

Ravi Prakash Singh

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4634395/publications.pdf>

Version: 2024-02-01

82
papers

9,868
citations

94433

37
h-index

62596

80
g-index

83
all docs

83
docs citations

83
times ranked

6890
citing authors

#	ARTICLE	IF	CITATIONS
1	A Putative ABC Transporter Confers Durable Resistance to Multiple Fungal Pathogens in Wheat. <i>Science</i> , 2009, 323, 1360-1363.	12.6	1,140
2	Genomic Selection in Plant Breeding: Methods, Models, and Perspectives. <i>Trends in Plant Science</i> , 2017, 22, 961-975.	8.8	1,004
3	Prediction of Genetic Values of Quantitative Traits in Plant Breeding Using Pedigree and Molecular Markers. <i>Genetics</i> , 2010, 186, 713-724.	2.9	664
4	The Emergence of Ug99 Races of the Stem Rust Fungus is a Threat to World Wheat Production. <i>Annual Review of Phytopathology</i> , 2011, 49, 465-481.	7.8	612
5	A recently evolved hexose transporter variant confers resistance to multiple pathogens in wheat. <i>Nature Genetics</i> , 2015, 47, 1494-1498.	21.4	575
6	Multiple wheat genomes reveal global variation in modern breeding. <i>Nature</i> , 2020, 588, 277-283.	27.8	513
7	Emergence and Spread of New Races of Wheat Stem Rust Fungus: Continued Threat to Food Security and Prospects of Genetic Control. <i>Phytopathology</i> , 2015, 105, 872-884.	2.2	393
8	Canopy Temperature and Vegetation Indices from High-Throughput Phenotyping Improve Accuracy of Pedigree and Genomic Selection for Grain Yield in Wheat. <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 2799-2808.	1.8	336
9	Disease Impact on Wheat Yield Potential and Prospects of Genetic Control. <i>Annual Review of Phytopathology</i> , 2016, 54, 303-322.	7.8	322
10	Increased Prediction Accuracy in Wheat Breeding Trials Using a Marker \times Environment Interaction Genomic Selection Model. <i>G3: Genes, Genomes, Genetics</i> , 2015, 5, 569-582.	1.8	266
11	Wheat improvement in India: present status, emerging challenges and future prospects. <i>Euphytica</i> , 2007, 157, 431-446.	1.2	262
12	Lr68: a new gene conferring slow rusting resistance to leaf rust in wheat. <i>Theoretical and Applied Genetics</i> , 2012, 124, 1475-1486.	3.6	248
13	New slow-rusting leaf rust and stripe rust resistance genes Lr67 and Yr46 in wheat are pleiotropic or closely linked. <i>Theoretical and Applied Genetics</i> , 2011, 122, 239-249.	3.6	224
14	Improving grain yield, stress resilience and quality of bread wheat using large-scale genomics. <i>Nature Genetics</i> , 2019, 51, 1530-1539.	21.4	216
15	Phenotypic and Genotypic Characterization of Race TKTF of <i>Puccinia graminis</i> f. sp. <i>tritici</i> that Caused a Wheat Stem Rust Epidemic in Southern Ethiopia in 2013-14. <i>Phytopathology</i> , 2015, 105, 917-928.	2.2	202
16	Application of unmanned aerial systems for high throughput phenotyping of large wheat breeding nurseries. <i>Plant Methods</i> , 2016, 12, 35.	4.3	200
17	A high density GBS map of bread wheat and its application for dissecting complex disease resistance traits. <i>BMC Genomics</i> , 2015, 16, 216.	2.8	188
18	Combining High-Throughput Phenotyping and Genomic Information to Increase Prediction and Selection Accuracy in Wheat Breeding. <i>Plant Genome</i> , 2018, 11, 170043.	2.8	175

