

# Gregory B Martin

## List of PR Articles by Year in descending order

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17,044

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7147

68

PR h-index

9399

128

g-index

181

documents

19259

doc citations

7912

72

h-index

14670

citing authors

#	ARTICLE	IF	PR CITATIONS
1	Helper <sc>NLRs</sc> Nrc2 and Nrc3 act codependently with Prf/Pto and activate MAPK signaling to induce immunity in tomato. <i>Plant Journal</i> , 2024, 117, 7-22.	6.2	20
2	Natural variation of immune epitopes reveals intrabacterial antagonism. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2024, 121, .	7.8	17
3	<sc>PP2C</sc> phosphatase Pic14 negatively regulates tomato Pto/Prfâ€ triggered immunity by inhibiting <sc>MAPK</sc> activation. <i>Plant Journal</i> , 2024, 119, 2622-2637.	6.2	3
4	Related type 2C protein phosphatases Pic3 and Pic12 negatively regulate immunity in tomato to <i>Pseudomonas syringae</i>. <i>Plant Physiology</i> , 2024, 196, 1997-2013.	5.5	2
5	Tomato receptor-like cytoplasmic kinase Fir1 is involved in flagellin signaling and preinvasion immunity. <i>Plant Physiology</i> , 2023, 192, 565-581.	5.5	12
6	Ptr1 is a CC-NLR immune receptor that mediates recognition of diverse bacterial effectors in multiple solanaceous plants. <i>Physiological and Molecular Plant Pathology</i> , 2023, 125, 101997.	3.5	4
7	Loss-of-function mutations in WRKY22 and WRKY25 impair stomatal-mediated immunity and PTI and ETI responses against <i>Pseudomonas syringae</i> pv. tomato. <i>Plant Molecular Biology</i> , 2023, 112, 161-177.	3.3	14
8	Ptr1 and <sc>ZAR1</sc> immune receptors confer overlapping and distinct bacterial pathogen effector specificities. <i>New Phytologist</i> , 2023, 239, 1935-1953.	8.2	31
9	A <i>Solanum lycopersicoides</i> reference genome facilitates insights into tomato specialized metabolism and immunity. <i>Plant Journal</i> , 2022, 110, 1791-1810.	6.2	30
10	The Emerging Role of PP2C Phosphatases in Tomato Immunity. <i>Molecular Plant-Microbe Interactions</i> , 2022, 35, 737-747.	3.3	16
11	Loss of function of the bHLH transcription factor Nrd1 in tomato enhances resistance to <i>Pseudomonas syringae</i>. <i>Plant Physiology</i> , 2022, 190, 1334-1348.	5.5	16
12	Spelling Changes and Fluorescent Tagging With Prime Editing Vectors for Plants. <i>Frontiers in Genome Editing</i> , 2021, 3, .	4.1	47
13	Integrative Proteomic and Phosphoproteomic Analyses of Pattern- and Effector-Trigged Immunity in Tomato. <i>Frontiers in Plant Science</i> , 2021, 12, .	4.1	25
14	Genome of <i>Solanum pimpinellifolium</i> provides insights into structural variants during tomato breeding. <i>Nature Communications</i> , 2020, 11, .	13.9	128
15	Tomato Wall-Associated Kinase SlWak1 Depends on Fls2/Fls3 to Promote Apoplastic Immune Responses to <i>Pseudomonas syringae</i>. <i>Plant Physiology</i> , 2020, 183, 1869-1882.	5.5	78
16	<i>Ptr1</i> evolved convergently with <i>RPS2</i> and <i>Mr5</i> to mediate recognition of AvrRpt2 in diverse solanaceous species. <i>Plant Journal</i> , 2020, 103, 1433-1445.	6.2	47
17	Molecular Characterization of Differences between the Tomato Immune Receptors Flagellin Sensing 3 and Flagellin Sensing 2. <i>Plant Physiology</i> , 2020, 183, 1825-1837.	5.5	37
18	WRKY22 and WRKY25 transcription factors are positive regulators of defense responses in <i>Nicotiana benthamiana</i> . <i>Plant Molecular Biology</i> , 2020, 105, 65-82.	3.3	46

