

# Antonio Di Pietro

## List of Publications by Year in descending order

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131  
papers

18,396  
citations

36303

51  
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15732

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140  
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140  
docs citations

140  
times ranked

25562  
citing authors

#	ARTICLE	IF	CITATIONS
1	In Vivo Monitoring of Cytosolic pH Using the Ratiometric pH Sensor pHluorin. <i>Methods in Molecular Biology</i> , 2022, 2391, 99-107.	0.9	3
2	Quantification and Isolation of Spontaneous Colony Growth Variants. <i>Methods in Molecular Biology</i> , 2022, 2391, 55-62.	0.9	1
3	Localisation of phosphoinositides in the grass endophyte <i>Epichloa festucae</i> and genetic and functional analysis of key components of their biosynthetic pathway in <i>E. festucae</i> symbiosis and <i>Fusarium oxysporum</i> pathogenesis. <i>Fungal Genetics and Biology</i> , 2022, 159, 103669.	2.1	2
4	<i>Marchantia polymorpha</i> model reveals conserved infection mechanisms in the vascular wilt fungal pathogen <i>Fusarium oxysporum</i> . <i>New Phytologist</i> , 2022, 234, 227-241.	7.3	22
5	Impairment of the cellulose degradation machinery enhances <i>Fusarium oxysporum</i> virulence but limits its reproductive fitness. <i>Science Advances</i> , 2022, 8, eabl9734.	10.3	8
6	Determinants of endophytic and pathogenic lifestyle in root colonizing fungi. <i>Current Opinion in Plant Biology</i> , 2022, 67, 102226.	7.1	23
7	Conserved secreted effectors contribute to endophytic growth and multihost plant compatibility in a vascular wilt fungus. <i>Plant Cell</i> , 2022, 34, 3214-3232.	6.6	20
8	Phylogenomic Analysis of a 55.1-kb 19-Gene Dataset Resolves a Monophyletic <i>Fusarium</i> that Includes the <i>Fusarium solani</i> Species Complex. <i>Phytopathology</i> , 2021, 111, 1064-1079.	2.2	107
9	Lineage-Specific Genes and Cryptic Sex: Parallels and Differences between Arbuscular Mycorrhizal Fungi and Fungal Pathogens. <i>Trends in Plant Science</i> , 2021, 26, 111-123.	8.8	25
10	Dual-specificity protein phosphatase Msg5 controls cell wall integrity and virulence in <i>Fusarium oxysporum</i> . <i>Fungal Genetics and Biology</i> , 2021, 146, 103486.	2.1	5
11	A diversity of resistance sources to <i>Fusarium oxysporum</i> f. sp. <i>pisii</i> found within grass pea germplasm. <i>Plant and Soil</i> , 2021, 463, 19-38.	3.7	12
12	TEfinder: A Bioinformatics Pipeline for Detecting New Transposable Element Insertion Events in Next-Generation Sequencing Data. <i>Genes</i> , 2021, 12, 224.	2.4	4
13	A 'Hydrolase Switch' for Vascular Specialization in Plant Pathogenic Bacteria. <i>Trends in Plant Science</i> , 2021, 26, 427-429.	8.8	3
14	Chemotropic Assay for Testing Fungal Response to Strigolactones and Strigolactone-Like Compounds. <i>Methods in Molecular Biology</i> , 2021, 2309, 105-111.	0.9	1
15	Rapid evolution in plant-microbe interactions – a molecular genomics perspective. <i>New Phytologist</i> , 2020, 225, 1134-1142.	7.3	96
16	A bacterial endophyte exploits chemotropism of a fungal pathogen for plant colonization. <i>Nature Communications</i> , 2020, 11, 5264.	12.8	41
17	No to <i>Neocosmospora</i> : Phylogenomic and Practical Reasons for Continued Inclusion of the <i>Fusarium solani</i> Species Complex in the Genus <i>Fusarium</i> . <i>MSphere</i> , 2020, 5, .	2.9	61
18	Chemotropism Assays for Plant Symbiosis and Mycoparasitism Related Compound Screening in <i>Trichoderma atroviride</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 601251.	3.5	27

