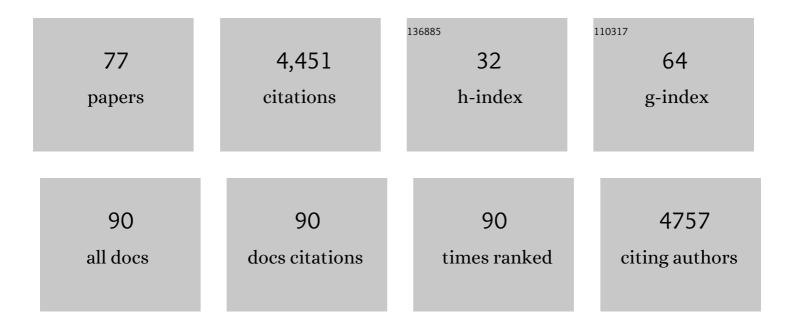
Michael James Wilkinson

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A new system of comparing PCR primers applied to ISSR fingerprinting of potato cultivars. Theoretical and Applied Genetics, 1999, 98, 107-112.	1.8	960
2	Developing an appropriate strategy to assess genetic variability in plant germplasm collections. Theoretical and Applied Genetics, 1999, 98, 1125-1131.	1.8	231
3	Selection of candidate coding DNA barcoding regions for use on land plants. Botanical Journal of the Linnean Society, 2009, 159, 1-11.	0.8	231
4	Environmental consequences of alternative practices for intensifying crop production. Agriculture, Ecosystems and Environment, 2002, 88, 279-290.	2.5	169
5	Hybridization Between Brassica napus and B. rapa on a National Scale in the United Kingdom. Science, 2003, 302, 457-459.	6.0	154
6	DNA Barcoding the Native Flowering Plants and Conifers of Wales. PLoS ONE, 2012, 7, e37945.	1.1	138
7	Low relative humidity triggers RNA-directed de novo DNA methylation and suppression of genes controlling stomatal development. Journal of Experimental Botany, 2012, 63, 3799-3813.	2.4	128
8	Risk assessment of GM plants: avoiding gridlock?. Trends in Plant Science, 2003, 8, 208-212.	4.3	110
9	Low probability of chloroplast movement from oilseed rape (Brassica napus) into wild Brassica rapa. Nature Biotechnology, 1999, 17, 390-392.	9.4	106
10	A direct regional scale estimate of transgene movement from genetically modified oilseed rape to its wild progenitors. Molecular Ecology, 2000, 9, 983-991.	2.0	106
11	Cytosine methylation regulates oviposition in the pathogenic blood fluke Schistosoma mansoni. Nature Communications, 2011, 2, 424.	5.8	103
12	A genetic linkage map of microsatellite, gene-specific and morphological markers in diploid Fragaria. Theoretical and Applied Genetics, 2004, 109, 1385-1391.	1.8	102
13	Progressive erosion of genetic and epigenetic variation in callusâ€derived cocoa (<i>Theobroma) Tj ETQq1 1 0.78</i>	4314 rgBT	Overlock 96
14	Polyploid speciation inHedera (Araliaceae): Phylogenetic and biogeographic insights based on chromosome counts and ITS sequences. Plant Systematics and Evolution, 1999, 219, 165-179.	0.3	84
15	Highâ€resolution melt analysis to identify and map sequenceâ€tagged site anchor points onto linkage maps: a white lupin (<i>Lupinus albus</i>) map as an exemplar. New Phytologist, 2008, 180, 594-607.	3.5	70
16	Transgene risk is low. Nature, 1998, 393, 320-320.	13.7	66
17	Biodiversity and biogeography of the cacao (Theobroma cacao) pathogen Moniliophthora roreri in tropical America. Plant Pathology, 2007, 56, 911-922.	1.2	66
18	Fitness of hybrids between rapeseed (<i>Brassica napus</i>) and wild <i>Brassica rapa</i> in natural habitats. Molecular Ecology, 2006, 15, 1175-1184.	2.0	65