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19	Generation and Molecular Characterization of CRISPR/Cas9-Induced Mutations in 63 Immunity-Associated Genes in Tomato Reveals Specificity and a Range of Gene Modifications. <i>Frontiers in Plant Science</i> , 2020, 11, .	4.1	66
20	Mai1 Protein Acts Between Host Recognition of Pathogen Effectors and Mitogen-Activated Protein Kinase Signaling. <i>Molecular Plant-Microbe Interactions</i> , 2019, 32, 1496-1507.	3.3	25
21	The tomato <i>Pto</i> gene confers resistance to <i>Pseudomonas floricida</i> , an emergent plant pathogen with just nine type III effectors. <i>Plant Pathology</i> , 2019, 68, 977-984.	2.6	5
22	PP2C phosphatase Pic1 negatively regulates the phosphorylation status of Pti1b kinase, a regulator of flagellin-triggered immunity in tomato. <i>Biochemical Journal</i> , 2019, 476, 1621-1635.	3.9	23
23	Transcriptome-based identification and validation of reference genes for plant-bacteria interaction studies using <i>Nicotiana benthamiana</i> . <i>Scientific Reports</i> , 2019, 9, .	3.5	48
24	Natural variation for unusual host responses and flagellin-mediated immunity against <i>Pseudomonas syringae</i> in genetically diverse tomato accessions. <i>New Phytologist</i> , 2019, 223, 447-461.	8.2	35
25	The <i>Ptr1</i> Locus of <i>Solanum lycopersoides</i> Confers Resistance to Race 1 Strains of <i>Pseudomonas syringae</i> pv. <i>tomato</i> and to <i>Ralstonia pseudosolanacearum</i> by Recognizing the Type III Effectors AvrRpt2 and RipBN. <i>Molecular Plant-Microbe Interactions</i> , 2019, 32, 949-960.	3.3	52
26	Virus-induced gene silencing database for phenomics and functional genomics in <i>Nicotiana benthamiana</i> . <i>Plant Direct</i> , 2018, 2, .	2.3	18
27	The Bacterial Effector AvrPto Targets the Regulatory Coreceptor SOBIR1 and Suppresses Defense Signaling Mediated by the Receptor-Like Protein Cf-4. <i>Molecular Plant-Microbe Interactions</i> , 2018, 31, 75-85.	3.3	18
28	<i>Pseudomonas syringae</i> pv. <i>tomato</i> Strains from New York Exhibit Virulence Attributes Intermediate Between Typical Race 0 and Race 1 Strains. <i>Plant Disease</i> , 2017, 101, 1442-1448.	2.4	11
29	A Subset of Ubiquitin-Conjugating Enzymes Is Essential for Plant Immunity. <i>Plant Physiology</i> , 2017, 173, 1371-1390.	5.5	60
30	Generation of a Collection of Mutant Tomato Lines Using Pooled CRISPR Libraries. <i>Plant Physiology</i> , 2017, 174, 2023-2037.	5.5	144
31	The Tomato Kinase Pti1 Contributes to Production of Reactive Oxygen Species in Response to Two Flagellin-Derived Peptides and Promotes Resistance to <i>Pseudomonas syringae</i> Infection. <i>Molecular Plant-Microbe Interactions</i> , 2017, 30, 725-738.	3.3	29
32	Use of RNA-seq data to identify and validate RT-qPCR reference genes for studying the tomato- <i>Pseudomonas</i> pathosystem. <i>Scientific Reports</i> , 2017, 7, .	3.5	97
33	Detecting the Interaction of Peptide Ligands with Plant Membrane Receptors. <i>Current Protocols in Plant Biology</i> , 2017, 2, 240-269.	1.0	2
34	Ser360 and Ser364 in the Kinase Domain of Tomato SIMAPKKK1 are Critical for Programmed Cell Death Associated with Plant Immunity. <i>Plant Pathology Journal</i> , 2017, 33, 163-169.	2.4	3
35	Detecting N-myristoylation and S-acylation of host and pathogen proteins in plants using click chemistry. <i>Plant Methods</i> , 2016, 12, .	4.0	25
36	iTAK: A Program for Genome-wide Prediction and Classification of Plant Transcription Factors, Transcriptional Regulators, and Protein Kinases. <i>Molecular Plant</i> , 2016, 9, 1667-1670.	19.2	1,029