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19	Structure of Fungal $\hat{\pm}$ Mating Pheromone in Membrane Mimetics Suggests a Possible Role for Regulation at the Water-Membrane Interface. <i>Frontiers in Microbiology</i> , 2020, 11, 1090.	3.5	5
20	The genome of opportunistic fungal pathogen <i>Fusarium oxysporum</i> carries a unique set of lineage-specific chromosomes. <i>Communications Biology</i> , 2020, 3, 50.	4.4	55
21	Nutritional factors modulating plant and fruit susceptibility to pathogens: BARD workshop, Haifa, Israel, February 25-26, 2018. <i>Phytoparasitica</i> , 2020, 48, 317-333.	1.2	0
22	Expansion Microscopy for Cell Biology Analysis in Fungi. <i>Frontiers in Microbiology</i> , 2020, 11, 574.	3.5	37
23	NADPH oxidase regulates chemotropic growth of the fungal pathogen <i>Fusarium oxysporum</i> towards the host plant. <i>New Phytologist</i> , 2019, 224, 1600-1612.	7.3	48
24	A Conserved Microbial Motif $\hat{\text{Traps}}^{\text{TM}}$ Protease Activation in Host Immunity. <i>Trends in Plant Science</i> , 2019, 24, 665-667.	8.8	0
25	The Role of Volatile Organic Compounds and Rhizosphere Competence in Mode of Action of the Non-pathogenic <i>Fusarium oxysporum</i> FO12 Toward <i>Verticillium</i> Wilt. <i>Frontiers in Microbiology</i> , 2019, 10, 1808.	3.5	27
26	Autocrine pheromone signalling regulates community behaviour in the fungal pathogen <i>Fusarium oxysporum</i> . <i>Nature Microbiology</i> , 2019, 4, 1443-1449.	13.3	54
27	Fusaric acid contributes to virulence of <i>Fusarium oxysporum</i> on plant and mammalian hosts. <i>Molecular Plant Pathology</i> , 2018, 19, 440-453.	4.2	105
28	Improved Assembly of Reference Genome <i>Fusarium oxysporum</i> f. sp. <i>lycopersici</i> Strain Fol4287. <i>Microbiology Resource Announcements</i> , 2018, 7, .	0.6	31
29	Adapt your shuttling proteins for virulence: a lesson from the corn smut fungus <i>Ustilago maydis</i> . <i>New Phytologist</i> , 2018, 220, 353-356.	7.3	2
30	Live-cell imaging of conidial anastomosis tube fusion during colony initiation in <i>Fusarium oxysporum</i> . <i>PLoS ONE</i> , 2018, 13, e0195634.	2.5	21
31	Characterization of a novel cysteine-rich antifungal protein from <i>Fusarium graminearum</i> with activity against maize fungal pathogens. <i>International Journal of Food Microbiology</i> , 2018, 283, 45-51.	4.7	11
32	Three <i>Fusarium oxysporum</i> mitogen-activated protein kinases (MAPKs) have distinct and complementary roles in stress adaptation and cross-kingdom pathogenicity. <i>Molecular Plant Pathology</i> , 2017, 18, 912-924.	4.2	77
33	Histomorphologic and ultrastructural recovery of myopathy in rats treated with low-level laser therapy. <i>Lasers in Medical Science</i> , 2017, 32, 841-849.	2.1	2
34	Stress Adaptation. <i>Microbiology Spectrum</i> , 2017, 5, .	3.0	46
35	Fungal pathogenesis: Combatting the oxidative burst. <i>Nature Microbiology</i> , 2017, 2, 17095.	13.3	12
36	Regulatory Mechanisms of a Highly Pectinolytic Mutant of <i>Penicillium occitanis</i> and Functional Analysis of a Candidate Gene in the Plant Pathogen <i>Fusarium oxysporum</i> . <i>Frontiers in Microbiology</i> , 2017, 8, 1627.	3.5	4

