

P Stephen Baenziger

List of Publications by Year in descending order

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300
papers

11,320
citations

30070

54
h-index

43889

91
g-index

306
all docs

306
docs citations

306
times ranked

7901
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide comparative diversity uncovers multiple targets of selection for improvement in hexaploid wheat landraces and cultivars. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 8057-8062.	7.1	1,065
2	Exploiting genetic diversity from landraces in wheat breeding for adaptation to climate change. <i>Journal of Experimental Botany</i> , 2015, 66, 3477-3486.	4.8	356
3	Drought Stress Tolerance in Wheat and Barley: Advances in Physiology, Breeding and Genetics Research. <i>International Journal of Molecular Sciences</i> , 2019, 20, 3137.	4.1	353
4	Management of Fusarium head blight of wheat and barley. <i>Crop Protection</i> , 2015, 73, 100-107.	2.1	236
5	Population- and genome-specific patterns of linkage disequilibrium and SNP variation in spring and winter wheat (<i>Triticum aestivum</i> L.). <i>BMC Genomics</i> , 2010, 11, 727.	2.8	234
6	GWAS: Fast-forwarding gene identification and characterization in temperate Cereals: lessons from Barley – A review. <i>Journal of Advanced Research</i> , 2020, 22, 119-135.	9.5	227
7	Genotype and Environment Effects on Quality Characteristics of Hard Red Winter Wheat. <i>Crop Science</i> , 1992, 32, 98-103.	1.8	221
8	A multi-sensor system for high throughput field phenotyping in soybean and wheat breeding. <i>Computers and Electronics in Agriculture</i> , 2016, 128, 181-192.	7.7	191
9	Genetic Diversity and Population Structure of F3:6 Nebraska Winter Wheat Genotypes Using Genotyping-By-Sequencing. <i>Frontiers in Genetics</i> , 2018, 9, 76.	2.3	183
10	Demarcating the gene-rich regions of the wheat genome. <i>Nucleic Acids Research</i> , 2004, 32, 3546-3565.	14.5	181
11	Transferability of SSR markers among wheat, rye, and triticale. <i>Theoretical and Applied Genetics</i> , 2004, 108, 1147-1150.	3.6	161
12	Comparison of phenotypic and molecular marker-based classifications of hard red winter wheat cultivars. <i>Euphytica</i> , 2005, 145, 133-146.	1.2	151
13	Identification of QTLs and Environmental Interactions Associated with Agronomic Traits on Chromosome 3A of Wheat. <i>Crop Science</i> , 2003, 43, 1493-1505.	1.8	139
14	Assessment of genetic diversity and relationship among a collection of US sweet sorghum germplasm by SSR markers. <i>Molecular Breeding</i> , 2008, 21, 497-509.	2.1	137
15	Introgression of Novel Traits from a Wild Wheat Relative Improves Drought Adaptation in Wheat. <i>Plant Physiology</i> , 2013, 161, 1806-1819.	4.8	124
16	Combining abilities and heritability of callus formation and plantlet regeneration in wheat (<i>Triticum</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	3.6	118
17	Improving predictions of developmental stages in winter wheat: a modified Wang and Engel model. <i>Agricultural and Forest Meteorology</i> , 2003, 115, 139-150.	4.8	113
18	Agronomic Effect of Wheat-Rye Translocation Carrying Rye Chromatin (1R) From Different Sources. <i>Crop Science</i> , 2004, 44, 1254-1258.	1.8	112