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37	Tomato receptor FLAGELLIN-SENSING 3 binds flgII-28 and activates the plant immune system. <i>Nature Plants</i> , 2016, 2, .	11.9	197
38	High-throughput CRISPR Vector Construction and Characterization of DNA Modifications by Generation of Tomato Hairy Roots. <i>Journal of Visualized Experiments</i> , 2016, , .	0.3	34
39	A novel method of transcriptome interpretation reveals a quantitative suppressive effect on tomato immune signaling by two domains in a single pathogen effector protein. <i>BMC Genomics</i> , 2016, 17, .	3.3	10
40	Natural Variation in Tomato Reveals Differences in the Recognition of AvrPto and AvrPtoB Effectors from <i>Pseudomonas syringae</i> . <i>Molecular Plant</i> , 2016, 9, 639-649.	19.2	13
41	Complete Genome Sequence of a Tomato-Infecting Tomato Mottle Mosaic Virus in New York. <i>Genome Announcements</i> , 2015, 3, .	0.7	11
42	Identification of a Candidate Gene in <i>Solanum habrochaites</i> for Resistance to a Race 1 Strain of <i>Pseudomonas syringae</i> pv. <i>tomato</i> . <i>Plant Genome</i> , 2015, 8, .	3.4	15
43	<i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000 Type III Secretion Effector Polymutants Reveal an Interplay between HopAD1 and AvrPtoB. <i>Cell Host and Microbe</i> , 2015, 17, 752-762.	15.5	148
44	The SGN VIGS Tool: User-Friendly Software to Design Virus-Induced Gene Silencing (VIGS) Constructs for Functional Genomics. <i>Molecular Plant</i> , 2015, 8, 486-488.	19.2	209
45	Greasy tactics in the plant-pathogen molecular arms race. <i>Journal of Experimental Botany</i> , 2015, 66, 1607-1616.	5.1	21
46	Acquisition of Iron Is Required for Growth of <i>Salmonella</i> spp. in Tomato Fruit. <i>Applied and Environmental Microbiology</i> , 2015, 81, 3663-3670.	3.5	21
47	Functional genomics of tomato for the study of plant immunity: Table 1. <i>Briefings in Functional Genomics</i> , 2015, 14, 291-301.	2.5	22
48	Five <i>Xanthomonas</i> type III effectors suppress cell death induced by components of immunity-associated MAP kinase cascades. <i>Plant Signaling and Behavior</i> , 2015, 10, e1064573.	3.0	23
49	Natural Variation for Responsiveness to flg22, flgII-28, and csp22 and <i>Pseudomonas syringae</i> pv. <i>tomato</i> in Heirloom Tomatoes. <i>PLoS ONE</i> , 2014, 9, e106119.	2.4	54
50	Transcriptomic analysis reveals tomato genes whose expression is induced specifically during effector-triggered immunity and identifies the Epk1 protein kinase which is required for the host response to three bacterial effector proteins. <i>Genome Biology</i> , 2014, 15, .	8.2	86
51	Pto Kinase Binds Two Domains of AvrPtoB and Its Proximity to the Effector E3 Ligase Determines if It Evades Degradation and Activates Plant Immunity. <i>PLoS Pathogens</i> , 2014, 10, e1004227.	4.5	60
52	Analysis of wild-species introgressions in tomato inbreds uncovers ancestral origins. <i>BMC Plant Biology</i> , 2014, 14, .	4.4	32
53	Transcriptomics-based screen for genes induced by flagellin and repressed by pathogen effectors identifies a cell wall-associated kinase involved in plant immunity. <i>Genome Biology</i> , 2013, 14, .	8.2	155
54	<i>Salmonella</i> colonization activates the plant immune system and benefits from association with plant pathogenic bacteria. <i>Environmental Microbiology</i> , 2013, 15, 2418-2430.	3.8	62

