

# Jennifer R Wortman

## List of Publications by Year in descending order

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Version: 2024-02-01

14  
papers

4,960  
citations

759233

12  
h-index

1199594

12  
g-index

14  
all docs

14  
docs citations

14  
times ranked

8654  
citing authors

#	ARTICLE	IF	CITATIONS
1	Improving the Arabidopsis genome annotation using maximal transcript alignment assemblies. <i>Nucleic Acids Research</i> , 2003, 31, 5654-5666.	14.5	1,597
2	The Genome Sequence of <i>Trypanosoma cruzi</i> , Etiologic Agent of Chagas Disease. <i>Science</i> , 2005, 309, 409-415.	12.6	1,273
3	Comparative Genomics of Trypanosomatid Parasitic Protozoa. <i>Science</i> , 2005, 309, 404-409.	12.6	713
4	Draft genome sequence of the oilseed species <i>Ricinus communis</i> . <i>Nature Biotechnology</i> , 2010, 28, 951-956.	17.5	449
5	Comprehensive annotation of secondary metabolite biosynthetic genes and gene clusters of <i>Aspergillus nidulans</i> , <i>A. fumigatus</i> , <i>A. niger</i> and <i>A. oryzae</i> . <i>BMC Microbiology</i> , 2013, 13, 91.	3.3	258
6	The Institute for Genomic Research Osa1 Rice Genome Annotation Database. <i>Plant Physiology</i> , 2005, 138, 18-26.	4.8	201
7	Annotation of the Arabidopsis Genome. <i>Plant Physiology</i> , 2003, 132, 461-468.	4.8	137
8	The Aspergillus Genome Database, a curated comparative genomics resource for gene, protein and sequence information for the Aspergillus research community. <i>Nucleic Acids Research</i> , 2010, 38, D420-D427.	14.5	111
9	Ergatis: a web interface and scalable software system for bioinformatics workflows. <i>Bioinformatics</i> , 2010, 26, 1488-1492.	4.1	83
10	Sybil: Methods and Software for Multiple Genome Comparison and Visualization. <i>Methods in Molecular Biology</i> , 2007, 408, 93-108.	0.9	55
11	Genome wide survey, discovery and evolution of repetitive elements in three <i>Entamoeba</i> species. <i>BMC Genomics</i> , 2008, 9, 595.	2.8	53
12	New resources for functional analysis of omics data for the genus <i>Aspergillus</i> . <i>BMC Genomics</i> , 2011, 12, 486.	2.8	28
13	Structural, Functional, and Comparative Annotation of Plant Genomes. , 0, , 373-395.		1
14	Unraveling the genomic diversity of small eukaryotes. <i>Genome Biology</i> , 2009, 10, 318.	9.6	1