Sara S Jawdy

List of Publications by Year in descending order

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201674 223800 2,265 48 27 46 citations h-index g-index papers 50 50 50 3382 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Forest stand and canopy development unaltered by 12Âyears of CO2 enrichment*. Tree Physiology, 2022, 42, 428-440.	3.1	12
2	Development of an Experimental Approach to Achieve Spatially Resolved Plant Root-Associated Metaproteomics Using an Agar-Plate System. Molecular Plant-Microbe Interactions, 2022, 35, 639-649.	2.6	3
3	Habitatâ€adapted microbial communities mediate <i>Sphagnum</i> peatmoss resilience to warming. New Phytologist, 2022, 234, 2111-2125.	7.3	18
4	Cultivating the Bacterial Microbiota of <i>Populus</i> Roots. MSystems, 2021, 6, e0130620.	3.8	17
5	Overexpression of a <i>Prefoldin \hat{l}^2</i> subunit gene reduces biomass recalcitrance in the bioenergy crop <i>Populus</i> . Plant Biotechnology Journal, 2020, 18, 859-871.	8.3	17
6	Identification of functional single nucleotide polymorphism of Populus trichocarpa PtrEPSPâ€₹F and determination of its transcriptional effect. Plant Direct, 2020, 4, e00178.	1.9	4
7	Mediation of plant–mycorrhizal interaction by a lectin receptor-like kinase. Nature Plants, 2019, 5, 676-680.	9.3	42
8	Relatively rare root endophytic bacteria drive plant resource allocation patterns and tissue nutrient concentration in unpredictable ways. American Journal of Botany, 2019, 106, 1423-1434.	1.7	9
9	Overexpression of a serine hydroxymethyltransferase increases biomass production and reduces recalcitrance in the bioenergy crop <i>Populus</i> Sustainable Energy and Fuels, 2019, 3, 195-207.	4.9	27
10	Identification of Populus Small RNAs Responsive to Mutualistic Interactions With Mycorrhizal Fungi, Laccaria bicolor and Rhizophagus irregularis. Frontiers in Microbiology, 2019, 10, 515.	3.5	17
11	The nature of the progression of drought stress drives differential metabolomic responses in Populus deltoides. Annals of Botany, 2019, 124, 617-626.	2.9	45
12	The Ectomycorrhizal Fungus <i>Laccaria bicolor</i> Produces Lipochitooligosaccharides and Uses the Common Symbiosis Pathway to Colonize <i>Populus</i> Roots. Plant Cell, 2019, 31, 2386-2410.	6.6	73
13	Abiotic Stresses Shift Belowground <i>Populus</i> -Associated Bacteria Toward a Core Stress Microbiome. MSystems, 2018, 3, .	3.8	89
14	Phytobiome and Transcriptional Adaptation of <i>Populus deltoides</i> to Acute Progressive Drought and Cyclic Drought. Phytobiomes Journal, 2018, 2, 249-260.	2.7	23
15	A New Calmodulin-Binding Protein Expresses in the Context of Secondary Cell Wall Biosynthesis and Impacts Biomass Properties in Populus. Frontiers in Plant Science, 2018, 9, 1669.	3.6	31
16	Association mapping, transcriptomics, and transient expression identify candidate genes mediating plant–pathogen interactions in a tree. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 11573-11578.	7.1	61
17	A Variable Polyglutamine Repeat Affects Subcellular Localization and Regulatory Activity of a <i>Populus</i> ANGUSTIFOLIA Protein. G3: Genes, Genomes, Genetics, 2018, 8, 2631-2641.	1.8	9
18	<scp>Genomeâ€wide association studies</scp> and expressionâ€based quantitative trait loci analyses reveal roles of <scp>HCT</scp> 2 in caffeoylquinic acid biosynthesis and its regulation by defenseâ€responsive transcription factors in <i>Populus</i> New Phytologist, 2018, 220, 502-516.	7.3	112