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55	Thymoquinone causes multiple effects, including cell death, on dividing plant cells. <i>Comptes Rendus - Biologies</i> , 2013, 336, 546-556.	0.4	4
56	Two leucines in the N-terminal MAPK docking site of tomato SIMKK2 are critical for interaction with a downstream MAPK to elicit programmed cell death associated with plant immunity. <i>FEBS Letters</i> , 2013, 587, 1460-1465.	2.8	15
57	Allelic variation in two distinct <i>Pseudomonas syringae</i> flagellin epitopes modulates the strength of plant immune responses but not bacterial motility. <i>New Phytologist</i> , 2013, 200, 847-860.	8.2	133
58	The Tomato Fni3 Lysine-63 Specific Ubiquitin-Conjugating Enzyme and Suv Ubiquitin E2 Variant Positively Regulate Plant Immunity. <i>Plant Cell</i> , 2013, 25, 3615-3631.	7.6	66
59	The Tomato Calcium Sensor Cbl10 and Its Interacting Protein Kinase Cipk6 Define a Signaling Pathway in Plant Immunity. <i>Plant Cell</i> , 2013, 25, 2748-2764.	7.6	147
60	Nonhost Resistance of Tomato to the Bean Pathogen <i>Pseudomonas syringae</i> pv. <i>syringae</i> B728a Is Due to a Defective E3 Ubiquitin Ligase Domain in AvrPtoB <sub>B728a</sub> . <i>Molecular Plant-Microbe Interactions</i> , 2013, 26, 387-397.	3.3	12
61	Type III Secretion and Effectors Shape the Survival and Growth Pattern of <i>Pseudomonas syringae</i> on Leaf Surfaces A A. <i>Plant Physiology</i> , 2012, 158, 1803-1818.	5.5	72
62	The Î²-Subunit of the SnRK1 Complex Is Phosphorylated by the Plant Cell Death Suppressor Adi3 A A. <i>Plant Physiology</i> , 2012, 159, 1277-1290.	5.5	38
63	Plant Programmed Cell Death Caused by an Autoactive Form of Prf Is Suppressed by Co-Expression of the Prf LRR Domain. <i>Molecular Plant</i> , 2012, 5, 1058-1067.	19.2	27
64	A Draft Genome Sequence of <i>Nicotiana benthamiana</i> to Enhance Molecular Plant-Microbe Biology Research. <i>Molecular Plant-Microbe Interactions</i> , 2012, 25, 1523-1530.	3.3	457
65	A tomato LysM receptor-like kinase promotes immunity and its kinase activity is inhibited by AvrPtoB. <i>Plant Journal</i> , 2012, 69, 92-103.	6.2	135
66	Structural Analysis of <i>Pseudomonas syringae</i> AvrPtoB Bound to Host BAK1 Reveals Two Similar Kinase-Interacting Domains in a Type III Effector. <i>Cell Host and Microbe</i> , 2011, 10, 616-626.	15.5	131
67	Effector-triggered immunity mediated by the Pto kinase. <i>Trends in Plant Science</i> , 2011, 16, 132-140.	12.1	118
68	Genetic disassembly and combinatorial reassembly identify a minimal functional repertoire of type III effectors in <i>Pseudomonas syringae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 2975-2980.	7.8	239
69	Tomato 14-3-3 Protein TFT7 Interacts with a MAP Kinase Kinase to Regulate Immunity-associated Programmed Cell Death Mediated by Diverse Disease Resistance Proteins. <i>Journal of Biological Chemistry</i> , 2011, 286, 14129-14136.	2.3	82
70	Two virulence determinants of type III effector AvrPto are functionally conserved in diverse <i>Pseudomonas syringae</i> pathovars. <i>New Phytologist</i> , 2010, 187, 969-982.	8.2	20
71	Phosphorylation of the <i>Pseudomonas syringae</i> effector AvrPto is required for FLS2/BAK1-independent virulence activity and recognition by tobacco. <i>Plant Journal</i> , 2010, 61, 16-24.	6.2	34
72	A secreted effector protein (SNE1) from <i>Phytophthora infestans</i> is a broadly acting suppressor of programmed cell death. <i>Plant Journal</i> , 2010, 62, 357-366.	6.2	125

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73	Tomato 14-3-3 Protein 7 Positively Regulates Immunity-Associated Programmed Cell Death by Enhancing Protein Abundance and Signaling Ability of MAPKKK Î±. <i>Plant Cell</i> , 2010, 22, 260-272.	7.6	147
74	Endosome-Associated CRT1 Functions Early in <i>Resistance</i> Gene-Mediated Defense Signaling in <i>Arabidopsis</i> and Tobacco. <i>Plant Cell</i> , 2010, 22, 918-936.	7.6	63
75	Identification of <i>Nicotiana benthamiana</i> Genes Involved in Pathogen-Associated Molecular Pattern-Triggered Immunity. <i>Molecular Plant-Microbe Interactions</i> , 2010, 23, 715-726.	3.3	75
76	Methods to Study PAMP-Triggered Immunity Using Tomato and <i>Nicotiana benthamiana</i>. <i>Molecular Plant-Microbe Interactions</i> , 2010, 23, 991-999.	3.3	207
77	The T-loop Extension of the Tomato Protein Kinase AvrPto-dependent Pto-interacting Protein 3 (Adi3) Directs Nuclear Localization for Suppression of Plant Cell Death. <i>Journal of Biological Chemistry</i> , 2010, 285, 17584-17594.	2.3	34
78	Deletions in the Repertoire of <i>Pseudomonas syringae</i> pv. tomato DC3000 Type III Secretion Effector Genes Reveal Functional Overlap among Effectors. <i>PLoS Pathogens</i> , 2009, 5, e1000388.	4.5	298
79	Crystal Structure of the Complex between <i>Pseudomonas</i> Effector AvrPtoB and the Tomato Pto Kinase Reveals Both a Shared and a Unique Interface Compared with AvrPto-Pto. <i>Plant Cell</i> , 2009, 21, 1846-1859.	7.6	82
80	Advances in experimental methods for the elucidation of <i>Pseudomonas syringae</i> effector function with a focus on AvrPtoB. <i>Molecular Plant Pathology</i> , 2009, 10, 777-793.	5.1	20
81	Virus-induced Gene Silencing (VIGS) in <i>Nicotiana benthamiana</i> and Tomato. <i>Journal of Visualized Experiments</i> , 2009, , .	0.3	133
82	<i>Xanthomonas</i> T3S Effector XopN Suppresses PAMP-Triggered Immunity and Interacts with a Tomato Atypical Receptor-Like Kinase and TFT1. <i>Plant Cell</i> , 2009, 21, 1305-1323.	7.6	189
83	A Draft Genome Sequence of <i>Pseudomonas syringae</i> pv. <i>tomato</i> T1 Reveals a Type III Effector Repertoire Significantly Divergent from That of <i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000. <i>Molecular Plant-Microbe Interactions</i> , 2009, 22, 52-62.	3.3	139
84	Assay for Pathogen-Associated Molecular Pattern (PAMP)-Triggered Immunity (PTI) in Plants. <i>Journal of Visualized Experiments</i> , 2009, , .	0.3	11
85	Bacterial Effectors Target the Common Signaling Partner BAK1 to Disrupt Multiple MAMP Receptor-Signaling Complexes and Impede Plant Immunity. <i>Cell Host and Microbe</i> , 2008, 4, 17-27.	15.5	537
86	<i>Pseudomonas syringae</i> Type III Effector AvrPtoB Is Phosphorylated in Plant Cells on Serine 258, Promoting Its Virulence Activity. <i>Journal of Biological Chemistry</i> , 2007, 282, 30737-30744.	2.3	41
87	Identification and Characterization of Plant Genes Involved in Agrobacterium-Mediated Plant Transformation by Virus-Induced Gene Silencing. <i>Molecular Plant-Microbe Interactions</i> , 2007, 20, 41-52.	3.3	80
88	Pto- and Prf-Mediated Recognition of AvrPto and AvrPtoB Restricts the Ability of Diverse <i>Pseudomonas syringae</i> Pathovars to Infect Tomato. <i>Molecular Plant-Microbe Interactions</i> , 2007, 20, 806-815.	3.3	67
89	An NB-LRR protein required for HR signalling mediated by both extra- and intracellular resistance proteins. <i>Plant Journal</i> , 2007, 50, 14-28.	6.2	189
90	A <i>Pseudomonas syringae</i> pv. tomato DC3000 mutant lacking the type III effector HopQ1-1 is able to cause disease in the model plant <i>Nicotiana benthamiana</i> . <i>Plant Journal</i> , 2007, 51, 32-46.	6.2	319

