

# Ramesh R Vetukuri

## List of Publications by Year in descending order

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59  
papers

1,740  
citations

471371

17  
h-index

302012

39  
g-index

73  
all docs

73  
docs citations

73  
times ranked

1975  
citing authors

#	ARTICLE	IF	CITATIONS
1	Plant Growth-Promoting Activity of <i>Pseudomonas aeruginosa</i> FG106 and Its Ability to Act as a Biocontrol Agent against Potato, Tomato and Taro Pathogens. <i>Biology</i> , 2022, 11, 140.	1.3	31
2	Optimization of Culture Conditions for Zinc Phosphate Solubilization by <i>Aspergillus</i> sp. Using Response Surface Methodology. <i>Journal of Soil Science and Plant Nutrition</i> , 2022, 22, 1009-1018.	1.7	3
3	Sorghum in dryland: morphological, physiological, and molecular responses of sorghum under drought stress. <i>Planta</i> , 2022, 255, 20.	1.6	55
4	Effect of Biochar and Microbial Inoculation on P, Fe, and Zn Bioavailability in a Calcareous Soil. <i>Processes</i> , 2022, 10, 343.	1.3	12
5	Optimization of Biofertilizer Formulation for Phosphorus Solubilizing by <i>Pseudomonas fluorescens</i> Ur21 via Response Surface Methodology. <i>Processes</i> , 2022, 10, 650.	1.3	9
6	Biodiversity of the Genus <i>Trichoderma</i> in the Rhizosphere of Coffee ( <i>Coffea arabica</i> ) Plants in Ethiopia and Their Potential Use in Biocontrol of Coffee Wilt Disease. <i>Crops</i> , 2022, 2, 120-141.	0.6	10
7	RNA-Seq Provides Novel Genomic Resources for Noug ( <i>Guizotia abyssinica</i> ) and Reveals Microsatellite Frequency and Distribution in Its Transcriptome. <i>Frontiers in Plant Science</i> , 2022, 13, .	1.7	9
8	Comparative Small RNA and Degradome Sequencing Provides Insights into Antagonistic Interactions in the Biocontrol Fungus <i>Clonostachys rosea</i> . <i>Applied and Environmental Microbiology</i> , 2022, 88, .	1.4	5
9	A Quantitative Luminol-Based Assay for ROS Burst Detection in Potato Leaves in Response to Biotic Stimuli. <i>Methods in Molecular Biology</i> , 2022, , 395-402.	0.4	1
10	Spray-Induced Gene Silencing to Study Gene Function in <i>Phytophthora</i> . <i>Methods in Molecular Biology</i> , 2022, , 459-474.	0.4	1
11	Characterization of Environmental Effects on Flowering and Plant Architecture in an Everbearing Strawberry F1-Hybrid by Meristem Dissection and Gene Expression Analysis. <i>Horticulturae</i> , 2022, 8, 626.	1.2	2
12	Spray-induced gene silencing: an innovative strategy for plant trait improvement and disease control. <i>Crop Breeding and Applied Biotechnology</i> , 2021, 21, .	0.1	16
13	Editorial: Genomics and Effectomics of Filamentous Plant Pathogens. <i>Frontiers in Genetics</i> , 2021, 12, 648690.	1.1	2
14	When is it biological control? A framework of definitions, mechanisms, and classifications. <i>Journal of Pest Science</i> , 2021, 94, 665-676.	1.9	86
15	Biological control of strawberry crown rot, root rot and grey mould by the beneficial fungus <i>Aureobasidium pullulans</i> . <i>BioControl</i> , 2021, 66, 535-545.	0.9	16
16	Exogenous melatonin-stimulated transcriptomic alterations of <i>Davidia involucreta</i> seedlings under drought stress. <i>Trees - Structure and Function</i> , 2021, 35, 1025-1038.	0.9	20
17	Larval response to frass and guaiacol: detection of an attractant produced by bacteria from <i>Spodoptera littoralis</i> frass. <i>Journal of Pest Science</i> , 2021, 94, 1105-1118.	1.9	10
18	Haustorium formation and a distinct biotrophic transcriptome characterize infection of <i>Nicotiana benthamiana</i> by the tree pathogen <i>Phytophthora kernoviae</i> . <i>Molecular Plant Pathology</i> , 2021, 22, 954-968.	2.0	5

