

Sarah L Lebeis

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2116082/publications.pdf>

Version: 2024-02-01

32
papers

5,936
citations

394421

19
h-index

526287

27
g-index

34
all docs

34
docs citations

34
times ranked

6632
citing authors

#	ARTICLE	IF	CITATIONS
1	Defining the core <i>Arabidopsis thaliana</i> root microbiome. <i>Nature</i> , 2012, 488, 86-90.	27.8	2,475
2	Salicylic acid modulates colonization of the root microbiome by specific bacterial taxa. <i>Science</i> , 2015, 349, 860-864.	12.6	957
3	Microbiota and Host Nutrition across Plant and Animal Kingdoms. <i>Cell Host and Microbe</i> , 2015, 17, 603-616.	11.0	628
4	Genomic features of bacterial adaptation to plants. <i>Nature Genetics</i> , 2018, 50, 138-150.	21.4	480
5	Disabling poxvirus pathogenesis by inhibition of Abl-family tyrosine kinases. <i>Nature Medicine</i> , 2005, 11, 731-739.	30.7	207
6	Soil indigenous microbiome and plant genotypes cooperatively modify soybean rhizosphere microbiome assembly. <i>BMC Microbiology</i> , 2019, 19, 201.	3.3	194
7	TLR Signaling Mediated by MyD88 Is Required for a Protective Innate Immune Response by Neutrophils to <i>Citrobacter rodentium</i> . <i>Journal of Immunology</i> , 2007, 179, 566-577.	0.8	162
8	The potential for give and take in plant-microbiome relationships. <i>Frontiers in Plant Science</i> , 2014, 5, 287.	3.6	106
9	Greater than the sum of their parts: characterizing plant microbiomes at the community-level. <i>Current Opinion in Plant Biology</i> , 2015, 24, 82-86.	7.1	93
10	Interleukin-1 Receptor Signaling Protects Mice from Lethal Intestinal Damage Caused by the Attaching and Effacing Pathogen <i>Citrobacter rodentium</i> . <i>Infection and Immunity</i> , 2009, 77, 604-614.	2.2	92
11	Giving back to the community: microbial mechanisms of plant-soil interactions. <i>Functional Ecology</i> , 2016, 30, 1043-1052.	3.6	89
12	Bacterial Production of Indole Related Compounds Reveals Their Role in Association Between Duckweeds and Endophytes. <i>Frontiers in Chemistry</i> , 2018, 6, 265.	3.6	75
13	Duckweed hosts a taxonomically similar bacterial assemblage as the terrestrial leaf microbiome. <i>PLoS ONE</i> , 2020, 15, e0228560.	2.5	51
14	Connexin26 35delG does not represent a mutational hotspot. <i>Human Genetics</i> , 2003, 113, 18-23.	3.8	46
15	Bridging the Gap Between Single-Strain and Community-Level Plant-Microbe Chemical Interactions. <i>Molecular Plant-Microbe Interactions</i> , 2020, 33, 124-134.	2.6	45
16	Overexpression of Strigolactone-Associated Genes Exerts Fine-Tuning Selection on Soybean Rhizosphere Bacterial and Fungal Microbiome. <i>Phytobiomes Journal</i> , 2020, 4, 239-251.	2.7	30
17	Aligning Antimicrobial Drug Discovery with Complex and Redundant Host-Pathogen Interactions. <i>Cell Host and Microbe</i> , 2009, 5, 114-122.	11.0	23
18	Genome-Resolved Proteomic Stable Isotope Probing of Soil Microbial Communities Using ¹³ CO ₂ and ¹³ C-Methanol. <i>Frontiers in Microbiology</i> , 2019, 10, 2706.	3.5	23

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19	Exercising influence: distinct biotic interactions shape root microbiomes. <i>Current Opinion in Plant Biology</i> , 2015, 26, 32-36.	7.1	18
20	Host-specific and tissue-dependent orchestration of microbiome community structure in traditional rice paddy ecosystems. <i>Plant and Soil</i> , 2020, 452, 379-395.	3.7	14
21	Root-Associated <i>Streptomyces</i> Isolates Harboring <i>mec</i> Genes Demonstrate Enhanced Plant Colonization. <i>Phytobiomes Journal</i> , 2019, 3, 165-176.	2.7	11
22	Bacterial communities of the <i>Salvia lyrata</i> rhizosphere explained by spatial structure and sampling grain. <i>Microbial Ecology</i> , 2020, 80, 846-858.	2.8	8
23	Plant Microbiome Identification and Characterization. <i>Current Protocols in Plant Biology</i> , 2017, 2, 135-146.	2.8	7
24	Distinguishing nutrient-dependent plant driven bacterial colonization patterns in alfalfa. <i>Environmental Microbiology Reports</i> , 2020, 12, 70-77.	2.4	7
25	Using the Microbiome Amplification Preference Tool (MAPT) to Reveal <i>Medicago sativa</i> -Associated Eukaryotic Microbes. <i>Phytobiomes Journal</i> , 2020, 4, 340-350.	2.7	3
26	Microbial Ecology: How to Fight the Establishment. <i>Current Biology</i> , 2019, 29, R1320-R1323.	3.9	2
27	Editorial overview: Biotic interactions: Inferring global implications for the molecular interface between plants and their biotic interactions across scales. <i>Current Opinion in Plant Biology</i> , 2017, 38, v-vii.	7.1	0
28	mSphere of Influence: Peering through a Keyhole into the Unseen World. <i>MSphere</i> , 2020, 5, .	2.9	0
29	Duckweed hosts a taxonomically similar bacterial assemblage as the terrestrial leaf microbiome. , 2020, 15, e0228560.		0
30	Duckweed hosts a taxonomically similar bacterial assemblage as the terrestrial leaf microbiome. , 2020, 15, e0228560.		0
31	Duckweed hosts a taxonomically similar bacterial assemblage as the terrestrial leaf microbiome. , 2020, 15, e0228560.		0
32	Duckweed hosts a taxonomically similar bacterial assemblage as the terrestrial leaf microbiome. , 2020, 15, e0228560.		0