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19	Variation for Grain Mineral Concentration in a Diversity Panel of Current and Historical Great Plains Hard Winter Wheat Germplasm. <i>Crop Science</i> , 2015, 55, 1035-1052.	1.8	112
20	Anther culture of wheat (<i>Triticum aestivum</i> L.) F1's and their reciprocal crosses. <i>Theoretical and Applied Genetics</i> , 1982, 62, 155-159.	3.6	110
21	Removing Spatial Variation from Wheat Yield Trials: A Comparison of Methods. <i>Crop Science</i> , 1994, 34, 62-66.	1.8	110
22	Molecular Mapping of Loci for Agronomic Traits on Chromosome 3A of Bread Wheat. <i>Crop Science</i> , 1999, 39, 1728-1732.	1.8	105
23	Genotyping-by-Sequencing Derived High-Density Linkage Map and its Application to QTL Mapping of Flag Leaf Traits in Bread Wheat. <i>Scientific Reports</i> , 2017, 7, 16394.	3.3	103
24	Designing crop technology for a future climate: An example using response surface methodology and the CERES-Wheat model. <i>Agricultural Systems</i> , 2006, 87, 63-79.	6.1	102
25	Earlier winter wheat heading dates and warmer spring in the U.S. Great Plains. <i>Agricultural and Forest Meteorology</i> , 2005, 135, 284-290.	4.8	97
26	Environmental modification of hard red winter wheat flour protein composition. <i>Journal of Cereal Science</i> , 1995, 22, 45-51.	3.7	96
27	The 1BL/1RS Translocation: Agronomic Performance of F ₃ Derived Lines from a Winter Wheat Cross. <i>Crop Science</i> , 1995, 35, 1051-1055.	1.8	94
28	Variation for nitrogen use efficiency traits in current and historical great plains hard winter wheat. <i>Euphytica</i> , 2017, 213, 1.	1.2	92
29	Haploid Plant Development from Anthers and In Vitro Embryo Culture of Wheat ¹ . <i>Crop Science</i> , 1979, 19, 697-702.	1.8	91
30	Genome-Wide Association Study Reveals Novel Genomic Regions for Grain Yield and Yield-Related Traits in Drought-Stressed Synthetic Hexaploid Wheat. <i>International Journal of Molecular Sciences</i> , 2018, 19, 3011.	4.1	90
31	Genetic improvement trends in agronomic performances and end-use quality characteristics among hard red winter wheat cultivars in Nebraska. <i>Euphytica</i> , 2005, 144, 187-198.	1.2	89
32	Genotypic and Environmental Modification of Wheat Flour Protein Composition in Relation to End-Use Quality. <i>Crop Science</i> , 1996, 36, 296-300.	1.8	86
33	Effect of Cultivar, Environment, and Their Interaction and Stability Analyses on Milling and Baking Quality of Soft Red Winter Wheat ¹ . <i>Crop Science</i> , 1985, 25, 5-8.	1.8	85
34	Cell Membrane Stability and Association Mapping for Drought and Heat Tolerance in a Worldwide Wheat Collection. <i>Sustainability</i> , 2017, 9, 1606.	3.2	85
35	Cultivar and cultivar x environment effects on the development of callus and polyhaploid plants from anther cultures of wheat. <i>Theoretical and Applied Genetics</i> , 1984, 67, 273-277.	3.6	80
36	Wheat Height Estimation Using LiDAR in Comparison to Ultrasonic Sensor and UAS. <i>Sensors</i> , 2018, 18, 3731.	3.8	80