#	ARTICLE	IF	CITATIONS
19	Wheat genetic resources enhancement by the International Maize and Wheat Improvement Center (CIMMYT). <i>Genetic Resources and Crop Evolution</i> , 2008, 55, 1095-1140.	1.6	155
20	Identification of genomic regions for grain yield and yield stability and their epistatic interactions. <i>Scientific Reports</i> , 2017, 7, 41578.	3.3	127
21	Genome-wide association mapping for resistance to leaf rust, stripe rust and tan spot in wheat reveals potential candidate genes. <i>Theoretical and Applied Genetics</i> , 2018, 131, 1405-1422.	3.6	101
22	Genomic and pedigree-based prediction for leaf, stem, and stripe rust resistance in wheat. <i>Theoretical and Applied Genetics</i> , 2017, 130, 1415-1430.	3.6	99
23	Strategic crossing of biomass and harvest index “source and sink” achieves genetic gains in wheat. <i>Euphytica</i> , 2017, 213, 1.	1.2	97
24	Hyperspectral Reflectance-Derived Relationship Matrices for Genomic Prediction of Grain Yield in Wheat. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 1231-1247.	1.8	96
25	Genetic Yield Gains In CIMMYT's International Elite Spring Wheat Yield Trials By Modeling The Genotype × Environment Interaction. <i>Crop Science</i> , 2017, 57, 789-801.	1.8	89
26	New Deep Learning Genomic-Based Prediction Model for Multiple Traits with Binary, Ordinal, and Continuous Phenotypes. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 1545-1556.	1.8	81
27	Integrating genomic-enabled prediction and high-throughput phenotyping in breeding for climate-resilient bread wheat. <i>Theoretical and Applied Genetics</i> , 2019, 132, 177-194.	3.6	78
28	Single-Step Genomic and Pedigree Genotype × Environment Interaction Models for Predicting Wheat Lines in International Environments. <i>Plant Genome</i> , 2017, 10, plantgenome2016.09.0089.	2.8	66
29	Prospects and Challenges of Applied Genomic Selection “A New Paradigm in Breeding for Grain Yield in Bread Wheat. <i>Plant Genome</i> , 2018, 11, 180017.	2.8	65
30	Fifty years of semi-dwarf spring wheat breeding at CIMMYT: Grain yield progress in optimum, drought and heat stress environments. <i>Field Crops Research</i> , 2020, 250, 107757.	5.1	64
31	Genetic Contribution of Synthetic Hexaploid Wheat to CIMMYT's Spring Bread Wheat Breeding Germplasm. <i>Scientific Reports</i> , 2019, 9, 12355.	3.3	62
32	Hybrid Wheat Prediction Using Genomic, Pedigree, and Environmental Covariables Interaction Models. <i>Plant Genome</i> , 2019, 12, 180051.	2.8	58
33	Genetic loci associated with high grain zinc concentration and pleiotropic effect on kernel weight in wheat (<i>Triticum aestivum</i> L.). <i>Molecular Breeding</i> , 2014, 34, 1893-1902.	2.1	56
34	A large-scale genomic association analysis identifies the candidate causal genes conferring stripe rust resistance under multiple field environments. <i>Plant Biotechnology Journal</i> , 2021, 19, 177-191.	8.3	54
35	Identification and characterization of pleiotropic and co-located resistance loci to leaf rust and stripe rust in bread wheat cultivar Sujata. <i>Theoretical and Applied Genetics</i> , 2015, 128, 549-561.	3.6	49
36	Comparison of Models and Whole-Genome Profiling Approaches for Genomic-Enabled Prediction of Septoria Tritici Blotch, Stagonospora Nodorum Blotch, and Tan Spot Resistance in Wheat. <i>Plant Genome</i> , 2017, 10, plantgenome2016.08.0082.	2.8	48

