

Crk Consortium

List of Publications by Year in descending order

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Version: 2024-02-01

103
papers

10,219
citations

44069

48
h-index

37204

96
g-index

122
all docs

122
docs citations

122
times ranked

9821
citing authors

#	ARTICLE	IF	CITATIONS
1	Transgenic approaches for plant disease control: Status and prospects 2021. <i>Plant Pathology</i> , 2022, 71, 207-225.	2.4	30
2	Regulation of Tomato Specialised Metabolism after Establishment of Symbiosis with the Endophytic Fungus <i>Serendipita indica</i> . <i>Microorganisms</i> , 2022, 10, 194.	3.6	8
3	The Fungal Endophyte <i>Penicillium olsonii</i> ML37 Reduces <i>Fusarium</i> Head Blight by Local Induced Resistance in Wheat Spikes. <i>Journal of Fungi (Basel, Switzerland)</i> , 2022, 8, 345.	3.5	8
4	Biological control of plant diseases – What has been achieved and what is the direction?. <i>Plant Pathology</i> , 2022, 71, 1024-1047.	2.4	78
5	Identification and Functional Characterisation of Two Oat UDP-Glucosyltransferases Involved in Deoxynivalenol Detoxification. <i>Toxins</i> , 2022, 14, 446.	3.4	5
6	Succession of the fungal endophytic microbiome of wheat is dependent on tissue-specific interactions between host genotype and environment. <i>Science of the Total Environment</i> , 2021, 759, 143804.	8.0	64
7	A Sesquiterpene Synthase from the Endophytic Fungus <i>Serendipita indica</i> Catalyzes Formation of <i>Viridiflorol</i> . <i>Biomolecules</i> , 2021, 11, 898.	4.0	12
8	<i>Fusarium</i> Head Blight Modifies Fungal Endophytic Communities During Infection of Wheat Spikes. <i>Microbial Ecology</i> , 2020, 79, 397-408.	2.8	56
9	Identification of two endophytic fungi that control <i>Septoria tritici</i> blotch in the field, using a structured screening approach. <i>Biological Control</i> , 2020, 141, 104128.	3.0	25
10	A 2-kb Mycovirus Converts a Pathogenic Fungus into a Beneficial Endophyte for Brassica Protection and Yield Enhancement. <i>Molecular Plant</i> , 2020, 13, 1420-1433.	8.3	113
11	Editorial: Plant Disease Management in the Post-genomic Era: From Functional Genomics to Genome Editing. <i>Frontiers in Microbiology</i> , 2020, 11, 107.	3.5	8
12	Selection of fungal endophytes with biocontrol potential against <i>Fusarium</i> head blight in wheat. <i>Biological Control</i> , 2020, 144, 104222.	3.0	82
13	Insights into the community structure and lifestyle of the fungal root endophytes of tomato by combining amplicon sequencing and isolation approaches with phytohormone profiling. <i>FEMS Microbiology Ecology</i> , 2020, 96, .	2.7	31
14	Race specificity and plant immunity.. , 2020, , 216-233.		3
15	Biological control of plant diseases.. , 2020, , 289-306.		1
16	Defining the twig fungal communities of <i>Fraxinus</i> species and <i>Fraxinus excelsior</i> genotypes with differences in susceptibility to ash dieback. <i>Fungal Ecology</i> , 2019, 42, 100859.	1.6	8
17	Searching for Novel Fungal Biological Control Agents for Plant Disease Control Among Endophytes. , 2019, , 25-51.		29
18	Transgenic crops and beyond: how can biotechnology contribute to the sustainable control of plant diseases?. <i>European Journal of Plant Pathology</i> , 2018, 152, 977-986.	1.7	10

