

# John B Ridenour

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7589185/publications.pdf>

Version: 2024-02-01

11  
papers

209  
citations

1307594

7  
h-index

1281871

11  
g-index

11  
all docs

11  
docs citations

11  
times ranked

301  
citing authors

#	ARTICLE	IF	CITATIONS
1	Regulation of Stomatal Tropism and Infection by Light in <i>Cercospora zea</i> - <i>maydis</i> : Evidence for Coordinated Host/Pathogen Responses to Photoperiod?. <i>PLoS Pathogens</i> , 2011, 7, e1002113.	4.7	65
2	HXX1 regulates carbon catabolism, sporulation, fumonisin B1 production and pathogenesis in <i>Fusarium verticillioides</i> . <i>Microbiology (United Kingdom)</i> , 2011, 157, 2658-2669.	1.8	33
3	Complementation of <i>CTB7</i> in the Maize Pathogen <i>Cercospora zeina</i> Overcomes the Lack of In Vitro Cercosporin Production. <i>Molecular Plant-Microbe Interactions</i> , 2017, 30, 710-724.	2.6	25
4	The novel fungal-specific gene FUG1 has a role in pathogenicity and fumonisin biosynthesis in <i>Fusarium verticillioides</i> . <i>Molecular Plant Pathology</i> , 2017, 18, 513-528.	4.2	24
5	The HAP complex in <i>Fusarium verticillioides</i> is a key regulator of growth, morphogenesis, secondary metabolism, and pathogenesis. <i>Fungal Genetics and Biology</i> , 2014, 69, 52-64.	2.1	20
6	<i>UBL1</i> of <i>Fusarium verticillioides</i> links the N-end rule pathway to extracellular sensing and plant pathogenesis. <i>Environmental Microbiology</i> , 2014, 16, 2004-2022.	3.8	17
7	<i>Agrobacterium tumefaciens</i> -mediated transformation of the soybean pathogen <i>Phomopsis longicolla</i> . <i>Journal of Microbiological Methods</i> , 2013, 92, 244-245.	1.6	10
8	Regulation of Pathogenesis by Light in <i>Cercospora zea</i> - <i>maydis</i> : An Updated Perspective. <i>Plant Pathology Journal</i> , 2011, 27, 103-109.	1.7	7
9	The HAP Complex Governs Fumonisin Biosynthesis and Maize Kernel Pathogenesis in <i>Fusarium verticillioides</i> . <i>Journal of Food Protection</i> , 2016, 79, 1498-1507.	1.7	6
10	Taxonomic Resolution of the Nematophagous Fungal Isolate ARF18 via Genome Sequencing. <i>Genome Announcements</i> , 2017, 5, .	0.8	1
11	Draft genome sequence of <i>Xylaria</i> sp., the causal agent of taproot decline of soybean in the southern United States. <i>Data in Brief</i> , 2018, 17, 129-133.	1.0	1