

# Vivianne G A A Vleeshouwers

## List of Publications by Year in descending order

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68  
papers

6,110  
citations

94433

37  
h-index

106344

65  
g-index

80  
all docs

80  
docs citations

80  
times ranked

3629  
citing authors

#	ARTICLE	IF	CITATIONS
1	Cleavage of a pathogen apoplastic protein by plant subtilases activates host immunity. <i>New Phytologist</i> , 2021, 229, 3424-3439.	7.3	24
2	Quantifying the Contribution to Virulence of <i>Phytophthora infestans</i> Effectors in Potato. <i>Methods in Molecular Biology</i> , 2021, 2354, 303-313.	0.9	0
3	A complex resistance locus in <i>Solanum americanum</i> recognizes a conserved <i>Phytophthora</i> effector. <i>Nature Plants</i> , 2021, 7, 198-208.	9.3	62
4	Evolutionarily distinct resistance proteins detect a pathogen effector through its association with different host targets. <i>New Phytologist</i> , 2021, 232, 1368-1381.	7.3	6
5	Qualitative and Quantitative Resistance against Early Blight Introgressed in Potato. <i>Biology</i> , 2021, 10, 892.	2.8	13
6	Identification of <i>Solanum</i> Immune Receptors by Bulk Segregant RNA-Seq and High-Throughput Recombinant Screening. <i>Methods in Molecular Biology</i> , 2021, 2354, 315-330.	0.9	3
7	Identification of <i>Avram1</i> from <i>Phytophthora infestans</i> using long read and cDNA pathogen-enrichment sequencing (PenSeq). <i>Molecular Plant Pathology</i> , 2020, 21, 1502-1512.	4.2	22
8	Divergent Evolution of PcF/SCR74 Effectors in Oomycetes Is Associated with Distinct Recognition Patterns in Solanaceous Plants. <i>MBio</i> , 2020, 11, .	4.1	11
9	Pathogen manipulation of chloroplast function triggers a light-dependent immune recognition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 9613-9620.	7.1	39
10	RLP/K enrichment sequencing; a novel method to identify receptor-like protein (RLP) and receptor-like kinase (RLK) genes. <i>New Phytologist</i> , 2020, 227, 1264-1276.	7.3	32
11	A rapid method to screen wild <i>Solanum</i> for resistance to early blight. <i>European Journal of Plant Pathology</i> , 2019, 154, 109-114.	1.7	12
12	<i>Phytophthora infestans</i> RXLR effectors act in concert at diverse subcellular locations to enhance host colonization. <i>Journal of Experimental Botany</i> , 2019, 70, 343-356.	4.8	66
13	The ELR-SOBIR1 Complex Functions as a Two-Component Receptor-Like Kinase to Mount Defense Against <i>Phytophthora infestans</i> . <i>Molecular Plant-Microbe Interactions</i> , 2018, 31, 795-802.	2.6	46
14	Gapless Genome Assembly of the Potato and Tomato Early Blight Pathogen <i>Alternaria solani</i> . <i>Molecular Plant-Microbe Interactions</i> , 2018, 31, 692-694.	2.6	48
15	Gene expression polymorphism underpins evasion of host immunity in an asexual lineage of the Irish potato famine pathogen. <i>BMC Evolutionary Biology</i> , 2018, 18, 93.	3.2	41
16	Effectoromics-Based Identification of Cell Surface Receptors in Potato. <i>Methods in Molecular Biology</i> , 2017, 1578, 337-353.	0.9	26
17	New Strategies Towards Durable Late Blight Resistance in Potato. <i>Compendium of Plant Genomes</i> , 2017, , 161-169.	0.5	6
18	Discovering Novel <i>Alternaria solani</i> Succinate Dehydrogenase Inhibitors by in Silico Modeling and Virtual Screening Strategies to Combat Early Blight. <i>Frontiers in Chemistry</i> , 2017, 5, 100.	3.6	16