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91	The N-terminal region of <i>Pseudomonas</i> type III effector AvrPtoB elicits Pto-dependent immunity and has two distinct virulence determinants. <i>Plant Journal</i> , 2007, 52, 595-614.	6.2	85
92	DspA/E, a type III effector of <i>Erwinia amylovora</i> , is required for early rapid growth in <i>Nicotiana benthamiana</i> and causes NbSGT1-dependent cell death. <i>Molecular Plant Pathology</i> , 2007, 8, 255-265.	5.1	34
93	A Bacterial Inhibitor of Host Programmed Cell Death Defenses Is an E3 Ubiquitin Ligase. <i>Science</i> , 2006, 311, 222-226.	37.0	320
94	Comparative Genomics of Host-Specific Virulence in <i>Pseudomonas syringae</i> . <i>Genetics</i> , 2006, 174, 1041-1056.	4.2	145
95	Specific Bacterial Suppressors of MAMP Signaling Upstream of MAPKKK in <i>Arabidopsis</i> Innate Immunity. <i>Cell</i> , 2006, 125, 563-575.	34.4	411
96	Whole-Genome Expression Profiling Defines the HrpL Regulon of <i>Pseudomonas syringae</i> pv. tomato DC3000, Allows de novo Reconstruction of the Hrp cis Element, and Identifies Novel Coregulated Genes. <i>Molecular Plant-Microbe Interactions</i> , 2006, 19, 1167-1179.	3.3	111
97	A novel link between tomato GRAS genes, plant disease resistance and mechanical stress response. <i>Molecular Plant Pathology</i> , 2006, 7, 593-604.	5.1	93
98	Bacterial elicitation and evasion of plant innate immunity. <i>Nature Reviews Molecular Cell Biology</i> , 2006, 7, 601-611.	79.0	382
99	Adi3 is a Pdk1-interacting AGC kinase that negatively regulates plant cell death. <i>EMBO Journal</i> , 2006, 25, 255-265.	7.5	80
100	Host-Mediated Phosphorylation of Type III Effector AvrPto Promotes <i>Pseudomonas</i> Virulence and Avirulence in Tomato. <i>Plant Cell</i> , 2006, 18, 502-514.	7.6	66
101	Type III effector AvrPtoB requires intrinsic E3 ubiquitin ligase activity to suppress plant cell death and immunity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 2851-2856.	7.8	217
102	Diverse AvrPtoB Homologs from Several <i>Pseudomonas syringae</i> Pathovars Elicit Pto-Dependent Resistance and Have Similar Virulence Activities. <i>Applied and Environmental Microbiology</i> , 2006, 72, 702-712.	3.5	67
103	Aconitase plays a role in regulating resistance to oxidative stress and cell death in <i>Arabidopsis</i> and <i>Nicotiana benthamiana</i> . <i>Plant Molecular Biology</i> , 2006, 63, 273-287.	3.3	158
104	An avrPto/avrPtoB Mutant of <i>Pseudomonas syringae</i> pv. tomato DC3000 Does Not Elicit Pto-Mediated Resistance and Is Less Virulent on Tomato. <i>Molecular Plant-Microbe Interactions</i> , 2005, 18, 43-51.	3.3	137
105	AvrPtoB: A bacterial type III effector that both elicits and suppresses programmed cell death associated with plant immunity. <i>FEMS Microbiology Letters</i> , 2005, 245, 1-8.	2.0	64
106	<i>Pseudomonas syringae</i> pv. tomato type III effectors AvrPto and AvrPtoB promote ethylene-dependent cell death in tomato. <i>Plant Journal</i> , 2005, 44, 139-154.	6.2	106
107	Role of mitogen-activated protein kinases in plant immunity. <i>Current Opinion in Plant Biology</i> , 2005, 8, 541-547.	7.2	285
108	Calmodulin-like Proteins from <i>Arabidopsis</i> and Tomato are Involved in Host Defense Against <i>Pseudomonas syringae</i> pv. tomato. <i>Plant Molecular Biology</i> , 2005, 58, 887-897.	3.3	142

