Ian Godwin

List of Publications by Year
in descending order

145 papers	6,074 citations	87888 38 h-index	88630 70 g-index
150 all docs	150 docs citations	150 times ranked	6226 citing authors

#	Article	IF	Citations
1	Beyond the gene: epigenetic and cis-regulatory targets offer new breeding potential for the future. Current Opinion in Biotechnology, 2022, 73, 88-94.	6.6	13
2	Epigenome guided crop improvement: current progress and future opportunities. Emerging Topics in Life Sciences, 2022, 6, 141-151.	2.6	5
3	Mining the Vavilov wheat diversity panel for new sources of adult plant resistance to stripe rust. Theoretical and Applied Genetics, 2022, 135, 1355-1373.	3.6	6
4	Endogenous U6 promoters improve CRISPR/Cas9 editing efficiencies in Sorghum bicolor and show potential for applications in other cereals. Plant Cell Reports, 2022, 41, 489-492.	5.6	16
5	Quantitative trait loci (QTL) for low temperature tolerance at the young microspore stage in rice (<i>Oryza sativa</i> L.) in Australian breeding material. Breeding Science, 2022, , .	1.9	1
6	Sorghum. , 2021, , 196-221.		9
7	Tunable crops are just a spray away. Nature Plants, 2021, 7, 102-103.	9.3	4
8	Stem vacuole-targetted sucrose isomerase enhances sugar content in sorghum. Biotechnology for Biofuels, 2021, 14, 53.	6.2	2
9	Current status and prospects of plant genome editing in Australia. In Vitro Cellular and Developmental Biology - Plant, 2021, 57, 574-583.	2.1	8
10	Genetic characterization of adult-plant resistance to tan spot (syn, yellow spot) in wheat. Theoretical and Applied Genetics, 2021, 134, 2823-2839.	3.6	8
11	Manipulating assimilate availability provides insight into the genes controlling grain size in sorghum. Plant Journal, 2021, 108, 231-243.	5.7	9
12	Sorghum as a novel biomass for the sustainable production of cellulose nanofibers. Industrial Crops and Products, 2021, 171, 113917.	5.2	20
13	Addressing the shortfalls of sorghum as a feed grain for chicken-meat production. World's Poultry Science Journal, 2021, 77, 29-41.	3.0	4
14	Hotter, drier, CRISPR: the latest edit on climate change. Theoretical and Applied Genetics, 2021, 134, 1691-1709.	3.6	40
15	Genetic Diversity and Population Structure of Field Isolates of Ganoderma Boninense from Oil Palm Plantation in Solomon Islands. Proceedings (mdpi), 2020, 36, .	0.2	1
16	How Do Crops Balance Water Supply and Demand when Water Is Limiting?. Proceedings (mdpi), 2020, 36, .	0.2	0
17	Trends in the production of cellulose nanofibers from non-wood sources. Cellulose, 2020, 27, 575-593.	4.9	151
18	Largeâ€scale GWAS in sorghum reveals common genetic control of grain size among cereals. Plant Biotechnology Journal, 2020, 18, 1093-1105.	8.3	72

