

Mengqiao Pan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4853262/publications.pdf>

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7
papers

2,034
citations

1306789

7
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1719596

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docs citations

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times ranked

1706
citing authors

#	ARTICLE	IF	CITATIONS
1	Sequencing of allotetraploid cotton (<i>Gossypium hirsutum</i> L. acc. TM-1) provides a resource for fiber improvement. <i>Nature Biotechnology</i> , 2015, 33, 531-537.	9.4	1,560
2	Genetic basis for glandular trichome formation in cotton. <i>Nature Communications</i> , 2016, 7, 10456.	5.8	130
3	Genomic insights into divergence and dual domestication of cultivated allotetraploid cottons. <i>Genome Biology</i> , 2017, 18, 33.	3.8	128
4	Small interfering RNA from bidirectional transcripts of <i>GhMML3_A12</i> regulate cotton fiber development. <i>New Phytologist</i> , 2016, 210, 1298-1310.	3.5	124
5	Parallel and Intertwining Threads of Domestication in Allopolyploid Cotton. <i>Advanced Science</i> , 2021, 8, 2003634.	5.6	45
6	The <i>Gossypium longicalyx</i> Genome as a Resource for Cotton Breeding and Evolution. <i>G3: Genes, Genomes, Genetics</i> , 2020, 10, 1457-1467.	0.8	32
7	Identification of centromeric regions on the linkage map of cotton using centromere-related repeats. <i>Genomics</i> , 2014, 104, 587-593.	1.3	15