <i>Empoasca fabae</i> in two shrub willow species and their hybrid progeny. <i>PLoS ONE</i> , 2020, 15, e0236586.	2.5	4
22	Differential Susceptibility of Diverse <i>Salix</i> spp. to <i>Melampsora americana</i> and <i>Melampsora paradoxa</i> . <i>Plant Disease</i> , 2020, 104, 2949-2957.	1.4	9
23	Developing Production Guidelines for Baby Leaf Hemp (<i>Cannabis sativa</i> L.) as an Edible Salad Green: Cultivar, Sowing Density and Seed Size. <i>Agriculture (Switzerland)</i> , 2020, 10, 617.	3.1	10
24	Genotypic diversity in willow (<i>Salix</i> spp.) is associated with chemical and morphological polymorphism, suggesting human-assisted dissemination in the Eastern Mediterranean. <i>Biochemical Systematics and Ecology</i> , 2020, 91, 104081.	1.3	6
25	A willow sex chromosome reveals convergent evolution of complex palindromic repeats. <i>Genome Biology</i> , 2020, 21, 38.	8.8	74
26	Pathways to sex determination in plants: how many roads lead to Rome?. <i>Current Opinion in Plant Biology</i> , 2020, 54, 61-68.	7.1	54
27	Development and validation of genetic markers for sex and cannabinoid chemotype in <i>Cannabis sativa</i> L.. <i>GCB Bioenergy</i> , 2020, 12, 213-222.	5.6	77
28	Genetic and Environmental Influences on First Rotation Shrub Willow (<i>Salix</i> spp.) Bark and Wood Elemental Composition. <i>Bioenergy Research</i> , 2020, 13, 797-809.	3.9	4
29	Cross-Infectivity of Powdery Mildew Isolates Originating from Hemp (<i>Cannabis sativa</i>) and Japanese Hop (<i>Humulus japonicus</i>) in New York. <i>Plant Health Progress</i> , 2020, 21, 47-53.	1.4	30
30	Genetic diversity and population structure of native, naturalized, and cultivated <i>Salix purpurea</i> . <i>Tree Genetics and Genomes</i> , 2019, 15, 1.	1.6	13
31	Joint linkage and association mapping of complex traits in shrub willow (<i>Salix purpurea</i> L.). <i>Annals of Botany</i> , 2019, 124, 701-715.	2.9	37
32	Tolerance of novel inter-specific shrub willow hybrids to water stress. <i>Trees - Structure and Function</i> , 2019, 33, 1015-1026.	1.9	12
33	Breeding progress and preparedness for mass-scale deployment of perennial lignocellulosic biomass crops switchgrass, miscanthus, willow and poplar. <i>GCB Bioenergy</i> , 2019, 11, 118-151.	5.6	116
34	Discovery of Geographically Robust Hybrid Poplar Clones. <i>Silvae Genetica</i> , 2019, 68, 101-110.	0.8	7
35	Differential growth response to fertilization of ten elite shrub willow (<i>Salix</i> spp.) bioenergy cultivars. <i>Trees - Structure and Function</i> , 2018, 32, 1061-1072.	1.9	8
36	Poplar and shrub willow energy crops in the United States: field trial results from the multiyear regional feedstock partnership and yield potential maps based on the PRISM-ELM model. <i>GCB Bioenergy</i> , 2018, 10, 735-751.	5.6	54

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37	Effects of nitrogen fertilization in shrub willow short rotation coppice production – a quantitative review. <i>GCB Bioenergy</i> , 2018, 10, 548-564.	5.6	34
38	Hardwood Tree Genomics: Unlocking Woody Plant Biology. <i>Frontiers in Plant Science</i> , 2018, 9, 1799.	3.6	50
39	Characterization of a large sex determination region in <i>Salix purpurea</i> L. (Salicaceae). <i>Molecular Genetics and Genomics</i> , 2018, 293, 1437-1452.	2.1	61
40	Genotype × environment interaction analysis of North American shrub willow yield trials confirms superior performance of triploid hybrids. <i>GCB Bioenergy</i> , 2017, 9, 445-459.	5.6	41
