

H-P Piepho

List of Publications by Year in descending order

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474
papers

18,070
citations

15504

65
h-index

29157

104
g-index

495
all docs

495
docs citations

495
times ranked

14440
citing authors

#	ARTICLE	IF	CITATIONS
1	Protein use efficiency and stability of baking quality in winter wheat based on the relation of loaf volume and grain protein content. Theoretical and Applied Genetics, 2022, , 1.	3.6	5
2	Contrasting drivers of belowground nitrogen cycling in a montane grassland exposed to a multifactorial global change experiment with elevated CO ₂ , warming, and drought. Global Change Biology, 2022, 28, 2425-2441.	9.5	25
3	Prediction of and for new environments: what's your model?. Molecular Plant, 2022, , .	8.3	7
4	Leveraging probability concepts for cultivar recommendation in multi-environment trials. Theoretical and Applied Genetics, 2022, 135, 1385-1399.	3.6	8
5	Two-dimensional P-spline smoothing for spatial analysis of plant breeding trials. Biometrical Journal, 2022, 64, 835-857.	1.0	8
6	Thyroid Hormone Concentrations in Testudo spp. by Season and Sex. Journal of Herpetological Medicine and Surgery, 2022, 32, .	0.4	0
7	Field investigation of topsoil moisture and temperature as drivers for decomposition or germination of sclerotia (<i>Sclerotinia sclerotiorum</i>) under winter-killed cover crops. Acta Agriculturae Scandinavica - Section B Soil and Plant Science, 2022, 72, 527-537.	0.6	2
8	Average semivariance directly yields accurate estimates of the genomic variance in complex trait analyses. G3: Genes, Genomes, Genetics, 2022, 12, .	1.8	7
9	Single-parent expression complementation contributes to phenotypic heterosis in maize hybrids. Plant Physiology, 2022, , .	4.8	6
10	Phenomics data processing: extracting dose-response curve parameters from high-resolution temperature courses and repeated field-based wheat height measurements. In Silico Plants, 2022, 4, .	1.9	9
11	How to observe the principle of concurrent control in an arm-based meta-analysis using <scp>SAS</scp> procedures <scp>GLIMMIX</scp> and <scp>BGLIMM</scp>. Research Synthesis Methods, 2022, 13, 821-828.	8.7	1
12	Assessing the response to genomic selection by simulation. Theoretical and Applied Genetics, 2022, 135, 2891-2905.	3.6	1
13	Linear mixed models and geostatistics for designed experiments in soil science: Two entirely different methods or two sides of the same coin?. European Journal of Soil Science, 2021, 72, 47-68.	3.9	11
14	Generating row-column field experimental designs with good neighbour balance and even distribution of treatment replications. Journal of Agronomy and Crop Science, 2021, 207, 745-753.	3.5	12
15	Missing association between nutrient concentrations in leaves and edible parts of food crops – A neglected food security issue. Food Chemistry, 2021, 345, 128723.	8.2	5
16	Genetic gain for rice yield in rainfed environments in India. Field Crops Research, 2021, 260, 107977.	5.1	37
17	Enviromics in breeding: applications and perspectives on envirotypic-assisted selection. Theoretical and Applied Genetics, 2021, 134, 95-112.	3.6	103
18	Optimizing the Allocation of Trials to Sub-regions in Multi-environment Crop Variety Testing. Journal of Agricultural, Biological, and Environmental Statistics, 2021, 26, 267-288.	1.4	3

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19	Early prediction of biomass in hybrid rye based on hyperspectral data surpasses genomic predictability in less-related breeding material. Theoretical and Applied Genetics, 2021, 134, 1409-1422.	3.6	15
20	Breeding progress of disease resistance and impact of disease severity under natural infections in winter wheat variety trials. Theoretical and Applied Genetics, 2021, 134, 1281-1302.	3.6	19
21	Methods of yield stability analysis in long-term field experiments. A review. Agronomy for Sustainable Development, 2021, 41, 1.	5.3	32
22	Projecting results of zoned multi-environment trials to new locations using environmental covariates with random coefficient models: accuracy and precision. Theoretical and Applied Genetics, 2021, 134, 1513-1530.	3.6	17
23	Influence of Cooling and Heating Systems on Pen Fouling, Lying Behavior, and Performance of Rearing Piglets. Agriculture (Switzerland), 2021, 11, 324.	3.1	4
24	Decline of seedling phosphorus use efficiency in the heterotic pool of flint maize breeding lines since the onset of hybrid breeding. Journal of Agronomy and Crop Science, 2021, 207, 857-872.	3.5	8
25	Importance of sources of variability, scales and experimental design: A case study about the effects of biochar and slurry application on soil properties in agricultural silty loam soils. European Journal of Soil Science, 2021, 72, 1954-1968.	3.9	0
26	Yield variability trends of winter wheat and spring barley grown during 1932â€“2019 in the Askov Long-term Experiment. Field Crops Research, 2021, 264, 108083.	5.1	13
27	Farmersâ€™ preferences for nature conservation compensation measures with a focus on eco-accounts according to the German Nature Conservation Act. Land Use Policy, 2021, 104, 105378.	5.6	9
28	Resolving the ambiguity of random-effects models with singular precision matrix. Statistica Neerlandica, 2021, 75, 482.	1.6	3
29	Regression models for order-of-addition experiments. Biometrical Journal, 2021, 63, 1673-1687.	1.0	1
30	High-throughput field phenotyping reveals genetic variation in photosynthetic traits in durum wheat under drought. Plant, Cell and Environment, 2021, 44, 2858-2878.	5.7	12
31	Assessing Spatial Variability of Barley Whole Crop Biomass Yield and Leaf Area Index in Silvoarable Agroforestry Systems Using UAV-Borne Remote Sensing. Remote Sensing, 2021, 13, 2751.	4.0	17
32	Appropriate sampling methods and statistics can tell apart fraud from pesticide drift in organic farming. Scientific Reports, 2021, 11, 14776.	3.3	4
33	Mineral-Ecological Cropping Systemsâ€”A New Approach to Improve Ecosystem Services by Farming without Chemical Synthetic Plant Protection. Agronomy, 2021, 11, 1710.	3.0	25
34	Average semivariance yields accurate estimates of the fraction of marker-associated genetic variance and heritability in complex trait analyses. PLoS Genetics, 2021, 17, e1009762.	3.5	12
35	Crafting for a better MAGIC: systematic design and test for Multiparental Advanced Generation Inter-Cross population. G3: Genes, Genomes, Genetics, 2021, 11, .	1.8	1
36	Back Cover Image. Plant, Cell and Environment, 2021, 44, .	5.7	0

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37	Effect of missing values in multi-environmental trials on variance component estimates. <i>Crop Science</i> , 2021, 61, 4087-4097.	1.8	11
38	Genome-enabled prediction for sparse testing in multi-environmental wheat trials. <i>Plant Genome</i> , 2021, 14, e20151.	2.8	15
39	Genetic variation for tolerance to the downy mildew pathogen <i>Peronospora variabilis</i> in genetic resources of quinoa (<i>Chenopodium quinoa</i>). <i>BMC Plant Biology</i> , 2021, 21, 41.	3.6	26
40	Long-term breeding progress of yield, yield-related, and disease resistance traits in five cereal crops of German variety trials. <i>Theoretical and Applied Genetics</i> , 2021, 134, 3805-3827.	3.6	14
41	Phenomics data processing: A plot-level model for repeated measurements to extract the timing of key stages and quantities at defined time points. <i>Field Crops Research</i> , 2021, 274, 108314.	5.1	18
42	Unraveling spatiotemporal variability of arbuscular mycorrhizal fungi in a temperate grassland plot. <i>Environmental Microbiology</i> , 2020, 22, 873-888.	3.8	27
43	Effects of converting a temperate short-rotation coppice to a silvo-arable alley cropping agroforestry system on soil quality indicators. <i>Agroforestry Systems</i> , 2020, 94, 389-400.	2.0	16
44	Choice of link and variance function for generalized linear mixed models: a case study with binomial response in proteomics. <i>Communications in Statistics - Theory and Methods</i> , 2020, 49, 4313-4332.	1.0	3
45	A Novel Model to Explain Extreme Feather Pecking Behavior in Laying Hens. <i>Behavior Genetics</i> , 2020, 50, 41-50.	2.1	19
46	Optimization of the extraction procedure for the determination of phenolic acids and flavonoids in the leaves of globe artichoke (<i>Cynara cardunculus</i> var. <i>scolymus</i> L.). <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2020, 177, 112879.	2.8	14
47	Stability Analysis of Tuber Yield and Starch Yield in Mid-Late and Late Maturing Starch Cultivars of Potato (<i>Solanum tuberosum</i>). <i>Potato Research</i> , 2020, 63, 179-197.	2.7	11
48	Growth responses of garden cress (<i>Lepidium sativum</i> L.) to biodynamic cow manure preparation in a bioassay. <i>Biological Agriculture and Horticulture</i> , 2020, 36, 16-34.	1.0	3
49	Impact of willow-based grassland alley cropping in relation to its plant species diversity on soil ecology of former arable land. <i>Applied Soil Ecology</i> , 2020, 147, 103373.	4.3	8
50	Analyzing designed experiments: Should we report standard deviations or standard errors of the mean or standard errors of the difference or what?. <i>Experimental Agriculture</i> , 2020, 56, 312-319.	0.9	5
51	Hyperspectral Reflectance Data and Agronomic Traits Can Predict Biomass Yield in Winter Rye Hybrids. <i>Bioenergy Research</i> , 2020, 13, 168-182.	3.9	10
52	Soil microbial community structure and function mainly respond to indirect effects in a multifactorial climate manipulation experiment. <i>Soil Biology and Biochemistry</i> , 2020, 142, 107704.	8.8	45
53	Novel strategies for genomic prediction of untested single-cross maize hybrids using unbalanced historical data. <i>Theoretical and Applied Genetics</i> , 2020, 133, 443-455.	3.6	22
54	Microbial growth and carbon use efficiency show seasonal responses in a multifactorial climate change experiment. <i>Communications Biology</i> , 2020, 3, 584.	4.4	30

