

Stephen I Wright

List of Publications by Year in descending order

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Version: 2024-02-01

127
papers

13,355
citations

31976

53
h-index

25787

108
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173
all docs

173
docs citations

173
times ranked

12987
citing authors

#	ARTICLE	IF	CITATIONS
1	Selective ancestral sorting and de novo evolution in the agricultural invasion of <i>Amaranthus tuberculatus</i> . <i>Evolution; International Journal of Organic Evolution</i> , 2022, 76, 70-85.	2.3	8
2	Repeated origins, widespread gene flow, and allelic interactions of target-site herbicide resistance mutations. <i>ELife</i> , 2022, 11, .	6.0	11
3	Recombination landscape dimorphism and sex chromosome evolution in the dioecious plant <i>Rumex hastatulus</i> . <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2022, 377, 20210226.	4.0	9
4	The relative role of plasticity and demographic history in <i>Capsella bursa-pastoris</i> : a common garden experiment in Asia and Europe. <i>AoB PLANTS</i> , 2022, 14, .	2.3	4
5	Effects of the neo-X chromosome on genomic signatures of hybridization in <i>Rumex hastatulus</i> . <i>Molecular Ecology</i> , 2022, 31, 3708-3721.	3.9	3
6	Widespread Recombination Suppression Facilitates Plant Sex Chromosome Evolution. <i>Molecular Biology and Evolution</i> , 2021, 38, 1018-1030.	8.9	42
7	Convergent Adaptation to Quantitative Host Resistance in a Major Plant Pathogen. <i>MBio</i> , 2021, 12, .	4.1	4
8	Deborah Charlesworth, winner of the society for the study of evolution's inaugural lifetime achievement award: Evolutionary biology for the genomics era. <i>Evolution; International Journal of Organic Evolution</i> , 2021, 75, 566-568.	2.3	0
9	The genetic architecture and population genomic signatures of glyphosate resistance in <i>Amaranthus tuberculatus</i> . <i>Molecular Ecology</i> , 2021, 30, 5373-5389.	3.9	22
10	New genomic resources and comparative analyses reveal differences in floral gene expression in selfing and outcrossing <i>Collinsia</i> sister species. <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	1.8	8
11	Patterns and Causes of Signed Linkage Disequilibria in Flies and Plants. <i>Molecular Biology and Evolution</i> , 2021, 38, 4310-4321.	8.9	12
12	Interacting evolutionary pressures drive mutation dynamics and health outcomes in aging blood. <i>Nature Communications</i> , 2021, 12, 4921.	12.8	11
13	Selection on Accessible Chromatin Regions in <i>Capsella grandiflora</i> . <i>Molecular Biology and Evolution</i> , 2021, 38, 5563-5575.	8.9	6
14	Ancestral and neo-Y sex chromosomes contribute to population divergence in a dioecious plant. <i>Evolution; International Journal of Organic Evolution</i> , 2020, 74, 256-269.	2.3	17
15	Evolutionary Genomics of Plant Gametophytic Selection. <i>Plant Communications</i> , 2020, 1, 100115.	7.7	28
16	Chromosome Evolution: Infectious Sex Chromosomes in the African Monarch Butterfly. <i>Current Biology</i> , 2020, 30, R657-R659.	3.9	0
17	The Evolutionary Forces Shaping Cis- and Trans-Regulation of Gene Expression within a Population of Outcrossing Plants. <i>Molecular Biology and Evolution</i> , 2020, 37, 2386-2393.	8.9	13
18	Estimation of the SNP Mutation Rate in Two Vegetatively Propagating Species of Duckweed. <i>G3: Genes, Genomes, Genetics</i> , 2020, 10, 4191-4200.	1.8	19