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37	Development and Utilization of SSRs to Estimate the Degree of Genetic Relationships in a Collection of Pearl Millet Germplasm. <i>Crop Science</i> , 2003, 43, 2284-2290.	1.8	77
38	Unlocking the novel genetic diversity and population structure of synthetic Hexaploid wheat. <i>BMC Genomics</i> , 2018, 19, 591.	2.8	76
39	Phenotypic Plasticity of Winter Wheat Heading Date and Grain Yield across the US Great Plains. <i>Crop Science</i> , 2016, 56, 2223-2236.	1.8	75
40	Seeding Rate and Genotype Effect on Agronomic Performance and End Use Quality of Winter Wheat. <i>Crop Science</i> , 2002, 42, 827-832.	1.8	74
41	Improving Lives: 50 Years of Crop Breeding, Genetics, and Cytology (Câ€¹). <i>Crop Science</i> , 2006, 46, 2230-2244.	1.8	74
42	Genomic Selection in Preliminary Yield Trials in a Winter Wheat Breeding Program. <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 2735-2747.	1.8	74
43	Genome-Wide Association Study Reveals Novel Genomic Regions Associated with 10 Grain Minerals in Synthetic Hexaploid Wheat. <i>International Journal of Molecular Sciences</i> , 2018, 19, 3237.	4.1	72
44	Registration of â€˜Maceâ€™™ Hard Red Winter Wheat. <i>Journal of Plant Registrations</i> , 2009, 3, 51-56.	0.5	71
45	Characterization of Genetic Variability Among Natural Populations of Wheat Streak Mosaic Virus. <i>Phytopathology</i> , 1996, 86, 1222.	2.2	70
46	Genome-Wide Association Study for Identification and Validation of Novel SNP Markers for Sr6 Stem Rust Resistance Gene in Bread Wheat. <i>Frontiers in Plant Science</i> , 2018, 9, 380.	3.6	68
47	A comparison between genotyping-by-sequencing and array-based scoring of SNPs for genomic prediction accuracy in winter wheat. <i>Plant Science</i> , 2018, 270, 123-130.	3.6	67
48	Functional properties of waxy wheat flours: genotypic and environmental effects. <i>Journal of Cereal Science</i> , 2003, 38, 69-76.	3.7	61
49	Evaluating canopy spectral reflectance vegetation indices to estimate nitrogen use traits in hard winter wheat. <i>Field Crops Research</i> , 2018, 217, 82-92.	5.1	61
50	Economic returns from fungicide application to control foliar fungal diseases in winter wheat. <i>Crop Protection</i> , 2011, 30, 685-692.	2.1	60
51	Root tip cell cycle synchronization and metaphase-chromosome isolation suitable for flow sorting in common wheat (<i>Triticum aestivum</i> L.). <i>Genome</i> , 1997, 40, 633-638.	2.0	58
52	Incorporating a Chronology Response into the Prediction of Leaf Appearance Rate in Winter Wheat. <i>Annals of Botany</i> , 2003, 92, 181-190.	2.9	58
53	Addition of Colchicine to Wheat Anther Culture Media to Increase Doubled Haploid Plant Production. <i>Plant Breeding</i> , 1994, 112, 192-198.	1.9	57
54	Effects of Powdery Mildew on Yield and Quality of Isogenic Lines of â€˜Chancellorâ€™™ Wheat 1. <i>Crop Science</i> , 1979, 19, 349-352.	1.8	56

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55	Prediction of genetic values of quantitative traits with epistatic effects in plant breeding populations. <i>Heredity</i> , 2012, 109, 313-319.	2.6	55
56	Understanding the Effect of Rye Chromatin in Bread Wheat. <i>Crop Science</i> , 2003, 43, 1643-1651.	1.8	53
57	Haploidy in Cultivated Wheats: Induction and Utility in Basic and Applied Research. <i>Crop Science</i> , 2009, 49, 737-755.	1.8	53
58	Prospects for Selecting Wheat with Increased Zinc and Decreased Cadmium Concentration in Grain. <i>Crop Science</i> , 2015, 55, 1712-1728.	1.8	52
59	Using Environmental Covariates to Explain Genotype \times Environment and QTL \times Environment Interactions for Agronomic Traits on Chromosome 3A of Wheat. <i>Crop Science</i> , 2004, 44, 620-627.	1.8	50
60	Effect of Sugars in Wheat Anther Culture Media. <i>Plant Breeding</i> , 1994, 112, 53-62.	1.9	49
61	Isolated wheat microspore culture. <i>Plant Cell, Tissue and Organ Culture</i> , 1995, 42, 207-213.	2.3	49
62	An Automated Near-Infrared System for Selecting Individual Kernels Based on Specific Quality Characteristics. <i>Cereal Chemistry</i> , 2006, 83, 537-543.	2.2	49
63	FR-H3: a new QTL to assist in the development of fall-sown barley with superior low temperature tolerance. <i>Theoretical and Applied Genetics</i> , 2013, 126, 335-347.	3.6	49
64	GWAS revealed effect of genotype \times environment interactions for grain yield of Nebraska winter wheat. <i>BMC Genomics</i> , 2021, 22, 2.	2.8	49
65	Chromosomal Location of Wheat Quantitative Trait Loci Affecting Agronomic Performance of Seven Traits, Using Reciprocal Chromosome Substitutions. <i>Crop Science</i> , 1992, 32, 621-627.	1.8	46
66	Agronomic Performance and End-Use Quality of 1B vs. 1BL/1RS Genotypes Derived from Winter Wheat "Rawhide"™. <i>Crop Science</i> , 1995, 35, 1607-1612.	1.8	46
67	Effect of growth stage on the relationship between tan spot and spot blotch severity and yield in winter wheat. <i>Crop Protection</i> , 2009, 28, 696-702.	2.1	46
68	Agronomic Performance of Wheat Doubled-Haploid Lines Derived from Cultivars by Anther Culture. <i>Plant Breeding</i> , 1989, 103, 101-109.	1.9	45
69	Breeding for end-use quality: Reflections on the Nebraska experience. <i>Euphytica</i> , 2001, 119, 95-100.	1.2	45
70	Quantification of Yield Loss Caused by <i>Triticum mosaic virus</i> and <i>Wheat streak mosaic virus</i> in Winter Wheat Under Field Conditions. <i>Plant Disease</i> , 2014, 98, 127-133.	1.4	45
71	Molecular genetic analysis of spring wheat core collection using genetic diversity, population structure, and linkage disequilibrium. <i>BMC Genomics</i> , 2020, 21, 434.	2.8	44
72	The Physical Environment in Relation to High Frequency Callus and Plantlet Development in Anther Cultures of Wheat (<i>Triticum aestivum</i> L.) cv. Chris. <i>Journal of Plant Physiology</i> , 1985, 121, 103-109.	3.5	43