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19	<a href="#">Solanum venturii</a> , a suitable model system for virus-induced gene silencing studies in potato reveals StMKK6 as an important player in plant immunity. <i>Plant Methods</i> , 2016, 12, 29.	4.3	10
20	Plant immunity switched from bacteria to virus. <i>Nature Biotechnology</i> , 2016, 34, 391-392.	17.5	7
21	Nine things to know about elicitors. <i>New Phytologist</i> , 2016, 212, 888-895.	7.3	84
22	Effector-driven marker development and cloning of resistance genes against <i>Phytophthora infestans</i> in potato breeding clone SW93-1015. <i>Theoretical and Applied Genetics</i> , 2016, 129, 105-115.	3.6	43
23	Elicitor recognition confers enhanced resistance to <i>Phytophthora infestans</i> in potato. <i>Nature Plants</i> , 2015, 1, 15034.	9.3	229
24	An updated conventional- and a novel GM potato late blight R gene differential set for virulence monitoring of <i>Phytophthora infestans</i> . <i>Euphytica</i> , 2015, 202, 219-234.	1.2	41
25	Effectors as Tools in Disease Resistance Breeding Against Biotrophic, Hemibiotrophic, and Necrotrophic Plant Pathogens. <i>Molecular Plant-Microbe Interactions</i> , 2015, 2015, 17-27.	2.6	4
26	Effectors as Tools in Disease Resistance Breeding Against Biotrophic, Hemibiotrophic, and Necrotrophic Plant Pathogens. <i>Molecular Plant-Microbe Interactions</i> , 2015, 2015, 40-50.	2.6	3
27	Effectors as Tools in Disease Resistance Breeding Against Biotrophic, Hemibiotrophic, and Necrotrophic Plant Pathogens. <i>Molecular Plant-Microbe Interactions</i> , 2014, 27, 196-206.	2.6	363
28	The Doâ€™s and Donâ€™ts of Effectoromics. <i>Methods in Molecular Biology</i> , 2014, 1127, 257-268.	0.9	17
29	Increased Difficulties to Control Late Blight in Tunisia Are Caused by a Genetically Diverse <i>Phytophthora infestans</i> Population Next to the Clonal Lineage NA-01. <i>Plant Disease</i> , 2014, 98, 898-908.	1.4	17
30	Agroinfiltration and PVX Agroinfection in Potato and <i>Nicotiana benthamiana</i> . <i>Journal of Visualized Experiments</i> , 2014, , e50971.	0.3	46
31	Functional analysis of potato genes involved in quantitative resistance to <i>Phytophthora infestans</i> . <i>Molecular Biology Reports</i> , 2013, 40, 957-967.	2.3	25
32	Characterisation of <i>Phytophthora infestans</i> Isolates Collected from Potato and Tomato Crops in Tunisia During 2006â€“2008. <i>Potato Research</i> , 2013, 56, 11-29.	2.7	13
33	Host Protein BSL1 Associates with <i>Phytophthora infestans</i> RXLR Effector AVR2 and the <i>Solanum demissum</i> Immune Receptor R2 to Mediate Disease Resistance. <i>Plant Cell</i> , 2012, 24, 3420-3434.	6.6	130
34	Genome Analyses of an Aggressive and Invasive Lineage of the Irish Potato Famine Pathogen. <i>PLoS Pathogens</i> , 2012, 8, e1002940.	4.7	321
35	Qualitative and Quantitative Late Blight Resistance in the Potato Cultivar Sarpo Mira Is Determined by the Perception of Five Distinct RXLR Effectors. <i>Molecular Plant-Microbe Interactions</i> , 2012, 25, 910-919.	2.6	162
36	High-Resolution Mapping of Two Broad-Spectrum Late Blight Resistance Genes from Two Wild Species of the <i>Solanum circaefolium</i> Group. <i>Potato Research</i> , 2012, 55, 109-123.	2.7	10

