

Nicolas Rispail

List of Publications by Year in descending order

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Version: 2024-02-01

39
papers

2,236
citations

279487

23
h-index

329751

37
g-index

40
all docs

40
docs citations

40
times ranked

2782
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative genomics of MAP kinase and calcium-calciuneurin signalling components in plant and human pathogenic fungi. <i>Fungal Genetics and Biology</i> , 2009, 46, 287-298.	0.9	302
2	Biotechnology approaches to overcome biotic and abiotic stress constraints in legumes. <i>Euphytica</i> , 2006, 147, 1-24.	0.6	214
3	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.	3.1	207
4	Achievements and Challenges in Legume Breeding for Pest and Disease Resistance. <i>Critical Reviews in Plant Sciences</i> , 2015, 34, 195-236.	2.7	153
5	Putative role of gamma -aminobutyric acid (GABA) as a long-distance signal in up-regulation of nitrate uptake in <i>Brassica napus</i> L.. <i>Plant, Cell and Environment</i> , 2004, 27, 1035-1046.	2.8	151
6	Genetics of Symbiosis in <i>Lotus japonicus</i> : Recombinant Inbred Lines, Comparative Genetic Maps, and Map Position of 35 Symbiotic Loci. <i>Molecular Plant-Microbe Interactions</i> , 2006, 19, 80-91.	1.4	94
7	Plant resistance to parasitic plants: molecular approaches to an old foe. <i>New Phytologist</i> , 2007, 173, 703-712.	3.5	89
8	<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.	1.4	87
9	Breeding approaches for crenate broomrape (<i>Orobanche crenata</i> Forsk.) management in pea (<i>Pisum sativum</i> L.). <i>Pest Management Science</i> , 2009, 65, 553-559.	1.7	71
10	Quantum Dot and Superparamagnetic Nanoparticle Interaction with Pathogenic Fungi: Internalization and Toxicity Profile. <i>ACS Applied Materials & Interfaces</i> , 2014, 6, 9100-9110.	4.0	71
11	Model legumes contribute to faba bean breeding. <i>Field Crops Research</i> , 2010, 115, 253-269.	2.3	64
12	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.	2.0	62
13	Physical and Chemical Barriers in Root Tissues Contribute to Quantitative Resistance to <i>Fusarium oxysporum</i> f. sp. <i>pisi</i> in Pea. <i>Frontiers in Plant Science</i> , 2018, 9, 199.	1.7	58
14	Genetic Diversity and Population Structure Among Oat Cultivars and Landraces. <i>Plant Molecular Biology Reporter</i> , 2013, 31, 1305-1314.	1.0	55
15	A detailed evaluation method to identify sources of quantitative resistance to <i>Fusarium oxysporum</i> f. sp. <i>pisi</i> race 2 within a <i>Pisum</i> spp. germplasm collection. <i>Plant Pathology</i> , 2012, 61, 532-542.	1.2	52
16	The homeodomain transcription factor Ste12. <i>Communicative and Integrative Biology</i> , 2010, 3, 327-332.	0.6	45
17	Genome-wide association study for crown rust (<i>Puccinia coronata</i> f. sp. <i>avenae</i>) and powdery mildew (<i>Blumeria graminis</i> f. sp. <i>avenae</i>) resistance in an oat (<i>Avena sativa</i>) collection of commercial varieties and landraces. <i>Frontiers in Plant Science</i> , 2015, 6, 103.	1.7	43
18	Genome-wide identification and comparison of legume MLO gene family. <i>Scientific Reports</i> , 2016, 6, 32673.	1.6	41

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19	Identification of the Main Toxins Isolated from <i>Fusarium oxysporum</i> f. sp. <i>pisi</i> Race 2 and Their Relation with Isolates' Pathogenicity. <i>Journal of Agricultural and Food Chemistry</i> , 2014, 62, 2574-2580.	2.4	40
20	Phenolic Compounds: extraction and analysis. , 2005, , 349-354.		32
21	Higher rust resistance and similar yield of oat landraces versus cultivars under high temperature and drought. <i>Agronomy for Sustainable Development</i> , 2017, 37, 1.	2.2	31
22	Legume breeding for rust resistance: lessons to learn from the model <i>Medicago truncatula</i> . <i>Euphytica</i> , 2011, 180, 89-98.	0.6	28
23	Identification of Sources of Quantitative Resistance to <i>Fusarium oxysporum</i> f. sp. <i>medicaginis</i> in <i>Medicago truncatula</i> . <i>Plant Disease</i> , 2014, 98, 667-673.	0.7	27
24	Salicylic acid regulates polyamine biosynthesis during drought responses in oat. <i>Plant Signaling and Behavior</i> , 2019, 14, e1651183.	1.2	24
25	Pisatin involvement in the variation of inhibition of <i>Fusarium oxysporum</i> f. sp. <i>pisi</i> spore germination by root exudates of <i>Pisum</i> spp. germplasm. <i>Plant Pathology</i> , 2018, 67, 1046-1054.	1.2	22
26	Parasitic plant infection is partially controlled through symbiotic pathways. <i>Weed Research</i> , 2010, 50, 76-82.	0.8	21
27	Deciphering Root Architectural Traits Involved to Cope With Water Deficit in Oat. <i>Frontiers in Plant Science</i> , 2019, 10, 1558.	1.7	19
28	Drought resistance in oat involves ABA-mediated modulation of transpiration and root hydraulic conductivity. <i>Environmental and Experimental Botany</i> , 2021, 182, 104333.	2.0	18
29	A nitrogen response pathway regulates virulence in plant pathogenic fungi. <i>Plant Signaling and Behavior</i> , 2010, 5, 1623-1625.	1.2	17
30	Resistance reaction of <i>Medicago truncatula</i> genotypes to <i>Fusarium oxysporum</i> : effect of plant age, substrate and inoculation method. <i>Crop and Pasture Science</i> , 2015, 66, 506.	0.7	17
31	Rapid and Efficient Estimation of Pea Resistance to the Soil-Borne Pathogen <i>Fusarium oxysporum</i> by Infrared Imaging. <i>Sensors</i> , 2015, 15, 3988-4000.	2.1	12
32	Multi-Environmental Trials Reveal Genetic Plasticity of Oat Agronomic Traits Associated With Climate Variable Changes. <i>Frontiers in Plant Science</i> , 2018, 9, 1358.	1.7	12
33	Secondary metabolite profiling of the model legume <i>Lotus japonicus</i> during its symbiotic interaction with <i>Mesorhizobium loti</i> . <i>Symbiosis</i> , 2010, 50, 119-128.	1.2	10
34	Population genomics of Mediterranean oat (<i>A. sativa</i>) reveals high genetic diversity and three loci for heading date. <i>Theoretical and Applied Genetics</i> , 2021, 134, 2063-2077.	1.8	10
35	Genomic regions associated with herbicide tolerance in a worldwide faba bean (<i>Vicia faba</i> L.) collection. <i>Scientific Reports</i> , 2022, 12, 158.	1.6	10
36	Editorial: Advances in Legume Research. <i>Frontiers in Plant Science</i> , 2018, 9, 501.	1.7	8

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37	Genetic Dissection of Heat Stress Tolerance in Faba Bean (<i>Vicia faba</i> L.) Using GWAS. <i>Plants</i> , 2022, 11, 1108.	1.6	7
38	Secondary metabolite profiling. , 2005, , 341-348.		5
39	Deciphering Main Climate and Edaphic Components Driving Oat Adaptation to Mediterranean Environments. <i>Frontiers in Plant Science</i> , 2021, 12, 780562.	1.7	3