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55	Soil and farm management effects on yield and nutrient concentrations of food crops in East Africa. <i>Science of the Total Environment</i> , 2020, 716, 137078.	8.0	14
56	Contrasting effects of cover crops on earthworms: Results from field monitoring and laboratory experiments on growth, reproduction and food choice. <i>European Journal of Soil Biology</i> , 2020, 100, 103225.	3.2	13
57	Nonproportional Hazards in Network Meta-Analysis: Efficient Strategies for Model Building and Analysis. <i>Value in Health</i> , 2020, 23, 918-927.	0.3	12
58	Integration of genotypic, hyperspectral, and phenotypic data to improve biomass yield prediction in hybrid rye. <i>Theoretical and Applied Genetics</i> , 2020, 133, 3001-3015.	3.6	34
59	Linear Variance, P-splines and Neighbour Differences for Spatial Adjustment in Field Trials: How are they Related?. <i>Journal of Agricultural, Biological, and Environmental Statistics</i> , 2020, 25, 676-698.	1.4	9
60	Guest Editorsâ€™ Introduction to the Special Issue on “Recent Advances in Design and Analysis of Experiments and Observational Studies in Agriculture” <i>Journal of Agricultural, Biological, and Environmental Statistics</i> , 2020, 25, 453-456.	1.4	2
61	Influence of A Cooled, Solid Lying Area on the Pen Fouling and Lying Behavior of Fattening Pigs. <i>Agriculture (Switzerland)</i> , 2020, 10, 307.	3.1	10
62	Long-term analysis from a cropping system perspective: Yield stability, environmental adaptability, and production risk of winter barley. <i>European Journal of Agronomy</i> , 2020, 117, 126056.	4.1	26
63	Augmented quasi-sudoku designs in field trials. <i>Computational Statistics and Data Analysis</i> , 2020, 150, 106988.	1.2	4
64	Determination of litter derived C and N in litterbags and soil using stable isotopes prevents overestimation of litter decomposition in alley cropping systems. <i>Pedobiologia</i> , 2020, 81-82, 150651.	1.2	4
65	The effects of cropping sequence, fertilization and straw management on the yield stability of winter wheat (1986â€“2017) in the Broadbalk Wheat Experiment, Rothamsted, UK. <i>Journal of Agricultural Science</i> , 2020, 158, 65-79.	1.3	17
66	Mapping Stem Rust (<i>Puccinia graminis</i> f. sp. <i>secalis</i>) Resistance in Self-Fertile Winter Rye Populations. <i>Frontiers in Plant Science</i> , 2020, 11, 667.	3.6	8
67	Long-term historical and projected herbivore population dynamics in Ngorongoro crater, Tanzania. <i>PLoS ONE</i> , 2020, 15, e0212530.	2.5	6
68	Aggregate formation and organo-mineral association affect characteristics of soil organic matter across soil horizons and parent materials in temperate broadleaf forest. <i>Biogeochemistry</i> , 2020, 148, 169-189.	3.5	6
69	Decoupling of impact factors reveals the response of German winter wheat yields to climatic changes. <i>Global Change Biology</i> , 2020, 26, 3601-3626.	9.5	35
70	Influence of Increased Light Intensity on the Acceptance of a Solid Lying Area and a Slatted Elimination Area in Fattening Pigs. <i>Agriculture (Switzerland)</i> , 2020, 10, 56.	3.1	10
71	Generating experimental designs for estimation of genetically related treatment effects using SAS. <i>Agronomy Journal</i> , 2020, 112, 3929-3940.	1.8	3
72	Influence of Anaerobic Digestion Processes on the Germination of Weed Seeds. <i>Gesunde Pflanzen</i> , 2020, 72, 181-194.	3.0	9

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73	Robust estimation of heritability and predictive accuracy in plant breeding: evaluation using simulation and empirical data. BMC Genomics, 2020, 21, 43.	2.8	6
74	Cross-validation of stagewise mixed-model analysis of Swedish variety trials with winter wheat and spring barley. Crop Science, 2020, 60, 2221-2240.	1.8	15
75	DRIFTS band areas as measured pool size proxy to reduce parameter uncertainty in soil organic matter models. Biogeosciences, 2020, 17, 1393-1413.	3.3	13
76	Mechanism of methane uptake in profiles of tropical soils converted from forest to rubber plantations. Soil Biology and Biochemistry, 2020, 145, 107796.	8.8	17
77	Heritability and Variability of Quality Parameters of Tomatoes in Outdoor Production. Research, 2020, 2020, 6707529.	5.7	13
78	Long-term field experiments in Germany: classification and spatial representation. Soil, 2020, 6, 579-596.	4.9	10
79	Does no-tillage decrease nitrate leaching compared to ploughing under a long-term crop rotation in Switzerland?. Soil and Tillage Research, 2020, 199, 104590.	5.6	18
80	Interactions between abiotic factors and the bioactivity of biodynamic horn manure on the growth of garden cress (<i>Lepidium sativum</i> L.) in a bioassay. Chemical and Biological Technologies in Agriculture, 2020, 7, .	4.6	0
81	DESIGNED EXPERIMENTS: DO YOU KNOW WHAT POPULATION YOU ARE SAMPLING FROM?. Experimental Agriculture, 2019, 55, 621-636.	0.9	3
82	Converting forests into rubber plantations weakened the soil CH ₄ sink in tropical uplands. Land Degradation and Development, 2019, 30, 2311-2322.	3.9	12
83	Heritability in Plant Breeding on a Genotype-Difference Basis. Genetics, 2019, 212, 991-1008.	2.9	94
84	Efficiency of Genomic Prediction of Nonassessed Testcrosses. Crop Science, 2019, 59, 2020-2027.	1.8	8
85	Error variance bias in neighbour balance and evenness of distribution designs. Australian and New Zealand Journal of Statistics, 2019, 61, 466-473.	0.9	6
86	Comparison of Weighted and Unweighted Stage-Wise Analysis for Genome-Wide Association Studies and Genomic Selection. Crop Science, 2019, 59, 2572-2584.	1.8	9
87	Trace gas fluxes from managed grassland soil subject to multifactorial climate change manipulation. Applied Soil Ecology, 2019, 137, 1-11.	4.3	14
88	Consequences of PCA graphs, SNP codings, and PCA variants for elucidating population structure. PLoS ONE, 2019, 14, e0218306.	2.5	26
89	Recent claim of declining climate resilience in European wheat is not supported by the statistics used. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 10625-10626.	7.1	9
90	Does fertilization impact production risk and yield stability across an entire crop rotation? Insights from a long-term experiment. Field Crops Research, 2019, 238, 82-92.	5.1	17

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91	Transcriptomic reprogramming of barley seminal roots by combined water deficit and salt stress. BMC Genomics, 2019, 20, 325.	2.8	42
92	Hypothesis Tests for Principal Component Analysis When Variables are Standardized. Journal of Agricultural, Biological, and Environmental Statistics, 2019, 24, 289-308.	1.4	31
93	Estimating Broadâ€Sense Heritability with Unbalanced Data from Agricultural Cultivar Trials. Crop Science, 2019, 59, 525-536.	1.8	56
94	A coefficient of determination (R^2) for generalized linear mixed models. Biometrical Journal, 2019, 61, 860-872.	1.0	77
95	Early diagnosis of ploidy status in doubled haploid production of maize by stomata length and flow cytometry measurements. Plant Breeding, 2019, 138, 266-276.	1.9	13
96	Testing multiplicative terms in AMMI and GGE models for multienvironment trials with replicates. Theoretical and Applied Genetics, 2019, 132, 2087-2096.	3.6	11
97	Modelling Spatio-Temporal Variation in Sparse Rainfall Data Using a Hierarchical Bayesian Regression Model. Journal of Agricultural, Biological, and Environmental Statistics, 2019, 24, 369-393.	1.4	9
98	A Crossâ€Validation of Statistical Models for Zonedâ€Based Prediction in Cultivar Testing. Crop Science, 2019, 59, 1544-1553.	1.8	11
99	Best Prediction of the Additive Genomic Variance in Random-Effects Models. Genetics, 2019, 213, 379-394.	2.9	8
100	Blocking and re-arrangement of pots in greenhouse experiments: which approach is more effective?. Plant Methods, 2019, 15, 143.	4.3	9
101	Do we need more drought for better nutrition? The effect of precipitation on nutrient concentration in East African food crops. Science of the Total Environment, 2019, 658, 405-415.	8.0	33
102	Analysing censored data in agricultural research: A review with examples and software tips. Annals of Applied Biology, 2019, 174, 3-13.	2.5	19
103	Mineral NPK and manure fertilisation affecting the yield stability of winter wheat: Results from a long-term field experiment. European Journal of Agronomy, 2019, 102, 14-22.	4.1	57
104	Similar spatial patterns of soil quality indicators in three poplar-based silvo-arable alley cropping systems in Germany. Biology and Fertility of Soils, 2019, 55, 1-14.	4.3	41
105	Simultaneous improvement of grain yield and protein content in durum wheat by different phenotypic indices and genomic selection. Theoretical and Applied Genetics, 2018, 131, 1315-1329.	3.6	87
106	Allowing for the structure of a designed experiment when estimating and testing trait correlations. Journal of Agricultural Science, 2018, 156, 59-70.	1.3	24
107	Estimating the variance for heterogeneity in armâ€based network metaâ€analysis. Pharmaceutical Statistics, 2018, 17, 264-277.	1.3	14
108	Weighted Estimation of AMMI and GGE Models. Journal of Agricultural, Biological, and Environmental Statistics, 2018, 23, 255-275.	1.4	8

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109	A tutorial on the statistical analysis of factorial experiments with qualitative and quantitative treatment factor levels. <i>Journal of Agronomy and Crop Science</i> , 2018, 204, 429-455.	3.5	59
110	Single-Parent Expression Is a General Mechanism Driving Extensive Complementation of Non-syntenic Genes in Maize Hybrids. <i>Current Biology</i> , 2018, 28, 431-437.e4.	3.9	50
111	Higher grain yield and higher grain protein deviation underline the potential of hybrid wheat for a sustainable agriculture. <i>Plant Breeding</i> , 2018, 137, 326-337.	1.9	46
112	What's normal anyway? Residual plots are more telling than significance tests when checking <scp>ANOVA</scp> assumptions. <i>Journal of Agronomy and Crop Science</i> , 2018, 204, 86-98.	3.5	227
113	An Evaluation of Error Variance Bias in Spatial Designs. <i>Journal of Agricultural, Biological, and Environmental Statistics</i> , 2018, 23, 83-91.	1.4	6
114	Factors controlling the variability of organic matter in the top- and subsoil of a sandy Dystric Cambisol under beech forest. <i>Geoderma</i> , 2018, 311, 37-44.	5.1	55
115	Efficiency of genomic prediction of non-assessed single crosses. <i>Heredity</i> , 2018, 120, 283-295.	2.6	17
116	Effects of biochar and slurry application as well as drying and rewetting on soil macroaggregate formation in agricultural silty loam soils. <i>Soil Use and Management</i> , 2018, 34, 575-583.	4.9	7
117	Phenotypic Selection in Ornamental Breeding: It's Better to Have the BLUPs Than to Have the BLUEs. <i>Frontiers in Plant Science</i> , 2018, 9, 1511.	3.6	17
118	Neighbor balance and evenness of distribution of treatment replications in row×column designs. <i>Biometrical Journal</i> , 2018, 60, 1172-1189.	1.0	19
119	Rainfall trends and variation in the Maasai Mara ecosystem and their implications for animal population and biodiversity dynamics. <i>PLoS ONE</i> , 2018, 13, e0202814.	2.5	61
120	Sowing Date in Egypt Affects Chia Seed Yield and Quality. <i>Agronomy Journal</i> , 2018, 110, 2310-2321.	1.8	6
121	Effects of chemical and physical grassland renovation on the temporal dynamics of organic carbon stocks and water-stable aggregate distribution in a sandy temperate grassland soil. <i>Soil Use and Management</i> , 2018, 34, 490-499.	4.9	2
122	Expected variance between seed germination test replicate results. <i>Seed Science and Technology</i> , 2018, 46, 197-209.	1.4	2
123	Genetic and phenotypic correlation for breeding relevant traits in <i>Dianthus caryophyllus</i> L.. <i>Postharvest Biology and Technology</i> , 2018, 143, 129-136.	6.0	6
124	Nonparametric Resampling Methods for Testing Multiplicative Terms in AMMI and GGE Models for Multi-environment Trials. <i>Crop Science</i> , 2018, 58, 752-761.	1.8	6
125	Predicting loaf volume for winter wheat by linear regression models based on protein concentration and sedimentation value using samples from VCU trials and mills. <i>Journal of Cereal Science</i> , 2018, 84, 132-141.	3.7	10
126	Letters in Mean Comparisons: What They Do and Don't Mean. <i>Agronomy Journal</i> , 2018, 110, 431-434.	1.8	45