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19	Opposing Evolutionary Pressures Drive Clonal Evolution and Health Outcomes in the Aging Blood System. <i>Blood</i> , 2020, 136, 37-37.	1.4	0
20	Population genomics of the facultatively asexual duckweed <i>Spirodela polyrhiza</i> . <i>New Phytologist</i> , 2019, 224, 1361-1371.	7.3	43
21	Hybridization and a loss of sex shape genome-wide diversity and the origin of species in the evening primroses (<i>Oenothera</i> , Onagraceae). <i>New Phytologist</i> , 2019, 224, 1372-1380.	7.3	16
22	Towards the new normal: Transcriptomic convergence and genomic legacy of the two subgenomes of an allopolyploid weed (<i>Capsella bursa-pastoris</i>). <i>PLoS Genetics</i> , 2019, 15, e1008131.	3.5	27
23	Transposable Elements Are Important Contributors to Standing Variation in Gene Expression in <i>Capsella Grandiflora</i> . <i>Molecular Biology and Evolution</i> , 2019, 36, 1734-1745.	8.9	34
24	Parental legacy, demography, and admixture influenced the evolution of the two subgenomes of the tetraploid <i>Capsella bursa-pastoris</i> (Brassicaceae). <i>PLoS Genetics</i> , 2019, 15, e1007949.	3.5	42
25	Multiple modes of convergent adaptation in the spread of glyphosate-resistant <i>Amaranthus tuberculatus</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 21076-21084.	7.1	98
26	Long-term balancing selection drives evolution of immunity genes in <i>Capsella</i> . <i>ELife</i> , 2019, 8, .	6.0	69
27	Genome-wide nucleotide diversity and associations with geography, ploidy level and glucosinolate profiles in <i>Aethionema arabicum</i> (Brassicaceae). <i>Plant Systematics and Evolution</i> , 2018, 304, 619-630.	0.9	13
28	Population Genomics of Herbicide Resistance: Adaptation via Evolutionary Rescue. <i>Annual Review of Plant Biology</i> , 2018, 69, 611-635.	18.7	80
29	A less selfish view of genome size evolution in maize. <i>PLoS Genetics</i> , 2018, 14, e1007249.	3.5	4
30	The effects of haploid selection on Y chromosome evolution in two closely related dioecious plants. <i>Evolution Letters</i> , 2018, 2, 368-377.	3.3	26
31	Coalescence and Linkage Disequilibrium in Facultatively Sexual Diploids. <i>Genetics</i> , 2018, 210, 683-701.	2.9	23
32	Homeologue-specific expression divergence in the recently formed tetraploid <i>Capsella bursa-pastoris</i> (Brassicaceae). <i>New Phytologist</i> , 2018, 220, 624-635.	7.3	10
33	The problem of estimating recent genetic connectivity in a changing world. <i>Conservation Biology</i> , 2017, 31, 126-135.	4.7	48
34	What can genome-wide association studies tell us about the evolutionary forces maintaining genetic variation for quantitative traits?. <i>New Phytologist</i> , 2017, 214, 21-33.	7.3	75
35	Purifying and Positive Selection Influence Patterns of Gene Loss and Gene Expression in the Evolution of a Plant Sex Chromosome System. <i>Molecular Biology and Evolution</i> , 2017, 34, 1140-1154.	8.9	50
36	The genetic architecture of tristylly and its breakdown to self-fertilization. <i>Molecular Ecology</i> , 2017, 26, 752-765.	3.9	9

