## Shiri Freilich

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4872655/publications.pdf Version: 2024-02-01



Shidi Edenich

#	Article	IF	CITATIONS
1	Challenges in microbial ecology: building predictive understanding of community function and dynamics. ISME Journal, 2016, 10, 2557-2568.	9.8	570
2	Competitive and cooperative metabolic interactions in bacterial communities. Nature Communications, 2011, 2, 589.	12.8	413
3	The large-scale organization of the bacterial network of ecological co-occurrence interactions. Nucleic Acids Research, 2010, 38, 3857-3868.	14.5	259
4	Prospects for Biological Soilborne Disease Control: Application of Indigenous Versus Synthetic Microbiomes. Phytopathology, 2017, 107, 256-263.	2.2	147
5	Modeling microbial communities from atrazine contaminated soils promotes the development of biostimulation solutions. ISME Journal, 2019, 13, 494-508.	9.8	119
6	Metabolic-network-driven analysis of bacterial ecological strategies. Genome Biology, 2009, 10, R61.	9.6	88
7	NetCooperate: a network-based tool for inferring host-microbe and microbe-microbe cooperation. BMC Bioinformatics, 2015, 16, 164.	2.6	82
8	Variations in the identity and complexity of endosymbiont combinations in whitefly hosts. Frontiers in Microbiology, 2014, 5, 310.	3.5	65
9	Global analysis of the apple fruit microbiome: are all apples the same?. Environmental Microbiology, 2021, 23, 6038-6055.	3.8	64
10	Analysis of Microbial Functions in the Rhizosphere Using a Metabolic-Network Based Framework for Metagenomics Interpretation. Frontiers in Microbiology, 2017, 8, 1606.	3.5	59
11	Effect of Washing, Waxing and Low-Temperature Storage on the Postharvest Microbiome of Apple. Microorganisms, 2020, 8, 944.	3.6	54
12	Compositional shifts in the strawberry fruit microbiome in response to near-harvest application of Metschnikowia fructicola, a yeast biocontrol agent. Postharvest Biology and Technology, 2021, 175, 111469.	6.0	50
13	To B or Not to B: Comparative Genomics Suggests Arsenophonus as a Source of B Vitamins in Whiteflies. Frontiers in Microbiology, 2018, 9, 2254.	3.5	49
14	NetCmpt: a network-based tool for calculating the metabolic competition between bacterial species. Bioinformatics, 2012, 28, 2195-2197.	4.1	47
15	Modeling trophic dependencies and exchanges among insects' bacterial symbionts in a host-simulated environment. BMC Genomics, 2018, 19, 402.	2.8	42
16	Decoupling Environment-Dependent and Independent Genetic Robustness across Bacterial Species. PLoS Computational Biology, 2010, 6, e1000690.	3.2	31
17	The pathobiome concept applied to postharvest pathology and its implication on biocontrol strategies. Postharvest Biology and Technology, 2022, 189, 111911.	6.0	16
18	NetMet: A Network-Based Tool for Predicting Metabolic Capacities of Microbial Species and their Interactions. Microorganisms, 2020, 8, 840.	3.6	13

Shiri Freilich

#	Article	IF	CITATIONS
19	Genome Analysis of Haplotype D of Candidatus Liberibacter Solanacearum. Frontiers in Microbiology, 2018, 9, 2933.	3.5	10
20	Stratification of co-evolving genomic groups using ranked phylogenetic profiles. BMC Bioinformatics, 2009, 10, 355.	2.6	8
21	The Apple Microbiome: Structure, Function, and Manipulation for Improved Plant Health. Compendium of Plant Genomes, 2021, , 341-382.	0.5	8
22	An eco-systems biology approach for modeling tritrophic networks reveals the influence of dietary amino acids on symbiont dynamics of <i>Bemisia tabaci</i> . FEMS Microbiology Ecology, 2021, 97, .	2.7	4
23	NetCom: A Network-Based Tool for Predicting Metabolic Activities of Microbial Communities Based on Interpretation of Metagenomics Data. Microorganisms, 2021, 9, 1838.	3.6	4