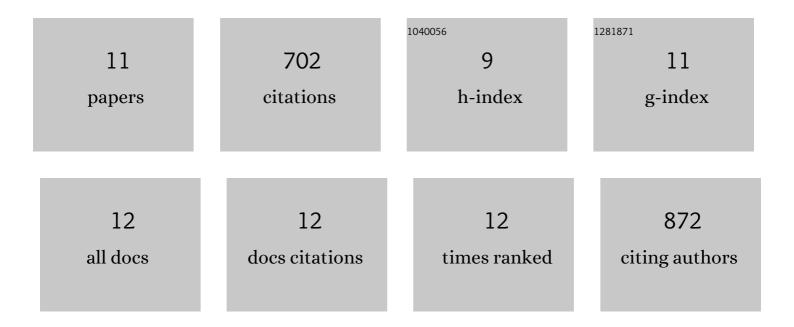
Surapareddy Sreenivasaprasad

List of Publications by Year in descending order

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SURAPAREDDY

#	Article	IF	CITATIONS
1	Genome Sequence of the Biocontrol Agent <i>Coniothyrium minitans</i> Conio (IMI 134523). Molecular Plant-Microbe Interactions, 2021, 34, 222-225.	2.6	2
2	Genomics Evolutionary History and Diagnostics of the Alternaria alternata Species Group Including Apple and Asian Pear Pathotypes. Frontiers in Microbiology, 2019, 10, 3124.	3.5	41
3	Genome Sequence of the Mycotoxigenic Crop Pathogen Fusarium proliferatum Strain ITEM 2341 from Date Palm. Microbiology Resource Announcements, 2018, 7, .	0.6	6
4	Gene family expansions and contractions are associated with host range in plant pathogens of the genus Colletotrichum. BMC Genomics, 2016, 17, 555.	2.8	151
5	Molecular Diversity of Anthracnose Pathogen Populations Associated with UK Strawberry Production Suggests Multiple Introductions of Three Different Colletotrichum Species. PLoS ONE, 2015, 10, e0129140.	2.5	81
6	Discrete lineages within Alternaria alternata species group: Identification using new highly variable loci and support from morphological characters. Fungal Biology, 2015, 119, 994-1006.	2.5	70
7	Draft Genome Sequence of <i>Colletotrichum acutatum Sensu Lato</i> (<i>Colletotrichum) Tj ETQq1 1 0.7843</i>	14 rgBT /(0.8	Overlock 10 Tf
8	The distinctive population structure ofColletotrichumspecies associated with olive anthracnose in the Algarve region of Portugal reflects a host–pathogen diversity hot spot. FEMS Microbiology Letters, 2009, 296, 31-38.	1.8	42
9	Disruption of the Coniothyrium minitans PIF1 DNA helicase gene impairs growth and capacity for sclerotial mycoparasitism. Microbiology (United Kingdom), 2008, 154, 1628-1636.	1.8	11
10	Analysis of cDNA transcripts from Coniothyrium minitans reveals a diverse array of genes involved in key processes during sclerotial mycoparasitism. Fungal Genetics and Biology, 2007, 44, 1262-1284.	2.1	31
11	A Novel Arabidopsis-Colletotrichum Pathosystem for the Molecular Dissection of Plant-Fungal Interactions. Molecular Plant-Microbe Interactions, 2004, 17, 272-282.	2.6	214