41	A mixed model approach for evaluating yield improvements in interspecific hybrids of shrub willow, a dedicated bioenergy crop. <i>Industrial Crops and Products</i> , 2017, 96, 57-70.	5.2	19
42	Contributions of environment and genotype to variation in shrub willow biomass composition. <i>Industrial Crops and Products</i> , 2017, 108, 149-161.	5.2	29
43	Dominance and Sexual Dimorphism Pervade the <i>Salix purpurea</i> L. Transcriptome. <i>Genome Biology and Evolution</i> , 2017, 9, 2377-2394.	2.5	35
44	Electrical capacitance as a predictor of root dry weight in shrub willow (<i>Salix</i> ; Salicaceae) parents and progeny. <i>Applications in Plant Sciences</i> , 2016, 4, 1600031.	2.1	10
45	Untapped Potential: Opportunities and Challenges for Sustainable Bioenergy Production from Marginal Lands in the Northeast USA. <i>Bioenergy Research</i> , 2015, 8, 482-501.	3.9	79
46	Variability in pyrolysis product yield from novel shrub willow genotypes. <i>Biomass and Bioenergy</i> , 2015, 72, 74-84.	5.7	13
47	Ploidy Level Affects Important Biomass Traits of Novel Shrub Willow (<i>Salix</i>) Hybrids. <i>Bioenergy Research</i> , 2015, 8, 259-269.	3.9	47
48	Whole-Genome Sequences of 13 Endophytic Bacteria Isolated from Shrub Willow (<i>Salix</i>) Grown in Geneva, New York. <i>Genome Announcements</i> , 2014, 2, .	0.8	25
49	Early selection of novel triploid hybrids of shrub willow with improved biomass yield relative to diploids. <i>BMC Plant Biology</i> , 2014, 14, 74.	3.6	50
50	Genetic evidence for three discrete taxa of <i>Melampsora</i> (Pucciniales) affecting willows (<i>Salix</i> spp.) in New York State. <i>Fungal Biology</i> , 2014, 118, 704-720.	2.5	12
51	Yield and Woody Biomass Traits of Novel Shrub Willow Hybrids at Two Contrasting Sites. <i>Bioenergy Research</i> , 2013, 6, 533-546.	3.9	92
52	Enzymatic saccharification of shrub willow genotypes with differing biomass composition for biofuel production. <i>Frontiers in Plant Science</i> , 2013, 4, 57.	3.6	39
53	Correlations of expression of cell wall biosynthesis genes with variation in biomass composition in shrub willow (<i>Salix</i> spp.) biomass crops. <i>Tree Genetics and Genomes</i> , 2012, 8, 775-788.	1.6	17
54	Biological conversion assay using <i>Clostridium</i> phytofermentans to estimate plant feedstock quality. <i>Biotechnology for Biofuels</i> , 2012, 5, 5.	6.2	28

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55	A Molecular and Fitness Evaluation of Commercially Available versus Locally Collected Blue Lupine (<i>Lupinus perennis</i> L. Seeds for Use in Ecosystem Restoration Efforts. Restoration Ecology, 2012, 20, 456-461.	2.9	10
56	Differential expression of genes encoding phosphate transporters contributes to arsenic tolerance and accumulation in shrub willow (Salix spp.). Environmental and Experimental Botany, 2012, 75, 248-257.	4.2	41
57	Shrub Willow. , 2012, , 687-708.		11
58	Analysis of Biomass Composition Using High-Resolution Thermogravimetric Analysis and Percent Bark Content for the Selection of Shrub Willow Bioenergy Crop Varieties. Bioenergy Research, 2009, 2, 1-9.	3.9	61
59	Population genetic structure of native versus naturalized sympatric shrub willows (<i>Salix</i>); Tj ETQq1 1 0.784314 rgBT /Overlock 10	1.7	36
60	High-resolution Thermogravimetric Analysis For Rapid Characterization of Biomass Composition and Selection of Shrub Willow Varieties. Applied Biochemistry and Biotechnology, 2008, 145, 3-11.	2.9	36
61	Quantitative Genetics of Traits Indicative of Biomass Production and Heterosis in 34 Full-sib F1 Salix eriocephala Families. Bioenergy Research, 2008, 1, 80-90.	3.9	21
