## Bo Liu

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5486326/publications.pdf

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#	Article	IF	CITATIONS
1	Three New Fungal Leaf Spot Diseases of Spinach in the United States and the Evaluation of Fungicide Efficacy for Disease Management. Plant Disease, 2021, 105, 316-323.	1.4	10
2	Phylogenetic Analysis, Vegetative Compatibility, Virulence, and Fungal Filtrates of Leaf Curl Pathogen <i>Colletotrichum fioriniae</i> from Celery. Phytopathology, 2021, 111, 751-760.	2.2	6
3	Long-term effect of non-irrigation and irrigation on soil Pythium, Fusarium, and Rhizoctonia communities and their relation with seed-rot, root-rot, and damping-off of soybean. European Journal of Plant Pathology, 2020, 158, 297-314.	1.7	5
4	Characterization of Leaf Spot Pathogens from Several Spinach Production Areas in the United States. Plant Disease, 2020, 104, 1994-2004.	1.4	9
5	Long-term effects of dryland and irrigation production systems on soil Fusarium communities in wheat. Canadian Journal of Plant Pathology, 2019, 41, 585-596.	1.4	1
6	Characterization of Foliar Web Blight of Spinach, Caused by Pythium aphanidermatum, in the Desert Southwest of the United States. Plant Disease, 2018, 102, 608-612.	1.4	4
7	Microbial communities in soils with different population densities of soybean cyst nematode. Canadian Journal of Plant Pathology, 2018, 40, 48-60.	1.4	2
8	Rhizoctonia communities in soybean fields and their relation with other microbes and nematode communities. European Journal of Plant Pathology, 2016, 144, 671-686.	1.7	13
9	Effect of organic, sustainable, and conventional management strategies in grower fields on soil physical, chemical, and biological factors and the incidence of Southern blight. Applied Soil Ecology, 2007, 37, 202-214.	<b>4.</b> 3	134
10	Characterization of Diversity in Colletotrichum acutatum sensu lato by Sequence Analysis of Two Gene Introns, mtDNA and Intron RFLPs, and Mating Compatibility. Mycologia, 2003, 95, 872.	1.9	133
11	Characterization of diversity in <i>Colletotrichum acutatumsensu lato</i> by sequence analysis of two gene introns, mtDNA and intron RFLPs, and mating compatibility. Mycologia, 2003, 95, 872-895.	1.9	233