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19	A cerato-platanin protein SsCP1 targets plant PR1 and contributes to virulence of <i>Sclerotinia sclerotiorum</i> . <i>New Phytologist</i> , 2018, 217, 739-755.	7.3	211
20	Fungal communities associated with species of <i>Fraxinus</i> tolerant to ash dieback, and their potential for biological control. <i>Fungal Biology</i> , 2018, 122, 110-120.	2.5	54
21	Endophytic fungi as biocontrol agents: elucidating mechanisms in disease suppression. <i>Plant Ecology and Diversity</i> , 2018, 11, 555-567.	2.4	159
22	Fusarium diseases: biology and management perspectives. <i>Burleigh Dodds Series in Agricultural Science</i> , 2018, , 23-45.	0.2	3
23	<i>Azadirachta indica</i> Reduces Black Sigatoka in East African Highland Banana by Direct Antimicrobial Effects against <i>Mycosphaerella fijiensis</i> without Inducing Resistance. <i>Journal of Agricultural Science</i> , 2017, 9, 61.	0.2	3
24	Large-Scale Phenomics Identifies Primary and Fine-Tuning Roles for CRKs in Responses Related to Oxidative Stress. <i>PLoS Genetics</i> , 2015, 11, e1005373.	3.5	167
25	Insights on the Evolution of Mycoparasitism from the Genome of <i>Clonostachys rosea</i> . <i>Genome Biology and Evolution</i> , 2015, 7, 465-480.	2.5	150
26	Activity-guided separation of <i>Chromolaena odorata</i> leaf extract reveals fractions with rice disease-reducing properties. <i>European Journal of Plant Pathology</i> , 2015, 143, 331-341.	1.7	3
27	Transcriptomic profiling to identify genes involved in Fusarium mycotoxin Deoxynivalenol and Zearalenone tolerance in the mycoparasitic fungus <i>Clonostachys rosea</i> . <i>BMC Genomics</i> , 2014, 15, 55.	2.8	61
28	Zearalenone detoxification by zearalenone hydrolase is important for the antagonistic ability of <i>Clonostachys rosea</i> against mycotoxigenic <i>Fusarium graminearum</i> . <i>Fungal Biology</i> , 2014, 118, 364-373.	2.5	99
29	The ash dieback crisis: genetic variation in resistance can prove a long-term solution. <i>Plant Pathology</i> , 2014, 63, 485-499.	2.4	191
30	The barley HvNAC6 transcription factor affects ABA accumulation and promotes basal resistance against powdery mildew. <i>Plant Molecular Biology</i> , 2013, 83, 577-590.	3.9	54
31	Fitness costs and trade-offs in plant disease. <i>Plant Pathology</i> , 2013, 62, 1-1.	2.4	18
32	Proteomic changes and endophytic micromycota during storage of organically and conventionally grown carrots. <i>Postharvest Biology and Technology</i> , 2013, 76, 26-33.	6.0	17
33	<i>Fusarium graminearum</i> and Its Interactions with Cereal Heads: Studies in the Proteomics Era. <i>Frontiers in Plant Science</i> , 2013, 4, 37.	3.6	84
34	The influence of the fungal pathogen <i>Mycocentrospora acerina</i> on the proteome and polyacetylenes and 6-methoxymellein in organic and conventionally cultivated carrots (<i>Daucus carota</i>) during post harvest storage. <i>Journal of Proteomics</i> , 2012, 75, 962-977.	2.4	18
35	Regulation of basal resistance by a powdery mildew-induced cysteine-rich receptor-like protein kinase in barley. <i>Molecular Plant Pathology</i> , 2012, 13, 135-147.	4.2	62
36	Secretomics identifies <i>Fusarium graminearum</i> proteins involved in the interaction with barley and wheat. <i>Molecular Plant Pathology</i> , 2012, 13, 445-453.	4.2	83