#	ARTICLE	IF	CITATIONS
37	The <i>Aegilops ventricosa</i> 2NvS segment in bread wheat: cytology, genomics and breeding. <i>Theoretical and Applied Genetics</i> , 2021, 134, 529-542.	3.6	48
38	Genetic structures of the CIMMYT international yield trial targeted to irrigated environments. <i>Molecular Breeding</i> , 2012, 29, 529-541.	2.1	41
39	Resistance to Spot Blotch in Two Mapping Populations of Common Wheat Is Controlled by Multiple QTL of Minor Effects. <i>International Journal of Molecular Sciences</i> , 2018, 19, 4054.	4.1	40
40	Genomic Bayesian functional regression models with interactions for predicting wheat grain yield using hyper-spectral image data. <i>Plant Methods</i> , 2017, 13, 62.	4.3	38
41	Strategies for Selecting Crosses Using Genomic Prediction in Two Wheat Breeding Programs. <i>Plant Genome</i> , 2017, 10, plantgenome2016.12.0128.	2.8	37
42	Increased ranking change in wheat breeding under climate change. <i>Nature Plants</i> , 2021, 7, 1207-1212.	9.3	37
43	Improving global integration of crop research. <i>Science</i> , 2017, 357, 359-360.	12.6	34
44	Grain yield genetic gains and changes in physiological related traits for CIMMYT's High Rainfall Wheat Screening Nursery tested across international environments. <i>Field Crops Research</i> , 2020, 249, 107742.	5.1	34
45	Regularized selection indices for breeding value prediction using hyper-spectral image data. <i>Scientific Reports</i> , 2020, 10, 8195.	3.3	32
46	Aerial high-throughput phenotyping enables indirect selection for grain yield at the early generation, seed-limited stages in breeding programs. <i>Crop Science</i> , 2020, 60, 3096-3114.	1.8	31
47	Resistance to stem rust Ug99 in six bread wheat cultivars maps to chromosome 6DS. <i>Theoretical and Applied Genetics</i> , 2014, 127, 231-239.	3.6	30
48	Haplotype-Based, Genome-Wide Association Study Reveals Stable Genomic Regions for Grain Yield in CIMMYT Spring Bread Wheat. <i>Frontiers in Genetics</i> , 2020, 11, 589490.	2.3	29
49	Cloning of the broadly effective wheat leaf rust resistance gene <i>Lr42</i> transferred from <i>Aegilops tauschii</i> . <i>Nature Communications</i> , 2022, 13, .	12.8	29
50	Identification and Validation of a Common Stem Rust Resistance Locus in Two Bi-parental Populations. <i>Frontiers in Plant Science</i> , 2018, 9, 1788.	3.6	28
51	Economic benefits of blast-resistant biofortified wheat in Bangladesh: The case of BARI Gom 33. <i>Crop Protection</i> , 2019, 123, 45-58.	2.1	28
52	Harnessing translational research in wheat for climate resilience. <i>Journal of Experimental Botany</i> , 2021, 72, 5134-5157.	4.8	28
53	Genome-wide association mapping for wheat blast resistance in CIMMYT's international screening nurseries evaluated in Bolivia and Bangladesh. <i>Scientific Reports</i> , 2020, 10, 15972.	3.3	27
54	Genomic Selection for Grain Yield in the CIMMYT Wheat Breeding Program—Status and Perspectives. <i>Frontiers in Plant Science</i> , 2020, 11, 564183.	3.6	27