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37	Structure-Activity Relationship of $\pm$ Mating Pheromone from the Fungal Pathogen <i>Fusarium oxysporum</i> . <i>Journal of Biological Chemistry</i> , 2017, 292, 3591-3602.	3.4	13
38	How alkalinization drives fungal pathogenicity. <i>PLoS Pathogens</i> , 2017, 13, e1006621.	4.7	73
39	Plant Defensins NaD1 and NaD2 Induce Different Stress Response Pathways in Fungi. <i>International Journal of Molecular Sciences</i> , 2016, 17, 1473.	4.1	8
40	A conserved chaperone is required for virulence in fungal plant pathogens. <i>New Phytologist</i> , 2016, 209, 1135-1148.	7.3	31
41	A fungal pathogen secretes plant alkalinizing peptides to increase infection. <i>Nature Microbiology</i> , 2016, 1, 16043.	13.3	249
42	Hyphal chemotropism in fungal pathogenicity. <i>Seminars in Cell and Developmental Biology</i> , 2016, 57, 69-75.	5.0	25
43	Editorial. <i>Seminars in Cell and Developmental Biology</i> , 2016, 57, 68.	5.0	0
44	Combined action of the major secreted exo- and endopolygalacturonases is required for full virulence of <i>Fusarium oxysporum</i> . <i>Molecular Plant Pathology</i> , 2016, 17, 339-353.	4.2	50
45	Guidelines for the use and interpretation of assays for monitoring autophagy (3rd edition). <i>Autophagy</i> , 2016, 12, 1-222.	9.1	4,701
46	Data quality aware analysis of differential expression in RNA-seq with NOISeq R/Bioc package. <i>Nucleic Acids Research</i> , 2015, 43, gkv711.	14.5	605
47	Autophagy contributes to regulation of nuclear dynamics during vegetative growth and hyphal fusion in <i>Fusarium oxysporum</i> . <i>Autophagy</i> , 2015, 11, 131-144.	9.1	47
48	Chemotropic sensing in fungus-plant interactions. <i>Current Opinion in Plant Biology</i> , 2015, 26, 135-140.	7.1	46
49	Fungal pathogen uses sex pheromone receptor for chemotropic sensing of host plant signals. <i>Nature</i> , 2015, 527, 521-524.	27.8	164
50	The transmembrane protein <i>S</i> ho1 cooperates with the mucin <i>M</i> sb2 to regulate invasive growth and plant infection in <i>Fusarium oxysporum</i> . <i>Molecular Plant Pathology</i> , 2015, 16, 593-603.	4.2	37
51	Murine Model for <i>Fusarium oxysporum</i> Invasive Fusariosis Reveals Organ-Specific Structures for Dissemination and Long-Term Persistence. <i>PLoS ONE</i> , 2014, 9, e89920.	2.5	14
52	Hyphal Growth of Phagocytosed <i>Fusarium oxysporum</i> Causes Cell Lysis and Death of Murine Macrophages. <i>PLoS ONE</i> , 2014, 9, e101999.	2.5	9
53	The Secreted Peptide PIP1 Amplifies Immunity through Receptor-Like Kinase 7. <i>PLoS Pathogens</i> , 2014, 10, e1004331.	4.7	186
54	Protein Kinases in Plant-Pathogenic Fungi: Conserved Regulators of Infection. <i>Annual Review of Phytopathology</i> , 2014, 52, 267-288.	7.8	199