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37	Interaction of barley powdery mildew effector candidate <sc>CSEP0055</sc> with the defence protein <sc>PR17c</sc>. <i>Molecular Plant Pathology</i> , 2012, 13, 1110-1119.	4.2	115
38	Fusarium Head Blight of Cereals in Denmark: Species Complex and Related Mycotoxins. <i>Phytopathology</i> , 2011, 101, 960-969.	2.2	152
39	Disease-Reducing Effect of <i>Chromolaena odorata</i> Extract on Sheath Blight and Other Rice Diseases. <i>Phytopathology</i> , 2011, 101, 231-240.	2.2	20
40	Engineering Pathogen Resistance in Crop Plants: Current Trends and Future Prospects. <i>Annual Review of Phytopathology</i> , 2010, 48, 269-291.	7.8	164
41	Investigation of the effect of nitrogen on severity of Fusarium Head Blight in barley. <i>Journal of Proteomics</i> , 2010, 73, 743-752.	2.4	49
42	Analysis of early events in the interaction between <i>Fusarium graminearum</i> and the susceptible barley (<i>Hordeum vulgare</i>) cultivar Scarlett. <i>Proteomics</i> , 2010, 10, 3748-3755.	2.2	55
43	Cell wall appositions: the first line of defence. <i>Journal of Experimental Botany</i> , 2009, 60, 351-352.	4.8	52
44	A cultivation independent, PCR-based protocol for the direct identification of plant pathogens in infected plant material. <i>European Journal of Plant Pathology</i> , 2009, 123, 473-476.	1.7	12
45	Identification and characterization of barley RNA-directed RNA polymerases. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2009, 1789, 375-385.	1.9	13
46	Effects of β -1,3-glucan from <i>Septoria tritici</i> on structural defence responses in wheat. <i>Journal of Experimental Botany</i> , 2009, 60, 4287-4300.	4.8	124
47	What are the prospects for genetically engineered, disease resistant plants?. <i>European Journal of Plant Pathology</i> , 2008, 121, 217-231.	1.7	77
48	How can we exploit functional genomics approaches for understanding the nature of plant defences? Barley as a case study. <i>European Journal of Plant Pathology</i> , 2008, 121, 257-266.	1.7	8
49	Roles of reactive oxygen species in interactions between plants and pathogens. <i>European Journal of Plant Pathology</i> , 2008, 121, 267-280.	1.7	262
50	Transcriptional regulation by an NAC (NAM-ATAF1, CUC2) transcription factor attenuates ABA signalling for efficient basal defence towards <i>Blumeria graminis</i> f. sp. <i>hordei</i> in <i>Arabidopsis</i> . <i>Plant Journal</i> , 2008, 56, 867-880.	5.7	210
51	Roles of reactive oxygen species in interactions between plants and pathogens. , 2008, , 267-280.		15
52	How can we exploit functional genomics approaches for understanding the nature of plant defences? Barley as a case study. , 2008, , 257-266.		1
53	Role of hydrogen peroxide during the interaction between the hemibiotrophic fungal pathogen <i>Septoria tritici</i> and wheat. <i>New Phytologist</i> , 2007, 174, 637-647.	7.3	220
54	The HvNAC6 transcription factor: a positive regulator of penetration resistance in barley and <i>Arabidopsis</i> . <i>Plant Molecular Biology</i> , 2007, 65, 137-150.	3.9	136

