

# Suomeng Dong

## List of Publications by Year in descending order

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66  
papers

5,266  
citations

101543

36  
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95266

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75  
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75  
docs citations

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times ranked

4061  
citing authors

#	ARTICLE	IF	CITATIONS
1	The two-speed genomes of filamentous pathogens: waltz with plants. <i>Current Opinion in Genetics and Development</i> , 2015, 35, 57-65.	3.3	503
2	Signatures of Adaptation to Obligate Biotrophy in the <i>Hyaloperonospora arabidopsidis</i> Genome. <i>Science</i> , 2010, 330, 1549-1551.	12.6	492
3	Transcriptional Programming and Functional Interactions within the <i>Phytophthora sojae</i> RXLR Effector Repertoire. <i>Plant Cell</i> , 2011, 23, 2064-2086.	6.6	455
4	A <i>Phytophthora sojae</i> Glycoside Hydrolase 12 Protein Is a Major Virulence Factor during Soybean Infection and Is Recognized as a PAMP. <i>Plant Cell</i> , 2015, 27, 2057-2072.	6.6	335
5	A paralogous decoy protects <i>Phytophthora sojae</i> apoplastic effector PsXEG1 from a host inhibitor. <i>Science</i> , 2017, 355, 710-714.	12.6	236
6	Effector Specialization in a Lineage of the Irish Potato Famine Pathogen. <i>Science</i> , 2014, 343, 552-555.	12.6	179
7	<i>Phytophthora sojae</i> Avirulence Effector Avr3b is a Secreted NADH and ADP-ribose Pyrophosphorylase that Modulates Plant Immunity. <i>PLoS Pathogens</i> , 2011, 7, e1002353.	4.7	169
8	The Basic Leucine Zipper Transcription Factor Moatf1 Mediates Oxidative Stress Responses and Is Necessary for Full Virulence of the Rice Blast Fungus <i>Magnaporthe oryzae</i> . <i>Molecular Plant-Microbe Interactions</i> , 2010, 23, 1053-1068.	2.6	156
9	Copy Number Variation and Transcriptional Polymorphisms of <i>Phytophthora sojae</i> RXLR Effector Genes Avr1a and Avr3a. <i>PLoS ONE</i> , 2009, 4, e5066.	2.5	151
10	The RxLR effector Avh241 from <i>Phytophthora sojae</i> requires plasma membrane localization to induce plant cell death. <i>New Phytologist</i> , 2012, 196, 247-260.	7.3	151
11	Leucine-rich repeat receptor-like gene screen reveals that <i>Nicotiana glauca</i> RXEG1 regulates glycoside hydrolase 12 MAMP detection. <i>Nature Communications</i> , 2018, 9, 594.	12.8	142
12	A <i>Phytophthora</i> Effector Manipulates Host Histone Acetylation and Reprograms Defense Gene Expression to Promote Infection. <i>Current Biology</i> , 2017, 27, 981-991.	3.9	120
13	Digital Gene Expression Profiling of the <i>Phytophthora sojae</i> Transcriptome. <i>Molecular Plant-Microbe Interactions</i> , 2011, 24, 1530-1539.	2.6	119
14	A <i>Phytophthora sojae</i> effector suppresses endoplasmic reticulum stress-mediated immunity by stabilizing plant Binding immunoglobulin Proteins. <i>Nature Communications</i> , 2016, 7, 11685.	12.8	119
15	The <i>Phytophthora sojae</i> Avirulence Locus Avr3c Encodes a Multi-Copy RXLR Effector with Sequence Polymorphisms among Pathogen Strains. <i>PLoS ONE</i> , 2009, 4, e5556.	2.5	116
16	The NLP Toxin Family in <i>Phytophthora sojae</i> Includes Rapidly Evolving Groups That Lack Necrosis-Inducing Activity. <i>Molecular Plant-Microbe Interactions</i> , 2012, 25, 896-909.	2.6	101
17	Two Host Cytoplasmic Effectors Are Required for Pathogenesis of <i>Phytophthora sojae</i> by Suppression of Host Defenses. <i>Plant Physiology</i> , 2011, 155, 490-501.	4.8	100
18	An oomycete plant pathogen reprograms host pre-mRNA splicing to subvert immunity. <i>Nature Communications</i> , 2017, 8, 2051.	12.8	84