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127	Biplots: Do Not Stretch Them!. Crop Science, 2018, 58, 1061-1069.	1.8	12
128	Chapter 6: Linear Regression Techniques. ACSESS Publications, 2018, , .	0.2	1
129	Ecosystem recovery indicators as decision criteria on potential reduction of fallow periods in swidden systems of Northern Thailand. Ecological Indicators, 2018, 95, 554-567.	6.3	5
130	Contribution to the discussion of "When should meta-analysis avoid making hidden normality assumptions?" Using general-purpose GLMM software for meta-analysis. Biometrical Journal, 2018, 60, 1059-1061.	1.0	0
131	More, Larger, Simpler: How Comparable Are On-Farm and On-Station Trials for Cultivar Evaluation?. Crop Science, 2018, 58, 1508-1518.	1.8	14
132	Modeling Spatially Correlated and Heteroscedastic Errors in Ethiopian Maize Trials. Crop Science, 2018, 58, 1575-1586.	1.8	8
133	Effects of fine root characteristics of beech on carbon turnover in the topsoil and subsoil of a sandy <sc>C</sc>ambisol. European Journal of Soil Science, 2017, 68, 177-188.	3.9	3
134	Comparative transcriptome analysis of vase life and carnation type in Dianthus caryophyllus L.. Scientia Horticulturae, 2017, 217, 61-72.	3.6	16
135	Selection for production-related traits in Pelargonium zonale: improved design and analysis make all the difference. Horticulture Research, 2017, 4, 17004.	6.3	13
136	Influence of elevated soil temperature and biochar application on organic matter associated with aggregate-size and density fractions in an arable soil. Agriculture, Ecosystems and Environment, 2017, 241, 79-87.	5.3	45
137	Efficient statistical design in two-phase experiments on vase life in carnations (Dianthus) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	6.0	16
138	The presence of extreme feather peckers in groups of laying hens. Animal, 2017, 11, 500-506.	3.3	11
139	Production objectives, trait and breed preferences of farmers keeping Nâ™Dama, Fulani Zebu and crossbred cattle and implications for breeding programs. Animal, 2017, 11, 687-695.	3.3	28
140	One Step at a Time: Stage-Wise Analysis of a Series of Experiments. Agronomy Journal, 2017, 109, 845-857.	1.8	71
141	Breeding progress, variation, and correlation of grain and quality traits in winter rye hybrid and population varieties and national on-farm progress in Germany over 26 years. Theoretical and Applied Genetics, 2017, 130, 981-998.	3.6	71
142	Genomic prediction in early selection stages using multi-year data in a hybrid rye breeding program. BMC Genetics, 2017, 18, 51.	2.7	31
143	Comparing the effectiveness profile of pharmacological interventions used for orthodontic pain relief: an arm-based multilevel network meta-analysis of longitudinal data. European Journal of Orthodontics, 2017, 39, 601-614.	2.4	9
144	Predicting biphasic responses in binary mixtures: Pelargonic acid versus glyphosate. Chemosphere, 2017, 178, 88-98.	8.2	33

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145	Cross-validation in AMMI and GGE Models: A Comparison of Methods. <i>Crop Science</i> , 2017, 57, 264-274.	1.8	12
146	Stability of Single-Parent Gene Expression Complementation in Maize Hybrids upon Water Deficit Stress. <i>Plant Physiology</i> , 2017, 173, 1247-1257.	4.8	36
147	A robust DF-REML framework for variance components estimation in genetic studies. <i>Bioinformatics</i> , 2017, 33, 3584-3594.	4.1	11
148	Designing an experiment with quantitative treatment factors to study the effects of climate change. <i>Journal of Agronomy and Crop Science</i> , 2017, 203, 584-592.	3.5	17
149	Breeding progress, genotypic and environmental variation and correlation of quality traits in malting barley in German official variety trials between 1983 and 2015. <i>Theoretical and Applied Genetics</i> , 2017, 130, 2411-2429.	3.6	35
150	Breeding progress, environmental variation and correlation of winter wheat yield and quality traits in German official variety trials and on-farm during 1983–2014. <i>Theoretical and Applied Genetics</i> , 2017, 130, 223-245.	3.6	133
151	Nondestructive Leaf Area Estimation for Chia. <i>Agronomy Journal</i> , 2017, 109, 1960-1969.	1.8	6
152	Quantifying uncertainty on sediment loads using bootstrap confidence intervals. <i>Hydrology and Earth System Sciences</i> , 2017, 21, 571-588.	4.9	9
153	Evaluating the Competitive Ability of Durum Wheat Varieties. <i>Agronomy Journal</i> , 2017, 109, 2606-2612.	1.8	0
154	Prediction Accuracy and Consistency in Cultivar Ranking for Factor-Analytic Linear Mixed Models for Winter Wheat Multienvironmental Trials. <i>Crop Science</i> , 2017, 57, 2506-2516.	1.8	9
155	Identifying Effective Design Approaches to Allocate Genotypes in Two-Phase Designs: A Case Study in <i>Pelargonium zonale</i> . <i>Frontiers in Plant Science</i> , 2017, 8, 2194.	3.6	6
156	Wildlife Population Dynamics in Human-Dominated Landscapes under Community-Based Conservation: The Example of Nakuru Wildlife Conservancy, Kenya. <i>PLoS ONE</i> , 2017, 12, e0169730.	2.5	42
157	Quantitative genetics theory for genomic selection and efficiency of genotypic value prediction in open-pollinated populations. <i>Scientia Agricola</i> , 2017, 74, 41-50.	1.2	11
158	Influence of wet heating and autoclaving on chemical composition and standardized ileal crude protein and amino acid digestibility in full-fat soybeans for pigs. <i>Journal of Animal Science</i> , 2017, 95, 779.	0.5	5
159	Farmer Participatory Early-Generation Yield Testing of Sorghum in West Africa: Possibilities to Optimize Genetic Gains for Yield in Farmers' Fields. <i>Crop Science</i> , 2016, 56, 2493-2505.	1.8	14
160	A Simulation-Based Approach for Evaluating the Efficiency of Multienvironment Trial Designs. <i>Crop Science</i> , 2016, 56, 2237-2250.	1.8	14
161	Diallel Analysis of Four Maize Traits and a Modified Heterosis Hypothesis. <i>Crop Science</i> , 2016, 56, 1115-1126.	1.8	10
162	Augmented Row-Column Designs for a Small Number of Checks. <i>Agronomy Journal</i> , 2016, 108, 2256-2262.	1.8	12

#	ARTICLE	IF	CITATIONS
163	Stability Analysis for a Countrywide Series of Wheat Trials in Pakistan. <i>Crop Science</i> , 2016, 56, 2465-2475.	1.8	16
164	Quantitative genetics theory for genomic selection and efficiency of breeding value prediction in open-pollinated populations. <i>Scientia Agricola</i> , 2016, 73, 243-251.	1.2	20
165	Extreme Wildlife Declines and Concurrent Increase in Livestock Numbers in Kenya: What Are the Causes?. <i>PLoS ONE</i> , 2016, 11, e0163249.	2.5	239
166	Agronomic improvements can make future cereal systems in South Asia far more productive and result in a lower environmental footprint. <i>Global Change Biology</i> , 2016, 22, 1054-1074.	9.5	70
167	Statistical Models and Methods for Network Meta-Analysis. <i>Phytopathology</i> , 2016, 106, 792-806.	2.2	81
168	Model training across multiple breeding cycles significantly improves genomic prediction accuracy in rye (<i>Secale cereale</i> L.). <i>Theoretical and Applied Genetics</i> , 2016, 129, 2043-2053.	3.6	84
169	Comparing the Predictive Abilities of Phenotypic and Marker-Assisted Selection Methods in a Biparental Lettuce Population. <i>Plant Genome</i> , 2016, 9, plantgenome2015.03.0014.	2.8	12
170	On generalized exponential transformations for proportions. <i>Communications in Statistics - Theory and Methods</i> , 2016, 45, 5857-5870.	1.0	4
171	Sediment-associated organic carbon and nitrogen inputs from erosion and irrigation to rice fields in a mountainous watershed in Northwest Vietnam. <i>Biogeochemistry</i> , 2016, 129, 93-113.	3.5	4
172	A Clustering-based Test for Nonadditivity in an Unreplicated Two-way Layout. <i>Communications in Statistics Part B: Simulation and Computation</i> , 2016, 45, 660-670.	1.2	5
173	Nonresolvable Row-Column Designs with an Even Distribution of Treatment Replications. <i>Journal of Agricultural, Biological, and Environmental Statistics</i> , 2016, 21, 227-242.	1.4	12
174	On a new family of shifted logarithmic transformations. <i>Journal of Statistical Computation and Simulation</i> , 2016, 86, 1697-1708.	1.2	1
175	Outlier detection methods for generalized lattices: a case study on the transition from ANOVA to REML. <i>Theoretical and Applied Genetics</i> , 2016, 129, 787-804.	3.6	80
176	Pitfalls in the use of middle-infrared spectroscopy: representativeness and ranking criteria for the estimation of soil properties. <i>Geoderma</i> , 2016, 268, 165-175.	5.1	20
177	Inferring relationships between Phosphorus utilization, feed per gain, and bodyweight gain in an F2 cross of Japanese quail using recursive models. <i>Poultry Science</i> , 2016, 95, 764-773.	3.4	21
178	Long-Term Experiments with cropping systems: Case studies on data analysis. <i>European Journal of Agronomy</i> , 2016, 77, 223-235.	4.1	48
179	Effects of soybean variety and <i>Bradyrhizobium</i> strains on yield, protein content and biological nitrogen fixation under cool growing conditions in Germany. <i>European Journal of Agronomy</i> , 2016, 72, 38-46.	4.1	104
180	Extensive tissue-specific transcriptomic plasticity in maize primary roots upon water deficit. <i>Journal of Experimental Botany</i> , 2016, 67, 1095-1107.	4.8	78