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55	What are the prospects for genetically engineered, disease resistant plants?. , 2007, , 217-231.		0
56	Defense-related genes expressed in Norway spruce roots after infection with the root rot pathogen <i>Ceratobasidium bicorne</i> (anamorph: <i>Rhizoctonia</i> sp.). <i>Tree Physiology</i> , 2005, 25, 1533-1543.	3.1	28
57	The molecular characterization of two barley proteins establishes the novel PR-17 family of pathogenesis-related proteins. <i>Molecular Plant Pathology</i> , 2002, 3, 135-144.	4.2	163
58	Mechanisms involved in control of <i>Blumeria graminis</i> f.sp. <i>hordei</i> in barley treated with mycelial extracts from cultured fungi. <i>Plant Pathology</i> , 2002, 51, 612-620.	2.4	7
59	Do 14-3-3 proteins and plasma membrane H ⁺ -ATPases interact in the barley epidermis in response to the barley powdery mildew fungus?. <i>Plant Molecular Biology</i> , 2002, 49, 137-147.	3.9	50
60	Post-translational modification of barley 14-3-3A is isoform-specific and involves removal of the hypervariable C-terminus. <i>Plant Molecular Biology</i> , 2002, 50, 535-542.	3.9	19
61	14-3-3 proteins and the response to abiotic and biotic stress. <i>Plant Molecular Biology</i> , 2002, 50, 1031-1039.	3.9	175
62	Control of <i>Blumeria graminis</i> f.sp. <i>hordei</i> by treatment with mycelial extracts from cultured fungi. <i>Plant Pathology</i> , 2001, 50, 552-560.	2.4	9
63	The Responses of Plants to Pathogens. , 2001, , 131-158.		1
64	Mechanical transmission of maize rayado fino marafivirus (MRFV) to maize and barley by means of the vascular puncture technique. <i>Plant Pathology</i> , 2000, 49, 302-307.	2.4	21
65	Proton extrusion is an essential signalling component in the HR of epidermal single cells in the barley-powdery mildew interaction. <i>Plant Journal</i> , 2000, 23, 245-254.	5.7	46
66	The Barley/ <i>Blumeria</i> (Syn. <i>Erysiphe</i>) <i>Graminis</i> Interaction. , 2000, , 77-100.		25
67	14-3-3 proteins: eukaryotic regulatory proteins with many functions. <i>Plant Molecular Biology</i> , 1999, 40, 545-554.	3.9	122
68	An epidermis/papilla-specific oxalate oxidase-like protein in the defence response of barley attacked by the powdery mildew fungus. <i>Plant Molecular Biology</i> , 1998, 36, 101-112.	3.9	134
69	A flavonoid 7-O-methyltransferase is expressed in barley leaves in response to pathogen attack. <i>Plant Molecular Biology</i> , 1998, 36, 219-227.	3.9	70
70	A chalcone synthase with an unusual substrate preference is expressed in barley leaves in response to UV light and pathogen attack. <i>Plant Molecular Biology</i> , 1998, 37, 849-857.	3.9	105
71	Molecular Characterization of the Oxalate Oxidase Involved in the Response of Barley to the Powdery Mildew Fungus1. <i>Plant Physiology</i> , 1998, 117, 33-41.	4.8	139
72	The 14-3-3 protein interacts directly with the C-terminal region of the plant plasma membrane H(+)-ATPase.. <i>Plant Cell</i> , 1997, 9, 1805-1814.	6.6	218

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73	Differential gene transcript accumulation in barley leaf epidermis and mesophyll in response to attack by <i>Blumeria graminis</i> f.sp. <i>hordei</i> (syn. <i>Erysiphe graminis</i> f.sp. <i>hordei</i>). <i>Physiological and Molecular Plant Pathology</i> , 1997, 51, 85-97.	2.5	93
74	PCR cloning, DNA sequencing and phylogenetic analysis of a xylanase gene from the phytopathogenic fungus <i>Ascochyta pisi</i> Lib.. <i>Physiological and Molecular Plant Pathology</i> , 1997, 51, 377-389.	2.5	14
75	Expression of a defence-related intercellular barley peroxidase in transgenic tobacco. <i>Plant Science</i> , 1997, 122, 173-182.	3.6	38
76	Subcellular localization of H ₂ O ₂ in plants. H ₂ O ₂ accumulation in papillae and hypersensitive response during the barley-powdery mildew interaction. <i>Plant Journal</i> , 1997, 11, 1187-1194.	5.7	2,406
77	Dimerization Characteristics of the 94-kDa Glucose-Regulated Protein. <i>Journal of Biochemistry</i> , 1996, 120, 249-256.	1.7	30
78	Ethanol increases sensitivity of oxalate oxidase assays and facilitates direct activity staining in SDS gels. <i>Plant Molecular Biology Reporter</i> , 1996, 14, 266-272.	1.8	35
79	Characterization of the transcript of a new class of retroposon-type repetitive element cloned from the powdery mildew fungus, <i>Erysiphe graminis</i> . <i>Molecular Genetics and Genomics</i> , 1996, 250, 477-482.	2.4	13
80	Germin-like oxalate oxidase, a H ₂ O ₂ -producing enzyme, accumulates in barley attacked by the powdery mildew fungus. <i>Plant Journal</i> , 1995, 8, 139-145.	5.7	192
81	A simple model based on known plant defence reactions is sufficient to explain most aspects of nodulation. <i>Journal of Experimental Botany</i> , 1995, 46, 1-18.	4.8	42
82	Nar-1 and Nar-2, Two Loci Required for Mla 12 -Specified Race-Specific Resistance to Powdery Mildew in Barley. <i>Plant Cell</i> , 1994, 6, 983.	6.6	65
83	A putative O-methyltransferase from barley is induced by fungal pathogens and UV light. <i>Plant Molecular Biology</i> , 1994, 26, 1797-1806.	3.9	39
84	Accumulation of a putative guanidine compound in relation to other early defence reactions in epidermal cells of barley and wheat exhibiting resistance to <i>Erysiphe graminis</i> f.sp. <i>hordei</i> . <i>Physiological and Molecular Plant Pathology</i> , 1994, 45, 469-484.	2.5	21
85	Induced resistance in sugar beet against <i>Cercospora beticola</i> : induction by dichloroisonicotinic acid is independent of chitinase and β -1,3-glucanase transcript accumulation. <i>Physiological and Molecular Plant Pathology</i> , 1994, 45, 89-99.	2.5	42
86	Purification, Characterization, and Molecular Cloning of Basic PR-1-Type Pathogenesis-Related Proteins from Barley. <i>Molecular Plant-Microbe Interactions</i> , 1994, 7, 267.	2.6	88
87	Plant chitinases. <i>Plant Journal</i> , 1993, 3, 31-40.	5.7	737
88	A pathogen-induced gene of barley encodes a HSP90 homologue showing striking similarity to vertebrate forms resident in the endoplasmic reticulum. <i>Plant Molecular Biology</i> , 1993, 21, 1097-1108.	3.9	77
89	Accumulation of defence-related transcripts and cloning of a chitinase mRNA from pea leaves (<i>Pisum</i>) Tj ETQq1 1 0,784314 rgBT /Ove	3.6	23
90	cDNA Cloning and Characterization of mRNAs Induced in Barley by the Fungal Pathogen, <i>Erysiphe Graminis</i> . <i>Developments in Plant Pathology</i> , 1993, , 304-307.	0.1	9