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37	Mycorrhizal Symbiosis: Ancient Signalling Mechanisms Co-opted. <i>Current Biology</i> , 2012, 22, R997-R999.	3.9	8
38	Understanding and Exploiting Late Blight Resistance in the Age of Effectors. <i>Annual Review of Phytopathology</i> , 2011, 49, 507-531.	7.8	369
39	Presence/absence, differential expression and sequence polymorphisms between <i>PiAVR2</i> and <i>PiAVR2-like</i> in <i>Phytophthora infestans</i> determine virulence on <i>R2</i> plants. <i>New Phytologist</i> , 2011, 191, 763-776.	7.3	142
40	High Resolution Mapping of a Novel Late Blight Resistance Gene <i>Rpi-avl1</i> , from the Wild Bolivian Species <i>Solanum avilesii</i> . <i>American Journal of Potato Research</i> , 2011, 88, 511-519.	0.9	13
41	SolRgene: an online database to explore disease resistance genes in tuber-bearing <i>Solanum</i> species. <i>BMC Plant Biology</i> , 2011, 11, 116.	3.6	38
42	Diversity, Distribution, and Evolution of <i>Solanum bulbocastanum</i> Late Blight Resistance Genes. <i>Molecular Plant-Microbe Interactions</i> , 2010, 23, 1206-1216.	2.6	69
43	A novel approach to locate <i>Phytophthora infestans</i> resistance genes on the potato genetic map. <i>Theoretical and Applied Genetics</i> , 2010, 120, 785-796.	3.6	49
44	In Planta Expression Screens of <i>Phytophthora infestans</i> RXLR Effectors Reveal Diverse Phenotypes, Including Activation of the <i>Solanum bulbocastanum</i> Disease Resistance Protein <i>Rpi-blb2</i> . <i>Plant Cell</i> , 2009, 21, 2928-2947.	6.6	376
45	<i>Phytophthora infestans</i> Isolates Lacking Class I <i>ipiO</i> Variants Are Virulent on <i>Rpi-blb1</i> Potato. <i>Molecular Plant-Microbe Interactions</i> , 2009, 22, 1535-1545.	2.6	118
46	Allele mining in <i>Solanum</i> : conserved homologues of <i>Rpi-blb1</i> are identified in <i>Solanum stoloniferum</i> . <i>Theoretical and Applied Genetics</i> , 2008, 116, 933-943.	3.6	117
47	AFLP analysis reveals a lack of phylogenetic structure within <i>Solanum</i> section <i>Petota</i> . <i>BMC Evolutionary Biology</i> , 2008, 8, 145.	3.2	52
48	Effector Genomics Accelerates Discovery and Functional Profiling of Potato Disease Resistance and <i>Phytophthora infestans</i> Avirulence Genes. <i>PLoS ONE</i> , 2008, 3, e2875.	2.5	361
49	Agroinfection-based high-throughput screening reveals specific recognition of INF elicitors in <i>Solanum</i> . <i>Molecular Plant Pathology</i> , 2006, 7, 499-510.	4.2	50
50	Differences in Intensity and Specificity of Hypersensitive Response Induction in <i>Nicotiana</i> spp. by INF1, INF2A, and INF2B of <i>Phytophthora infestans</i> . <i>Molecular Plant-Microbe Interactions</i> , 2005, 18, 183-193.	2.6	56
51	The Late Blight Resistance Locus <i>Rpi-blb3</i> from <i>Solanum bulbocastanum</i> Belongs to a Major Late Blight R Gene Cluster on Chromosome 4 of Potato. <i>Molecular Plant-Microbe Interactions</i> , 2005, 18, 722-729.	2.6	133
52	Comparative genomics enabled the isolation of the R3a late blight resistance gene in potato. <i>Plant Journal</i> , 2005, 42, 251-261.	5.7	355
53	Dissection of foliage and tuber late blight resistance in mapping populations of potato. <i>Euphytica</i> , 2005, 143, 75-83.	1.2	41
54	High-resolution Mapping and Analysis of the Resistance Locus <i>Rpi-abpt</i> Against <i>Phytophthora infestans</i> in Potato. <i>Molecular Breeding</i> , 2005, 16, 33-43.	2.1	66

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55	An Accurate In Vitro Assay for High-Throughput Disease Testing of Phytophthora infestans in Potato. Plant Disease, 2005, 89, 1263-1267.	1.4	30
56	A G1± subunit controls zoospore motility and virulence in the potato late blight pathogen Phytophthora infestans. Molecular Microbiology, 2004, 51, 925-936.	2.5	130
57	The R3 Resistance to Phytophthora infestans in Potato is Conferred by Two Closely Linked R Genes with Distinct Specificities. Molecular Plant-Microbe Interactions, 2004, 17, 428-435.	2.6	121
58	Active defence responses associated with non-host resistance of Arabidopsis thaliana to the oomycete pathogen Phytophthora infestans. Molecular Plant Pathology, 2003, 4, 487-500.	4.2	90
59	Ancient Diversification of the Pto Kinase Family Preceded Speciation in Solanum. Molecular Plant-Microbe Interactions, 2001, 14, 996-1005.	2.6	23
60	The hypersensitive response is associated with host and nonhost resistance to Phytophthora infestans. Planta, 2000, 210, 853-864.	3.2	217
61	Does basal PR gene expression in Solanum species contribute to non-specific resistance to Phytophthora infestans ?. Physiological and Molecular Plant Pathology, 2000, 57, 35-42.	2.5	73
62	Title is missing!. European Journal of Plant Pathology, 1999, 105, 241-250.	1.7	146
63	Resistance to oomycetes: a general role for the hypersensitive response?. Trends in Plant Science, 1999, 4, 196-200.	8.8	183
64	Resistance of Nicotiana benthamiana to Phytophthora infestans Is Mediated by the Recognition of the Elicitor Protein INF1. Plant Cell, 1998, 10, 1413-1425.	6.6	371
65	A Gene Encoding a Protein Elicitor of Phytophthora infestans Is Down-Regulated During Infection of Potato. Molecular Plant-Microbe Interactions, 1997, 10, 13-20.	2.6	233
66	Production of the AVR9 elicitor from the fungal pathogen Cladosporium fulvum in transgenic tobacco and tomato plants. Plant Molecular Biology, 1995, 29, 909-920.	3.9	39
67	Effectors as Tools in Disease Resistance Breeding Against Biotrophic, Hemibiotrophic, and Necrotrophic Plant Pathogens. Molecular Plant-Microbe Interactions, 0, , MPMI-10-13-0313.	2.6	1
68	Effectors as Tools in Disease Resistance Breeding Against Biotrophic, Hemibiotrophic, and Necrotrophic Plant Pathogens. Molecular Plant-Microbe Interactions, 0, , MPMI-10-13-0313.	2.6	0