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19	A Hypomethylated population of Brassica rapa for forward and reverse Epi-genetics. BMC Plant Biology, 2012, 12, 193.	1.6	64
20	Assembling spatially explicit landscape models of pollen and spore dispersal by wind for risk assessment. Proceedings of the Royal Society B: Biological Sciences, 2006, 273, 1705-1713.	1.2	63
21	Frosty Pod of Cacao: A Disease with a Limited Geographic Range but Unlimited Potential for Damage. Phytopathology, 2007, 97, 1644-1647.	1.1	63
22	A new taxonomic treatment of the Festuca ovina L. aggregate (Poaceae) in the British Isles. Botanical Journal of the Linnean Society, 1991, 106, 347-397.	0.8	61
23	Feulgen Staining of Intact Plant Tissues for Confocal Microscopy. Biotechnic and Histochemistry, 1996, 71, 84-87.	0.7	55
24	Epi-fingerprinting and epi-interventions for improved crop production and food quality. Frontiers in Plant Science, 2015, 6, 397.	1.7	52
25	Transgenerational, Dynamic Methylation of Stomata Genes in Response to Low Relative Humidity. International Journal of Molecular Sciences, 2013, 14, 6674-6689.	1.8	51
26	Cytological and molecular observations on Solanum phureja-induced dihaploid potatoes. Theoretical and Applied Genetics, 1991, 82, 545-551.	1.8	49
27	Evaluation of cacao (Theobroma cacao) clones against seven Colombian isolates of Moniliophthora roreri from four pathogen genetic groups. Plant Pathology, 2005, 54, 483-490.	1.2	48
28	Spontaneous gene flow from rapeseed (Brassica napus) to wild Brassica oleracea. Proceedings of the Royal Society B: Biological Sciences, 2006, 273, 3111-3115.	1.2	46
29	Replacing Sanger with Next Generation Sequencing to improve coverage and quality of reference DNA barcodes for plants. Scientific Reports, 2017, 7, 46040.	1.6	45
30	Epigenetic rather than genetic factors may explain phenotypic divergence between coastal populations of diploid and tetraploid Limonium spp. (Plumbaginaceae) in Portugal. BMC Plant Biology, 2013, 13, 205.	1.6	41
31	Meristem micropropagation of cassava (Manihot esculenta) evokes genome-wide changes in DNA methylation. Frontiers in Plant Science, 2015, 6, 590.	1.7	40
32	Direct Detection and Quantification of Methylation in Nucleic Acid Sequences Using High-Resolution Melting Analysis. Analytical Chemistry, 2010, 82, 9100-9108.	3.2	39
33	Detection and quantification of in vitro-culture induced chimerism using simple sequence repeat (SSR) analysis in Theobroma cacao (L.). Theoretical and Applied Genetics, 2004, 110, 157-166.	1.8	36
34	Differential Effect of Three Base Modifications on DNA Thermostability Revealed by High Resolution Melting. Analytical Chemistry, 2012, 84, 7336-7342.	3.2	35
35	Active and adaptive plasticity in a changing climate. Trends in Plant Science, 2022, 27, 717-728.	4.3	35
36	The Structure of Interrupted Human AC Microsatellites. Molecular Biology and Evolution, 2003, 20, 453-459.	3.5	32

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37	The first genetic maps of cashew (Anacardium occidentale L.). Euphytica, 2007, 157, 131-143.	0.6	31
38	Title is missing!. Euphytica, 2002, 124, 139-145.	0.6	30
39	Allelic Size Standards and Reference Genotypes to Unify International Cocoa (Theobroma cacao L.) Microsatellite Data. Genetic Resources and Crop Evolution, 2006, 53, 1643-1652.	0.8	30
40	Fitness and beyond: Preparing for the arrival of GM crops with ecologically important novel characters. Environmental Biosafety Research, 2009, 8, 1-14.	1.1	30
41	What's in a name: Crinipellis, the final resting place for the frosty pod rot pathogen of cocoa?. The Mycologist, 2002, 16, .	0.5	29
42	Quantitative and Qualitative Differences in Morphological Traits Revealed between Diploid Fragaria Species. Annals of Botany, 2004, 94, 787-796.	1.4	29
43	Rapeseed cytoplasm gives advantage in wild relatives and complicates genetically modified crop biocontainment. New Phytologist, 2009, 183, 1201-1211.	3.5	27
44	Characterization and PCR multiplexing of polymorphic microsatellite loci in cashew (Anacardium) Tj ETQq0 0 0 r	gBT /Overl 1.7	ock_{25} 10 Tf 50
45	Production of haploids and doubled haploids in oil palm. BMC Plant Biology, 2010, 10, 218.	1.6	25
46	Solanum phureja genes are expressed in the leaves and tubers of aneusomatic potato dihaploids. Euphytica, 1993, 69, 1-6.	0.6	23
47	Pre-conditioning the epigenetic response to high vapor pressure deficit increases the drought tolerance of <i>Arabidopsis thaliana</i> . Plant Signaling and Behavior, 2013, 8, e25974.	1.2	23
48	Salt Stress Induces Non-CG Methylation in Coding Regions of Barley Seedlings (Hordeum vulgare). Epigenomes, 2018, 2, 12.	0.8	21
49	Spontaneous capture of oilseed rape (Brassica napus) chloroplasts by wild B. rapa: implications for the use of chloroplast transformation for biocontainment. Current Genetics, 2009, 55, 139-150.	0.8	20
50	Conserving marginal populations of the food plant (Impatiens noli-tangere) of an endangered moth (Eustroma reticulatum) in a changing climate. Biological Conservation, 2004, 116, 305-317.	1.9	18
51	Detection of somaclonal variation during cocoa somatic embryogenesis characterised using cleaved amplified polymorphic sequence and the new freeware Artbio. Molecular Breeding, 2010, 25, 501-516.	1.0	18

52	Functional Genomics of Cacao. Advances in Botanical Research, 2010, 55, 119-177.	0.5	17
53	METHODOLOGICAL INSIGHTS: The role of satellite image-processing for national-scale estimates of gene flow from genetically modified crops: rapeseed in the UK as a model. Journal of Applied Ecology, 2004, 41, 1174-1184.	1.9	16
54	High throughput, high resolution selection of polymorphic microsatellite loci for multiplex analysis.	1.9	15

Plant Methods, 2005, 1, 3.