#	ARTICLE	IF	CITATIONS
181	Underdispersion of replicate results in germination tests is species and laboratory specific. <i>Seed Science and Technology</i> , 2016, 44, 281-297.	1.4	4
182	Is it necessary to split nitrogen fertilization for winter wheat? On-farm research on Luvisols in South-West Germany – CORRIGENDUM. <i>Journal of Agricultural Science</i> , 2015, 153, 1149-1149.	1.3	3
183	Phosphorous Efficiency and Tolerance Traits for Selection of Sorghum for Performance in Phosphorous-Limited Environments. <i>Crop Science</i> , 2015, 55, 1152-1162.	1.8	16
184	Beyond Latin Squares: A Brief Tour of Row-Column Designs. <i>Agronomy Journal</i> , 2015, 107, 2263-2270.	1.8	25
185	Controlling Misclassification Rates in Identification of Haploid Seeds from Induction Crosses in Maize with High-Oil Inducers. <i>Crop Science</i> , 2015, 55, 1076-1086.	1.8	10
186	Analysis of Series of Cultivar Trials with Perennial Grasses for Subdivided Target Regions. <i>Crop Science</i> , 2015, 55, 597-609.	1.8	5
187	Effect of substrate quality on the N uptake routes of soil microorganisms in different soil depths. <i>Pedobiologia</i> , 2015, 58, 211-218.	1.2	8
188	Optimality and Contrasts in Block Designs with Unequal Treatment Replication. <i>Australian and New Zealand Journal of Statistics</i> , 2015, 57, 203-209.	0.9	13
189	How Rainfall Variation Influences Reproductive Patterns of African Savanna Ungulates in an Equatorial Region Where Photoperiod Variation Is Absent. <i>PLoS ONE</i> , 2015, 10, e0133744.	2.5	20
190	How Genetic Variance and Number of Genotypes and Markers Influence Estimates of Genomic Prediction Accuracy in Plant Breeding. <i>Crop Science</i> , 2015, 55, 1911-1924.	1.8	5
191	Is it necessary to split nitrogen fertilization for winter wheat? On-farm research on Luvisols in South-West Germany. <i>Journal of Agricultural Science</i> , 2015, 153, 575-587.	1.3	32
192	Multiplicative interaction in network meta-analysis. <i>Statistics in Medicine</i> , 2015, 34, 582-594.	1.6	9
193	Segregation of non-target-site resistance to herbicides in multiple-resistant <i>Alopecurus myosuroides</i> plants. <i>Weed Research</i> , 2015, 55, 298-308.	1.7	5
194	Treatment comparisons in agricultural field trials accounting for spatial correlation. <i>Journal of Agricultural Science</i> , 2015, 153, 1187-1207.	1.3	16
195	Statistical modeling of the hormetic dose zone and the toxic potency completes the quantitative description of hormetic dose responses. <i>Environmental Toxicology and Chemistry</i> , 2015, 34, 1169-1177.	4.3	27
196	A High-Resolution Tissue-Specific Proteome and Phosphoproteome Atlas of Maize Primary Roots Reveals Functional Gradients along the Root Axes. <i>Plant Physiology</i> , 2015, 168, 233-246.	4.8	64
197	Inter-block information: to recover or not to recover it?. <i>Theoretical and Applied Genetics</i> , 2015, 128, 1541-1554.	3.6	18
198	Assessing the relative importance of dairy products to family nutrition in mixed crop-livestock production systems of Ethiopia. <i>Food Security</i> , 2015, 7, 1003-1015.	5.3	8

#	ARTICLE	IF	CITATIONS
199	Response of soil fertility indices to long-term application of biogas and raw slurry under organic farming. <i>Applied Soil Ecology</i> , 2015, 96, 99-107.	4.3	47
200	Yield performance and stability of CMS-based triticale hybrids. <i>Theoretical and Applied Genetics</i> , 2015, 128, 291-301.	3.6	8
201	The Elucidation of the Interactome of 16 Arabidopsis bZIP Factors Reveals Three Independent Functional Networks. <i>PLoS ONE</i> , 2015, 10, e0139884.	2.5	25
202	Deducing Hybrid Performance from Parental Metabolic Profiles of Young Primary Roots of Maize by Using a Multivariate Diallel Approach. <i>PLoS ONE</i> , 2014, 9, e85435.	2.5	19
203	Reproductive seasonality in African ungulates in relation to rainfall. <i>Wildlife Research</i> , 2014, 41, 323.	1.4	21
204	Genetic and non-genetic long-term trends of 12 different crops in German official variety performance trials and on-farm yield trends. <i>Theoretical and Applied Genetics</i> , 2014, 127, 2599-2617.	3.6	111
205	Regularized group regression methods for genomic prediction: Bridge, MCP, SCAD, group bridge, group lasso, sparse group lasso, group MCP and group SCAD. <i>BMC Proceedings</i> , 2014, 8, S7.	1.6	37
206	Transcriptomic complexity in young maize primary roots in response to low water potentials. <i>BMC Genomics</i> , 2014, 15, 741.	2.8	69
207	Nonsynthetic Genes Drive Highly Dynamic Complementation of Gene Expression in Maize Hybrids. <i>Plant Cell</i> , 2014, 26, 3939-3948.	6.6	80
208	Analysis of series of variety trials with perennial crops. <i>Grass and Forage Science</i> , 2014, 69, 431-440.	2.9	35
209	An Evaluation of Super-Valid Restricted Randomization. <i>Journal of Agricultural, Biological, and Environmental Statistics</i> , 2014, 19, 470-478.	1.4	2
210	Influence of Outliers on Accuracy Estimation in Genomic Prediction in Plant Breeding. <i>G3: Genes, Genomes, Genetics</i> , 2014, 4, 2317-2328.	1.8	19
211	Diversity of Stability, Localization, Interaction and Control of Downstream Gene Activity in the Maize Aux/IAA Protein Family. <i>PLoS ONE</i> , 2014, 9, e107346.	2.5	14
212	Effect of feeding level on ileal and total tract digestibility of nutrients and energy from soybean meal-based diets for piglets. <i>Journal of Animal Physiology and Animal Nutrition</i> , 2014, 98, 1154-1165.	2.2	7
213	Interspecies Variability of Plant Hormesis by the Antiauxin PCIB in a Laboratory Bioassay. <i>Journal of Plant Growth Regulation</i> , 2014, 33, 499-512.	5.1	15
214	Genome-based prediction of maize hybrid performance across genetic groups, testers, locations, and years. <i>Theoretical and Applied Genetics</i> , 2014, 127, 1375-1386.	3.6	90
215	Efficiency of augmented p-rep designs in multi-environmental trials. <i>Theoretical and Applied Genetics</i> , 2014, 127, 1049-1060.	3.6	45
216	Dissecting genetic and non-genetic sources of long-term yield trend in German official variety trials. <i>Theoretical and Applied Genetics</i> , 2014, 127, 1009-1018.	3.6	67

#	ARTICLE	IF	CITATIONS
217	Large herbivore responses to surface water and land use in an East African savanna: implications for conservation and human-wildlife conflicts. <i>Biodiversity and Conservation</i> , 2014, 23, 573-596.	2.6	50
218	Yield stability of hybrids versus lines in wheat, barley, and triticale. <i>Theoretical and Applied Genetics</i> , 2014, 127, 309-316.	3.6	130
219	Another Look at Bayesian Analysis of AMMI Models for Genotype-Environment Data. <i>Journal of Agricultural, Biological, and Environmental Statistics</i> , 2014, 19, 240.	1.4	21
220	Parametric bootstrap methods for testing multiplicative terms in GGE and AMMI models. <i>Biometrics</i> , 2014, 70, 639-647.	1.4	30
221	Impacts of climate change and variability on cattle production in southern Ethiopia: Perceptions and empirical evidence. <i>Agricultural Systems</i> , 2014, 130, 23-34.	6.1	49
222	The importance of phenotypic data analysis for genomic prediction - a case study comparing different spatial models in rye. <i>BMC Genomics</i> , 2014, 15, 646.	2.8	51
223	Exploitation of yield stability in barley. <i>Theoretical and Applied Genetics</i> , 2014, 127, 1949-1962.	3.6	32
224	Livestock Diversification: an Adaptive Strategy to Climate and Rangeland Ecosystem Changes in Southern Ethiopia. <i>Human Ecology</i> , 2014, 42, 509-520.	1.4	55
225	Genetic parameters for feather pecking and aggressive behavior in a large F2-cross of laying hens using generalized linear mixed models. <i>Poultry Science</i> , 2014, 93, 810-817.	3.4	46
226	Network-meta analysis made easy: detection of inconsistency using factorial analysis-of-variance models. <i>BMC Medical Research Methodology</i> , 2014, 14, 61.	3.1	48
227	A turbidity-based method to continuously monitor sediment, carbon and nitrogen flows in mountainous watersheds. <i>Journal of Hydrology</i> , 2014, 513, 45-57.	5.4	37
228	Herbivore Dynamics and Range Contraction in Kajiado County Kenya: Climate and Land Use Changes, Population Pressures, Governance, Policy and Human-wildlife Conflicts. <i>Open Ecology Journal</i> , 2014, 7, 9-31.	2.0	44
229	Variability of hormetic dose responses of the antiauxin <i>PCIB</i> on <i>Lactuca sativa</i> in a plant bioassay. <i>Weed Research</i> , 2013, 53, 418-428.	1.7	25
230	A Comparison of Spatial Designs for Field Variety Trials. <i>Australian and New Zealand Journal of Statistics</i> , 2013, 55, 253-258.	0.9	12
231	Evaluation of approaches for estimating the accuracy of genomic prediction in plant breeding. <i>BMC Genomics</i> , 2013, 14, 860.	2.8	38
232	Energy crop production in double-cropping systems: Results from an experiment at seven sites. <i>European Journal of Agronomy</i> , 2013, 51, 120-129.	4.1	68
233	Japanese Bindweed (<i>Calystegia hederacea</i>) Abundance and Response to Winter Wheat Seeding Rate and Nitrogen Fertilization in the North China Plain. <i>Weed Technology</i> , 2013, 27, 768-777.	0.9	8
234	Comparisons of single-stage and two-stage approaches to genomic selection. <i>Theoretical and Applied Genetics</i> , 2013, 126, 69-82.	3.6	51