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37	The Relationship between Selection, Network Connectivity, and Regulatory Variation within a Population of <i>Capsella grandiflora</i> . <i>Genome Biology and Evolution</i> , 2017, 9, 1099-1109.	2.5	41
38	Hill-Robertson Interference Reduces Genetic Diversity on a Young Plant Y-Chromosome. <i>Genetics</i> , 2017, 207, 685-695.	2.9	30
39	Coevolution between transposable elements and recombination. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2017, 372, 20160458.	4.0	214
40	Genomic Loss and Silencing on the Y Chromosomes of <i>Rumex</i> . <i>Genome Biology and Evolution</i> , 2017, 9, 3345-3355.	2.5	16
41	Mutation Accumulation in an Asexual Relative of <i>Arabidopsis</i> . <i>PLoS Genetics</i> , 2017, 13, e1006550.	3.5	54
42	The influence of population structure on gene expression and flowering time variation in the ubiquitous weed <i>Capsella bursa-pastoris</i> (Brassicaceae). <i>Molecular Ecology</i> , 2016, 25, 1106-1121.	3.9	36
43	Spontaneous Chloroplast Mutants Mostly Occur by Replication Slippage and Show a Biased Pattern in the Plastome of <i>Oenothera</i> . <i>Plant Cell</i> , 2016, 28, 911-929.	6.6	49
44	Recent mating system evolution in <i>Eichhornia</i> is accompanied by cis-regulatory divergence. <i>New Phytologist</i> , 2016, 211, 697-707.	7.3	7
45	Charlesworth <i>et al.</i> on Background Selection and Neutral Diversity. <i>Genetics</i> , 2016, 204, 829-832.	2.9	2
46	Repeated Inactivation of the First Committed Enzyme Underlies the Loss of Benzaldehyde Emission after the Selfing Transition in <i>Capsella</i> . <i>Current Biology</i> , 2016, 26, 3313-3319.	3.9	43
47	Transposable element evolution in the allotetraploid <i>Capsella bursa-pastoris</i> . <i>American Journal of Botany</i> , 2016, 103, 1197-1202.	1.7	27
48	Standing genetic variation in a tissue-specific enhancer underlies selfing-syndrome evolution in <i>Capsella</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 13911-13916.	7.1	50
49	Genomic signature of successful colonization of Eurasia by the allopolyploid shepherd's purse (<i>Capsella bursa-pastoris</i>). <i>Molecular Ecology</i> , 2016, 25, 616-629.	3.9	48
50	The Limits of Natural Selection in a Nonequilibrium World. <i>Trends in Genetics</i> , 2016, 32, 201-210.	6.7	98
51	Coalescent Times and Patterns of Genetic Diversity in Species with Facultative Sex: Effects of Gene Conversion, Population Structure, and Heterogeneity. <i>Genetics</i> , 2016, 202, 297-312.	2.9	35
52	On the Trail of Linked Selection. <i>PLoS Genetics</i> , 2016, 12, e1006240.	3.5	4
53	No evidence that sex and transposable elements drive genome size variation in evening primroses. <i>Evolution; International Journal of Organic Evolution</i> , 2015, 69, 1053-1062.	2.3	40
54	The Evolution of Selfing Is Accompanied by Reduced Efficacy of Selection and Purging of Deleterious Mutations. <i>Genetics</i> , 2015, 199, 817-829.	2.9	100

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55	Recurrent Loss of Sex Is Associated with Accumulation of Deleterious Mutations in <i>Oenothera</i> . <i>Molecular Biology and Evolution</i> , 2015, 32, 896-905.	8.9	82
56	Hybrid origins and the earliest stages of diploidization in the highly successful recent polyploid <i>Capsella bursa-pastoris</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 2806-2811.	7.1	128
57	The butterfly plant arms-race escalated by gene and genome duplications. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 8362-8366.	7.1	458
58	Selfish genetic elements and plant genome size evolution. <i>Trends in Plant Science</i> , 2015, 20, 195-196.	8.8	16
59	Divergent sorting of a balanced ancestral polymorphism underlies the establishment of gene-flow barriers in <i>Capsella</i> . <i>Nature Communications</i> , 2015, 6, 7960.	12.8	81
60	Association mapping reveals the role of purifying selection in the maintenance of genomic variation in gene expression. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 15390-15395.	7.1	92
61	Genome-Wide Patterns of Genetic Variation within and among Alternative Selective Regimes. <i>PLoS Genetics</i> , 2014, 10, e1004527.	3.5	49
62	Evidence for Widespread Positive and Negative Selection in Coding and Conserved Noncoding Regions of <i>Capsella grandiflora</i> . <i>PLoS Genetics</i> , 2014, 10, e1004622.	3.5	128
63	MUTATIONISM 2.0: VIEWING EVOLUTION THROUGH MUTATION'S LENS. <i>Evolution; International Journal of Organic Evolution</i> , 2014, 68, 1225-1227.	2.3	0
64	Limited genomic consequences of mixed mating in the recently derived sister species pair, <i>C. ollinsia concolor</i> and <i>C. ollinsia parryi</i> . <i>Journal of Evolutionary Biology</i> , 2014, 27, 1400-1412.	1.7	9
65	Chromosomal Distribution of Cytonuclear Genes in a Dioecious Plant with Sex Chromosomes. <i>Genome Biology and Evolution</i> , 2014, 6, 2439-2443.	2.5	9
66	Genetic degeneration of old and young Y chromosomes in the flowering plant <i>Rumex hastatulus</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 7713-7718.	7.1	120
67	Mating system shifts and transposable element evolution in the plant genus <i>Capsella</i> . <i>BMC Genomics</i> , 2014, 15, 602.	2.8	61
68	The demography and population genomics of evolutionary transitions to self-fertilization in plants. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2014, 369, 20130344.	4.0	86
69	Repeated Evolutionary Changes of Leaf Morphology Caused by Mutations to a Homeobox Gene. <i>Current Biology</i> , 2014, 24, 1880-1886.	3.9	105
70	An atlas of over 90,000 conserved noncoding sequences provides insight into crucifer regulatory regions. <i>Nature Genetics</i> , 2013, 45, 891-898.	21.4	350
71	Patterns of Selection in Plant Genomes. <i>Annual Review of Ecology, Evolution, and Systematics</i> , 2013, 44, 31-49.	8.3	53
72	The <i>Capsella rubella</i> genome and the genomic consequences of rapid mating system evolution. <i>Nature Genetics</i> , 2013, 45, 831-835.	21.4	374