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55	Fungal model systems and the elucidation of pathogenicity determinants. <i>Fungal Genetics and Biology</i> , 2014, 70, 42-67.	2.1	133
56	Combinatorial function of velvet and AreA in transcriptional regulation of nitrate utilization and secondary metabolism. <i>Fungal Genetics and Biology</i> , 2014, 62, 78-84.	2.1	34
57	The velvet complex governs mycotoxin production and virulence of <i>Fusarium oxysporum</i> on plant and mammalian hosts. <i>Molecular Microbiology</i> , 2013, 87, 49-65.	2.5	132
58	One Fungus, One Name: Defining the Genus <i>Fusarium</i> in a Scientifically Robust Way That Preserves Longstanding Use. <i>Phytopathology</i> , 2013, 103, 400-408.	2.2	219
59	Targeting Iron Acquisition Blocks Infection with the Fungal Pathogens <i>Aspergillus fumigatus</i> and <i>Fusarium oxysporum</i> . <i>PLoS Pathogens</i> , 2013, 9, e1003436.	4.7	101
60	Iron competition in fungus-plant interactions. <i>Plant Signaling and Behavior</i> , 2013, 8, e23012.	2.4	9
61	The CAP protein superfamily: function in sterol export and fungal virulence. <i>Biomolecular Concepts</i> , 2013, 4, 519-525.	2.2	61
62	HapX-Mediated Iron Homeostasis Is Essential for Rhizosphere Competence and Virulence of the Soilborne Pathogen <i>Fusarium oxysporum</i> . <i>Plant Cell</i> , 2012, 24, 3805-3822.	6.6	138
63	A PR-1-like Protein of <i>Fusarium oxysporum</i> Functions in Virulence on Mammalian Hosts. <i>Journal of Biological Chemistry</i> , 2012, 287, 21970-21979.	3.4	52
64	Role of new bacterial surfactins in the antifungal interaction between <i>Bacillus amyloliquefaciens</i> and <i>Fusarium oxysporum</i> . <i>Plant Pathology</i> , 2012, 61, 689-699.	2.4	63
65	The Top 10 fungal pathogens in molecular plant pathology. <i>Molecular Plant Pathology</i> , 2012, 13, 414-430.	4.2	3,270
66	The Top 10 fungal pathogens in molecular plant pathology. <i>Molecular Plant Pathology</i> , 2012, 13, 804-804.	4.2	72
67	The Top 10 fungal pathogens in molecular plant pathology. <i>Molecular Plant Pathology</i> , 2012, , no-no.	4.2	22
68	Signaling of Infectious Growth in <i>Fusarium oxysporum</i> . <i>Topics in Current Genetics</i> , 2012, , 61-79.	0.7	0
69	<i>Galleria mellonella</i> as model host for the trans-kingdom pathogen <i>Fusarium oxysporum</i> . <i>Fungal Genetics and Biology</i> , 2011, 48, 1124-1129.	2.1	47
70	Antifungal Lipopeptides from <i>Bacillus amyloliquefaciens</i> Strain BO7. <i>Journal of Natural Products</i> , 2011, 74, 145-151.	3.0	49
71	The Membrane Mucin Msb2 Regulates Invasive Growth and Plant Infection in <i>Fusarium oxysporum</i> . <i>Plant Cell</i> , 2011, 23, 1171-1185.	6.6	97
72	The Molecular Pathogenicity of <i>Fusarium Keratitis</i> . <i>Cornea</i> , 2010, 29, 1440-1444.	1.7	16

