Simon G Krattinger

List of Publications by Year in descending order

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58 papers

5,243 citations

147801 31 h-index 53 g-index

64 all docs

64
docs citations

64 times ranked 4254 citing authors

#	Article	IF	CITATIONS
1	The long road to engineering durable disease resistance in wheat. Current Opinion in Biotechnology, 2022, 73, 270-275.	6.6	14
2	Long-read genome sequencing of bread wheat facilitates disease resistance gene cloning. Nature Genetics, 2022, 54, 227-231.	21.4	63
3	Oat genome — sequence of a superfood. Nature Plants, 2022, 8, 602-603.	9.3	5
4	A membrane-bound ankyrin repeat protein confers race-specific leaf rust disease resistance in wheat. Nature Communications, 2021, 12, 956.	12.8	63
5	Population genomics and haplotype analysis in spelt and bread wheat identifies a gene regulating glume color. Communications Biology, 2021, 4, 375.	4.4	11
6	Alleles of a wallâ€associated kinase gene account for three of the major northern corn leaf blight resistance loci in maize. Plant Journal, 2021, 106, 526-535.	5.7	23
7	Genome-wide association study for septoria tritici blotch resistance reveals the occurrence and distribution of Stb6 in a historic Swiss landrace collection. Euphytica, 2021, 217, 1.	1.2	3
8	Comparative Transcriptome Analysis of Wheat Lines in the Field Reveals Multiple Essential Biochemical Pathways Suppressed by Obligate Pathogens. Frontiers in Plant Science, 2021, 12, 720462.	3.6	14
9	Expression of the wheat disease resistance gene Lr34 in transgenic barley leads to accumulation of abscisic acid at the leaf tip. Plant Physiology and Biochemistry, 2021, 166, 950-957.	5.8	10
10	Multiple wheat genomes reveal global variation in modern breeding. Nature, 2020, 588, 277-283.	27.8	513
11	Fonio millet genome unlocks African orphan crop diversity for agriculture in a changing climate. Nature Communications, 2020, 11, 4488.	12.8	63
12	The NLR-Annotator Tool Enables Annotation of the Intracellular Immune Receptor Repertoire. Plant Physiology, 2020, 183, 468-482.	4.8	147
13	Impedimetric Plant Biosensor Based on Minimally Invasive and Flexible Microneedle Electrodes. , 2020, ,		2
14	A new player contributing to durable Fusarium resistance. Nature Genetics, 2019, 51, 1070-1071.	21.4	36
15	Orthologous receptor kinases quantitatively affect the host status of barley to leaf rust fungi. Nature Plants, 2019, 5, 1129-1135.	9.3	37
16	Updated inventory, evolutionary and expression analyses of G (PDR) type ABC transporter genes of rice. Plant Physiology and Biochemistry, 2019, 142, 429-439.	5.8	17
17	Abscisic acid is a substrate of the <scp>ABC</scp> transporter encoded by the durable wheat disease resistance gene <i>Lr34</i> . New Phytologist, 2019, 223, 853-866.	7.3	102
18	Fungal resistance mediated by maize wallâ€associated kinase Zm <scp>WAK</scp> â€ <scp>RLK</scp> 1 correlates with reduced benzoxazinoid content. New Phytologist, 2019, 221, 976-987.	7.3	71

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19	Point Inoculation Method for Measuring Adult Plant Response of Wheat to Stripe Rust Infection. Plant Disease, 2019, 103, 1228-1233.	1.4	2
20	Genebank genomics highlights the diversity of a global barley collection. Nature Genetics, 2019, 51, 319-326.	21.4	322
21	Rapid gene cloning in cereals. Theoretical and Applied Genetics, 2019, 132, 699-711.	3.6	26
22	Precision Phenotyping Reveals Novel Loci for Quantitative Resistance to Septoria Tritici Blotch. Plant Phenomics, 2019, 2019, 3285904.	5.9	37
23	A new player in race-specific resistance. Nature Plants, 2018, 4, 197-198.	9.3	11
24	Pathogenâ€inducible <i>Ta</i> â€ <i>Lr34res</i> expression in heterologous barley confers disease resistance without negative pleiotropic effects. Plant Biotechnology Journal, 2018, 16, 245-253.	8.3	39
25	Transcriptional profiling reveals no response of fungal pathogens to the durable, quantitative <i>Lr34</i> disease resistance gene of wheat. Plant Pathology, 2018, 67, 792-798.	2.4	5
26	Unlocking the diversity of genebanks: whole-genome marker analysis of Swiss bread wheat and spelt. Theoretical and Applied Genetics, 2018, 131, 407-416.	3.6	47
27	Advances in Wheat and Pathogen Genomics: Implications for Disease Control. Annual Review of Phytopathology, 2018, 56, 67-87.	7.8	66
28	Chromosome-scale comparative sequence analysis unravels molecular mechanisms of genome dynamics between two wheat cultivars. Genome Biology, 2018, 19, 104.	8.8	54
29	Resistance: Double gain with one gene. Nature Plants, 2017, 3, 17019.	9.3	6
30	Rapid cloning of genes in hexaploid wheat using cultivar-specific long-range chromosome assembly. Nature Biotechnology, 2017, 35, 793-796.	17.5	218
31	Genomic compartments in barley. Nature, 2017, 544, 424-425.	27.8	18
32	The wheat Lr34 multipathogen resistance gene confers resistance to anthracnose and rust in sorghum. Plant Biotechnology Journal, 2017, 15, 1387-1396.	8.3	52
33	The <i>Lr34</i> adult plant rust resistance gene provides seedling resistance in durum wheat without senescence. Plant Biotechnology Journal, 2017, 15, 894-905.	8.3	56
34	Rapid Identification of Rust Resistance Genes Through Cultivar-Specific De Novo Chromosome Assemblies. Methods in Molecular Biology, 2017, 1659, 245-255.	0.9	2
35	Combined GC- and UHPLC-HR-MS Based Metabolomics to Analyze Durable Anti-fungal Resistance Processes in Cereals. Chimia, 2017, 71, 156-159.	0.6	6
36	The durable wheat disease resistance gene <i>Lr34</i> confers common rust and northern corn leaf blight resistance in maize. Plant Biotechnology Journal, 2017, 15, 489-496.	8.3	75