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19	A 5-Enolpyruvylshikimate 3-Phosphate Synthase Functions as a Transcriptional Repressor in <i>Populus</i>). Plant Cell, 2018, 30, 1645-1660.	6.6	56
20	Quantitative proteome profile of water deficit stress responses in eastern cottonwood (Populus) Tj ETQq0 0 C	rgBT ₂ /Overl	ock 10 Tf 50
21	Overexpression of a Domain of Unknown Function 266-containing protein results in high cellulose content, reduced recalcitrance, and enhanced plant growth in the bioenergy crop Populus. Biotechnology for Biofuels, 2017, 10, 74.	6.2	22
22	Effects of Biomass Accessibility and Klason Lignin Contents during Consolidated Bioprocessing in <i>Populus trichocarpa</i> . ACS Sustainable Chemistry and Engineering, 2017, 5, 5075-5081.	6.7	20
23	Populus trichocarpa encodes small, effector-like secreted proteins that are highly induced during mutualistic symbiosis. Scientific Reports, 2017, 7, 382.	3.3	36
24	Overexpression of a Domain of Unknown Function 231-containing protein increases O-xylan acetylation and cellulose biosynthesis in Populus. Biotechnology for Biofuels, 2017, 10, 311.	6.2	26
25	A Carotenoid-Deficient Mutant in Pantoea sp. YR343, a Bacteria Isolated from the Rhizosphere of Populus deltoides, Is Defective in Root Colonization. Frontiers in Microbiology, 2016, 7, 491.	3.5	48
26	Two Poplar-Associated Bacterial Isolates Induce Additive Favorable Responses in a Constructed Plant-Microbiome System. Frontiers in Plant Science, 2016, 7, 497.	3.6	113
27	Down-Regulation of KORRIGAN-Like Endo- \hat{l}^2 -1,4-Glucanase Genes Impacts Carbon Partitioning, Mycorrhizal Colonization and Biomass Production in Populus. Frontiers in Plant Science, 2016, 7, 1455.	3.6	32
28	Knockdown of a laccase in <i>Populus deltoides</i> confers altered cell wall chemistry and increased sugar release. Plant Biotechnology Journal, 2016, 14, 2010-2020.	8.3	64
29	Genome-wide analysis of lectin receptor-like kinases in Populus. BMC Genomics, 2016, 17, 699.	2.8	72
30	Simultaneous knockdown of six non-family genes using a single synthetic RNAi fragment in Arabidopsis thaliana. Plant Methods, 2016, 12, 16.	4.3	12
31	Root bacterial endophytes alter plant phenotype, but not physiology. PeerJ, 2016, 4, e2606.	2.0	64
32	Scaling nitrogen and carbon interactions: what are the consequences of biological buffering?. Ecology and Evolution, 2015, 5, 2839-2850.	1.9	4
33	Metabolic functions of Pseudomonas fluorescens strains from Populus deltoides depend on rhizosphere or endosphere isolation compartment. Frontiers in Microbiology, 2015, 6, 1118.	3.5	60
34	High-resolution genetic mapping of allelic variants associated with cell wall chemistry in Populus. BMC Genomics, 2015, 16, 24.	2.8	106
35	Metabolic profiling reveals altered sugar and secondary metabolism in response to UGPase overexpression in Populus. BMC Plant Biology, 2014, 14, 265.	3.6	61
36	Leaf endophytes and <i>Populus</i> genotype affect severity of damage from the necrotrophic leaf pathogen, <i>Drepanopeziza populi</i> Ecosphere, 2013, 4, 1-12.	2.2	35

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37	Initial characterization of shade avoidance response suggests functional diversity between <i>Populus</i> phytochrome B genes. New Phytologist, 2012, 196, 726-737.	7.3	25
38	<i>Pseudomonas fluorescens</i> Induces Strain-Dependent and Strain-Independent Host Plant Responses in Defense Networks, Primary Metabolism, Photosynthesis, and Fitness. Molecular Plant-Microbe Interactions, 2012, 25, 765-778.	2.6	100
39	Chemical, ultrastructural and supramolecular analysis of tension wood in Populus tremula x alba as a model substrate for reduced recalcitrance. Energy and Environmental Science, 2011, 4, 4962.	30.8	61
40	Comparative physiology and transcriptional networks underlying the heat shock response in <i>Populus trichocarpa</i> , <i>Arabidopsis thaliana</i> and <i>Glycine max</i> . Plant, Cell and Environment, 2011, 34, 1488-1506.	5.7	71
41	Discovery and annotation of small proteins using genomics, proteomics, and computational approaches. Genome Research, 2011, 21, 634-641.	5.5	105
42	Auxin Signaling and Response Mechanisms and Roles in Plant Growth and Development., 2011,, 231-254.		3
43	Bioinformatics-Based Identification of Candidate Genes from QTLs Associated with Cell Wall Traits in Populus. Bioenergy Research, 2010, 3, 172-182.	3.9	25
44	Apoplast proteome reveals that extracellular matrix contributes to multistress response in poplar. BMC Genomics, 2010, 11, 674.	2.8	70
45	Genome-wide identification of lineage-specific genes in Arabidopsis, Oryza and Populus. Genomics, 2009, 93, 473-480.	2.9	50
46	The F-Box Gene Family Is Expanded in Herbaceous Annual Plants Relative to Woody Perennial Plants Â. Plant Physiology, 2008, 148, 1189-1200.	4.8	125
47	Genetic and physical mapping of Melampsora rust resistance genes in Populus and characterization of linkage disequilibrium and flanking genomic sequence. New Phytologist, 2004, 164, 95-105.	7.3	66
48	Leaf dynamics of a deciduous forest canopy: no response to elevated CO 2. Oecologia, 2003, 136, 574-584.	2.0	106