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19	Development of a loop-mediated isothermal amplification assay for detection of <i>Phytophthora sojae</i> . <i>FEMS Microbiology Letters</i> , 2012, 334, 27-34.	1.8	83
20	The role of vacuolar processing enzyme (VPE) from <i>Nicotiana benthamiana</i> in the elicitor-triggered hypersensitive response and stomatal closure. <i>Journal of Experimental Botany</i> , 2010, 61, 3799-3812.	4.8	76
21	Sequence Variants of the <i>Phytophthora sojae</i> RXLR Effector Avr3a/5 Are Differentially Recognized by Rps3a and Rps5 in Soybean. <i>PLoS ONE</i> , 2011, 6, e20172.	2.5	76
22	Distinct regions of the <i>Phytophthora</i> essential effector Avh238 determine its function in cell death activation and plant immunity suppression. <i>New Phytologist</i> , 2017, 214, 361-375.	7.3	67
23	The Activation of <i>Phytophthora</i> Effector Avr3b by Plant Cyclophilin is Required for the Nudix Hydrolase Activity of Avr3b. <i>PLoS Pathogens</i> , 2015, 11, e1005139.	4.7	66
24	The <i>Phytophthora sojae</i> RXLR effector Avh238 destabilizes soybean Type2 GmACSs to suppress ethylene biosynthesis and promote infection. <i>New Phytologist</i> , 2019, 222, 425-437.	7.3	63
25	<i>Phytophthora</i> methylomes are modulated by 6mA methyltransferases and associated with adaptive genome regions. <i>Genome Biology</i> , 2018, 19, 181.	8.8	61
26	<i>Phytophthora sojae</i> Effector PsAvh240 Inhibits Host Aspartic Protease Secretion to Promote Infection. <i>Molecular Plant</i> , 2019, 12, 552-564.	8.3	60
27	A <i>Phytophthora</i> effector recruits a host cytoplasmic transacetylase into nuclear speckles to enhance plant susceptibility. <i>ELife</i> , 2018, 7, .	6.0	60
28	Extracellular proteolytic cascade in tomato activates immune protease Rcr3. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 17409-17417.	7.1	55
29	N-glycosylation shields <i>Phytophthora sojae</i> apoplastic effector PsXEG1 from a specific host aspartic protease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 27685-27693.	7.1	51
30	Distribution, Pathotypes, and Metalaxyl Sensitivity of <i>Phytophthora sojae</i> from Heilongjiang and Fujian Provinces in China. <i>Plant Disease</i> , 2010, 94, 881-884.	1.4	50
31	<i>Phytophthora</i> Effectors Modulate Genome-wide Alternative Splicing of Host mRNAs to Reprogram Plant Immunity. <i>Molecular Plant</i> , 2020, 13, 1470-1484.	8.3	49
32	Effector gene silencing mediated by histone methylation underpins host adaptation in an oomycete plant pathogen. <i>Nucleic Acids Research</i> , 2020, 48, 1790-1799.	14.5	47
33	PsSAK1, a Stress-Activated MAP Kinase of <i>Phytophthora sojae</i> , Is Required for Zoospore Viability and Infection of Soybean. <i>Molecular Plant-Microbe Interactions</i> , 2010, 23, 1022-1031.	2.6	45
34	The <i>Nicotiana benthamiana</i> Mitogen-Activated Protein Kinase Cascade and WRKY Transcription Factor Participate in Nep1 <sub>Mo</sub> -Triggered Plant Responses. <i>Molecular Plant-Microbe Interactions</i> , 2012, 25, 1639-1653.	2.6	43
35	The <i>Phytophthora sojae</i> Avr1d Gene Encodes an RxLR-dEER Effector with Presence and Absence Polymorphisms Among Pathogen Strains. <i>Molecular Plant-Microbe Interactions</i> , 2013, 26, 958-968.	2.6	43
36	Nudix Effectors: A Common Weapon in the Arsenal of Plant Pathogens. <i>PLoS Pathogens</i> , 2016, 12, e1005704.	4.7	43