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109	Transcriptome and Selected Metabolite Analyses Reveal Multiple Points of Ethylene Control during Tomato Fruit Development. <i>Plant Cell</i> , 2005, 17, 2954-2965.	7.6	492
110	Suppression of pathogen-inducible NO synthase (iNOS) activity in tomato increases susceptibility to <i>Pseudomonas syringae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 8239-8244.	7.8	17
111	PeerGAD: a peer-review-based and community-centric web application for viewing and annotating prokaryotic genome sequences. <i>Nucleic Acids Research</i> , 2004, 32, 3124-3135.	15.8	15
112	Identification of MAPKs and Their Possible MAPK Kinase Activators Involved in the Pto-mediated Defense Response of Tomato. <i>Journal of Biological Chemistry</i> , 2004, 279, 49229-49235.	2.3	116
113	Silencing of subfamily I of protein phosphatase 2A catalytic subunits results in activation of plant defense responses and localized cell death. <i>Plant Journal</i> , 2004, 38, 563-577.	6.2	130
114	Applications and advantages of virus-induced gene silencing for gene function studies in plants. <i>Plant Journal</i> , 2004, 39, 734-746.	6.2	712
115	Comprehensive EST analysis of tomato and comparative genomics of fruit ripening. <i>Plant Journal</i> , 2004, 40, 47-59.	6.2	210
116	MAPKKK1 $\pm$ is a positive regulator of cell death associated with both plant immunity and disease. <i>EMBO Journal</i> , 2004, 23, 3072-3082.	7.5	321
117	The Solution Structure of Type III Effector Protein AvrPto Reveals Conformational and Dynamic Features Important for Plant Pathogenesis. <i>Structure</i> , 2004, 12, 1257-1268.	3.9	52
118	Strategies used by bacterial pathogens to suppress plant defenses. <i>Current Opinion in Plant Biology</i> , 2004, 7, 356-364.	7.2	222
119	Strategies used by bacterial pathogens to suppress plant defenses. <i>Current Opinion in Plant Biology</i> , 2004, , .	7.2	12
120	Identification and Expression Profiling of Tomato Genes Differentially Regulated During a Resistance Response to <i>Xanthomonas campestris</i> pv. <i>vesicatoria</i> . <i>Molecular Plant-Microbe Interactions</i> , 2004, 17, 1212-1222.	3.3	55
121	Molecular Mechanisms Involved in Bacterial Speck Disease Resistance of Tomato. <i>Plant Pathology Journal</i> , 2004, 20, 7-12.	2.4	17
122	<i>Pseudomonas</i> type III effector AvrPtoB induces plant disease susceptibility by inhibition of host programmed cell death. <i>EMBO Journal</i> , 2003, 22, 60-69.	7.5	373
123	UNDERSTANDING THE FUNCTIONS OF PLANT DISEASE RESISTANCE PROTEINS. <i>Annual Review of Plant Biology</i> , 2003, 54, 23-61.	24.7	864
124	Two MAPK cascades, NPR1, and TGA transcription factors play a role in Pto-mediated disease resistance in tomato. <i>Plant Journal</i> , 2003, 36, 905-917.	6.2	337
125	MOLECULAR BASIS OF PTO-MEDIATED RESISTANCE TO BACTERIAL SPECK DISEASE IN TOMATO. <i>Annual Review of Phytopathology</i> , 2003, 41, 215-243.	10.4	315
126	The complete genome sequence of the <i>Arabidopsis</i> and tomato pathogen <i>Pseudomonas syringae</i> pv. <i>tomato</i>	7.8	820