#	ARTICLE	IF	CITATIONS
235	Hippopotamus and livestock grazing: influences on riparian vegetation and facilitation of other herbivores in the Mara Region of Kenya. <i>Landscape and Ecological Engineering</i> , 2013, 9, 47-58.	1.5	46
236	Choice of shrinkage parameter and prediction of genomic breeding values in elite maize breeding populations. <i>Plant Breeding</i> , 2013, 132, 99-106.	1.9	20
237	Performance of empirical BLUP and Bayesian prediction in small randomized complete block experiments. <i>Journal of Agricultural Science</i> , 2013, 151, 381-395.	1.3	12
238	Heterosis-associated proteome analyses of maize (<i>Zea mays</i> L.) seminal roots by quantitative label-free LC-MS. <i>Journal of Proteomics</i> , 2013, 93, 295-302.	2.4	31
239	Visual Scorings of Drought Stress Intensity as Covariates for Improved Variety Trial Analysis. <i>Journal of Agronomy and Crop Science</i> , 2013, 199, 321-330.	3.5	5
240	Genomic selection allowing for marker-by-environment interaction. <i>Plant Breeding</i> , 2013, 132, 532-538.	1.9	51
241	Why Randomize Agricultural Experiments?. <i>Journal of Agronomy and Crop Science</i> , 2013, 199, 374-383.	3.5	27
242	Comparative Transcriptome Profiling of Maize Coleoptilar Nodes during Shoot-Borne Root Initiation. <i>Plant Physiology</i> , 2013, 163, 419-430.	4.8	25
243	Response to "Controlling type 1 error rates in genome-wide association studies in plants" by Andrew W George. <i>Heredity</i> , 2013, 111, 88-88.	2.6	1
244	Assessment of variability in biomass yield and quality: what is an adequate size of sampling area for miscanthus?. <i>GCB Bioenergy</i> , 2013, 5, 572-579.	5.6	22
245	Using Genome-Wide Association Analysis to Characterize Environmental Sensitivity of Milk Traits in Dairy Cattle. <i>G3: Genes, Genomes, Genetics</i> , 2013, 3, 1085-1093.	1.8	16
246	Responses of phenology, synchrony and fecundity of breeding by African ungulates to interannual variation in rainfall. <i>Wildlife Research</i> , 2013, 40, 698.	1.4	13
247	Multiplicative Models for Specific Combining Ability in Half-Diallels with Parents. <i>Crop Science</i> , 2013, 53, 2321-2331.	1.8	1
248	Comparison of the Performance of Best Linear Unbiased Estimation and Best Linear Unbiased Prediction of Genotype Effects from Zoned Indian Maize Data. <i>Crop Science</i> , 2013, 53, 1384-1391.	1.8	31
249	Efficient Computation of Ridge Regression Best Linear Unbiased Prediction in Genomic Selection in Plant Breeding. <i>Crop Science</i> , 2012, 52, 1093-1104.	1.8	42
250	Strategies to Subdivide a Target Population of Environments: Results from the CIMMYT-Led Maize Hybrid Testing Programs in Africa. <i>Crop Science</i> , 2012, 52, 2143-2152.	1.8	49
251	Genomic Selection using Multiple Populations. <i>Crop Science</i> , 2012, 52, 2453-2461.	1.8	68
252	Crop yield evaluation under controlled drainage in Ohio, United States. <i>Journal of Soils and Water Conservation</i> , 2012, 67, 465-473.	1.6	42

#	ARTICLE	IF	CITATIONS
253	QTL detection power of multi-parental RIL populations in <i>Arabidopsis thaliana</i> . <i>Heredity</i> , 2012, 108, 626-632.	2.6	27
254	Selection Strategy for Sorghum Targeting Phosphorusâ€Limited Environments in West Africa: Analysis of Multiâ€Environment Experiments. <i>Crop Science</i> , 2012, 52, 2517-2527.	1.8	41
255	Drought yield index to select high yielding rice lines under different drought stress severities. <i>Rice</i> , 2012, 5, 31.	4.0	92
256	Occurrence and Distribution of 13 Trichothecene Toxins in Naturally Contaminated Maize Plants in Germany. <i>Toxins</i> , 2012, 4, 778-787.	3.4	38
257	Humanâ€hippo conflicts in Kenya during 1997â€2008: vulnerability of a megaherbivore to anthropogenic land use changes. <i>Journal of Land Use Science</i> , 2012, 7, 395-406.	2.2	25
258	The Use of Twoâ€Way Linear Mixed Models in Multitreatment Metaâ€Analysis. <i>Biometrics</i> , 2012, 68, 1269-1277.	1.4	58
259	Ostrich recruitment dynamics in relation to rainfall in the Maraâ€Serengeti ecosystem. <i>Ostrich</i> , 2012, 83, 119-136.	1.1	3
260	Checking Normality and Homoscedasticity in the General Linear Model Using Diagnostic Plots. <i>Communications in Statistics Part B: Simulation and Computation</i> , 2012, 41, 141-154.	1.2	28
261	The Area Under the Disease Progress Stairs: Calculation, Advantage, and Application. <i>Phytopathology</i> , 2012, 102, 381-389.	2.2	288
262	A stageâ€wise approach for the analysis of multiâ€environment trials. <i>Biometrical Journal</i> , 2012, 54, 844-860.	1.0	145
263	Modeling Effective Dosages in Hormetic Dose-Response Studies. <i>PLoS ONE</i> , 2012, 7, e33432.	2.5	58
264	Standardized ileal digestibilities of crude protein, amino acids, and contents of antinutritional factors, mycotoxins, and isoflavones of European soybean meal imports fed to piglets1. <i>Journal of Animal Science</i> , 2012, 90, 4883-4895.	0.5	17
265	Effect of feed intake level on ileal digestibilities of crude protein and amino acids in diets for piglets. <i>Journal of the Science of Food and Agriculture</i> , 2012, 92, 1261-1266.	3.5	8
266	Comparative changes in density and demography of large herbivores in the Masai Mara Reserve and its surrounding human-dominated pastoral ranches in Kenya. <i>Biodiversity and Conservation</i> , 2012, 21, 1509-1530.	2.6	67
267	Using precision farming technology to quantify yield effects attributed to weed competition and herbicide application. <i>Weed Research</i> , 2012, 52, 6-15.	1.7	28
268	High-yielding, drought-tolerant, stable rice genotypes for the shallow rainfed lowland drought-prone ecosystem. <i>Field Crops Research</i> , 2012, 133, 37-47.	5.1	48
269	Residual analysis of linear mixed models using a simulation approach. <i>Computational Statistics and Data Analysis</i> , 2012, 56, 1405-1416.	1.2	22
270	The distribution of large herbivore hotspots in relation to environmental and anthropogenic correlates in the Mara region of Kenya. <i>Journal of Animal Ecology</i> , 2012, 81, 1268-1287.	2.8	55

#	ARTICLE	IF	CITATIONS
271	Getting the Most Out of Sorghum Lowâ€Input Field Trials in West Africa Using Spatial Adjustment. Journal of Agronomy and Crop Science, 2012, 198, 349-359.	3.5	21
272	Genomic selection using regularized linear regression models: ridge regression, lasso, elastic net and their extensions. BMC Proceedings, 2012, 6, S10.	1.6	245
273	Production objectives and breeding goals of Sahiwal cattle keepers in Kenya and implications for a breeding programme. Tropical Animal Health and Production, 2012, 44, 519-530.	1.4	32
274	Dynamics of ungulates in relation to climatic and land use changes in an insularized African savanna ecosystem. Biodiversity and Conservation, 2012, 21, 1033-1053.	2.6	55
275	Statistical aspects of on-farm experimentation. Crop and Pasture Science, 2011, 62, 721.	1.5	46
276	Land-use intensity modifies spatial distribution and function of soil microorganisms in grasslands. Pedobiologia, 2011, 54, 341-351.	1.2	29
277	Combining phenotypic data from ordinal rating scales in multiple plant experiments. Trends in Plant Science, 2011, 16, 235-237.	8.8	18
278	Evaluation of basic and alternative breeding programs for Sahiwal cattle genetic resources in Kenya. Animal Production Science, 2011, 51, 682.	1.3	8
279	REMLâ€Based Diallel Analysis. Crop Science, 2011, 51, 470-478.	1.8	47
280	Genetic architecture of plant height in winter rye introgression libraries. Plant Breeding, 2011, 130, 209-216.	1.9	28
281	Corn hybrids display lower metabolite variability and complex metabolite inheritance patterns. Plant Journal, 2011, 68, 326-336.	5.7	75
282	Influence of leaching on the chemical composition of grassland biomass for combustion. Grass and Forage Science, 2011, 66, 464-473.	2.9	20
283	Continuing wildlife population declines and range contraction in the Mara region of Kenya during 1977â€2009. Journal of Zoology, 2011, 285, 99-109.	1.7	191
284	Influence of land-use intensity on the spatial distribution of N-cycling microorganisms in grassland soils. FEMS Microbiology Ecology, 2011, 77, 95-106.	2.7	70
285	A general method for controlling the genome-wide type I error rate in linkage and association mapping experiments in plants. Heredity, 2011, 106, 825-831.	2.6	32
286	Chemical composition and standardised ileal digestibilities of crude protein and amino acids in grain legumes for growing pigs. Livestock Science, 2011, 138, 229-243.	1.6	47
287	Integrating a simple shading algorithm into CERES-wheat and CERES-maize with particular regard to a changing microclimate within a relay-intercropping system. Field Crops Research, 2011, 121, 274-285.	5.1	44
288	Stability analysis of farmer participatory trials for conservation agriculture using mixed models. Field Crops Research, 2011, 121, 450-459.	5.1	34