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55	Processes affecting genetic structure and conservation: a case study of wild and cultivated Brassica rapa. Genetic Resources and Crop Evolution, 2009, 56, 189-200.	0.8	15
56	Confocal observations of late-acting self-incompatibility in Theobroma cacao L. Sexual Plant Reproduction, 2012, 25, 169-183.	2.2	14
57	Evidence that genes from the male parent may influence the morphology of potato dihaploids. Theoretical and Applied Genetics, 1997, 94, 241-248.	1.8	13
58	Juvenile-mature wood transition in pine: correlation between wood properties and candidate gene expression profiles. Euphytica, 2009, 166, 341.	0.6	13
59	Adaptive divergence and speciation among sexual and pseudoviviparous populations of Festuca. Heredity, 2011, 106, 854-861.	1.2	12
60	Factors influencing the utility of gametic microprotoplasts for partial genome transfer in potato. Plant Cell Reports, 1999, 18, 786-790.	2.8	10
61	Genetic Variation in a Chilean Endangered Endemic: Gomortega keule (molina) Baillon. Biodiversity and Conservation, 2005, 14, 2871-2881.	1.2	10
62	Development and characterisation of microsatellite markers for the fungus Lasiodiplodia theobromae. Summa Phytopathologica, 2008, 34, 55-57.	0.3	9
63	Dual-locus DNA metabarcoding reveals southern hairy-nosed wombats (Lasiorhinus latifrons Owen) have a summer diet dominated by toxic invasive plants. PLoS ONE, 2020, 15, e0229390.	1.1	9
64	Botrytis cinerea Loss and Restoration of Virulence during In Vitro Culture Follows Flux in Global DNA Methylation. International Journal of Molecular Sciences, 2022, 23, 3034.	1.8	8
65	The partial stability of additional chromosomes in Solanum tuberosum cv. Torridon. Euphytica, 1992, 60, 115-122.	0.6	7
66	A simple, high throughput method to locate single copy sequences from Bacterial Artificial Chromosome (BAC) libraries using High Resolution Melt analysis. BMC Genomics, 2010, 11, 301.	1.2	7
67	BAC-HAPPY Mapping (BAP Mapping): A New and Efficient Protocol for Physical Mapping. PLoS ONE, 2010, 5, e9089.	1.1	6
68	Assessing the value of imperfect biocontainment nationally: rapeseed in the U nited K ingdom as an exemplar. New Phytologist, 2015, 205, 1342-1349.	3.5	6
69	Ranking the value of germplasm: new oil palm (<i><scp>E</scp>laeis guineensis</i>) breeding stocks as a case study. Annals of Applied Biology, 2012, 160, 145-156.	1.3	5
70	GM risk assessment: Pollen carriage from Brassica napus to B. rapa varies widely between pollinators. Basic and Applied Ecology, 2017, 19, 36-44.	1.2	5
71	The potential of aerosol eDNA sampling for the characterisation of commercial seed lots. PLoS ONE, 2018, 13, e0201617.	1.1	5
72	Greenhouse Spatial Effects Detected in the Barley (Hordeum vulgare L.) Epigenome Underlie Stochasticity of DNA Methylation. Frontiers in Plant Science, 2020, 11, 553907.	1.7	5

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73	Abandoning â€~responsive' GM risk assessment. Trends in Biotechnology, 2004, 22, 438-439.	4.9	3
74	DNA Barcoding Simplifies Environmental Risk Assessment of Genetically Modified Crops in Biodiverse Regions. PLoS ONE, 2012, 7, e35929.	1.1	3
75	Atlas of Age- and Tissue-Specific DNA Methylation during Early Development of Barley (Hordeum) Tj ETQq1 1 0.78	34314 rgB	T /Overlock
76	Caps DNA Barcoding for Field Laboratory Identification of Grass Species (British Grasses as a Model). Agriculture, 2020, 66, 74-86.	0.2	1
77	The love life of a rose. A commentary on: â€~Asymmetrical canina meiosis is accompanied by the expansion of a pericentric satellite in non-recombining univalent chromosomes'. Annals of Botany, 2020, 125, v-vi.	1.4	0