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127	The Tomato Transcription Factor Pti4 Regulates Defense-Related Gene Expression via GCC Box and Non-GCC Box cis Elements[W]. <i>Plant Cell</i> , 2003, 15, 3033-3050.	7.6	273
128	Overexpression of the Disease Resistance Gene Pto in Tomato Induces Gene Expression Changes Similar to Immune Responses in Human and Fruitfly Å. <i>Plant Physiology</i> , 2003, 132, 1901-1912.	5.5	57
129	The tobacco salicylic acid-binding protein 3 (SABP3) is the chloroplast carbonic anhydrase, which exhibits antioxidant activity and plays a role in the hypersensitive defense response. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 11640-11645.	7.8	359
130	Genomewide identification of <i>Pseudomonas syringae</i> pv. tomato DC3000 promoters controlled by the HrpL alternative sigma factor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 2275-2280.	7.8	295
131	Tomato Transcription Factors Pti4, Pti5, and Pti6 Activate Defense Responses When Expressed in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2002, 14, 817-831.	7.6	416
132	Two Distinct <i>Pseudomonas</i> Effector Proteins Interact with the Pto Kinase and Activate Plant Immunity. <i>Cell</i> , 2002, 109, 589-598.	34.4	274
133	Location and activity of members of a family of virPphA homologues in pathovars of <i>Pseudomonas syringae</i> and <i>P. savastanoi</i> . <i>Molecular Plant Pathology</i> , 2002, 3, 205-216.	5.1	38
134	Comprehensive transcript profiling of Pto- and Prf-mediated host defense responses to infection by <i>Pseudomonas syringae</i> pv. tomato. <i>Plant Journal</i> , 2002, 32, 299-315.	6.2	130
135	Title is missing!. <i>Journal of Biomolecular NMR</i> , 2002, 23, 247-248.	1.6	3
136	Title is missing!. <i>Genome Biology</i> , 2001, 2, reviews1003.1.	12.7	22
137	Ancient origin of pathogen recognition specificity conferred by the tomato disease resistance gene Pto. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 2059-2064.	7.8	65
138	Innate immunity in plants. <i>Current Opinion in Immunology</i> , 2001, 13, 55-62.	5.3	134
139	The major site of the Pti1 kinase phosphorylated by the Pto kinase is located in the activation domain and is required for Pto-Pti1 physical interaction. <i>FEBS Journal</i> , 2000, 267, 171-178.	0.2	37
140	Signal recognition and transduction mediated by the tomato Pto kinase: a paradigm of innate immunity in plants. <i>Microbes and Infection</i> , 2000, 2, 1591-1597.	2.4	32
141	Thr38 and Ser198 are Pto autophosphorylation sites required for the AvrPto-mediated hypersensitive response. <i>EMBO Journal</i> , 2000, 19, 2257-2269.	7.5	97
142	AvrPto-dependent Pto-interacting proteins and AvrPto-interacting proteins in tomato. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 8836-8840.	7.8	124
143	The <i>Pseudomonas</i> AvrPto Protein Is Differentially Recognized by Tomato and Tobacco and Is Localized to the Plant Plasma Membrane. <i>Plant Cell</i> , 2000, 12, 2323.	7.6	1
144	Pti4 Is Induced by Ethylene and Salicylic Acid, and Its Product Is Phosphorylated by the Pto Kinase. <i>Plant Cell</i> , 2000, 12, 771.	7.6	8

