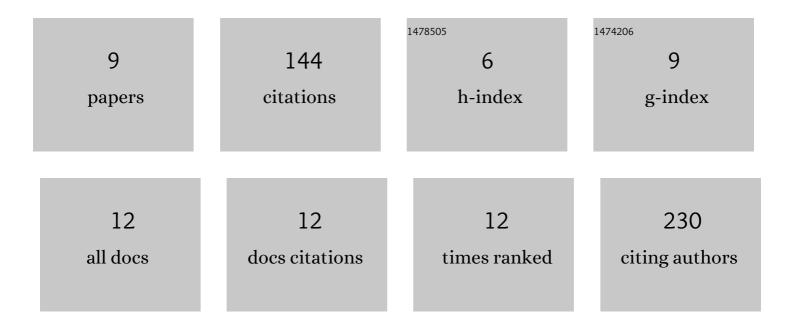
## Mengmeng Lu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11210/publications.pdf Version: 2024-02-01



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#	Article	IF	CITATIONS
1	Haploid, diploid, and pooled exome capture recapitulate features of biology and paralogy in two nonâ€model tree species. Molecular Ecology Resources, 2022, 22, 225-238.	4.8	3
2	Evaluating the accuracy of variant calling methods using the frequency of parentâ€offspring genotype mismatch. Molecular Ecology Resources, 2022, , .	4.8	1
3	Comparative Gene Expression Analysis Reveals Mechanism of <i>Pinus contorta</i> Response to the Fungal Pathogen <i>Dothistroma septosporum</i> . Molecular Plant-Microbe Interactions, 2021, 34, 397-409.	2.6	10
4	Detecting the genetic basis of local adaptation in loblolly pine ( Pinus taeda L.) using whole exomeâ€wide genotyping and an integrative landscape genomics analysis approach. Ecology and Evolution, 2019, 9, 6798-6809.	1.9	25
5	Predicting Adaptive Genetic Variation of Loblolly Pine (Pinus taeda L.) Populations Under Projected Future Climates Based on Multivariate Models. Journal of Heredity, 2019, 110, 857-865.	2.4	12
6	Purifying selection does not drive signatures of convergent local adaptation of lodgepole pine and interior spruce. BMC Evolutionary Biology, 2019, 19, 110.	3.2	1
7	Exploring the genetic basis of gene transcript abundance and metabolite levels in loblolly pine (Pinus) Tj ETQq1 1	0,784314 2.7	l rgBT /Over
8	Association genetics of growth and adaptive traits in loblolly pine (Pinus taeda L.) using whole-exome-discovered polymorphisms. Tree Genetics and Genomes, 2017, 13, 1.	1.6	29
9	Exome genotyping, linkage disequilibrium and population structure in loblolly pine (Pinus taeda L.). BMC Genomics, 2016, 17, 730.	2.8	53