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37	Pathogen manipulation of chloroplast function triggers a light-dependent immune recognition. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 9613-9620.	7.1	39
38	Natural allelic variations provide insights into host adaptation of <i>Phytophthora</i> avirulence effector PsAvr3c. New Phytologist, 2019, 221, 1010-1022.	7.3	37
39	A Myb Transcription Factor of <i>Phytophthora sojae</i> , Regulated by MAP Kinase PsSAK1, Is Required for Zoospore Development. PLoS ONE, 2012, 7, e40246.	2.5	33
40	The LCB <sub>2</sub> subunit of the sphingolip biosynthesis enzyme serine palmitoyltransferase can function as an attenuator of the hypersensitive response and Bax-induced cell death. New Phytologist, 2009, 181, 127-146.	7.3	32
41	Long transposon-rich centromeres in an oomycete reveal divergence of centromere features in Stramenopila-Alveolata-Rhizaria lineages. PLoS Genetics, 2020, 16, e1008646.	3.5	29
42	The MADS-box Transcription Factor PsMAD1 Is Involved in Zoosporogenesis and Pathogenesis of <i>Phytophthora sojae</i> . Frontiers in Microbiology, 2018, 9, 2259.	3.5	26
43	Chitin synthase is involved in vegetative growth, asexual reproduction and pathogenesis of <i>Phytophthora capsici</i> and <i>Phytophthora sojae</i> . Environmental Microbiology, 2019, 21, 4537-4547.	3.8	25
44	Cleavage of a pathogen apoplastic protein by plant subtilases activates host immunity. New Phytologist, 2021, 229, 3424-3439.	7.3	24
45	Analysis of polymorphism and transcription of the effector gene <i>Avr1b</i> in <i>Phytophthora sojae</i> isolates from China virulent to <i>Rps1b</i> . Molecular Plant Pathology, 2012, 13, 114-122.	4.2	23
46	How to win a tug-of-war: the adaptive evolution of <i>Phytophthora</i> effectors. Current Opinion in Plant Biology, 2021, 62, 102027.	7.1	22
47	PsAAT3, an oomycete-specific aspartate aminotransferase, is required for full pathogenicity of the oomycete pathogen <i>Phytophthora sojae</i> . Fungal Biology, 2016, 120, 620-630.	2.5	20
48	Rapid detection of potato late blight using a loop-mediated isothermal amplification assay. Journal of Integrative Agriculture, 2020, 19, 1274-1282.	3.5	18
49	The <i>Phytophthora</i> effector Avh241 interacts with host NDR1-like proteins to manipulate plant immunity. Journal of Integrative Plant Biology, 2021, 63, 1382-1396.	8.5	16
50	Editing of an effector gene promoter sequence impacts plant- <i>Phytophthora</i> interaction. Journal of Integrative Plant Biology, 2020, 62, 378-392.	8.5	15
51	<i>Phytophthora sojae</i> apoplastic effector AEP1 mediates sugar uptake by mutarotation of extracellular aldose and is recognized as a MAMP. Plant Physiology, 2021, 187, 321-335.	4.8	15
52	Specific interaction of an RNA-binding protein with the 3'-UTR of its target mRNA is critical to oomycete sexual reproduction. PLoS Pathogens, 2021, 17, e1010001.	4.7	13
53	Rapid detection of <i>Colletotrichum gloeosporioides</i> using a loop-mediated isothermal amplification assay. Australasian Plant Pathology, 2017, 46, 493-498.	1.0	12
54	Silent control: microbial plant pathogens evade host immunity without coding sequence changes. FEMS Microbiology Reviews, 2021, 45, .	8.6	12

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55	Mammalian pro-apoptotic bax gene enhances tobacco resistance to pathogens. <i>Plant Cell Reports</i> , 2008, 27, 1559-1569.	5.6	11
56	Genome Analysis of Two Newly Emerged Potato Late Blight Isolates Sheds Light on Pathogen Adaptation and Provides Tools for Disease Management. <i>Phytopathology</i> , 2021, 111, 96-107.	2.2	9
57	Decoding co-/post-transcriptional complexities of plant transcriptomes and epitranscriptome using next-generation sequencing technologies. <i>Biochemical Society Transactions</i> , 2020, 48, 2399-2414.	3.4	9
58	Polymorphism in natural alleles of the avirulence gene Avr1c is associated with the host adaptation of <i>Phytophthora sojae</i> . <i>Phytopathology Research</i> , 2019, 1, .	2.4	8
59	The bZIP transcription factor PsBZP32 is involved in cyst germination, oxidative stress response, and pathogenicity of <i>Phytophthora sojae</i> . <i>Phytopathology Research</i> , 2021, 3, .	2.4	8
60	Functional Analysis of PsAvr3c Effector Family From <i>Phytophthora</i> Provides Probes to Dissect SKRP Mediated Plant Susceptibility. <i>Frontiers in Plant Science</i> , 2018, 9, 1105.	3.6	6
61	An Improved Method for the Identification of Soybean Resistance to <i>Phytophthora sojae</i> Applied to Germplasm Resources from the Huanghuaihai and Dongbei Regions of China. <i>Plant Disease</i> , 2020, 104, 408-413.	1.4	5
62	Structures of plant resistosome reveal how NLR immune receptors are activated. <i>ABIOTECH</i> , 2020, 1, 147-150.	3.9	5
63	ATAC-seq reveals the landscape of open chromatin and cis-regulatory elements in the <i>Phytophthora sojae</i> genome. <i>Molecular Plant-Microbe Interactions</i> , 2022, , .	2.6	5
64	A new roadmap for the breeding of disease-resistant and high-yield crops. <i>Stress Biology</i> , 2021, 1, 1.	3.1	4
65	Green fluorescent protein (GFP) as a vital marker for studying the interaction of <i>Phytophthora sojae</i> and soybean. <i>Science Bulletin</i> , 2009, 54, 2822-2829.	9.0	2
66	Plant biotic interactions: From fundamental research toward sustainable agriculture. <i>Journal of Integrative Plant Biology</i> , 2021, 63, 275-276.	8.5	1