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73	A Generalized Vernalization Response Function for Winter Wheat. <i>Agronomy Journal</i> , 2003, 95, 155.	1.8	43
74	Fusarium Head Blight Resistance in U.S. Winter Wheat Cultivars and Elite Breeding Lines. <i>Crop Science</i> , 2013, 53, 2006-2013.	1.8	43
75	Genetic variation in drought tolerance at seedling stage and grain yield in low rainfall environments in wheat (<i>Triticum aestivum</i> L.). <i>Euphytica</i> , 2018, 214, 1.	1.2	43
76	Inheritance of Multiple Transgenes in Wheat. <i>Crop Science</i> , 2000, 40, 1133-1141.	1.8	42
77	Linkage mapping of powdery mildew and greenbug resistance genes on recombinant 1RS from 'Amigo' and 'Kavkaz' wheat-rye translocations of chromosome 1RS.1AL. <i>Genome</i> , 2004, 47, 292-298.	2.0	42
78	Transgenic expression of lactoferrin imparts enhanced resistance to head blight of wheat caused by <i>Fusarium graminearum</i> . <i>BMC Plant Biology</i> , 2012, 12, 33.	3.6	42
79	Evaluation and Association Mapping of Resistance to Tan Spot and <i>Stagonospora Nodorum</i> Blotch in Adapted Winter Wheat Germplasm. <i>Plant Disease</i> , 2015, 99, 1333-1341.	1.4	42
80	Influence of soil water status and atmospheric vapor pressure deficit on leaf gas exchange in field-grown winter wheat. <i>Environmental and Experimental Botany</i> , 2004, 51, 167-179.	4.2	41
81	High-density mapping and comparative analysis of agronomically important traits on wheat chromosome 3A. <i>Genomics</i> , 2006, 88, 74-87.	2.9	41
82	Mapping QTL for Agronomic Traits on Wheat Chromosome 3A and a Comparison of Recombinant Inbred Chromosome Line Populations. <i>Crop Science</i> , 2011, 51, 553-566.	1.8	40
83	Genetic Dissection of Yield and Its Component Traits Using High-Density Composite Map of Wheat Chromosome 3A: Bridging Gaps between QTLs and Underlying Genes. <i>PLoS ONE</i> , 2013, 8, e70526.	2.5	40
84	Validation of QTL for Grain Yield-Related Traits on Wheat Chromosome 3A Using Recombinant Inbred Chromosome Lines. <i>Crop Science</i> , 2012, 52, 1622-1632.	1.8	39
85	Seeding Rate, Genotype, and Topdressed Nitrogen Effects on Yield and Agronomic Characteristics of Winter Wheat. <i>Crop Science</i> , 2017, 57, 951-963.	1.8	38
86	Seeding Rate and Genotype Effect on Agronomic Performance and End-Use Quality of Winter Wheat. <i>Crop Science</i> , 2002, 42, 827.	1.8	38
87	Registration of 'NE01643'™ Wheat. <i>Journal of Plant Registrations</i> , 2008, 2, 36-42.	0.5	38
88	Winter Wheat Cultivar Characteristics Affect Annual Weed Suppression. <i>Weed Technology</i> , 2004, 18, 988-998.	0.9	37
89	Genetic architecture of common bunt resistance in winter wheat using genome-wide association study. <i>BMC Plant Biology</i> , 2018, 18, 280.	3.6	37
90	Yield and Grain Quality Responses of Soft Red Winter Wheat Exposed to Ozone During Anthesis 1. <i>Agronomy Journal</i> , 1986, 78, 593-600.	1.8	36