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73	Genomic Identification of Founding Haplotypes Reveals the History of the Selfing Species <i>Capsella rubella</i> . <i>PLoS Genetics</i> , 2013, 9, e1003754.	3.5	86
74	Evolutionary consequences of self-fertilization in plants. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2013, 280, 20130133.	2.6	346
75	Pollen-Specific, but Not Sperm-Specific, Genes Show Stronger Purifying Selection and Higher Rates of Positive Selection Than Sporophytic Genes in <i>Capsella grandiflora</i> . <i>Molecular Biology and Evolution</i> , 2013, 30, 2475-2486.	8.9	90
76	Impact of Sampling Schemes on Demographic Inference: An Empirical Study in Two Species with Different Mating Systems and Demographic Histories. <i>G3: Genes, Genomes, Genetics</i> , 2012, 2, 803-814.	1.8	10
77	The <i>ARC1</i> E3 Ligase Gene Is Frequently Deleted in Self-Compatible Brassicaceae Species and Has a Conserved Role in <i>Arabidopsis lyrata</i> Self-Pollen Rejection. <i>Plant Cell</i> , 2012, 24, 4607-4620.	6.6	94
78	Signatures of balancing selection are maintained at disease resistance loci following mating system evolution and a population bottleneck in the genus <i>Capsella</i> . <i>BMC Evolutionary Biology</i> , 2012, 12, 152.	3.2	32
79	COMPARATIVE POPULATION GENOMICS IN COLLINSIA SISTER SPECIES REVEALS EVIDENCE FOR REDUCED EFFECTIVE POPULATION SIZE, RELAXED SELECTION, AND EVOLUTION OF BIASED GENE CONVERSION WITH AN ONGOING MATING SYSTEM SHIFT. <i>Evolution; International Journal of Organic Evolution</i> , 2012, 67, no-no.	2.3	36
80	Coalescent-Based Analysis Distinguishes between Allo- and Autopolyploid Origin in Shepherd's Purse (<i>Capsella bursa-pastoris</i>). <i>Molecular Biology and Evolution</i> , 2012, 29, 1721-1733.	8.9	29
81	Recipient of the 2011 <i>Molecular Ecology</i> Prize: Deborah Charlesworth. <i>Molecular Ecology</i> , 2012, 21, 23-25.	3.9	1
82	GENETIC ARCHITECTURE AND ADAPTIVE SIGNIFICANCE OF THE SELFING SYNDROME IN <i>CAPSELLA</i> . <i>Evolution; International Journal of Organic Evolution</i> , 2012, 66, 1360-1374.	2.3	79
83	Does Speciation between <i>Arabidopsis halleri</i> and <i>Arabidopsis lyrata</i> Coincide with Major Changes in a Molecular Target of Adaptation?. <i>PLoS ONE</i> , 2011, 6, e26872.	2.5	87
84	Reduced Efficacy of Natural Selection on Codon Usage Bias in Selfing <i>Arabidopsis</i> and <i>Capsella</i> Species. <i>Genome Biology and Evolution</i> , 2011, 3, 868-880.	2.5	85
85	Co-evolution between transposable elements and their hosts: a major factor in genome size evolution?. <i>Chromosome Research</i> , 2011, 19, 777-786.	2.2	77
86	Genomic Determinants of Protein Evolution and Polymorphism in <i>Arabidopsis</i> . <i>Genome Biology and Evolution</i> , 2011, 3, 1210-1219.	2.5	98
87	Sizing up <i>Arabidopsis</i> genome evolution. <i>Heredity</i> , 2011, 107, 509-510.	2.6	13
88	The population genomics of plant adaptation. <i>New Phytologist</i> , 2010, 188, 313-332.	7.3	105
89	RECONSTRUCTING ORIGINS OF LOSS OF SELF-INCOMPATIBILITY AND SELFING IN NORTH AMERICAN <i>ARABIDOPSIS LYRATA</i> : A POPULATION GENETIC CONTEXT. <i>Evolution; International Journal of Organic Evolution</i> , 2010, 64, 3495-3510.	2.3	101
90	Sex-Linked Inheritance in Macaque Monkeys: Implications for Effective Population Size and Dispersal to Sulawesi. <i>Genetics</i> , 2010, 185, 923-937.	2.9	19

