

# Takako Mochizuki

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10234198/publications.pdf>

Version: 2024-02-01

12  
papers

383  
citations

1307594

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h-index

1372567

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13  
times ranked

624  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome evolution of a nonparasitic secondary heterotroph, the diatom <i>Nitzschia putrida</i> . Science Advances, 2022, 8, eabi5075.	10.3	9
2	Identification of the sex-determining factor in the liverwort <i>Marchantia polymorpha</i> reveals unique evolution of sex chromosomes in a haploid system. Current Biology, 2021, 31, 5522-5532.e7.	3.9	36
3	Chromatin Organization in Early Land Plants Reveals an Ancestral Association between H3K27me3, Transposons, and Constitutive Heterochromatin. Current Biology, 2020, 30, 573-588.e7.	3.9	160
4	Draft Sequencing of the Heterozygous Diploid Genome of Satsuma ( <i>Citrus unshiu</i> Marc.) Using a Hybrid Assembly Approach. Frontiers in Genetics, 2017, 8, 180.	2.3	49
5	DNApod: DNA polymorphism annotation database from next-generation sequence read archives. PLoS ONE, 2017, 12, e0172269.	2.5	2
6	OryzaGenome: Genome Diversity Database of Wild <i>Oryza</i> Species. Plant and Cell Physiology, 2016, 57, e1-e1.	3.1	34
7	Hybrid Origins of Citrus Varieties Inferred from DNA Marker Analysis of Nuclear and Organelle Genomes. PLoS ONE, 2016, 11, e0166969.	2.5	76
8	Be happy with your Perl scripts. Ikushugaku Kenkyu, 2016, 18, 27-33.	0.3	0
9	Global expression differences and tissue specific expression differences in rice evolution result in two contrasting types of differentially expressed genes. BMC Genomics, 2015, 16, 1099.	2.8	5
10	Introduction of NIG supercomputer with command line-based NGS analysis. Ikushugaku Kenkyu, 2015, 17, 88-93.	0.3	0
11	Draft Genome Sequence of <i>Weissella oryzae</i> SG25 T, Isolated from Fermented Rice Grains. Genome Announcements, 2014, 2, .	0.8	11
12	Draft Genome Sequence of <i>Lactobacillus oryzae</i> Strain SG293 T. Genome Announcements, 2014, 2, .	0.8	1