Huamin Chen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8926559/publications.pdf

Version: 2024-02-01

566801 433756 2,137 29 15 31 citations h-index g-index papers 34 34 34 3470 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Diguanylate Cyclase GdpX6 with c-di-GMP Binding Activity Involved in the Regulation of Virulence Expression in Xanthomonas oryzae pv. oryzae. Microorganisms, 2021, 9, 495.	1.6	3
2	Transcriptome Analysis Revealed Overlapping and Special Regulatory Roles of RpoN1 and RpoN2 in Motility, Virulence, and Growth of Xanthomonas oryzae pv. oryzae. Frontiers in Microbiology, 2021, 12, 653354.	1.5	3
3	<scp>BLB8</scp> , an antiviral protein from <i>Brevibacillus laterosporus</i> strain <scp>B8</scp> , inhibits <i>Tobacco mosaic virus</i> infection by triggering immune response in tobacco. Pest Management Science, 2021, 77, 4383-4392.	1.7	5
4	The Regulatory Functions of $\ddot{l}f54$ Factor in Phytopathogenic Bacteria. International Journal of Molecular Sciences, 2021, 22, 12692.	1.8	8
5	Identification of the Regulatory Components Mediated by the Cyclic di-GMP Receptor Filp and Its Interactor PilZX3 and Functioning in Virulence of Xanthomonas oryzae pv. oryzae. Molecular Plant-Microbe Interactions, 2020, 33, 1196-1208.	1.4	3
6	The RpoN2â€PilRX regulatory system governs type IV pilus gene transcription and is required for bacterial motility and virulence in ⟨i⟩Xanthomonas oryzae⟨ i⟩ pv. ⟨i⟩oryzae⟨ i⟩. Molecular Plant Pathology, 2020, 21, 652-666.	2.0	10
7	<i>Xanthomonas oryzae</i> pv. <i>oryzae</i> Response Regulator TriP Regulates Virulence and Exopolysaccharide Production Via Interacting With c-di-GMP Phosphodiesterase PdeR. Molecular Plant-Microbe Interactions, 2019, 32, 729-739.	1.4	8
8	Overexpression of miR169o, an Overlapping MicroRNA in Response to Both Nitrogen Limitation and Bacterial Infection, Promotes Nitrogen Use Efficiency and Susceptibility to Bacterial Blight in Rice. Plant and Cell Physiology, 2018, 59, 1234-1247.	1.5	46
9	A ten geneâ€containing genomic island determines flagellin glycosylation: implication for its regulatory role in motility and virulence of ⟨i>Xanthomonas oryzae⟨ i> pv. ⟨i>oryzae⟨ i>. Molecular Plant Pathology, 2018, 19, 579-592.	2.0	15
10	Phosphodiesterase EdpX1 Promotes Xanthomonas oryzae pv. oryzae Virulence, Exopolysaccharide Production, and Biofilm Formation. Applied and Environmental Microbiology, 2018, 84, .	1.4	19
11	The <i>Xanthomonas</i> effector XopK harbours E3 ubiquitinâ€ligase activity that is required for virulence. New Phytologist, 2018, 220, 219-231.	3.5	47
12	Identification of phenolic compounds that suppress the virulence of <i>Xanthomonas oryzae</i> on rice via the type III secretion system. Molecular Plant Pathology, 2017, 18, 555-568.	2.0	67
13	RpoN2- and FliA-regulated fliTX is indispensible for flagellar motility and virulence in Xanthomonas oryzae pv. oryzae. BMC Microbiology, 2017, 17, 171.	1.3	3
14	The GGDEF-domain protein GdpX1 attenuates motility, exopolysaccharide production and virulence in <i>Xanthomonas oryzae</i> pv.Â <i>oryzae</i> . Journal of Applied Microbiology, 2016, 120, 1646-1657.	1.4	25
15	OxyR-regulated catalase CatB promotes the virulence in rice via detoxifying hydrogen peroxide in Xanthomonas oryzae pv. oryzae. BMC Microbiology, 2016, 16, 269.	1.3	29
16	XA21-specific induction of stress-related genes following <i>Xanthomonas</i> i>infection of detached rice leaves. PeerJ, 2016, 4, e2446.	0.9	9
17	The Arabidopsis Transcription Factor BRASSINOSTEROID INSENSITIVE1-ETHYL METHANESULFONATE-SUPPRESSOR1 Is a Direct Substrate of MITOGEN-ACTIVATED PROTEIN KINASE6 and Regulates Immunity. Plant Physiology, 2015, 167, 1076-1086.	2.3	87
18	PXO_00987, a putative acetyltransferase, is required for flagellin glycosylation, and regulates flagellar motility, exopolysaccharide production, and biofilm formation in Xanthomonas oryzae pv. oryzae. Microbial Pathogenesis, 2015, 85, 50-57.	1.3	15

#	Article	IF	CITATIONS
19	The rice immune receptor XA21 recognizes a tyrosine-sulfated protein from a Gram-negative bacterium. Science Advances, 2015, 1, e1500245.	4.7	209
20	The Xanthomonas oryzae pv. oryzae PilZ Domain Proteins Function Differentially in Cyclic di-GMP Binding and Regulation of Virulence and Motility. Applied and Environmental Microbiology, 2015, 81, 4358-4367.	1.4	47
21	Identification of differentially-expressed genes of rice in overlapping responses to bacterial infection by Xanthomonas oryzae pv. oryzae and nitrogen deficiency. Journal of Integrative Agriculture, 2015, 14, 888-899.	1.7	1
22	Alternative sigma factor RpoN2 is required for flagellar motility and full virulence of Xanthomonas oryzae pv. oryzae. Microbiological Research, 2015, 170, 177-183.	2.5	52
23	The Degenerate EAL-GGDEF Domain Protein Filp Functions as a Cyclic di-GMP Receptor and Specifically Interacts with the PilZ-Domain Protein PXO_02715 to Regulate Virulence in <i>Xanthomonas oryzae</i> pv. <i>oryzae</i> . Molecular Plant-Microbe Interactions, 2014, 27, 578-589.	1.4	65
24	Mutation of alkyl hydroperoxide reductase gene ahpC of Xanthomonas oryzae pv. oryzae affects hydrogen peroxide accumulation during the rice–pathogen interaction. Research in Microbiology, 2014, 165, 605-611.	1.0	2
25	Differentially-expressed genes in rice infected by Xanthomonas oryzae pv. oryzae relative to a flagellin-deficient mutant reveal potential functions of flagellin in host–pathogen interactions. Rice, 2014, 7, 20.	1.7	18
26	A Novel Two-Component System PdeK/PdeR Regulates c-di-GMP Turnover and Virulence of <i>Xanthomonas oryzae</i> pv. <i>oryzae</i> Molecular Plant-Microbe Interactions, 2012, 25, 1361-1369.	1.4	78
27	ETHYLENE INSENSITIVE3 and ETHYLENE INSENSITIVE3-LIKE1 Repress <i>SALICYLIC ACID INDUCTION DEFICIENT2</i> Expression to Negatively Regulate Plant Innate Immunity in <i>Arabidopsis</i> ÂÂ. Plant Cell, 2009, 21, 2527-2540.	3.1	267
28	Reporter-based screen for Arabidopsis mutants compromised in nonhost resistance. Science Bulletin, 2008, 53, 1027-1034.	4.3	3
29	Firefly Luciferase Complementation Imaging Assay for Protein-Protein Interactions in Plants. Plant Physiology, 2008, 146, 323-324.	2.3	989