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73	The two-component histidine kinase Fhk1 controls stress adaptation and virulence of <i>Fusarium oxysporum</i> . <i>Molecular Plant Pathology</i> , 2010, 11, 395-407.	4.2	62
74	Comparative genomics reveals mobile pathogenicity chromosomes in <i>Fusarium</i> . <i>Nature</i> , 2010, 464, 367-373.	27.8	1,442
75	Nuclear Dynamics during Germination, Conidiation, and Hyphal Fusion of <i>Fusarium oxysporum</i> . <i>Eukaryotic Cell</i> , 2010, 9, 1216-1224.	3.4	60
76	A Nitrogen Response Pathway Regulates Virulence Functions in <i>Fusarium oxysporum</i> via the Protein Kinase TOR and the bZIP Protein MeaB. <i>Plant Cell</i> , 2010, 22, 2459-2475.	6.6	207
77	The homeodomain transcription factor Ste12. <i>Communicative and Integrative Biology</i> , 2010, 3, 327-332.	1.4	45
78	A nitrogen response pathway regulates virulence in plant pathogenic fungi. <i>Plant Signaling and Behavior</i> , 2010, 5, 1623-1625.	2.4	17
79	The <i>Fusarium oxysporum</i> cell wall proteome under adhesion-inducing conditions. <i>Proteomics</i> , 2009, 9, 4755-4769.	2.2	34
80	Identification of virulence genes in <i>Fusarium oxysporum</i> f. sp. <i>lycopersici</i> by large-scale transposon tagging. <i>Molecular Plant Pathology</i> , 2009, 10, 95-107.	4.2	40
81	Possible involvement of C-proteins and cAMP in the induction of progesterone hydroxylating enzyme system in the vascular wilt fungus <i>Fusarium oxysporum</i> . <i>Journal of Steroid Biochemistry and Molecular Biology</i> , 2009, 113, 241-247.	2.5	2
82	From Tools of Survival to Weapons of Destruction: The Role of Cell Wall-Degrading Enzymes in Plant Infection. , 2009, , 181-200.		9
83	Comparative genomics of MAP kinase and calcium-calcieneurin signalling components in plant and human pathogenic fungi. <i>Fungal Genetics and Biology</i> , 2009, 46, 287-298.	2.1	302
84	<i>Fusarium oxysporum</i> Ste12 Controls Invasive Growth and Virulence Downstream of the Fmk1 MAPK Cascade. <i>Molecular Plant-Microbe Interactions</i> , 2009, 22, 830-839.	2.6	87
85	Differential toxicity of antifungal protein AFP against mutants of <i>Fusarium oxysporum</i> . <i>International Microbiology</i> , 2009, 12, 115-21.	2.4	12
86	Rho1 has distinct functions in morphogenesis, cell wall biosynthesis and virulence of <i>Fusarium oxysporum</i> . <i>Cellular Microbiology</i> , 2008, 10, 1339-1351.	2.1	75
87	Ctf1, a transcriptional activator of cutinase and lipase genes in <i>Fusarium oxysporum</i> is dispensable for virulence. <i>Molecular Plant Pathology</i> , 2008, 9, 293-304.	4.2	33
88	Isolation and expression analysis of a cobalamin-independent methionine synthase gene from the parasitic plant <i>Orobanche ramosa</i> . <i>Scientia Horticulturae</i> , 2008, 116, 337-341.	3.6	2
89	Regulatory elements mediating expression of xylanase genes in <i>Fusarium oxysporum</i> . <i>Fungal Genetics and Biology</i> , 2008, 45, 28-34.	2.1	8
90	The <i>Fusarium oxysporum</i> sti35 gene functions in thiamine biosynthesis and oxidative stress response. <i>Fungal Genetics and Biology</i> , 2008, 45, 6-16.	2.1	23