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91	Evaluating the Genetic Diversity of Triticale with Wheat and Rye SSR Markers. <i>Crop Science</i> , 2006, 46, 1692-1700.	1.8	36
92	Genotype, environment, seeding rate, and top-dressed nitrogen effects on end-use quality of modern Nebraska winter wheat. <i>Journal of the Science of Food and Agriculture</i> , 2017, 97, 5311-5318.	3.5	36
93	The effects of interactions of culture environment with genotype on wheat (<i>Triticum aestivum</i>) anther culture response. <i>Plant Cell Reports</i> , 1990, 8, 525-529.	5.6	34
94	Chromosomal Location of Wheat Quantitative Trait Loci Affecting Stability of Six Traits, Using Reciprocal Chromosome Substitutions. <i>Crop Science</i> , 1992, 32, 628-633.	1.8	34
95	High-yielding winter synthetic hexaploid wheats resistant to multiple diseases and pests. <i>Plant Genetic Resources: Characterisation and Utilisation</i> , 2018, 16, 273-278.	0.8	34
96	Distribution of Cadmium, Iron, and Zinc in Millstreams of Hard Winter Wheat (<i>Triticum</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 542 Td	5.2	33
97	Structuring an Efficient Organic Wheat Breeding Program. <i>Sustainability</i> , 2011, 3, 1190-1205.	3.2	32
98	Plant Height Response of Semidwarf and Nonsemidwarf Wheats to the Environment. <i>Crop Science</i> , 1995, 35, 447-451.	1.8	32
99	Genotypic variation of gas exchange parameters and carbon isotope discrimination in winter wheat. <i>Journal of Plant Physiology</i> , 2002, 159, 891-898.	3.5	31
100	Effects of Single and Double Infections of Winter Wheat by <i>Triticum mosaic virus</i> and <i>Wheat streak mosaic virus</i> on Yield Determinants. <i>Plant Disease</i> , 2012, 96, 859-864.	1.4	31
101	Genome-Wide Association Study for Multiple Biotic Stress Resistance in Synthetic Hexaploid Wheat. <i>International Journal of Molecular Sciences</i> , 2019, 20, 3667.	4.1	31
102	Analysis of Genotype x Environment Interaction in Wheat Using a Structural Equation Model and Chromosome Substitution Lines. <i>Crop Science</i> , 2007, 47, 477-484.	1.8	30
103	Principal variable selection to explain grain yield variation in winter wheat from features extracted from UAV imagery. <i>Plant Methods</i> , 2019, 15, 123.	4.3	30
104	Production, morphology, and cytogenetic analysis of <i>Elymus caninus</i> (<i>Agropyron caninum</i>) x <i>Triticum aestivum</i> F1 hybrids and backcross-1 derivatives. <i>Theoretical and Applied Genetics</i> , 1986, 71, 750-756.	3.6	29
105	Impact of wheat bran physical properties and chemical composition on whole grain flour mixing and baking properties. <i>Journal of Cereal Science</i> , 2019, 89, 102790.	3.7	29
106	Nuclear Genome Diversity and Relationships among Naturally Occurring Buffalograss Genotypes Determined by Sequence-related Amplified Polymorphism Markers. <i>Hortscience: A Publication of the American Society for Horticultural Science</i> , 2005, 40, 537-541.	1.0	28
107	Predicting phenological development in winter wheat. <i>Climate Research</i> , 2004, 25, 243-252.	1.1	28
108	Inheritance of the blue aleurone trait in diverse wheat crosses. <i>Genome</i> , 1990, 33, 525-529.	2.0	27