62	Cuticular wax composition of Salix varieties in relation to biomass productivity. Phytochemistry, 2008, 69, 396-402.	2.9	17
63	Hydroponic Screening of Shrub Willow (<i>Salix</i> Spp.) for Arsenic Tolerance and Uptake. International Journal of Phytoremediation, 2008, 10, 515-528.	3.1	44
64	Genetic Improvement of Willow (Salix spp.) as a Dedicated Bioenergy Crop. , 2008, , 377-396.		26
65	Increased Accumulation of Cuticular Wax and Expression of Lipid Transfer Protein in Response to Periodic Drying Events in Leaves of Tree Tobacco. Plant Physiology, 2006, 140, 176-183.	4.8	357
66	A second member of the Nicotiana glauca lipid transfer protein gene family, NgLTP2, encodes a divergent and differentially expressed protein. Functional Plant Biology, 2006, 33, 141.	2.1	6
67	The development of short-rotation willow in the northeastern United States for bioenergy and bioproducts, agroforestry and phytoremediation. Biomass and Bioenergy, 2006, 30, 715-727.	5.7	237
68	Collection and storage of pollen from (<i>Salix</i>) (Salicaceae). American Journal of Botany, 2002, 89, 248-252.	1.7	19
69	Predicting within-family variability in juvenile height growth of Salix based upon similarity among parental AFLP fingerprints. Theoretical and Applied Genetics, 2002, 105, 106-112.	3.6	21
70	Diversity of cuticular wax among Salix species and Populus species hybrids. Phytochemistry, 2002, 60, 715-725.	2.9	52
71	MIP Genes are Down-regulated Under Drought Stress in Nicotiana glauca. Plant and Cell Physiology, 2001, 42, 686-693.	3.1	134
72	Isolation of genes predominantly expressed in guard cells and epidermal cells of Nicotiana glauca. Plant Molecular Biology, 2000, 42, 857-869.	3.9	16

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73	Isolation of RNA and Protein from Guard Cells of <i>Nicotiana glauca</i> . <i>Plant Molecular Biology Reporter</i> , 1999, 17, 371-383.	1.8	4
74	Genes Involved in Osmoregulation during Turgor-Driven Cell Expansion of Developing Cotton Fibers Are Differentially Regulated ¹ . <i>Plant Physiology</i> , 1998, 116, 1539-1549.	4.8	177
75	Absence of PsaC subunit allows assembly of photosystem I core but prevents the binding of PsaD and PsaE in <i>Synechocystis</i> sp. PCC6803. <i>Plant Molecular Biology</i> , 1995, 29, 331-342.	3.9	70
76	A mixed-ligand iron-sulfur cluster (C556SPaB or C565SPsaB) in the Fx-binding site leads to a decreased quantum efficiency of electron transfer in photosystem I. <i>Biophysical Journal</i> , 1995, 69, 1544-1553.	0.5	42
77	Genetic inactivation of the <i>psaB</i> gene in <i>Synechocystis</i> sp. PCC 6803 disrupts assembly of photosystem I. <i>Plant Molecular Biology</i> , 1993, 21, 177-180.	3.9	31
78	Site-directed conversion of cysteine-565 to serine in PsaB of photosystem I results in the assembly of iron-sulfur [3Fe-4S] and iron-sulfur [4Fe-4S] clusters in Fx. A mixed-ligand iron-sulfur [4Fe-4S] cluster is capable of electron transfer to FA and FB. <i>Biochemistry</i> , 1993, 32, 4411-4419.	2.5	49
79	Mutational analysis of the structure and biogenesis of the photosystem I reaction center in the cyanobacterium <i>Synechocystis</i> sp. PCC 6803.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1993, 90, 1132-1136.	7.1	54
80	Expression of photosynthesis genes in the cyanobacterium <i>Synechocystis</i> sp. PCC 6803: <i>psaA-psaB</i> and <i>psbA</i> transcripts accumulate in dark-grown cells. <i>Plant Molecular Biology</i> , 1991, 17, 959-971.	3.9	55