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19	Spray-Induced Gene Silencing as a Potential Tool to Control Potato Late Blight Disease. <i>Phytopathology</i> , 2021, 111, 2168-2175.	1.1	32
20	Comparison of two commercial recirculated aquacultural systems and their microbial potential in plant disease suppression. <i>BMC Microbiology</i> , 2021, 21, 205.	1.3	6
21	Draft genome assemblies for tree pathogens <i>Phytophthora pseudosyringae</i> and <i>Phytophthora boehmeriae</i> . <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	0.8	4
22	Optimization of Culture Conditions and Production of Bio-Fungicides from <i>Trichoderma</i> Species under Solid-State Fermentation Using Mathematical Modeling. <i>Microorganisms</i> , 2021, 9, 1675.	1.6	23
23	A fast, nondestructive method for the detection of disease-related lesions and wounded leaves. <i>BioTechniques</i> , 2021, 71, 425-430.	0.8	5
24	Identification of Unique Peptides for SARS-CoV-2 Diagnostics and Vaccine Development by an In Silico Proteomics Approach. <i>Frontiers in Immunology</i> , 2021, 12, 725240.	2.2	12
25	Role of Dicer-Dependent RNA Interference in Regulating Mycoparasitic Interactions. <i>Microbiology Spectrum</i> , 2021, 9, e0109921.	1.2	12
26	Interactions between Biochar and Compost Treatment and Mycorrhizal Fungi to Improve the Qualitative Properties of a Calcareous Soil under Rhizobox Conditions. <i>Agriculture (Switzerland)</i> , 2021, 11, 993.	1.4	7
27	Harnessing the Potential of Symbiotic Endophytic Fungi and Plant Growth-Promoting Rhizobacteria to Enhance Soil Quality in Saline Soils. <i>Processes</i> , 2021, 9, 1810.	1.3	5
28	Effect of RNA silencing suppression activity of chrysanthemum virus B p12 protein on small RNA species. <i>Archives of Virology</i> , 2020, 165, 2953-2959.	0.9	3
29	Differential Gene Expression Analysis of Wheat Breeding Lines Reveal Molecular Insights in Yellow Rust Resistance under Field Conditions. <i>Agronomy</i> , 2020, 10, 1888.	1.3	8
30	Horizontal Gene Transfer and Tandem Duplication Shape the Unique CAZyme Complement of the Mycoparasitic Oomycetes <i>Pythium oligandrum</i> and <i>Pythium periplocum</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 581698.	1.5	10
31	Dominance of Mating Type A1 and Indication of Epigenetic Effects During Early Stages of Mating in <i>Phytophthora infestans</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 252.	1.5	9
32	Monitoring and discrimination of Pandemis moths in apple orchards using semiochemicals, wing pattern morphology and DNA barcoding. <i>Crop Protection</i> , 2020, 132, 105110.	1.0	5
33	The presence of <i>Phytophthora infestans</i> in the rhizosphere of a wild <i>Solanum</i> species may contribute to off-season survival and pathogenicity. <i>Applied Soil Ecology</i> , 2020, 148, 103475.	2.1	7
34	Transient expression and purification of Î²-caryophyllene synthase in <i>Nicotiana benthamiana</i> to produce Î²-caryophyllene in vitro. <i>PeerJ</i> , 2020, 8, e8904.	0.9	9
35	Lignocellulolytic and Chitinolytic Glycoside Hydrolases: Structure, Catalytic Mechanism, Directed Evolution and Industrial Implementation. , 2020, , 97-127.		0
36	Efficient RNA silencing suppression activity of Potato Mop-Top Virus 8K protein is driven by variability and positive selection. <i>Virology</i> , 2019, 535, 111-121.	1.1	8