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91	14-3-3 proteins: a highly conserved, widespread family of eukaryotic proteins. Trends in Biochemical Sciences, 1992, 17, 498-501.	7.5	478
92	cDNA cloning and characterization of two barley peroxidase transcripts induced differentially by the powdery mildew fungus Erysiphe graminis. Physiological and Molecular Plant Pathology, 1992, 40, 395-409.	2.5	98
93	Cloning and characterization of a pathogen-induced chitinase in Brassica napus. Plant Molecular Biology, 1992, 20, 277-287.	3.9	75
94	A pathogen-induced gene of barley encodes a protein showing high similarity to a protein kinase regulator. Plant Journal, 1992, 2, 815-820.	5.7	53
95	Induction, purification and characterization of chitinase isolated from pea leaves inoculated with Ascochyta pisi. Planta, 1991, 184, 24-9.	3.2	35
96	Early induction of new mRNAs accompanies the resistance reaction of barley to the wheat pathogen, Erysiphe graminis f.sp. tritici. Physiological and Molecular Plant Pathology, 1990, 36, 471-481.	2.5	20
97	Plant gene expression in response to pathogens. Plant Molecular Biology, 1987, 9, 389-410.	3.9	215
98	Gene expression in Brassica campestris showing a hypersensitive response to the incompatible pathogen Xanthomonas campestris pv. vitians. Plant Molecular Biology, 1987, 8, 405-414.	3.9	50
99	The inheritance of cyanoglucoside content in Trifolium repens L.. Biochemical Genetics, 1984, 22, 139-151.	1.7	16
100	Evidence that linamarin and lotaustralin, the two cyanogenic glucosides of Trifolium repens L., are synthesized by a single set of microsomal enzymes controlled by the Ac/ac locus. Plant Science Letters, 1984, 34, 119-125.	1.8	22
101	Developmental and Physiological Studies on the Cyanogenic Glucosides of White Clover, Trifolium repens L.. Journal of Experimental Botany, 1982, 33, 154-161.	4.8	27
102	In vitro characterization of the Ac locus in white clover (Trifolium repens L.). Archives of Biochemistry and Biophysics, 1982, 218, 38-45.	3.0	29
103	The Status and Prospects for Biotechnological Approaches for Attaining Sustainable Disease Resistance. , 0, .		1