#	ARTICLE	IF	PR CITATIONS
145	The <i>Pseudomonas</i> AvrPto Protein Is Differentially Recognized by Tomato and Tobacco and Is Localized to the Plant Plasma Membrane. <i>Plant Cell</i> , 2000, 12, 2323-2337.	7.6	166
146	Pti4 Is Induced by Ethylene and Salicylic Acid, and Its Product Is Phosphorylated by the Pto Kinase. <i>Plant Cell</i> , 2000, 12, 771-785.	7.6	284
147	Overexpression of Pto Activates Defense Responses and Confers Broad Resistance. <i>Plant Cell</i> , 1999, 11, 15-29.	7.6	256
148	<i>Pseudomonas syringae</i> pv tomato induces the expression of tomato EREBP-like genes Pti4 and Pti5 independent of ethylene, salicylate and jasmonate. <i>Plant Journal</i> , 1999, 20, 475-483.	6.2	99
149	Title is missing!. , 1999, 40, 455-465.		64
150	Functional analysis of plant disease resistance genes and their downstream effectors. <i>Current Opinion in Plant Biology</i> , 1999, 2, 273-279.	7.2	158
151	Overexpression of Pto Activates Defense Responses and Confers Broad Resistance. <i>Plant Cell</i> , 1999, 11, 15.	7.6	19
152	Gene discovery for crop improvement. <i>Current Opinion in Biotechnology</i> , 1998, 9, 220-226.	7.0	19
153	Recognition Specificity for the Bacterial Avirulence Protein AvrPto Is Determined by Thr-204 in the Activation Loop of the Tomato Pto Kinase. <i>Molecular Cell</i> , 1998, 2, 241-245.	13.7	125
154	Molecular mechanisms involved in bacterial speck disease resistance of tomato. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 1998, 353, 1455-1461.	3.7	17
155	A Nitrilase-Like Protein Interacts with GCC Box DNA-Binding Proteins Involved in Ethylene and Defense Responses. <i>Plant Physiology</i> , 1998, 118, 867-874.	5.5	52
156	Biochemical Properties of Two Protein Kinases Involved in Disease Resistance Signaling in Tomato. <i>Journal of Biological Chemistry</i> , 1998, 273, 15860-15865.	2.3	39
157	The Myristylation Motif of Pto Is Not Required for Disease Resistance. <i>Molecular Plant-Microbe Interactions</i> , 1998, 11, 572-576.	3.3	40
158	Alleles of Pto and Fen Occur in Bacterial Speck-Susceptible and Fenthion-Insensitive Tomato Cultivars and Encode Active Protein Kinases. <i>Plant Cell</i> , 1997, 9, 61.	7.6	16
159	The Pto kinase conferring resistance to tomato bacterial speck disease interacts with proteins that bind a cis-element of pathogenesis-related genes. <i>EMBO Journal</i> , 1997, 16, 3207-3218.	7.5	447
160	The Pto kinase mediates a signaling pathway leading to the oxidative burst in tomato. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1996, 93, 13393-13397.	7.8	86
161	Expression of the Tomato Pto Gene in Tobacco Enhances Resistance to <i>Pseudomonas syringae</i> pv tabaci Expressing avrPto. <i>Plant Cell</i> , 1995, 7, 1529.	7.6	41
162	Chromosome landing: a paradigm for map-based gene cloning in plants with large genomes. <i>Trends in Genetics</i> , 1995, 11, 63-68.	10.0	407

#	ARTICLE	IF	PR CITATIONS
163	Molecular genetic analysis of the ripening-inhibitor and non-ripening loci of tomato: A first step in genetic map-based cloning of fruit ripening genes. <i>Molecular Genetics and Genomics</i> , 1995, 248, 195-206.	0.5	55
164	The tomato gene <i>Pti1</i> encodes a serine/threonine kinase that is phosphorylated by <i>Pto</i> and is involved in the hypersensitive response. <i>Cell</i> , 1995, 83, 925-935.	34.4	403
165	Map-based cloning in crop plants: tomato as a model system II. Isolation and characterization of a set of overlapping yeast artificial chromosomes encompassing the <i>jointless</i> locus. <i>Molecular Genetics and Genomics</i> , 1994, 244, 613-621.	0.5	32
166	Chromosome 2-specific DNA clones from flow-sorted chromosomes of tomato. <i>Molecular Genetics and Genomics</i> , 1994, 242, 551-558.	0.5	73
167	Analysis of the molecular basis of <i>Pseudomonas syringae</i> pv. <i>tomato</i> resistance in tomato. <i>Euphytica</i> , 1994, 79, 187-193.	1.5	4
168	A Member of the Tomato <i>Pto</i> Gene Family Confers Sensitivity to Fenthion Resulting in Rapid Cell Death. <i>Plant Cell</i> , 1994, 6, 1543.	7.6	41
169	High-Resolution Linkage Analysis and Physical Characterization of the <i>Pto</i> Bacterial Resistance Locus in Tomato. <i>Molecular Plant-Microbe Interactions</i> , 1993, 6, 26.	3.3	75
170	Construction of a yeast artificial chromosome library of tomato and identification of cloned segments linked to two disease resistance loci. <i>Molecular Genetics and Genomics</i> , 1992, 233, 25-32.	0.5	124
171	<i>Bradyrhizobium japonicum</i> <i>ntrBC/glnA</i> and <i>nifA/glnA</i> Mutants: Further Evidence that Separate Regulatory Pathways Govern <i>nif</i> Expression in Free-Living and Symbiotic Cells. <i>Molecular Plant-Microbe Interactions</i> , 1991, 4, 254.	3.3	5
172	Landraces of <i>Phaseolus vulgaris</i> (Fabaceae) in Northern Malawi. I. Regional variation. <i>Economic Botany</i> , 1987, 41, 190-203.	1.2	56
173	Landraces of <i>Phaseolus vulgaris</i> (Fabaceae) in Northern Malawi. II. Generation and maintenance of variability. <i>Economic Botany</i> , 1987, 41, 204-215.	1.2	40
174	Comparative genomics and phylogenetic discordance of cultivated tomato and close wild relatives. <i>PeerJ</i> , 0, 3, e793.	0.0	27