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109	Impact of Pre-Anthesis Water Deficit on Yield and Yield Components in Barley (<i>Hordeum vulgare</i> L.) Plants Grown under Controlled Conditions. <i>Agronomy</i> , 2016, 6, 33.	3.0	27
110	A simple wheat haploid and doubled haploid production system using anther culture. <i>In Vitro Cellular and Developmental Biology - Plant</i> , 2005, 41, 22-27.	2.1	26
111	Agronomic and quality effects in winter wheat of a gene conditioning resistance to wheat streak mosaic virus. <i>Euphytica</i> , 2006, 152, 41-49.	1.2	26
112	Characterization of Stem Rust Resistance in Wheat Cultivar Gage. <i>Crop Science</i> , 2015, 55, 229-239.	1.8	26
113	Genes Conditioning Resistance of <i>Hordeum spontaneum</i> to <i>Erysiphe graminis</i> f. sp. <i>hordei</i> 1. <i>Crop Science</i> , 1981, 21, 229-232.	1.8	25
114	The Effects of Genes Controlling Barley Leaf and Sheath Waxes on Agronomic Performance in Irrigated and Dryland Environments 1. <i>Crop Science</i> , 1983, 23, 116-120.	1.8	25
115	Putting genes into genetic coefficients. <i>Field Crops Research</i> , 2004, 90, 133-143.	5.1	25
116	Detailed Genetic Analysis for Identifying QTLs Associated with Drought Tolerance at Seed Germination and Seedling Stages in Barley. <i>Plants</i> , 2020, 9, 1425.	3.5	25
117	The Significance of Doubled Haploid Variation. <i>Stadler Genetics Symposia Series</i> , 1984, , 385-414.	0.0	25
118	Identification and Characterization of the Gene Conditioning Powdery Mildew Resistance in "Amigo"™ Wheat 1. <i>Crop Science</i> , 1984, 24, 129-132.	1.8	24
119	Genetic diversity and genetic variation in morpho-physiological traits to improve heat tolerance in Spring barley. <i>Molecular Biology Reports</i> , 2018, 45, 2441-2453.	2.3	24
120	Investigation of Heat-Induced Changes in the Grain Yield and Grains Metabolites, with Molecular Insights on the Candidate Genes in Barley. <i>Agronomy</i> , 2020, 10, 1730.	3.0	24
121	Evidence for microspore embryogenesis in wheat anther culture. <i>In Vitro Cellular and Developmental Biology - Plant</i> , 1991, 27, 168-174.	2.1	23
122	Identifying Winter Forage Triticale (<i>Triticosecale</i> Wittmack) Strains for the Central Great Plains. <i>Crop Science</i> , 2008, 48, 2040-2048.	1.8	23
123	Genome-wide association study reveals favorable alleles associated with common bunt resistance in synthetic hexaploid wheat. <i>Euphytica</i> , 2018, 214, 1.	1.2	23
124	Estimation of heterosis and combining abilities of U.S. winter wheat germplasm for hybrid development in Texas. <i>Crop Science</i> , 2020, 60, 788-803.	1.8	23
125	Registration of "Arapahoe"™ Wheat. <i>Crop Science</i> , 1989, 29, 832-832.	1.8	23
126	A better way to construct recombinant chromosome lines and their controls. <i>Genome</i> , 1992, 35, 827-830.	2.0	22