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19	Ganoderma infection of oil palm â€“ a persistent problem in Papua New Guinea and Solomon Islands. Australasian Plant Pathology, 2020, 49, 69-77.	1.0	9
20	Tracking seasonal changes in diversity of pollen allergen exposure: Targeted metabarcoding of a subtropical aerobiome. Science of the Total Environment, 2020, 747, 141189.	8.0	23
21	Genomic interventions for sustainable agriculture. Plant Biotechnology Journal, 2020, 18, 2388-2405.	8.3	71
22	Transgenic expression of flavanone 3â€hydroxylase redirects flavonoid biosynthesis and alleviates anthracnose susceptibility in sorghum. Plant Biotechnology Journal, 2020, 18, 2170-2172.	8.3	30
23	Biolistic DNA Delivery and Its Applications in Sorghum bicolor. Methods in Molecular Biology, 2020, 2124, 197-215.	0.9	4
24	Genetic variation for leaf carbon isotope discrimination and its association with transpiration efficiency in canola (Brassica napus). Functional Plant Biology, 2020, 47, 355.	2.1	3
25	Upâ€regulation of lipid biosynthesis increases the oil content in leaves of Sorghum bicolor. Plant Biotechnology Journal, 2019, 17, 220-232.	8.3	75
26	The Role of Pullulanase in Starch Biosynthesis, Structure, and Thermal Properties by Studying Sorghum with Increased Pullulanase Activity. Starch/Staerke, 2019, 71, 1900072.	2.1	9
27	A SWEET solution to rice blight. Nature Biotechnology, 2019, 37, 1280-1282.	17.5	20
28	Breeding crops to feed 10 billion. Nature Biotechnology, 2019, 37, 744-754.	17.5	577
29	Technological perspectives for plant breeding. Theoretical and Applied Genetics, 2019, 132, 555-557.	3.6	27
30	Increasing protein content and digestibility in sorghum grain with a synthetic biology approach. Journal of Cereal Science, 2019, 85, 27-34.	3.7	19
31	Genome Editing by CRISPR/Cas9 in Sorghum Through Biolistic Bombardment. Methods in Molecular Biology, 2019, 1931, 169-183.	0.9	53
32	Discovering new alleles for yellow spot resistance in the Vavilov wheat collection. Theoretical and Applied Genetics, 2019, 132, 149-162.	3.6	21
33	Sterile leucaena becomes a reality?. Tropical Grasslands - Forrajes Tropicales, 2019, 7, 74-79.	0.5	4
34	Vavilov wheat accessions provide useful sources of resistance to tan spot (syn. yellow spot) of wheat. Plant Pathology, 2018, 67, 1076-1087.	2.4	15
35	VERNALIZATION1 Modulates Root System Architecture in Wheat and Barley. Molecular Plant, 2018, 11, 226-229.	8.3	118
36	Outlook: Sorghum as a feed grain for Australian chicken-meat production. Animal Nutrition, 2018, 4, 17-30.	5.1	24

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37	Applications and potential of genome editing in crop improvement. Genome Biology, 2018, 19, 210.	8.8	286
38	Effects of the growth environment on the yield and material properties of nanocellulose derived from the Australian desert grass Triodia. Industrial Crops and Products, 2018, 126, 238-249.	5.2	7
39	Highâ€throughput detection and screening of plants modified by gene editing using quantitative realâ€time polymerase chain reaction. Plant Journal, 2018, 95, 557-567.	5.7	39
40	Into the vault of the Vavilov wheats: old diversity for new alleles. Genetic Resources and Crop Evolution, 2017, 64, 531-544.	1.6	41
41	Post-anthesis nitrate uptake is critical to yield and grain protein content in Sorghum bicolor. Journal of Plant Physiology, 2017, 216, 118-124.	3.5	20
42	The wheat Lr34 multipathogen resistance gene confers resistance to anthracnose and rust in sorghum. Plant Biotechnology Journal, 2017, 15, 1387-1396.	8.3	52
43	The vegetative nitrogen response of sorghum lines containing different alleles for nitrate reductase and glutamate synthase. Molecular Breeding, 2017, 37, 1.	2.1	3
44	The functionality of Î±-kafirin promoter and Î±-kafirin signal peptide. Plant Cell, Tissue and Organ Culture, 2017, 128, 133-143.	2.3	6
45	Expression patterns of the native Shrunken-2 promoter in Sorghum bicolor visualised through use of the GFP reporter gene. Plant Cell Reports, 2017, 36, 1689-1700.	5.6	4
46	Whole-Genome Analysis of Candidate genes Associated with Seed Size and Weight in Sorghum bicolor Reveals Signatures of Artificial Selection and Insights into Parallel Domestication in Cereal Crops. Frontiers in Plant Science, 2017, 8, 1237.	3.6	59
47	Whole Genome Sequencing Reveals Potential New Targets for Improving Nitrogen Uptake and Utilization in Sorghum bicolor. Frontiers in Plant Science, 2016, 7, 1544.	3.6	39
48	Domestication and the storage starch biosynthesis pathway: signatures of selection from a whole sorghum genome sequencing strategy. Plant Biotechnology Journal, 2016, 14, 2240-2253.	8.3	38
49	Rice-cold tolerance across reproductive stages. Crop and Pasture Science, 2016, 67, 823.	1.5	14
50	Genomic Approaches for Improving Grain Quality of Sorghum. Compendium of Plant Genomes, 2016, , 189-205.	0.5	4
51	Resistance to yellow spot in wheat grown under accelerated growth conditions. Euphytica, 2016, 209, 693-707.	1.2	43
52	Heterosis in locally adapted sorghum genotypes and potential of hybrids for increased productivity in contrasting environments in Ethiopia. Crop Journal, 2016, 4, 479-489.	5.2	26
53	Microsatellite Analysis of Oil Palms and their Progenies Bred in Papua New Guinea. Tropical Plant Biology, 2016, 9, 280-289.	1.9	1
54	Regional and seasonal variation in airborne grass pollen levels between cities of Australia and New Zealand. Aerobiologia, 2016, 32, 289-302.	1.7	34