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91	Genome-Wide Evidence for Efficient Positive and Purifying Selection in <i>Capsella grandiflora</i> , a Plant Species with a Large Effective Population Size. <i>Molecular Biology and Evolution</i> , 2010, 27, 1813-1821.	8.9	153
92	The Long-Term Benefits of Self-Rejection. <i>Science</i> , 2010, 330, 459-460.	12.6	27
93	Mating-System Variation, Demographic History and Patterns of Nucleotide Diversity in the Tristylous Plant <i>Eichhornia paniculata</i> . <i>Genetics</i> , 2010, 184, 381-392.	2.9	79
94	Using DNA Sequence Diversity to Test for Selection in <i>Silene</i> . <i>International Journal of Plant Sciences</i> , 2010, 171, 1072-1082.	1.3	2
95	Signature of Diversifying Selection on Members of the Pentatricopeptide Repeat Protein Family in <i>Arabidopsis lyrata</i> . <i>Genetics</i> , 2009, 183, 663-672.	2.9	15
96	Recent speciation associated with the evolution of selfing in <i>Capsella</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 5241-5245.	7.1	245
97	Analysis of Site Frequency Spectra from <i>Arabidopsis</i> with Context-Dependent Corrections for Ancestral Misinference. <i>Plant Physiology</i> , 2009, 149, 616-624.	4.8	8
98	Claudin-8 and -27 tight junction proteins in puffer fish <i>Tetraodon nigroviridis</i> acclimated to freshwater and seawater. <i>Journal of Comparative Physiology B: Biochemical, Systemic, and Environmental Physiology</i> , 2009, 179, 419-431.	1.5	42
99	Conditional neutrality at two adjacent NBS-LRR disease resistance loci in natural populations of <i>Arabidopsis lyrata</i> . <i>Molecular Ecology</i> , 2008, 17, 4953-4962.	3.9	19
100	The Impact of Natural Selection on the Genome: Emerging Patterns in <i>Drosophila</i> and <i>Arabidopsis</i> . <i>Annual Review of Ecology, Evolution, and Systematics</i> , 2008, 39, 193-213.	8.3	97
101	Genomic Consequences of Outcrossing and Selfing in Plants. <i>International Journal of Plant Sciences</i> , 2008, 169, 105-118.	1.3	198
102	Claudin-3 tight junction proteins in <i>Tetraodon nigroviridis</i> : cloning, tissue-specific expression, and a role in hydromineral balance. <i>American Journal of Physiology - Regulatory Integrative and Comparative Physiology</i> , 2008, 294, R1638-R1647.	1.8	59
103	High DNA Sequence Diversity in Pericentromeric Genes of the Plant <i>Arabidopsis lyrata</i> . <i>Genetics</i> , 2008, 179, 985-995.	2.9	22
104	Selection on Amino Acid Substitutions in <i>Arabidopsis</i> . <i>Molecular Biology and Evolution</i> , 2008, 25, 1375-1383.	8.9	71
105	Contrasting Patterns of Transposable-Element Insertion Polymorphism and Nucleotide Diversity in Autotetraploid and Allotetraploid <i>Arabidopsis</i> Species. <i>Genetics</i> , 2008, 179, 581-592.	2.9	29
106	Effective population size and tests of neutrality at cytoplasmic genes in <i>Arabidopsis</i> . <i>Genetical Research</i> , 2008, 90, 119-128.	0.9	31
107	Patterns of Polymorphism and Demographic History in Natural Populations of <i>Arabidopsis lyrata</i> . <i>PLoS ONE</i> , 2008, 3, e2411.	2.5	163
108	Genomic Screening for Artificial Selection during Domestication and Improvement in Maize. <i>Annals of Botany</i> , 2007, 100, 967-973.	2.9	70