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91	Vegetative Hyphal Fusion Is Not Essential for Plant Infection by <i>Fusarium oxysporum</i> . <i>Eukaryotic Cell</i> , 2008, 7, 162-171.	3.4	98
92	Role of the Transcriptional Activator XlnR of <i>Fusarium oxysporum</i> in Regulation of Xylanase Genes and Virulence. <i>Molecular Plant-Microbe Interactions</i> , 2007, 20, 977-985.	2.6	73
93	The <i>Fusarium graminearum</i> Genome Reveals a Link Between Localized Polymorphism and Pathogen Specialization. <i>Science</i> , 2007, 317, 1400-1402.	12.6	837
94	Identification and expression analysis of a MYB family transcription factor in the parasitic plant <i>Orobanche ramosa</i> . <i>Annals of Applied Biology</i> , 2007, 150, 123-130.	2.5	5
95	Distinct signalling pathways coordinately contribute to virulence of <i>Fusarium oxysporum</i> on mammalian hosts. <i>Microbes and Infection</i> , 2006, 8, 2825-2831.	1.9	20
96	A peroxidase gene expressed during early developmental stages of the parasitic plant <i>Orobanche ramosa</i> . <i>Journal of Experimental Botany</i> , 2006, 57, 185-192.	4.8	19
97	Nitrogen-responsive genes are differentially regulated in planta during <i>Fusarium oxysporum</i> f. sp. <i>lycopersici</i> infection. <i>Molecular Plant Pathology</i> , 2005, 6, 459-470.	4.2	43
98	<i>Fusarium oxysporum</i> gas1 Encodes a Putative $\beta$ -1, 3-Glucanoyltransferase Required for Virulence on Tomato Plants. <i>Molecular Plant-Microbe Interactions</i> , 2005, 18, 1140-1147.	2.6	62
99	An Improved Axenic System for Studying Pre-infection Development of the Parasitic Plant <i>Orobanche ramosa</i> . <i>Annals of Botany</i> , 2005, 96, 1121-1127.	2.9	18
100	G-protein $\gamma$ subunit Fgb1 regulates hyphal growth, development, and virulence through multiple signalling pathways. <i>Fungal Genetics and Biology</i> , 2005, 42, 61-72.	2.1	61
101	<i>Fusarium oxysporum</i> as a Multihost Model for the Genetic Dissection of Fungal Virulence in Plants and Mammals. <i>Infection and Immunity</i> , 2004, 72, 1760-1766.	2.2	164
102	Class V chitin synthase determines pathogenesis in the vascular wilt fungus <i>Fusarium oxysporum</i> and mediates resistance to plant defence compounds. <i>Molecular Microbiology</i> , 2003, 47, 257-266.	2.5	139
103	The pH signalling transcription factor PacC controls virulence in the plant pathogen <i>Fusarium oxysporum</i> . <i>Molecular Microbiology</i> , 2003, 48, 765-779.	2.5	196
104	<i>Fusarium oxysporum</i> : exploring the molecular arsenal of a vascular wilt fungus. <i>Molecular Plant Pathology</i> , 2003, 4, 315-325.	4.2	360
105	<i>Fusarium</i> as a model for studying virulence in soilborne plant pathogens. <i>Physiological and Molecular Plant Pathology</i> , 2003, 62, 87-98.	2.5	123
106	pH Response Transcription Factor PacC Controls Salt Stress Tolerance and Expression of the P-Type Na <sup>+</sup> -ATPase <i>Ena1</i> in <i>Fusarium oxysporum</i> . <i>Eukaryotic Cell</i> , 2003, 2, 1246-1252.	3.4	76
107	Role in Pathogenesis of Two Endo- $\beta$ -1,4-xylanase Genes from the Vascular Wilt Fungus <i>Fusarium oxysporum</i> . <i>Fungal Genetics and Biology</i> , 2002, 35, 213-222.	2.1	96
108	Molecular Characterization of a Subtilase from the Vascular Wilt Fungus <i>Fusarium oxysporum</i> . <i>Molecular Plant-Microbe Interactions</i> , 2001, 14, 653-662.	2.6	41