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55	SorGSD: a sorghum genome SNP database. Biotechnology for Biofuels, 2016, 9, 6.	6.2	44
56	Differences in grass pollen allergen exposure across Australia. Australian and New Zealand Journal of Public Health, 2015, 39, 51-55.	1.8	42
57	Diurnal changes in Sorghum leaf starch molecular structure. Plant Science, 2015, 239, 147-154.	3.6	29
58	Characterization of the time evolution of starch structure from rice callus. Carbohydrate Polymers, 2015, 127, 116-123.	10.2	12
59	A robust tissue culture system for sorghum [Sorghum bicolor (L.) Moench]. South African Journal of Botany, 2015, 98, 157-160.	2.5	32
60	Genetic differentiation analysis for the identification of complementary parental pools for sorghum hybrid breeding in Ethiopia. Theoretical and Applied Genetics, 2015, 128, 1765-1775.	3.6	14
61	Total transcriptome, proteome, and allergome of Johnson grass pollen, which is important for allergic rhinitis in subtropical regions. Journal of Allergy and Clinical Immunology, 2015, 135, 133-142.	2.9	36
62	The plasticity of NBS resistance genes in sorghum is driven by multiple evolutionary processes. BMC Plant Biology, 2014, 14, 253.	3.6	49
63	Impacts of Kafirin Allelic Diversity, Starch Content, and Protein Digestibility on Ethanol Conversion Efficiency in Grain Sorghum. Cereal Chemistry, 2014, 91, 218-227.	2.2	24
64	Grain Sorghum Proteomics: Integrated Approach toward Characterization of Endosperm Storage Proteins in Kafirin Allelic Variants. Journal of Agricultural and Food Chemistry, 2014, 62, 9819-9831.	5.2	40
65	Sorghum Genetic Transformation by Particle Bombardment. Methods in Molecular Biology, 2014, 1099, 219-234.	0.9	24
66	The Macroecology of Airborne Pollen in Australian and New Zealand Urban Areas. PLoS ONE, 2014, 9, e97925.	2.5	58
67	Whole-genome sequencing reveals untapped genetic potential in Africaâ€™s indigenous cereal crop sorghum. Nature Communications, 2013, 4, 2320.	12.8	405
68	Allelic variation at a single gene increases food value in a drought-tolerant staple cereal. Nature Communications, 2013, 4, 1483.	12.8	41
69	Additive effects of three auxins and copper on sorghum in vitro root induction. In Vitro Cellular and Developmental Biology - Plant, 2013, 49, 191-197.	2.1	9
70	Insights into Sorghum Starch Biosynthesis from Structure Changes Induced by Different Growth Temperatures. Cereal Chemistry, 2013, 90, 223-230.	2.2	24
71	Potential of Australian bermudagrasses (Cynodon spp.) for pasture in subtropical Australia. Tropical Grasslands - Forrajes Tropicales, 2013, 1, 81.	0.5	2
72	Factors Controlling Selfâ€Fertility in Sunflower: The Role of GCA/SCA Effects, S Alleles, and Floret Characteristics. Crop Science, 2012, 52, 128-135.	1.8	5