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37	Transcriptomic Responses of Dove Tree ( <i>Davidia involucrata</i> Baill.) to Heat Stress at the Seedling Stage. <i>Forests</i> , 2019, 10, 656.	0.9	9
38	Transcriptional stimulation of rate-limiting components of the autophagic pathway improves plant fitness. <i>Journal of Experimental Botany</i> , 2018, 69, 1415-1432.	2.4	120
39	Genome Sequence Resource for the Oomycete Taro Pathogen <i>Phytophthora colocasiae</i> . <i>Molecular Plant-Microbe Interactions</i> , 2018, 31, 903-905.	1.4	8
40	Screening of alternative products for integrated pest management of cucurbit powdery mildew in Sweden. <i>European Journal of Plant Pathology</i> , 2018, 150, 127-138.	0.8	22
41	Draft Genome Sequence for the Tree Pathogen <i>Phytophthora plurivora</i> . <i>Genome Biology and Evolution</i> , 2018, 10, 2432-2442.	1.1	19
42	Draft Genome Sequence of the Mycoparasitic Oomycete <i>Pythium periplocum</i> Strain CBS 532.74. <i>Genome Announcements</i> , 2017, 5, .	0.8	12
43	Draft genome of the oomycete pathogen <i>Phytophthora cactorum</i> strain LV007 isolated from European beech ( <i>Fagus sylvatica</i> ). <i>Genomics Data</i> , 2017, 12, 155-156.	1.3	18
44	<i>Phytophthora infestans</i> effector Pi14054 is a novel candidate suppressor of host silencing mechanisms. <i>European Journal of Plant Pathology</i> , 2017, 149, 771-777.	0.8	17
45	Draft Genome Sequence of the Mycoparasitic Oomycete <i>Pythium oligandrum</i> Strain CBS 530.74. <i>Genome Announcements</i> , 2017, 5, .	0.8	18
46	<i>Phytophthora infestans</i> Argonaute 1 binds microRNA and small RNA from effector genes and transposable elements. <i>New Phytologist</i> , 2016, 211, 993-1007.	3.5	41
47	The occurrence of pathogen suppressive soils in Sweden in relation to soil biota, soil properties, and farming practices. <i>Applied Soil Ecology</i> , 2016, 107, 57-65.	2.1	78
48	Real-time PCR for detection and quantification, and histological characterization of <i>Neonectria ditissima</i> in apple trees. <i>Trees - Structure and Function</i> , 2016, 30, 1111-1125.	0.9	17
49	Importin- $\beta$ -Mediated Nucleolar Localization of Potato Mop-Top Virus TRIPLE GENE BLOCK1 (TGB1) Protein Facilitates Virus Systemic Movement, Whereas TGB1 Self-Interaction Is Required for Cell-to-Cell Movement in <i>Nicotiana benthamiana</i> . <i>Plant Physiology</i> , 2015, 167, 738-752.	2.3	35
50	Plant-mediated gene silencing restricts growth of the potato late blight pathogen <i>Phytophthora infestans</i> . <i>Journal of Experimental Botany</i> , 2015, 66, 2785-2794.	2.4	124
51	Retromer Contributes to Immunity-Associated Cell Death in Arabidopsis. <i>Plant Cell</i> , 2015, 27, 463-479.	3.1	67
52	Fragmentation of tRNA in <i>Phytophthora infestans</i> asexual life cycle stages and during host plant infection. <i>BMC Microbiology</i> , 2014, 14, 308.	1.3	24
53	A viral transcription factor exhibits antiviral RNA silencing suppression activity independent of its nuclear localization. <i>Journal of General Virology</i> , 2014, 95, 2831-2837.	1.3	16
54	Phenotypic diversification by gene silencing in <i>Phytophthora</i> plant pathogens. <i>Communicative and Integrative Biology</i> , 2013, 6, e25890.	0.6	9

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55	Can silencing of transposons contribute to variation in effector gene expression in <i>Phytophthora infestans</i> ? <i>Mobile Genetic Elements</i> , 2012, 2, 110-114.	1.8	43
56	Evidence for Small RNAs Homologous to Effector-Encoding Genes and Transposable Elements in the Oomycete <i>Phytophthora infestans</i> . <i>PLoS ONE</i> , 2012, 7, e51399.	1.1	79
57	Silencing of the PiAvr3a effector-encoding gene from <i>Phytophthora infestans</i> by transcriptional fusion to a short interspersed element. <i>Fungal Biology</i> , 2011, 115, 1225-1233.	1.1	18
58	Evidence for involvement of Dicer-like, Argonaute and histone deacetylase proteins in gene silencing in <i>Phytophthora infestans</i> . <i>Molecular Plant Pathology</i> , 2011, 12, 772-785.	2.0	64
59	<i>Phytophthora infestans</i> effector AVR3a is essential for virulence and manipulates plant immunity by stabilizing host E3 ligase CMPG1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 9909-9914.	3.3	412