

Tiffany M Lowe-Power

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

20
papers

527
citations

13
h-index

22
g-index

22
ext. papers

794
ext. citations

6.9
avg, IF

3.92
L-index

#	Paper	IF	Citations
20	Meta-analysis of the species complex (RSSC) based on comparative evolutionary genomics and reverse ecology.. <i>Microbial Genomics</i> , 2022 , 8,	4.4	3
19	Genome-Wide Identification of Tomato Xylem Sap Fitness Factors for Three Plant-Pathogenic Species. <i>MSystems</i> , 2021 , e0122921	7.6	3
18	A generation of junior faculty is at risk from the impacts of COVID-19. <i>PLoS Biology</i> , 2021 , 19, e3001266	9.7	0
17	A Plant Pathogen Type III Effector Protein Subverts Translational Regulation to Boost Host Polyamine Levels. <i>Cell Host and Microbe</i> , 2019 , 26, 638-649.e5	23.4	27
16	Plant-like bacterial expansins play contrasting roles in two tomato vascular pathogens. <i>Molecular Plant Pathology</i> , 2018 , 19, 1210-1221	5.7	19
15	Plant Assays for Quantifying Virulence. <i>Bio-protocol</i> , 2018 , 8, e3028	0.9	13
14	Metabolomics of tomato xylem sap during bacterial wilt reveals <i>Ralstonia solanacearum</i> produces abundant putrescine, a metabolite that accelerates wilt disease. <i>Environmental Microbiology</i> , 2018 , 20, 1330-1349	5.2	59
13	How <i>Ralstonia solanacearum</i> Exploits and Thrives in the Flowing Plant Xylem Environment. <i>Trends in Microbiology</i> , 2018 , 26, 929-942	12.4	58
12	A Single Regulator Mediates Strategic Switching between Attachment/Spread and Growth/Virulence in the Plant Pathogen. <i>MBio</i> , 2017 , 8,	7.8	38
11	Functional Identification of Putrescine C- and N-Hydroxylases. <i>ACS Chemical Biology</i> , 2016 , 11, 2782-2789	4.9	17
10	Interactions between Bacteria And Aspen Defense Chemicals at the Phyllosphere - Herbivore Interface. <i>Journal of Chemical Ecology</i> , 2016 , 42, 193-201	2.7	19
9	<i>Ralstonia solanacearum</i> lipopeptide induces chlamydospore development in fungi and facilitates bacterial entry into fungal tissues. <i>ISME Journal</i> , 2016 , 10, 2317-30	11.9	73
8	In planta comparative transcriptomics of host-adapted strains of <i>Ralstonia solanacearum</i> . <i>PeerJ</i> , 2016 , 4, e1549	3.1	23
7	Degradation of the Plant Defense Signal Salicylic Acid Protects <i>Ralstonia solanacearum</i> from Toxicity and Enhances Virulence on Tobacco. <i>MBio</i> , 2016 , 7,	7.8	44
6	Insights into the environmental reservoir of pathogenic <i>Vibrio parahaemolyticus</i> using comparative genomics. <i>Frontiers in Microbiology</i> , 2015 , 6, 204	5.7	23
5	Comparative genomic analysis of <i>Ralstonia solanacearum</i> reveals candidate genes for host specificity. <i>BMC Genomics</i> , 2015 , 16, 270	4.5	56
4	Hydroxycinnamic Acid Degradation, a Broadly Conserved Trait, Protects <i>Ralstonia solanacearum</i> from Chemical Plant Defenses and Contributes to Root Colonization and Virulence. <i>Molecular Plant-Microbe Interactions</i> , 2015 , 28, 286-97	3.6	44

- 3 A Meta-analysis of the known Global Distribution and Host Range of the *Ralstonia* Species Complex 4
- 2 Genome-wide identification of tomato xylem sap fitness factors for *Ralstonia pseudosolanacearum* and *Ralstonia syzygii* 2
- 1 Revisiting the source of wilt symptoms: X-ray microcomputed tomography provides direct evidence that *Ralstonia* biomass clogs xylem vessels 2