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73	Maximizing Genetic, Morphological, and Geographic Diversity in a Core Collection of Australian Bermudagrass. Crop Science, 2012, 52, 879-889.	1.8	28
74	Expression Pattern of the Alpha-Kafirin Promoter Coupled with a Signal Peptide fromSorghum bicolorL. Moench. Journal of Biomedicine and Biotechnology, 2012, 2012, 1-8.	3.0	9
75	Phylogenetic analysis reveals multiple introductions of Cynodon species in Australia. Molecular Phylogenetics and Evolution, 2012, 65, 390-396.	2.7	21
76	Characterisation of grain quality in diverse sorghum germplasm using a Rapid Visco-Analyzer and near infrared reflectance spectroscopy. Journal of the Science of Food and Agriculture, 2012, 92, 1402-1410.	3.5	16
77	Highly efficient sorghum transformation. Plant Cell Reports, 2012, 31, 999-1007.	5.6	96
78	Rapid phenotyping for adultâ€plant resistance to stripe rust in wheat. Plant Breeding, 2012, 131, 54-61.	1.9	63
79	Occurrence of LINE, gypsy-like, and copia-like retrotransposons in the clonally propagated sweet potato (Ipomoea batatas L.). Genome, 2011, 54, 603-609.	2.0	1
80	Lack of Low Frequency Variants Masks Patterns of Non-Neutral Evolution following Domestication. PLoS ONE, 2011, 6, e23041.	2.5	17
81	Cultivar-specific effects of pathogen testing on storage root yield of sweetpotato, Ipomoea batatas. Annals of Applied Biology, 2011, 158, 288-296.	2.5	8
82	Effect of a gibberellin-biosynthesis inhibitor treatment on the physicochemical properties of sorghum starch. Journal of Cereal Science, 2011, 53, 328-334.	3.7	51
83	IRAP, a retrotransposon-based marker system for the detection of somaclonal variation in barley. Molecular Breeding, 2011, 27, 193-206.	2.1	36
84	Allelic variation of the Î²-, Î³- and Î´-kafirin genes in diverse Sorghum genotypes. Theoretical and Applied Genetics, 2010, 121, 1227-1237.	3.6	39
85	Characterization, inheritance, and molecular study of opaque leaf mutant in mungbean (Vigna radiata) Tj ETQq1 1 0.784314 rgBT /Ov	1.5	2
86	A QTL controlling low temperature induced spikelet sterility at booting stage in rice. Euphytica, 2010, 176, 291-301.	1.2	44
87	Characterization and multiplexing of ESTâ€SSR primers in Cynodon (Poaceae) species¹. American Journal of Botany, 2010, 97, e99-e101.	1.7	21
88	Development and characterization of microsatellite loci for Khaya senegalensis (Meliaceae)¹. American Journal of Botany, 2010, 97, e111-3.	1.7	15
89	Transgenic Plants for Abiotic Stress Resistance. , 2010, , 67-132.		90
90	Cold tolerance in rice varieties at different growth stages. Crop and Pasture Science, 2009, 60, 328.	1.5	69

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91	Multifunctional grains for the future: genetic engineering for enhanced and novel cereal quality. In Vitro Cellular and Developmental Biology - Plant, 2009, 45, 383-399.	2.1	10
92	Identification of QTL for sugar-related traits in a sweetÂÃ—Âgrain sorghum (Sorghum bicolor L. Moench) recombinant inbred population. Molecular Breeding, 2008, 22, 367-384.	2.1	138
93	Assessment and rationalization of genetic diversity of Papua New Guinea taro (Colocasia esculenta) using SSR DNA fingerprinting. Genetic Resources and Crop Evolution, 2008, 55, 811-822.	1.6	21
94	QTL analysis of ergot resistance in sorghum. Theoretical and Applied Genetics, 2008, 117, 369-382.	3.6	46
95	Iron-fortified parboiled rice â€“ A novel solution to high iron density in rice-based diets. Food Chemistry, 2008, 110, 390-398.	8.2	38
96	Genetic erosion and changes in distribution of sorghum (Sorghum bicolor L. (Moench)) landraces in north-eastern Ethiopia. Plant Genetic Resources: Characterisation and Utilisation, 2008, 6, 1-10.	0.8	22
97	Effect of Grain Morphology on Degree of Milling and Iron Loss in Rice. Cereal Chemistry, 2007, 84, 384-388.	2.2	26
98	Genotypic variation of iron partitioning in rice grain. Journal of the Science of Food and Agriculture, 2007, 87, 2049-2054.	3.5	38
99	An assessment of the genetic relationship between sweet and grain sorghums, within Sorghum bicolor ssp. bicolor (L.) Moench, using AFLP markers. Euphytica, 2007, 157, 161-176.	1.2	83
100	Genetic Components of Variance and the Role of Pollen Traits in Sorghum Ergot Resistance. Crop Science, 2006, 46, 2387-2395.	1.8	15
101	Rationalization of taro germplasm collections in the Pacific Island region using simple sequence repeat (SSR) markers. Plant Genetic Resources: Characterisation and Utilisation, 2006, 4, 210-220.	0.8	19
102	Comparison of identity by descent and identity by state for detecting genetic regions under selection in a sorghum pedigree breeding program. Molecular Breeding, 2005, 14, 441-454.	2.1	21
103	Molecular Characterization of a Novel Methionine-Rich Î´-Kafirin Seed Storage Protein Gene in Sorghum (Sorghum bicolorL.). Cereal Chemistry, 2005, 82, 706-710.	2.2	29
104	Identification of quantitative trait loci for resistance to two species of root-lesion nematode (Pratylenchus thornei and P. neglectus) in wheat. Australian Journal of Agricultural Research, 2005, 56, 345.	1.5	45
105	Segregation Distortion for Seed Testa Color in Mungbean (Vigna radiata L. Wilcek). Journal of Heredity, 2004, 95, 532-535.	2.4	35
106	Genetic analysis of resistance to root-lesion nematode (Pratylenchus thornei) in wheat. Plant Breeding, 2004, 123, 209-212.	1.9	32
107	Sequence diversity in the coat protein coding region of the genome RNA of Johnsongrass mosaic virus in Australia. Archives of Virology, 2004, 149, 1633-41.	2.1	5
108	Homologues of the maize rust resistance gene Rp1-D are genetically associated with a major rust resistance QTL in sorghum. Theoretical and Applied Genetics, 2004, 109, 875-883.	3.6	29