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109	Recombination: an underappreciated factor in the evolution of plant genomes. <i>Nature Reviews Genetics</i> , 2007, 8, 77-84.	16.3	223
110	Neutral Evolution of Synonymous Base Composition in the Brassicaceae. <i>Journal of Molecular Evolution</i> , 2007, 64, 136-141.	1.8	21
111	Testing for Effects of Recombination Rate on Nucleotide Diversity in Natural Populations of <i>Arabidopsis lyrata</i> . <i>Genetics</i> , 2006, 174, 1421-1430.	2.9	64
112	Selective Constraints on Codon Usage of Nuclear Genes from <i>Arabidopsis thaliana</i> . <i>Molecular Biology and Evolution</i> , 2006, 24, 122-129.	8.9	43
113	The map-based sequence of the rice genome. <i>Nature</i> , 2005, 436, 793-800.	27.8	3,365
114	Molecular Population Genetics and the Search for Adaptive Evolution in Plants. <i>Molecular Biology and Evolution</i> , 2005, 22, 506-519.	8.9	301
115	The Effects of Artificial Selection on the Maize Genome. <i>Science</i> , 2005, 308, 1310-1314.	12.6	742
116	Effects of Gene Expression on Molecular Evolution in <i>Arabidopsis thaliana</i> and <i>Arabidopsis lyrata</i> . <i>Molecular Biology and Evolution</i> , 2004, 21, 1719-1726.	8.9	132
117	The HKA Test Revisited. <i>Genetics</i> , 2004, 168, 1071-1076.	2.9	179
118	Subdivision and haplotype structure in natural populations of <i>Arabidopsis lyrata</i> . <i>Molecular Ecology</i> , 2003, 12, 1247-1263.	3.9	131
119	Effects of Recombination Rate and Gene Density on Transposable Element Distributions in <i>Arabidopsis thaliana</i> . <i>Genome Research</i> , 2003, 13, 1897-1903.	5.5	186
120	Rates and Patterns of Molecular Evolution in Inbred and Outbred <i>Arabidopsis</i> . <i>Molecular Biology and Evolution</i> , 2002, 19, 1407-1420.	8.9	180
121	Breeding systems and genome evolution. <i>Current Opinion in Genetics and Development</i> , 2001, 11, 685-690.	3.3	329
122	Genome evolution: Sex and the transposable element. <i>Current Biology</i> , 2001, 11, R296-R299.	3.9	89
123	Population Dynamics of an <i>Ac</i> -like Transposable Element in Self- and Cross-Pollinating <i>Arabidopsis</i> . <i>Genetics</i> , 2001, 158, 1279-1288.	2.9	116
124	Transposon diversity in <i>Arabidopsis thaliana</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 7376-7381.	7.1	232
125	<i>Mutator</i> -like Elements in <i>Arabidopsis thaliana</i> : Structure, Diversity and Evolution. <i>Genetics</i> , 2000, 156, 2019-2031.	2.9	106
126	Size-dependent gender modification in a hermaphroditic perennial herb. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 1999, 266, 225-232.	2.6	62

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127	Transposon dynamics and the breeding system. , 1999, 107, 139-148.		107