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109	Molecular characterization of a novel endo- $\beta$ -1,4-xylanase gene from the vascular wilt fungus <i>Fusarium oxysporum</i> . <i>Current Genetics</i> , 2001, 40, 268-275.	1.7	62
110	Molecular Characterization of an Endopolygalacturonase from <i>Fusarium oxysporum</i> Expressed during Early Stages of Infection. <i>Applied and Environmental Microbiology</i> , 2001, 67, 2191-2196.	3.1	84
111	A MAP kinase of the vascular wilt fungus <i>Fusarium oxysporum</i> is essential for root penetration and pathogenesis. <i>Molecular Microbiology</i> , 2001, 39, 1140-1152.	2.5	378
112	A MAP kinase of the vascular wilt fungus <i>Fusarium oxysporum</i> is essential for root penetration and pathogenesis. <i>Molecular Microbiology</i> , 2001, 39, 1140-1152.	2.5	9
113	Cloning and Disruption of <i>pgx4</i> Encoding an In Planta Expressed Exopolygalacturonase from <i>Fusarium oxysporum</i> . <i>Molecular Plant-Microbe Interactions</i> , 2000, 13, 359-365.	2.6	73
114	Genetic variability in callus formation and regeneration of garlic ( <i>Allium sativum</i> L.). <i>Plant Cell Reports</i> , 1999, 18, 434-437.	5.6	28
115	Cloning and characterization of <i>pl1</i> encoding an in planta-secreted pectate lyase of <i>Fusarium oxysporum</i> . <i>Current Genetics</i> , 1999, 35, 36-40.	1.7	55
116	Two xylanase genes of the vascular wilt pathogen <i>Fusarium oxysporum</i> are differentially expressed during infection of tomato plants. <i>Molecular Genetics and Genomics</i> , 1999, 261, 530-536.	2.4	57
117	An efficient method for the in vitro management of multiple garlic accessions. <i>In Vitro Cellular and Developmental Biology - Plant</i> , 1999, 35, 466-469.	2.1	12
118	Cross protection provides evidence for race-specific avirulence factors in <i>Fusarium oxysporum</i> . <i>Physiological and Molecular Plant Pathology</i> , 1999, 54, 63-72.	2.5	32
119	An Efficient Method for Callus Culture and Shoot Regeneration of Garlic ( <i>Allium sativum</i> L.). <i>Hortscience: A Publication of the American Society for Horticultural Science</i> , 1999, 34, 348-349.	1.0	25
120	Biolistic transfer and expression of a <i>uidA</i> reporter gene in different tissues of <i>Allium sativum</i> L.. <i>Plant Cell Reports</i> , 1998, 17, 737-741.	5.6	39
121	Cloning, Expression, and Role in Pathogenicity of <i>pg1</i> Encoding the Major Extracellular Endopolygalacturonase of the Vascular Wilt Pathogen <i>Fusarium oxysporum</i> . <i>Molecular Plant-Microbe Interactions</i> , 1998, 11, 91-98.	2.6	268
122	Endopolygalacturonase PG1 in Different Formae Speciales of <i>Fusarium oxysporum</i> . <i>Applied and Environmental Microbiology</i> , 1998, 64, 1967-1971.	3.1	19
123	Purification and characterization of an acidic endo- $\beta$ -1,4-xylanase from the tomato vascular pathogen <i>Fusarium oxysporum</i> f. sp. <i>lycopersici</i> . <i>FEMS Microbiology Letters</i> , 1997, 148, 75-82.	1.8	18
124	Purification and characterization of a novel exopolygalacturonase from <i>Fusarium oxysporum</i> f.sp. <i>lycopersici</i> . <i>FEMS Microbiology Letters</i> , 1997, 154, 37-43.	1.8	28
125	Purification and characterization of a pectate lyase from <i>Fusarium oxysporum</i> f.sp. <i>lycopersici</i> produced on tomato vascular tissue. <i>Physiological and Molecular Plant Pathology</i> , 1996, 49, 177-185.	2.5	24
126	Purification and characterization of an exo-polygalacturonase from the tomato vascular wilt pathogen <i>Fusarium oxysporum</i> f.sp. <i>lycopersici</i> . <i>FEMS Microbiology Letters</i> , 1996, 145, 295-299.	1.8	57



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127	Occurrence of a retrotransposon-like sequence among different formae speciales and races of <i>Fusarium oxysporum</i> . <i>Mycological Research</i> , 1994, 98, 993-996.	2.5	16
128	Isolation and sequence of an endochitinase-encoding gene from a cDNA library of <i>Trichoderma harzianum</i> . <i>Gene</i> , 1994, 138, 143-148.	2.2	127
129	Biolistic transformation of <i>Trichoderma harzianum</i> and <i>Gliocladium virens</i> using plasmid and genomic DNA. <i>Current Genetics</i> , 1993, 24, 349-356.	1.7	90
130	Stress Adaptation. , 0, , 463-485.		9
131	CHAPTER 3: Host/Pathogen Interactions. , 0, , 21-38.		0