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109	Prediction of hybrid performance in grain sorghum using RFLP markers. Theoretical and Applied Genetics, 2003, 106, 559-567.	3.6	109
110	Development and characterization of polymorphic microsatellite markers in taro (Colocasia) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 702 T	2.0	41
111	A rapid PCR protocol for marker assisted detection of heterozygotes in segregating generations involving 1BL/1RS translocation and normal wheat lines. Australian Journal of Agricultural Research, 2002, 53, 931.	1.5	8
112	The investigation of optimal bombardment parameters for transient and stable transgene expression in Sorghum. In Vitro Cellular and Developmental Biology - Plant, 2001, 37, 341-348.	2.1	53
113	Genetic mapping of the Lablab purpureus genome suggests the presence of â€™cuckooâ€™ gene(s) in this species. Theoretical and Applied Genetics, 2000, 100, 866-871.	3.6	25
114	Two genetic linkage maps of mungbean using RFLP and RAPD markers. Australian Journal of Agricultural Research, 2000, 51, 415.	1.5	64
115	A New Technique for Coconut (Cocos nucifera) Germplasm Collection from Remote Sites: Culturability of Embryos Following Low-temperature Incubation. Australian Journal of Botany, 1999, 47, 69.	0.6	4
116	Title is missing!. Euphytica, 1998, 102, 1-7.	1.2	43
117	USE OF A SCAR-BASED MARKER FOR THE EARLY DETECTION OF DWARF OFF-TYPES IN MICROPROPAGATED CAVENDISH BANANAS. Acta Horticulturae, 1998, , 157-164.	0.2	11
118	CONTROL OF ETHYLENE AND USE OF POLYAMINES CAN OPTIMISE THE CONDITIONS FOR SOMATIC EMBRYOGENESIS IN COCONUT (COCOS NUCIFERA L.) AND PAPAYA (CARICA PAPAYA L.). Acta Horticulturae, 1998, , 459-466.	0.2	8
119	An efficient interspecific hybridisation protocol for Carica papaya L. Ã— C. cauliflora Jacq.. Australian Journal of Experimental Agriculture, 1998, 38, 523.	1.0	20
120	IDENTIFICATION AND CHARACTERISATION OF DWARF OFF-TYPES FROM MICROPROPAGATED CAVENDISH BANANAS. Acta Horticulturae, 1998, , 79-84.	0.2	13
121	RANDOMLY AMPLIFIED POLYMORPHIC DNA MARKERS FOR A CARICA INTERSPECIFIC HYBRID. Acta Horticulturae, 1998, , 133-140.	0.2	3
122	THE EFFECT OF EXOGENOUS POLYAMINES ON SOMATIC EMBRYOGENESIS AND PLANT REGENERATION FROM SORGHUM BICOLOR AND SACCHARUM SPP.. Acta Horticulturae, 1998, , 451-458.	0.2	3
123	Effect of ethylene and culture environment on development of papaya nodal cultures. Plant Cell, Tissue and Organ Culture, 1997, 49, 93-100.	2.3	21
124	Morphological, molecular and cytological analyses of Carica papayaÃ—C. cauliflora interspecific hybrids. Theoretical and Applied Genetics, 1997, 95, 224-229.	3.6	42
125	Screening Carica papaya â€ƒÃ—â€ƒ C. cauliflora hybrids for resistance to papaya ringspot virusâ€type P. Plant Pathology, 1997, 46, 837-841.	2.4	32
126	Application of inter simple sequence repeat (ISSR) markers to plant genetics. Electrophoresis, 1997, 18, 1524-1528.	2.4	329