#	ARTICLE	IF	CITATIONS
289	Dynamics of births and juvenile recruitment in Maraâ€“Serengeti ungulates in relation to climatic and land use changes. <i>Population Ecology</i> , 2011, 53, 195-213.	1.2	15
290	A comparison of random forests, boosting and support vector machines for genomic selection. <i>BMC Proceedings</i> , 2011, 5, S11.	1.6	169
291	Pre-selection of markers for genomic selection. <i>BMC Proceedings</i> , 2011, 5, S12.	1.6	25
292	Gene set analysis for longitudinal gene expression data. <i>BMC Bioinformatics</i> , 2011, 12, 273.	2.6	11
293	Augmented p-rep designs. <i>Biometrical Journal</i> , 2011, 53, 19-27.	1.0	109
294	Influence of coastal vegetation on the 2004 tsunami wave impact in west Aceh. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 18612-18617.	7.1	86
295	On Estimation of Genotypic Correlations and Their Standard Errors by Multivariate REML using the MIXED Procedure of the SAS System. <i>Crop Science</i> , 2011, 51, 2449-2454.	1.8	21
296	Population structure and phenotypic variation of a spring barley world collection set up for association studies. <i>Plant Breeding</i> , 2010, 129, 271-279.	1.9	47
297	Generation Means Analysis Using Mixed Models. <i>Crop Science</i> , 2010, 50, 1674-1680.	1.8	21
298	Molecular marker assisted broadening of the Central European heterotic groups in rye with Eastern European germplasm. <i>Theoretical and Applied Genetics</i> , 2010, 120, 291-299.	3.6	39
299	Optimum allocation of resources for QTL detection using a nested association mapping strategy in maize. <i>Theoretical and Applied Genetics</i> , 2010, 120, 553-561.	3.6	14
300	Heterotic patterns of sugar and amino acid components in developing maize kernels. <i>Theoretical and Applied Genetics</i> , 2010, 120, 369-381.	3.6	25
301	Heterosis in early seed development: a comparative study of F1 embryo and endosperm tissues 6Âdays after fertilization. <i>Theoretical and Applied Genetics</i> , 2010, 120, 389-400.	3.6	78
302	Prediction of hybrid performance in maize using molecular markers and joint analyses of hybrids and parental inbreds. <i>Theoretical and Applied Genetics</i> , 2010, 120, 451-461.	3.6	70
303	Background correction of two-colour cDNA microarray data using spatial smoothing methods. <i>Theoretical and Applied Genetics</i> , 2010, 120, 475-490.	3.6	3
304	Rainfall extremes explain interannual shifts in timing and synchrony of calving in topi and warthog. <i>Population Ecology</i> , 2010, 52, 89-102.	1.2	30
305	Genetic and economic evaluation of a basic breeding programme for Kenya Boran cattle. <i>Tropical Animal Health and Production</i> , 2010, 42, 327-340.	1.4	11
306	DNA polymorphisms and haplotype patterns of transcription factors involved in barley endosperm development are associated with key agronomic traits. <i>BMC Plant Biology</i> , 2010, 10, 5.	3.6	34

#	ARTICLE	IF	CITATIONS
307	Regulation of the maize (<i>Zea mays</i> L.) embryo proteome by RTCS which controls seminal root initiation. <i>European Journal of Cell Biology</i> , 2010, 89, 242-249.	3.6	17
308	Series of randomized complete block experiments with non-normal data. <i>Computational Statistics and Data Analysis</i> , 2010, 54, 1840-1857.	1.2	8
309	Current statistical issues in <i>Weed Research</i> . <i>Weed Research</i> , 2010, 50, 5-24.	1.7	118
310	Variation explained in mixed-model association mapping. <i>Heredity</i> , 2010, 105, 333-340.	2.6	133
311	Linear variance models for plant breeding trials. <i>Plant Breeding</i> , 2010, 129, 1-8.	1.9	50
312	Model selection and its consequences for different split-plot designs with spatial covariance and trend. <i>Plant Breeding</i> , 2010, 129, 590-598.	1.9	11
313	Arrangement of check plots in augmented block designs when spatial analysis is used. <i>Plant Breeding</i> , 2010, 129, 581-589.	1.9	16
314	Comparison of Spatial Models for Sugar Beet and Barley Trials. <i>Crop Science</i> , 2010, 50, 794-802.	1.8	31
315	Extension and Evaluation of Intercropping Field Trials Using Spatial Models. <i>Agronomy Journal</i> , 2010, 102, 1023-1031.	1.8	11
316	Correlation between per se and Testcross Performance in Rye (<i>Secale cereale</i> L.) Introgression Lines Estimated with a Bivariate Mixed Linear Model. <i>Crop Science</i> , 2010, 50, 1863-1873.	1.8	8
317	Development of Heterotic Groups in Triticale. <i>Crop Science</i> , 2010, 50, 584-590.	1.8	76
318	A Modeling Approach to Simulate Effects of Intercropping and Interspecific Competition in Arable Crops. <i>International Journal of Information Systems and Social Change</i> , 2010, 1, 44-65.	0.1	10
319	Genome-wide selection by mixed model ridge regression and extensions based on geostatistical models. <i>BMC Proceedings</i> , 2010, 4, S8.	1.6	17
320	Large herbivore responses to water and settlements in savannas. <i>Ecological Monographs</i> , 2010, 80, 241-266.	5.4	52
321	Nonadditive Protein Accumulation Patterns in Maize (<i>Zea mays</i> L.) Hybrids during Embryo Development. <i>Journal of Proteome Research</i> , 2010, 9, 6511-6522.	3.7	42
322	Estimates of dietary threshold levels for crude protein and amino acids to obtain plateau values of apparent ileal crude protein and amino acid digestibilities in newly weaned pigs. <i>Archives of Animal Nutrition</i> , 2010, 64, 357-372.	1.8	15
323	Comparison of Weighting in Two-Stage Analysis of Plant Breeding Trials. <i>Crop Science</i> , 2009, 49, 1977-1988.	1.8	183
324	Impact of Genetic Divergence on the Ratio of Variance Due to Specific vs. General Combining Ability in Winter Triticale. <i>Crop Science</i> , 2009, 49, 2119-2122.	1.8	14

#	ARTICLE	IF	CITATIONS
325	Standardised ileal crude protein and amino acid digestibilities in protein supplements for piglets. Archives of Animal Nutrition, 2009, 63, 356-378.	1.8	24
326	Specification of Cortical Parenchyma and Stele of Maize Primary Roots by Asymmetric Levels of Auxin, Cytokinin, and Cytokinin-Regulated Proteins. Plant Physiology, 2009, 152, 4-18.	4.8	56
327	Unraveling Epistasis With Triple Testcross Progenies of Near-Isogenic Lines. Genetics, 2009, 181, 247-257.	2.9	28
328	Validation of candidate genes putatively associated with resistance to SCMV and MDMV in maize (Zea mays L.) overlock 1000. Theoretical and Applied Genetics, 2009, 118, 259-273.	3.6	96
329	Taste aversion learning to eliminate feather pecking in laying hens, Gallus gallus domesticus. Animal Behaviour, 2009, 78, 485-490.	1.9	20
330	Quantitative trait loci mapping for biomass yield traits in a Lolium inbred line derived F2 population. Euphytica, 2009, 170, 99-107.	1.2	30
331	Association mapping reveals gene action and interactions in the determination of flowering time in barley. Theoretical and Applied Genetics, 2009, 118, 259-273.	3.6	96
332	Molecular marker-based prediction of hybrid performance in maize using unbalanced data from multiple experiments with factorial crosses. Theoretical and Applied Genetics, 2009, 118, 741-751.	3.6	76
333	Testcross performance of rye introgression lines developed by marker-assisted backcrossing using an Iranian accession as donor. Theoretical and Applied Genetics, 2009, 118, 1225-1238.	3.6	42
334	Dynamics of Serengeti ungulates in relation to land use changes. Journal of Zoology, 2009, 278, 1-14.	1.7	129
335	Modelling of a recording scheme for market-oriented smallholder pig producers in Northwest Vietnam. Livestock Science, 2009, 123, 241-248.	1.6	20
336	Stated preferences of llama keeping functions in Bolivia. Livestock Science, 2009, 124, 119-125.	1.6	24
337	Tissue Specific Control of the Maize (Zea mays L.) Embryo, Cortical Parenchyma, and Stele Proteomes by RUM1 Which Regulates Seminal and Lateral Root Initiation. Journal of Proteome Research, 2009, 8, 2285-2297.	3.7	21
338	Ridge Regression and Extensions for Genomewide Selection in Maize. Crop Science, 2009, 49, 1165-1176.	1.8	234
339	Data Transformation in Statistical Analysis of Field Trials with Changing Treatment Variance. Agronomy Journal, 2009, 101, 865-869.	1.8	93
340	Determination of standardized ileal crude protein and amino acid digestibilities in protein supplements for piglets. Animal, 2009, 3, 1696-1705.	3.3	11
341	An on-farm approach to quantify yield variation and to derive decision rules for site-specific weed management. Precision Agriculture, 2008, 9, 133-146.	6.0	33
342	BLUP for phenotypic selection in plant breeding and variety testing. Euphytica, 2008, 161, 209-228.	1.2	569

#	ARTICLE	IF	CITATIONS
343	Assessing the importance of genotype—environment interaction for root traits in rice using a mapping population III: QTL analysis by mixed models. <i>Euphytica</i> , 2008, 161, 229-240.	1.2	5
344	Comparing the performance of cereal varieties in organic and non-organic cropping systems in different European countries. <i>Euphytica</i> , 2008, 163, 417-433.	1.2	80
345	REML approach for adjusting the Fusarium head blight rating to a phenological date in inoculated selection experiments of wheat. <i>Theoretical and Applied Genetics</i> , 2008, 117, 65-73.	3.6	48
346	Multi-trait association mapping in sugar beet (<i>Beta vulgaris</i> L.). <i>Theoretical and Applied Genetics</i> , 2008, 117, 947-954.	3.6	57
347	Association mapping in multiple segregating populations of sugar beet (<i>Beta vulgaris</i> L.). <i>Theoretical and Applied Genetics</i> , 2008, 117, 1167-1179.	3.6	28
348	Analysis of nonadditive protein accumulation in young primary roots of a maize (<i>Zea mays</i> L.) F1-Hybrid and Parental Inbred Line Primary Root Transcriptomes Suggests Organ-Specific Patterns of Nonadditive Gene Expression and Conserved Expression Trends. <i>Genetics</i> , 2008, 179, 1275-1283.	2.2	57
349	Estimates of basal ileal endogenous losses of amino acids by regression analysis and determination of standardised ileal amino acid digestibilities from casein in newly weaned pigs. <i>Journal of the Science of Food and Agriculture</i> , 2008, 88, 641-651.	3.5	24
350	Nearest Neighbour Adjustment and Linear Variance Models in Plant Breeding Trials. <i>Biometrical Journal</i> , 2008, 50, 164-189.	1.0	53
351	Variance Component Estimation for Mixed Model Analysis of cDNA Microarray Data. <i>Biometrical Journal</i> , 2008, 50, 927-939.	1.0	3
352	Effect of dietary crude protein level on basal ileal endogenous losses and standardized ileal digestibilities of crude protein and amino acids in newly weaned pigs. <i>Journal of Animal Physiology and Animal Nutrition</i> , 2008, 92, 578-590.	2.2	29
353	Trends in genetic variance components during 30 years of hybrid maize breeding at the University of Hohenheim. <i>Plant Breeding</i> , 2008, 127, 446-451.	1.9	44
354	El Niño—Southern Oscillation, rainfall, temperature and Normalized Difference Vegetation Index fluctuations in the Mara—Serengeti ecosystem. <i>African Journal of Ecology</i> , 2008, 46, 132-143.	0.9	88
355	Rainfall influences on ungulate population abundance in the Mara—Serengeti ecosystem. <i>Journal of Animal Ecology</i> , 2008, 77, 814-829.	2.8	115
356	Erratum to “Meta-Analysis of the Relative Efficiency of Methionine-Hydroxy-Analogue-Free-Acid Compared with dl-Methionine in Broilers Using Nonlinear Mixed Models” (Poult Sci. 87:2023–2031). <i>Poultry Science</i> , 2008, 87, 2690.	3.4	0
357	Comparison of Mixed-Model Approaches for Association Mapping. <i>Genetics</i> , 2008, 178, 1745-1754.	2.9	273
358	Meta-Analysis of the Relative Efficiency of Methionine-Hydroxy-Analogue-Free-Acid Compared with dl-Methionine in Broilers Using Nonlinear Mixed Models. <i>Poultry Science</i> , 2008, 87, 2023-2031.	3.4	55
359	Comparison of Maize (<i>Zea mays</i> L.) F1-Hybrid and Parental Inbred Line Primary Root Transcriptomes Suggests Organ-Specific Patterns of Nonadditive Gene Expression and Conserved Expression Trends. <i>Genetics</i> , 2008, 179, 1275-1283.	2.9	111
360	A note on the bias of genetic distances in linkage maps based on small samples for backcrosses and intercrosses with complete dominance. <i>Genome</i> , 2008, 51, 1054-1061.	2.0	2

