

Vinita Joardar

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11341206/publications.pdf>

Version: 2024-02-01

11

papers

2,449

citations

840776

11

h-index

1281871

11

g-index

11

all docs

11

docs citations

11

times ranked

3324

citing authors

#	ARTICLE	IF	CITATIONS
1	Consensus coding sequence (CCDS) database: a standardized set of human and mouse protein-coding regions supported by expert curation. <i>Nucleic Acids Research</i> , 2018, 46, D221-D228.	14.5	97
2	Mobile elements and mitochondrial genome expansion in the soil fungus and potato pathogen <i>Rhizoctonia solani</i> AG-3. <i>FEMS Microbiology Letters</i> , 2014, 352, 165-173.	1.8	143
3	A Rickettsia Genome Overrun by Mobile Genetic Elements Provides Insight into the Acquisition of Genes Characteristic of an Obligate Intracellular Lifestyle. <i>Journal of Bacteriology</i> , 2012, 194, 376-394.	2.2	152
4	Sequencing of mitochondrial genomes of nine <i>Aspergillus</i> and <i>Penicillium</i> species identifies mobile introns and accessory genes as main sources of genome size variability. <i>BMC Genomics</i> , 2012, 13, 698.	2.8	131
5	Genome sequencing and analysis of the filamentous fungus <i>Penicillium chrysogenum</i> . <i>Nature Biotechnology</i> , 2008, 26, 1161-1168.	17.5	427
6	Bioinformatics-Enabled Identification of the HrpL Regulon and Type III Secretion System Effector Proteins of <i>Pseudomonas syringae</i> pv. <i>phaseolicola</i> 1448A. <i>Molecular Plant-Microbe Interactions</i> , 2006, 19, 1193-1206.	2.6	81
7	Lineage-specific regions in <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000. <i>Molecular Plant Pathology</i> , 2005, 6, 53-64.	4.2	14
8	Whole-Genome Sequence Analysis of <i>Pseudomonas syringae</i> pv. <i>phaseolicola</i> 1448A Reveals Divergence among Pathovars in Genes Involved in Virulence and Transposition. <i>Journal of Bacteriology</i> , 2005, 187, 6488-6498.	2.2	301
9	The complete genome sequence of the <i>Arabidopsis</i> and tomato pathogen <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 10181-10186.	7.1	785
10	Identification of <i>Pseudomonas syringae</i> pv. <i>tomato</i> genes induced during infection of <i>Arabidopsis thaliana</i> . <i>Molecular Microbiology</i> , 2002, 44, 73-88.	2.5	180
11	Identification of novel <i>hrp</i> -regulated genes through functional genomic analysis of the <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 genome. <i>Molecular Microbiology</i> , 2002, 45, 1207-1218.	2.5	138