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127	Genetic analysis of preharvest sprouting tolerance in three wheat crosses. Australian Journal of Agricultural Research, 1997, 48, 215.	1.5	12
128	Micropropagated dwarf off-type Cavendish bananas (Musa spp., AAA) show improved tolerance to suboptimal temperatures. Australian Journal of Agricultural Research, 1997, 48, 377.	1.5	12
129	RAPD polymorphisms among variant and phenotypically normal rice (Oryza sativa var. indica) somaclonal progenies. Plant Cell Reports, 1997, 16, 320-324.	5.6	15
130	An Improved Embryo-Rescue Protocol for a Carica Interspecific Hybrid. Australian Journal of Botany, 1996, 44, 343.	0.6	36
131	Random amplified polymorphic DNA (RAPD) detection of dwarf off-types in micropropagated Cavendish (Musa spp. AAA) bananas. Plant Cell Reports, 1996, 16, 118-123.	5.6	94
132	Comparison of DNA Marker Technologies in Characterizing Plant Genome Diversity: Variability in Chinese Sorghums. Crop Science, 1996, 36, 1669-1676.	1.8	152
133	Phylogenetic position of hoop pine (Araucaria cunninghamii). Australian Systematic Botany, 1996, 9, 893.	0.9	11
134	Gibberellic acid detection of dwarf offtypes in micropropagated Cavendish bananas. Australian Journal of Experimental Agriculture, 1996, 36, 237.	1.0	22
135	The hypersensitive reaction to bacterial canker in Mirabilis jalapa is simply inherited. Australasian Plant Pathology, 1996, 25, 64.	1.0	0
136	Random amplified polymorphic DNA (RAPD) detection of dwarf off-types in micropropagated Cavendish (Musa spp. AAA) bananas. Plant Cell Reports, 1996, 16, 118-123.	5.6	1
137	Somaclonal Variation in Rice Ìµ2 Drought Tolerance and Other Agronomic Characters. Australian Journal of Botany, 1995, 43, 201.	0.6	66
138	An Efficient in vitro Regeneration System for Australian-Grown Chickpea (Cicer arietinum) Cultivars. Australian Journal of Botany, 1995, 43, 491.	0.6	9
139	Assessment of genome origins and genetic diversity in the genusEleusinewith DNA markers. Genome, 1995, 38, 757-763.	2.0	179
140	High-Efficiency Plant Regeneration From Callus Induced on Mature Indica Rice Caryopses. Australian Journal of Botany, 1995, 43, 337.	0.6	7
141	Interpretation of randomly amplified polymorphic DNA marker data for fingerprinting sweet potato (Ipomoea batatas L.) genotypes. Theoretical and Applied Genetics, 1994, 88-88, 332-336.	3.6	40
142	In vitro Approaches to Extending the Host-Range of Agrobacterium for Plant Transformation. Australian Journal of Botany, 1992, 40, 751.	0.6	32
143	The effects of acetosyringone and pH on Agrobacterium-mediated transformation vary according to plant species. Plant Cell Reports, 1991, 9, 671-675.	5.6	168
144	Variation among somaclonal progenies from three species of Stylosanthes. Australian Journal of Agricultural Research, 1990, 41, 645.	1.5	6

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145	Plant regeneration from leaf-derived callus cultures of the tropical pasture legume Stylosanthes scabra Vog.. Plant Cell, Tissue and Organ Culture, 1987, 9, 3-8.	2.3	8