#	ARTICLE	IF	CITATIONS
361	Statistical Analysis of Yield Trials by AMMI and GGE: Further Considerations. Crop Science, 2008, 48, 866-889.	1.8	347
362	Effect of Ileal Infusion of Short-Chain Fatty Acids on Pancreatic Prandial Secretion and Gastrointestinal Hormones in Pigs. Pancreas, 2008, 37, 196-202.	1.1	13
363	The Role of Epistasis in the Manifestation of Heterosis: A Systems-Oriented Approach. Genetics, 2007, 177, 1815-1825.	2.9	125
364	Comment on the article on "Within-field wheat yield prediction from IKONOS data, a new matrix approach"™ by E.Á. Enclona, P.Á. Thenkabail, D. Celis and J. Diekmann, International Journal of Remote Sensing, 25, 377-388 (2004). International Journal of Remote Sensing, 2007, 28, 1907-1911.	2.9	1
365	Simple State-Space Models in a Mixed Model Framework. American Statistician, 2007, 61, 224-232.	1.6	20
366	Analysis of a Triple Testcross Design With Recombinant Inbred Lines Reveals a Significant Role of Epistasis in Heterosis for Biomass-Related Traits in Arabidopsis. Genetics, 2007, 175, 2009-2017.	2.9	65
367	Genetic Basis of Heterosis for Growth-Related Traits in Arabidopsis Investigated by Testcross Progenies of Near-Isogenic Lines Reveals a Significant Role of Epistasis. Genetics, 2007, 177, 1827-1837.	2.9	95
368	Power to Detect Higher-Order Epistatic Interactions in a Metabolic Pathway Using a New Mapping Strategy. Genetics, 2007, 176, 563-570.	2.9	43
369	Heterosis for Biomass-Related Traits in Arabidopsis Investigated by Quantitative Trait Loci Analysis of the Triple Testcross Design With Recombinant Inbred Lines. Genetics, 2007, 177, 1839-1850.	2.9	55
370	Computing Heritability and Selection Response From Unbalanced Plant Breeding Trials. Genetics, 2007, 177, 1881-1888.	2.9	483
371	Algorithms for compact letter displays: Comparison and evaluation. Computational Statistics and Data Analysis, 2007, 52, 725-736.	1.2	28
372	The Effect of Preceding Crop and Pre-Sprouting on Crop Growth, N Use and Tuber Yield of Maincrop Potatoes for Processing Under Conditions of N Stress. Journal of Agronomy and Crop Science, 2007, 193, 270-291.	3.5	11
373	Temporal changes in allele frequencies in two European F2 flint maize populations under modified recurrent full-sib selection. Theoretical and Applied Genetics, 2007, 114, 765-776.	3.6	20
374	Prediction of single-cross hybrid performance in maize using haplotype blocks associated with QTL for grain yield. Theoretical and Applied Genetics, 2007, 114, 1345-1355.	3.6	33
375	Linkage disequilibrium in two European F2 flint maize populations under modified recurrent full-sib selection. Theoretical and Applied Genetics, 2007, 115, 289-297.	3.6	6
376	Potential causes of linkage disequilibrium in a European maize breeding program investigated with computer simulations. Theoretical and Applied Genetics, 2007, 115, 529-536.	3.6	20
377	Are ordinal rating scales better than percent ratings? a statistical and "psychological" view. Euphytica, 2007, 155, 15-26.	1.2	21
378	Multiplicative main cultivar effects in Czech official winter wheat trials 1976-2005. Czech Journal of Genetics and Plant Breeding, 2007, 43, 117-124.	0.8	1

#	ARTICLE	IF	CITATIONS
379	A Cautionary Note on Appropriate Statistical Methods to Compare Dose Responses of Methionine Sources. Poultry Science, 2006, 85, 1511-1512.	3.4	6
380	Selection in Cultivar Trials—Is It Ignorable?. Crop Science, 2006, 46, 192-201.	1.8	85
381	Subacute effects of moderate feed loads of isolated Fusarium toxin deoxynivalenol on selected parameters of metabolism in weaned growing piglets. Journal of Animal Physiology and Animal Nutrition, 2006, 90, 421-428.	2.2	19
382	On the use of multiple lattice designs and λ -designs in plant breeding trials. Plant Breeding, 2006, 125, 523-528.	1.9	28
383	Efficiency of strip- and line-transect surveys of African savanna mammals. Journal of Zoology, 2006, 269, 060303002124001-???	1.7	59
384	Application of mark-recapture methods to lions: satisfying assumptions by using covariates to explain heterogeneity. Journal of Zoology, 2006, 269, 060423083931002-???	1.7	22
385	Comparative expression profiling in meristems of inbred-hybrid triplets of maize based on morphological investigations of heterosis for plant height. Plant Molecular Biology, 2006, 63, 21-34.	3.9	97
386	Analysis of Genebank Evaluation Data by using Geostatistical Methods. Genetic Resources and Crop Evolution, 2006, 53, 737-751.	1.6	2
387	A simple note on how to save money in linkage analysis. Molecular Breeding, 2006, 18, 291-300.	2.1	3
388	Manifestation of heterosis during early maize (<i>Zea mays</i> L.) root development. Theoretical and Applied Genetics, 2006, 112, 421-429.	3.6	104
389	Assessing the importance of genotype \times environment interaction for root traits in rice using a mapping population. I: a soil-filled box screen. Theoretical and Applied Genetics, 2006, 113, 977-986.	3.6	37
390	Assessing the importance of genotype \times environment interaction for root traits in rice using a mapping population II: conventional QTL analysis. Theoretical and Applied Genetics, 2006, 113, 953-964.	3.6	113
391	A new test for family-based association mapping with inbred lines from plant breeding programs. Theoretical and Applied Genetics, 2006, 113, 1121-1130.	3.6	33
392	ZmGrp3: identification of a novel marker for root initiation in maize and development of a robust assay to quantify allele-specific contribution to gene expression in hybrids. Theoretical and Applied Genetics, 2006, 113, 1305-1315.	3.6	8
393	A comparison of experimental designs for selection in breeding trials with nested treatment structure. Theoretical and Applied Genetics, 2006, 113, 1505-1513.	3.6	34
394	Combining signals from spotted cDNA microarrays obtained at different scanning intensities. Bioinformatics, 2006, 22, 802-807.	4.1	11
395	The Effect of Feather Eating on Feed Passage in Laying Hens. Poultry Science, 2006, 85, 21-25.	3.4	43
396	Best Linear Unbiased Prediction of Cultivar Effects for Subdivided Target Regions. Crop Science, 2005, 45, 1151-1159.	1.8	81

#	ARTICLE	IF	CITATIONS
397	A simulation study on tests of hypotheses and confidence intervals for fixed effects in mixed models for blocked experiments with missing data. <i>Journal of Agricultural, Biological, and Environmental Statistics</i> , 2005, 10, 374-389.	1.4	50
398	Analysis of Unbalanced Data by Mixed Linear Models Using the mixed Procedure of the SAS System. <i>Journal of Agronomy and Crop Science</i> , 2005, 191, 47-54.	3.5	97
399	Statistical tests for QTL and QTL-by-environment effects in segregating populations derived from line crosses. <i>Theoretical and Applied Genetics</i> , 2005, 110, 561-566.	3.6	39
400	Designing a microarray experiment to estimate dominance in maize (<i>Zea mays</i> L.). <i>Theoretical and Applied Genetics</i> , 2005, 111, 57-64.	3.6	10
401	Permutation tests for the correlation among genetic distances and measures of heterosis. <i>Theoretical and Applied Genetics</i> , 2005, 111, 95-99.	3.6	7
402	Spatial models as a tool to identify spatial patterns of surficial sediment composition and their contributing factors in the littoral zone of Lake Constance (Germany). <i>Aquatic Sciences</i> , 2005, 67, 326-336.	1.5	3
403	Is heterosis an artefact governed by the choice of scale?. <i>Euphytica</i> , 2005, 145, 113-121.	1.2	5
404	A Threshold Model for Multiyear Genebank Data Based on Different Rating Scales. <i>Crop Science</i> , 2005, 45, 1045-1051.	1.8	5
405	Optimal Allocation in Designs for Assessing Heterosis From cDNA Gene Expression Data. <i>Genetics</i> , 2005, 171, 359-364.	2.9	9
406	Simultaneous Confidence Intervals for Two Estimable Functions and Their Ratio Under a Linear Model. <i>American Statistician</i> , 2005, 59, 292-300.	1.6	12
407	Effect of supplementation of xylanase and phospholipase to a wheat-based diet for weanling pigs on nutrient digestibility and concentrations of microbial metabolites in ileal digesta and feces ¹ . <i>Journal of Animal Science</i> , 2004, 82, 2647-2656.	0.5	102
408	A Mixed Modelling Approach for Randomized Experiments with Repeated Measures. <i>Journal of Agronomy and Crop Science</i> , 2004, 190, 230-247.	3.5	154
409	Mixed modelling for QTL \times environment interaction analysis. <i>Euphytica</i> , 2004, 137, 147-153.	1.2	22
410	Legume rotation effects on early growth and rhizosphere microbiology of sorghum in West African soils. <i>Plant and Soil</i> , 2004, 264, 325-334.	3.7	19
411	Approximating the degrees of freedom for contrasts of genotypes laid out as subplots in an alpha-design in a split-plot experiment. <i>Plant Breeding</i> , 2004, 123, 193-197.	1.9	10
412	Inbreeding coefficients for stochastically varying small population sizes—bias of calculation based on effective numbers. <i>Journal of Theoretical Biology</i> , 2004, 226, 467-475.	1.7	2
413	Serum IgA-Promoting Effects Induced by Feed Loads Containing Isolated Deoxynivalenol (DON) in Growing Piglets. <i>Journal of Toxicology and Environmental Health - Part A: Current Issues</i> , 2004, 67, 1051-1067.	2.3	29
414	Transformations in mixed models: Application to risk analysis for a multienvironment trial. <i>Journal of Agricultural, Biological, and Environmental Statistics</i> , 2004, 9, 123-137.	1.4	15

