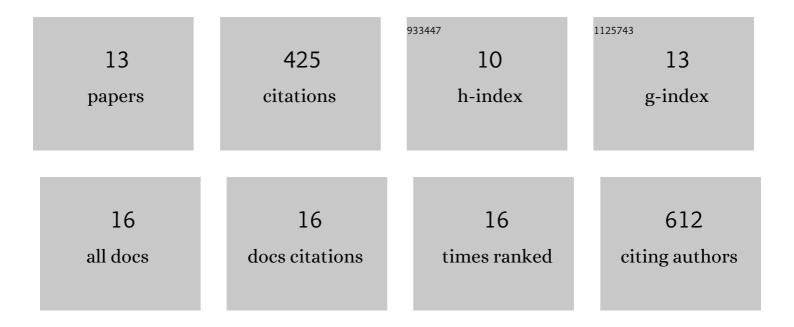
## Heba M M Ibrahim

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7469930/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Transcriptional response to host chemical cues underpins the expansion of host range in a fungal plant pathogen lineage. ISME Journal, 2022, 16, 138-148.	9.8	17
2	Genomeâ€wide alternative splicing profiling in the fungal plant pathogen <i>Sclerotinia sclerotiorum</i> during the colonization of diverse host families. Molecular Plant Pathology, 2021, 22, 31-47.	4.2	25
3	Pathogen Effectors: Exploiting the Promiscuity of Plant Signaling Hubs. Trends in Plant Science, 2021, 26, 780-795.	8.8	36
4	Efficient CRISPR-mediated base editing in <i>Agrobacterium</i> spp Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	38
5	Surfactin Stimulated by Pectin Molecular Patterns and Root Exudates Acts as a Key Driver of the <i>Bacillus</i> -Plant Mutualistic Interaction. MBio, 2021, 12, e0177421.	4.1	25
6	A Short-Read Genome Assembly Resource for Leveillula taurica Causing Powdery Mildew Disease of Sweet Pepper (Capsicum annuum). Molecular Plant-Microbe Interactions, 2020, 33, 782-786.	2.6	11
7	A Chromosome-Scale Genome Assembly Resource for <i>Myriosclerotinia sulcatula</i> Infecting Sedge Grass ( <i>Carex</i> sp.). Molecular Plant-Microbe Interactions, 2020, 33, 880-883.	2.6	2
8	Effective approaches to study the plant-root knot nematode interaction. Plant Physiology and Biochemistry, 2019, 141, 332-342.	5.8	32
9	In vitro evaluation of antioxidant, anticancer, and antiviral activities of exopolysaccharide from Streptomyces hirsutus NRC2018. Journal of Applied Pharmaceutical Science, 2019, 9, 10-18.	1.0	6
10	Production and Assessment of Antioxidant Activity of Exopolysaccharide from Marine Streptomyces globisporus BU2018. Egyptian Journal of Botany, 2019, .	0.2	3
11	Transcriptome analyses reveal SR45 to be a neutral splicing regulator and a suppressor of innate immunity in Arabidopsis thaliana. BMC Genomics, 2017, 18, 772.	2.8	64
12	Post-transcriptional gene silencing of root-knot nematode in transformed soybean roots. Experimental Parasitology, 2011, 127, 90-99.	1.2	85
13	Analysis of Gene expression in soybean (Glycine max) roots in response to the root knot nematode Meloidogyne incognita using microarrays and KEGG pathways. BMC Genomics, 2011, 12, 220.	2.8	78