

Venkatachalam Lakshmanan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10702508/publications.pdf>

Version: 2024-02-01

17
papers

1,383
citations

516710

16
h-index

888059

17
g-index

17
all docs

17
docs citations

17
times ranked

2159
citing authors

#	ARTICLE	IF	CITATIONS
1	Microbe to Microbiome: A Paradigm Shift in the Application of Microorganisms for Sustainable Agriculture. <i>Frontiers in Microbiology</i> , 2020, 11, 622926.	3.5	88
2	Toward a Resilient, Functional Microbiome: Drought Tolerance-Alleviating Microbes for Sustainable Agriculture. <i>Methods in Molecular Biology</i> , 2017, 1631, 69-84.	0.9	26
3	Interplant Aboveground Signaling Prompts Upregulation of Auxin Promoter and Malate Transporter as Part of Defensive Response in the Neighboring Plants. <i>Frontiers in Plant Science</i> , 2017, 8, 595.	3.6	10
4	Impact of Seed Exudates on Growth and Biofilm Formation of <i>Bacillus amyloliquefaciens</i> ALB629 in Common Bean. <i>Frontiers in Microbiology</i> , 2017, 8, 2631.	3.5	31
5	Killing Two Birds with One Stone: Natural Rice Rhizospheric Microbes Reduce Arsenic Uptake and Blast Infections in Rice. <i>Frontiers in Plant Science</i> , 2016, 7, 1514.	3.6	19
6	<i>Bacillus subtilis</i> Early Colonization of <i>Arabidopsis thaliana</i> Roots Involves Multiple Chemotaxis Receptors. <i>MBio</i> , 2016, 7, .	4.1	189
7	A perspective on inter-kingdom signaling in plantâ€“beneficial microbe interactions. <i>Plant Molecular Biology</i> , 2016, 90, 537-548.	3.9	97
8	Crucial Roles of Abscisic Acid Biogenesis in Virulence of Rice Blast Fungus <i>Magnaporthe oryzae</i> . <i>Frontiers in Plant Science</i> , 2015, 6, 1082.	3.6	74
9	Root Microbiome Assemblage is Modulated by Plant Host Factors. <i>Advances in Botanical Research</i> , 2015, 75, 57-79.	1.1	28
10	A natural rice rhizospheric bacterium abates arsenic accumulation in rice (<i>Oryza sativa</i> L.). <i>Planta</i> , 2015, 242, 1037-1050.	3.2	63
11	Functional Soil Microbiome: Belowground Solutions to an Aboveground Problem Â. <i>Plant Physiology</i> , 2014, 166, 689-700.	4.8	299
12	Root transcriptome analysis of <i>Arabidopsis thaliana</i> exposed to beneficial <i>Bacillus subtilis</i> FB17 rhizobacteria revealed genes for bacterial recruitment and plant defense independent of malate efflux. <i>Planta</i> , 2013, 238, 657-668.	3.2	84
13	Characterization of the Complex Regulation of <i>AtALMT1</i> Expression in Response to Phytohormones and Other Inducers Â Â. <i>Plant Physiology</i> , 2013, 162, 732-740.	4.8	77
14	Overexpression of <i>AtALMT1</i> in the <i>Arabidopsis thaliana</i> ecotype Columbia results in enhanced Al-activated malate excretion and beneficial bacterium recruitment. <i>Plant Signaling and Behavior</i> , 2013, 8, e25565.	2.4	21
15	Factors other than root secreted malic acid that contributes toward <i>Bacillus subtilis</i> FB17 colonization on <i>Arabidopsis</i> roots. <i>Plant Signaling and Behavior</i> , 2013, 8, e27277.	2.4	22
16	Rhizobacteria <i>Bacillus subtilis</i> restricts foliar pathogen entry through stomata. <i>Plant Journal</i> , 2012, 72, 694-706.	5.7	98
17	Microbe-Associated Molecular Patterns-Triggered Root Responses Mediate Beneficial Rhizobacterial Recruitment in <i>Arabidopsis</i> Â Â Â. <i>Plant Physiology</i> , 2012, 160, 1642-1661.	4.8	157