#	ARTICLE	IF	CITATIONS
415	An Algorithm for a Letter-Based Representation of All-Pairwise Comparisons. Journal of Computational and Graphical Statistics, 2004, 13, 456-466.	1.7	313
416	Digesta characteristics of dorsal, middle and ventral rumen of cows fed with different hay qualities and concentrate levels. Archives of Animal Nutrition, 2004, 58, 325-342.	1.8	45
417	Model-based mean adjustment in quantitative germplasm evaluation data. Genetic Resources and Crop Evolution, 2003, 50, 281-290.	1.6	5
418	Responses of different arabica coffee (Coffea arabica L.) clones to varied environmental conditions. Euphytica, 2003, 129, 175-182.	1.2	13
419	Zur Beschreibung des Fichtenwachstums mit Hilfe nichtlinearer Regression. European Journal of Forest Research, 2003, 122, 359-367.	0.3	0
420	Determining the sample size for co-dominant molecular marker-assisted linkage detection for a monogenic qualitative trait by controlling the type-I and type-II errors in a segregating F2 population. Theoretical and Applied Genetics, 2003, 106, 840-845.	3.6	11
421	The folded exponential transformation for proportions. Journal of the Royal Statistical Society: Series D (the Statistician), 2003, 52, 575-589.	0.2	33
422	A Hitchhiker's Guide to Mixed Models for Randomized Experiments. Journal of Agronomy and Crop Science, 2003, 189, 310-322.	3.5	282
423	Threshold models with fixed and random effects for ordered categorical data. Food Quality and Preference, 2003, 14, 343-357.	4.6	10
424	MULTI-SITE TIME-TREND ANALYSIS OF SOIL FERTILITY MANAGEMENT EFFECTS ON CROP PRODUCTION IN SUB-SAHARAN WEST AFRICA. Experimental Agriculture, 2002, 38, 163-183.	0.9	38
425	A simple mixed model for trend analysis in wildlife populations. Journal of Agricultural, Biological, and Environmental Statistics, 2002, 7, 350-360.	1.4	28
426	Estimating the product-moment correlation in samples with censoring on both variables. Biometrical Journal, 2002, 44, 657-670.	1.0	7
427	Can the sample variance estimator be improved by using a covariate?. Journal of Agricultural, Biological, and Environmental Statistics, 2002, 7, 157-175.	1.4	1
428	Efficient phosphorus application strategies for increased crop production in sub-Saharan West Africa. Field Crops Research, 2001, 72, 1-15.	5.1	104
429	Significance testing for QTL mapping by marker difference regression. Theoretical and Applied Genetics, 2001, 102, 1099-1102.	3.6	0
430	Exploiting quantitative information in the analysis of dominant markers. Theoretical and Applied Genetics, 2001, 103, 462-468.	3.6	6
431	A Quick Method for Computing Approximate Thresholds for Quantitative Trait Loci Detection. Genetics, 2001, 157, 425-432.	2.9	150
432	Marker Pair Selection for Mapping Quantitative Trait Loci. Genetics, 2001, 157, 433-444.	2.9	70

#	ARTICLE	IF	CITATIONS
433	Multiple Treatment Comparisons in Linear Models When the Standard Error of a Difference is not Constant. <i>Biometrical Journal</i> , 2000, 42, 823-835.	1.0	13
434	Optimal marker density for interval mapping in a backcross population. <i>Heredity</i> , 2000, 84, 437-440.	2.6	42
435	Exact Confidence Limits for Covariate-Dependent Risk in Cultivar Trials. <i>Journal of Agricultural, Biological, and Environmental Statistics</i> , 2000, 5, 202.	1.4	11
436	Codominant Analysis of Banding Data From a Dominant Marker System by Normal Mixtures. <i>Genetics</i> , 2000, 155, 1459-1468.	2.9	47
437	A Mixed-Model Approach to Mapping Quantitative Trait Loci in Barley on the Basis of Multiple Environment Data. <i>Genetics</i> , 2000, 156, 2043-2050.	2.9	91
438	Stability Analysis Using the SAS System. <i>Agronomy Journal</i> , 1999, 91, 154-160.	1.8	116
439	Analysing disease incidence data from designed experiments by generalized linear mixed models. <i>Plant Pathology</i> , 1999, 48, 668-674.	2.4	42
440	Fitting A Regression Model for Genotype-By-Environment Data on Heading Dates in Grasses by Methods for Nonlinear Mixed Models. <i>Biometrics</i> , 1999, 55, 1120-1128.	1.4	9
441	Methods for Comparing the Yield Stability of Cropping Systems. <i>Journal of Agronomy and Crop Science</i> , 1998, 180, 193-213.	3.5	221
442	Empirical best linear unbiased prediction in cultivar trials using factor-analytic variance-covariance structures. <i>Theoretical and Applied Genetics</i> , 1998, 97, 195-201.	3.6	152
443	An algorithm for fitting the shefted multiplicative model. <i>Journal of Statistical Computation and Simulation</i> , 1998, 62, 29-43.	1.2	0
444	Predicting Cultivar Differences Using Covariates. <i>Journal of Agricultural, Biological, and Environmental Statistics</i> , 1998, 3, 151.	1.4	23
445	Distribution-Free Tests for One-Way Homoscedasticity in a Two-Way Classification. <i>Biometrics</i> , 1997, 53, 340.	1.4	9
446	Analyzing Genotype-Environment Data by Mixed Models with Multiplicative Terms. <i>Biometrics</i> , 1997, 53, 761.	1.4	170
447	Analysis of a Randomized Block Design with Unequal Subclass Numbers. <i>Agronomy Journal</i> , 1997, 89, 718-723.	1.8	14
448	Modelling expectation and variance for genotype by environment data. <i>Heredity</i> , 1997, 79, 162-171.	2.6	58
449	Modelling expectation and variance for genotype by environment data. <i>Heredity</i> , 1997, 79, 162-171.	2.6	7
450	A Monte Carlo Test for Variance Homogeneity in Linear Models. <i>Biometrical Journal</i> , 1996, 38, 461-473.	1.0	6

#	ARTICLE	IF	CITATIONS
451	Weighted estimates of interlaboratory consensus values. Computational Statistics and Data Analysis, 1996, 22, 471-479.	1.2	3
452	Comparing cultivar means in multilocation trials when the covariance structure is not circular. Heredity, 1996, 76, 198-203.	2.6	6
453	A Simplified Procedure for Comparing the Stability of Cropping Systems. Biometrics, 1996, 52, 315.	1.4	15
454	A Robust Test for Homoscedasticity in a Two-Way Layout. Biometrical Journal, 1995, 37, 151-160.	1.0	5
455	A simple procedure for yield component analysis. Euphytica, 1995, 84, 43-48.	1.2	33
456	Robustness of statistical tests for multiplicative terms in the additive main effects and multiplicative interaction model for cultivar trials. Theoretical and Applied Genetics, 1995, 90, 438-443.	3.6	74
457	Implication of correlations among some common stability statistics " a Monte Carlo simulations. Theoretical and Applied Genetics, 1995, 90, 457-461.	3.6	4
458	The use of multilocation trials to select cultivars that are better than a control. Plant Breeding, 1995, 114, 337-340.	1.9	10
459	Assessing Cultivar Adaptability by Multiple Comparison with the Best. Agronomy Journal, 1995, 87, 1225-1227.	1.8	5
460	Detecting and handling heteroscedasticity in yield trial data. Communications in Statistics Part B: Simulation and Computation, 1995, 24, 243-274.	1.2	8
461	Remark on a Desirability Index for Selecting Genotypes. Crop Science, 1995, 35, 1498-1500.	1.8	0
462	Partitioning Genotype×Environmental Interaction in Regional Yield Trials via A Generalized Stability Variance. Crop Science, 1994, 34, 1682-1685.	1.8	6
463	C410. a distribution-free test for homoscedasticity in a two-way layout. Journal of Statistical Computation and Simulation, 1994, 49, 223-225.	1.2	4
464	Best Linear Unbiased Prediction (BLUP) for regional yield trials: a comparison to additive main effects and multiplicative interaction (AMMI) analysis. Theoretical and Applied Genetics, 1994, 89, 647-654.	3.6	119
465	Relationships Between Kendall's Coefficient of Concordance and a Nonparametric Measure of Phenotypic Stability with Implications for the Consistency in Rankings as Affected by Variance Components. Biometrical Journal, 1994, 36, 719-727.	1.0	10
466	Application of a generalized Grubbs' model in the analysis of genotype-environment interaction. Heredity, 1994, 73, 113-116.	2.6	4
467	Missing observations in the analysis of stability. Heredity, 1994, 72, 141-145.	2.6	11
468	ON TESTS FOR INTERACTION IN A NONREPLICATED TWO×WAY LAYOUT. The Australian Journal of Statistics, 1994, 36, 363-369.	0.2	9

#	ARTICLE	IF	CITATIONS
469	A Comparison of the Ecovalence and the Variance of Relative Yield as Measures of Stability. Journal of Agronomy and Crop Science, 1994, 173, 1-4.	3.5	9
470	Relationships between genotype x environment interactions and rank orders for a set of genotypes tested in different environments. Theoretical and Applied Genetics, 1993, 86, 943-950.	3.6	13
471	Note on bias in estimates of the regression coefficient in the analysis of genotype-environmental interaction. Heredity, 1993, 70, 98-100.	2.6	3
472	Rank correlation among parametric and nonparametric measures of phenotypic stability. Euphytica, 1992, 64, 221-225.	1.2	33
473	Linear Regression Techniques. Assa, Cssa and Sssa, 0, , 107-176.	0.6	3
474	Assessing the between-country genetic correlation in maize yield using German and Polish official variety trials. Theoretical and Applied Genetics, 0, , .	3.6	0