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127	DNA content of wheat monosomics at interphase estimated by flow cytometry. <i>Theoretical and Applied Genetics</i> , 1997, 95, 1300-1304.	3.6	22
128	Registration of 'Goodstreak'™ Wheat. <i>Crop Science</i> , 2004, 44, 1473-1474.	1.8	22
129	Genotype Imputation in Winter Wheat Using First-Generation Haplotype Map SNPs Improves Genome-Wide Association Mapping and Genomic Prediction of Traits. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 125-133.	1.8	22
130	The Scientific Grand Challenges of the 21st Century for the Crop Science Society of America. <i>Crop Science</i> , 2012, 52, 1003-1010.	1.8	21
131	Populations of doubled haploids for genetic mapping in hexaploid winter triticale. <i>Molecular Breeding</i> , 2018, 38, 46.	2.1	21
132	Evaluation of a global spring wheat panel for stripe rust: Resistance loci validation and novel resources identification. <i>PLoS ONE</i> , 2019, 14, e0222755.	2.5	21
133	Registration of 'NE06545'™ (Husker Genetics Brand Freeman) Hard Red Winter Wheat. <i>Journal of Plant Registrations</i> , 2014, 8, 279-284.	0.5	20
134	Effects of fungicide chemical class, fungicide application timing, and environment on Fusarium head blight in winter wheat. <i>European Journal of Plant Pathology</i> , 2020, 158, 667-679.	1.7	20
135	Combined GWAS and QTL mapping revealed candidate genes and SNP network controlling recovery and tolerance traits associated with drought tolerance in seedling winter wheat. <i>Genomics</i> , 2022, 114, 110358.	2.9	20
136	Cytogenetic characteristics of wheat plants regenerated from anther calli of 'Centurk'. <i>Genome</i> , 1983, 25, 513-517.	0.7	19
137	Genetic Analyses of Agronomic Traits Controlled by Wheat Chromosome 3A. <i>Crop Science</i> , 1999, 39, 1016-1021.	1.8	19
138	Perspectives on Low Temperature Tolerance and Vernalization Sensitivity in Barley: Prospects for Facultative Growth Habit. <i>Frontiers in Plant Science</i> , 2020, 11, 585927.	3.6	19
139	Registration of 'NH03614 CL'™ Wheat. <i>Journal of Plant Registrations</i> , 2011, 5, 75-80.	0.5	19
140	Registration of 'Mattern'™ Waxy (Amylose-free) Winter Wheat. <i>Journal of Plant Registrations</i> , 2014, 8, 43-48.	0.5	19
141	Using DArT Markers to Monitor Genetic Diversity throughout Selection: A Case Study in Nebraska's Winter Wheat Breeding Nurseries. <i>Crop Science</i> , 2013, 53, 2363-2373.	1.8	18
142	Native Fusarium head blight resistance from winter wheat cultivars 'Lyman', 'Overland', 'Ernie' and 'Freedom' mapped and pyramided onto 'Wesley'-Fhb1 backgrounds. <i>Molecular Breeding</i> , 2015, 35, 1-11.	2.1	18
143	Marker-Trait Associations for Enhancing Agronomic Performance, Disease Resistance, and Grain Quality in Synthetic and Bread Wheat Accessions in Western Siberia. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 4209-4222.	1.8	18
144	Baking quality of hard winter wheat: Response of cultivars to environment in the Great Plains. <i>Developments in Plant Breeding</i> , 1997, , 223-228.	0.2	18

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145	Grain Yield Performance and Stability of Cultivar Blends vs. Component Cultivars of Hard Winter Wheat in Nebraska. <i>Crop Science</i> , 2010, 50, 617-623.	1.8	17
146	Genetic diversity and population structure analysis of synthetic and bread wheat accessions in Western Siberia. <i>Journal of Applied Genetics</i> , 2019, 60, 283-289.	1.9	17
147	<i>Agrobacterium tumefaciens</i> -Mediated Wheat Transformation. <i>Cereal Research Communications</i> , 2003, 31, 9-16.	1.6	17
148	Registration of "Infinity CL"™ Wheat. <i>Crop Science</i> , 2006, 46, 975-977.	1.8	16
149	Identification of markers linked to genes for sprouting tolerance (independent of grain color) in hard white winter wheat (HWWW). <i>Theoretical and Applied Genetics</i> , 2016, 129, 419-430.	3.6	16
150	Variation in asparagine concentration in Nebraska wheat. <i>Cereal Chemistry</i> , 2018, 95, 264-273.	2.2	16
151	Yield and Quality in Purple-Grained Wheat Isogenic Lines. <i>Agronomy</i> , 2020, 10, 86.	3.0	16
152	The Effect of Gelling Agents on Wheat Anther and Immature Embryo Culture. <i>Plant Breeding</i> , 1992, 109, 211-217.	1.9	15
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256	Chromosomal locations of genes that control major RNA-degrading activities in common wheat (<i>Triticum aestivum</i> L.). <i>Theoretical and Applied Genetics</i> , 1996, 93, 645-648.	3.6	3
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277	Registration of "Hallam"™ Wheat. <i>Crop Science</i> , 2006, 46, 977-979.	1.8	1
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