

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