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37	Characterization of Lr75: a partial, broad-spectrum leaf rust resistance gene in wheat. Theoretical and Applied Genetics, 2017, 130, 1-12.	3.6	130
38	Large-scale Maize Seedling Infection with Exserohilum turcicum in the Greenhouse. Bio-protocol, 2017, 7, e2567.	0.4	3
39	The wheat durable, multipathogen resistance gene <i>Lr34</i> confers partial blast resistance in rice. Plant Biotechnology Journal, 2016, 14, 1261-1268.	8.3	98
40	Trapping the intruder â€" immune receptor domain fusions provide new molecular leads for improving disease resistance in plants. Genome Biology, 2016, 17, 23.	8.8	7
41	Molecular genetics and evolution of disease resistance in cereals. New Phytologist, 2016, 212, 320-332.	7.3	99
42	Rapid gene isolation in barley and wheat by mutant chromosome sequencing. Genome Biology, 2016, 17, 221.	8.8	265
43	TheLr34adult plant rust resistance gene provides seedling resistance in durum wheat without senescence. Plant Biotechnology Journal, 2016, , .	8.3	0
44	The wheat resistance gene <i>Lr34</i> results in the constitutive induction of multiple defense pathways in transgenic barley. Plant Journal, 2015, 84, 202-215.	5.7	45
45	The maize disease resistance gene <i>Htn1</i> against northern corn leaf blight encodes a wall-associated receptor-like kinase. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 8780-8785.	7.1	302
46	Genomic Approaches Towards Durable Fungal Disease Resistance in Wheat., 2015,, 369-375.		2
47	Recent emergence of the wheat Lr34 multi-pathogen resistance: insights from haplotype analysis in wheat, rice, sorghum and Aegilops tauschii. Theoretical and Applied Genetics, 2013, 126, 663-672.	3.6	79
48	The wheat <i><scp>L</scp>r34</i> gene provides resistance against multiple fungal pathogens in barley. Plant Biotechnology Journal, 2013, 11, 847-854.	8.3	116
49	Comment on â€ ⁻ In Turkish wheat cultivars the resistance allele of LR34 is ineffective against leaf rust'. Journal of Plant Diseases and Protection, 2013, 120, 3-3.	2.9	1
50	Functional variability of the <i>Lr34</i> durable resistance gene in transgenic wheat. Plant Biotechnology Journal, 2012, 10, 477-487.	8.3	99
51	<i>Lr34</i> multiâ€pathogen resistance ABC transporter: molecular analysis of homoeologous and orthologous genes in hexaploid wheat and other grass species. Plant Journal, 2011, 65, 392-403.	5.7	79
52	Relationships among the A Genomes of Triticum L. Species as Evidenced by SSR Markers, in Iran. International Journal of Molecular Sciences, 2010, 11, 4309-4325.	4.1	11
53	Gene-specific markers for the wheat gene Lr34/Yr18/Pm38 which confers resistance to multiple fungal pathogens. Theoretical and Applied Genetics, 2009, 119, 889-898.	3.6	342
54	Analysis of Intraspecies Diversity in Wheat and Barley Genomes Identifies Breakpoints of Ancient Haplotypes and Provides Insight into the Structure of Diploid and Hexaploid Triticeae Gene Pools Â. Plant Physiology, 2009, 149, 258-270.	4.8	38

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55	A Putative ABC Transporter Confers Durable Resistance to Multiple Fungal Pathogens in Wheat. Science, 2009, 323, 1360-1363.	12.6	1,140
56	Map-Based Cloning of Genes in Triticeae (Wheat and Barley)., 2009,, 337-357.		33
57	Development of simple sequence repeat markers specific for the Lr34 resistance region of wheat using sequence information from rice and Aegilops tauschii. Theoretical and Applied Genetics, 2006, 113, 1049-1062.	3.6	82
58	Unconventional R proteins in the botanical tribe Triticeae. Essays in Biochemistry, 0, , .	4.7	3