## Amanda R De La Torre

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/788471/publications.pdf

Version: 2024-02-01

24 papers 1,071 citations

623734 14 h-index 677142 22 g-index

27 all docs

27 docs citations

times ranked

27

1489 citing authors

#	Article	IF	CITATIONS
1	Insights into Conifer Giga-Genomes. Plant Physiology, 2014, 166, 1724-1732.	4.8	164
2	Contrasting Rates of Molecular Evolution and Patterns of Selection among Gymnosperms and Flowering Plants. Molecular Biology and Evolution, 2017, 34, 1363-1377.	8.9	164
3	Functional and evolutionary genomic inferences in <i>Populus</i> through genome and population sequencing of American and European aspen. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E10970-E10978.	7.1	84
4	Single-Copy Genes as Molecular Markers for Phylogenomic Studies in Seed Plants. Genome Biology and Evolution, 2017, 9, 1130-1147.	2.5	75
5	Adaptation and exogenous selection in a <i><scp>P</scp>icea glauca</i> $\tilde{A}$ — <i><scp>P</scp>icea engelmannii</i> hybrid zone: implications for forest management under climate change. New Phytologist, 2014, 201, 687-699.	7.3	74
6	Novel Insights into Tree Biology and Genome Evolution as Revealed Through Genomics. Annual Review of Plant Biology, 2017, 68, 457-483.	18.7	64
7	Genomeâ€wide admixture and ecological niche modelling reveal the maintenance of species boundaries despite long history of interspecific gene flow. Molecular Ecology, 2014, 23, 2046-2059.	3.9	63
8	Genome-Wide Analysis Reveals Diverged Patterns of Codon Bias, Gene Expression, and Rates of Sequence Evolution in Picea Gene Families. Genome Biology and Evolution, 2015, 7, 1002-1015.	2.5	63
9	Genomic architecture of complex traits in loblolly pine. New Phytologist, 2019, 221, 1789-1801.	7.3	60
10	Environmental Genome-Wide Association Reveals Climate Adaptation Is Shaped by Subtle to Moderate Allele Frequency Shifts in Loblolly Pine. Genome Biology and Evolution, 2019, 11, 2976-2989.	2.5	54
11	Functional and morphological evolution in gymnosperms: A portrait of implicated gene families. Evolutionary Applications, 2020, 13, 210-227.	3.1	32
12	Genomic basis of white pine blister rust quantitative disease resistance and its relationship with qualitative resistance. Plant Journal, 2020, 104, 365-376.	5.7	32
13	Assembled and annotated 26.5 Gbp coast redwood genome: a resource for estimating evolutionary adaptive potential and investigating hexaploid origin. G3: Genes, Genomes, Genetics, 2022, 12, .	1.8	28
14	Fine-scale environmental variation contributes to introgression in a three-species spruce hybrid complex. Tree Genetics and Genomes, $2015, 11, 1$ .	1.6	25
15	Genetic Variation Related to High Elevation Adaptation Revealed by Common Garden Experiments in Pinus yunnanensis. Frontiers in Genetics, 2019, 10, 1405.	2.3	17
16	Genomeâ€wide association identifies candidate genes for drought tolerance in coast redwood and giant sequoia. Plant Journal, 2022, 109, 7-22.	5.7	17
17	Dissecting the Polygenic Basis of Cold Adaptation Using Genome-Wide Association of Traits and Environmental Data in Douglas-fir. Genes, 2021, 12, 110.	2.4	14

Finding loci associated to partial resistance to white pine blister rust in sugar pine (Pinus lambertiana) Tj ETQq $0.0\,^\circ$ 11 Overlock 10 Tf 11 Overlock 10

#	Article	IF	CITATIONS
19	Selective Sweeps and Polygenic Adaptation Drive Local Adaptation along Moisture and Temperature Gradients in Natural Populations of Coast Redwood and Giant Sequoia. Genes, 2021, 12, 1826.	2.4	7
20	Transcriptome-based single-nucleotide polymorphism markers between Pinus brutia and Pinus halepensis and the analysis of their hybrids. Tree Genetics and Genomes, 2021, 17, 1.	1.6	4
21	Maritime Pine Genomics in Focus. Compendium of Plant Genomes, 2022, , 67-123.	0.5	4
22	Genomics of Climate Adaptation in Pinus Lambertiana. Compendium of Plant Genomes, 2022, , 51-65.	0.5	1
23	Comparative Genomics of Spruce and Other Gymnosperms. Compendium of Plant Genomes, 2020, , 97-105.	0.5	O
24	Prospects: The Spruce Genome, a Model for Understanding Gymnosperm Evolution and Supporting Tree Improvement Efforts. Compendium of Plant Genomes, 2020, , 215-218.	0.5	0