#	ARTICLE	IF	CITATIONS
55	Target Population of Environments for Wheat Breeding in India: Definition, Prediction and Genetic Gains. <i>Frontiers in Plant Science</i> , 2021, 12, 638520.	3.6	26
56	Sources of the highly expressed wheat bread making (wbm) gene in CIMMYT spring wheat germplasm and its effect on processing and bread-making quality. <i>Euphytica</i> , 2016, 209, 689-692.	1.2	24
57	A Bayesian Genomic Multi-output Regressor Stacking Model for Predicting Multi-trait Multi-environment Plant Breeding Data. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 3381-3393.	1.8	22
58	Characterization of QTLs for Seedling Resistance to Tan Spot and Septoria Nodorum Blotch in the PBW343/Kenya Nyangumi Wheat Recombinant Inbred Lines Population. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5432.	4.1	17
59	Screening and Mapping for Head Blast Resistance in a Panel of CIMMYT and South Asian Bread Wheat Germplasm. <i>Frontiers in Genetics</i> , 2021, 12, 679162.	2.3	16
60	Genome-wide association analysis for arabinoxylan content in common wheat (<i>T. Aestivum</i> L.) flour. <i>Journal of Cereal Science</i> , 2021, 98, 103166.	3.7	14
61	Genomic Selection for Wheat Blast in a Diversity Panel, Breeding Panel and Full-Sibs Panel. <i>Frontiers in Plant Science</i> , 2021, 12, 745379.	3.6	13
62	Elucidating the genetics of grain yield and stress-resilience in bread wheat using a large-scale genome-wide association mapping study with 55,568 lines. <i>Scientific Reports</i> , 2021, 11, 5254.	3.3	11
63	Multi-trait selection of bread wheat ideotypes for adaptation to early sown condition. <i>Crop Science</i> , 2022, 62, 67-82.	1.8	10
64	Refined mapping of stripe rust resistance gene YrP10090 within a desirable haplotype for wheat improvement on chromosome 6A. <i>Theoretical and Applied Genetics</i> , 2021, 134, 2005-2021.	3.6	9
65	Genome-Wide Association Mapping Indicates Quantitative Genetic Control of Spot Blotch Resistance in Bread Wheat and the Favorable Effects of Some Spot Blotch Loci on Grain Yield. <i>Frontiers in Plant Science</i> , 2022, 13, 835095.	3.6	9
66	Genomic selection for spot blotch in bread wheat breeding panels, full-sibs and half-sibs and index-based selection for spot blotch, heading and plant height. <i>Theoretical and Applied Genetics</i> , 2022, , 1.	3.6	9
67	A singular value decomposition Bayesian multiple-trait and multiple-environment genomic model. <i>Heredity</i> , 2019, 122, 381-401.	2.6	8
68	Genome-wide mapping and allelic fingerprinting provide insights into the genetics of resistance to wheat stripe rust in India, Kenya and Mexico. <i>Scientific Reports</i> , 2020, 10, 10908.	3.3	8
69	Improving Wheat Yield Prediction Using Secondary Traits and High-Density Phenotyping Under Heat-Stressed Environments. <i>Frontiers in Plant Science</i> , 2021, 12, 633651.	3.6	8
70	Bayesian multitrait kernel methods improve multi-environment genome-based prediction. <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	1.8	8
71	Juvenile Heat Tolerance in Wheat for Attaining Higher Grain Yield by Shifting to Early Sowing in October in South Asia. <i>Genes</i> , 2021, 12, 1808.	2.4	8
72	Retrospective Quantitative Genetic Analysis and Genomic Prediction of Global Wheat Yields. <i>Frontiers in Plant Science</i> , 2020, 11, 580136.	3.6	7

#	ARTICLE	IF	CITATIONS
73	Breeding increases grain yield, zinc, and iron, supporting enhanced wheat biofortification. <i>Crop Science</i> , 2022, 62, 1912-1925.	1.8	7
74	Combined linkage and association mapping reveals two major QTL for stripe rust adult plant resistance in Shaanmai 155 and their haplotype variation in common wheat germplasm. <i>Crop Journal</i> , 2022, 10, 783-792.	5.2	5
75	Quantitative trait loci mapping reveals the complexity of adult plant resistance to leaf rust in spring wheat "Copio"™. <i>Crop Science</i> , 2022, 62, 1037-1050.	1.8	5
76	Identification and mapping of two adult plant leaf rust resistance genes in durum wheat. <i>Molecular Breeding</i> , 2019, 39, 1.	2.1	4
77	Pre-emptive Breeding Against Karnal Bunt Infection in Common Wheat: Combining Genomic and Agronomic Information to Identify Suitable Parents. <i>Frontiers in Plant Science</i> , 2021, 12, 675859.	3.6	4
78	Plant breeding increases spring wheat yield potential in Afghanistan. <i>Crop Science</i> , 2022, 62, 167-177.	1.8	3
79	Genome-Wide Association Mapping Identifies Key Genomic Regions for Grain Zinc and Iron Biofortification in Bread Wheat. <i>Frontiers in Plant Science</i> , 0, 13, .	3.6	3
80	Molecular Characterization of Genomic Regions for Adult Plant Resistance to Stem Rust in a Spring Wheat Mapping Population. <i>Plant Disease</i> , 2022, 106, 439-450.	1.4	1
81	Achieving Genetic Gains in Practice. , 2022, , 97-123.		1
82	Dataset of historic and modern bread and durum wheat cultivar performance under conventional and reduced tillage with full and reduced irrigation. <i>Data in Brief</i> , 2022, 43, 108439.	1.0	0