John-James Wilson

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

59 papers 1,294 citations 18 h-index g-index

66 1,605 2.8 4.7 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
59	Integration of DNA barcoding into an ongoing inventory of complex tropical biodiversity. <i>Molecular Ecology Resources</i> , 2009 , 9 Suppl s1, 1-26	8.4	268
58	DNA metabarcoding of insects and allies: an evaluation of primers and pipelines. <i>Bulletin of Entomological Research</i> , 2015 , 105, 717-27	1.7	90
57	DNA barcodes for insects. <i>Methods in Molecular Biology</i> , 2012 , 858, 17-46	1.4	66
56	Compositional heterogeneity in true bug mitochondrial phylogenomics. <i>Molecular Phylogenetics and Evolution</i> , 2018 , 118, 135-144	4.1	63
55	Factors driving changes in freshwater mussel (Bivalvia, Unionida) diversity and distribution in Peninsular Malaysia. <i>Science of the Total Environment</i> , 2016 , 571, 1069-78	10.2	60
54	Preying on commercial fisheries and accumulating paralytic shellfish toxins: a dietary analysis of invasive Dosidicus gigas (Cephalopoda Ommastrephidae) stranded in Pacific Canada. <i>Marine Biology</i> , 2012 , 159, 25-31	2.5	57
53	When species matches are unavailable are DNA barcodes correctly assigned to higher taxa? An assessment using sphingid moths. <i>BMC Ecology</i> , 2011 , 11, 18	2.7	57
52	Diagnosing mitochondrial DNA diversity: applications of a sentinel gene approach. <i>Journal of Molecular Evolution</i> , 2008 , 66, 362-7	3.1	37
51	Higher-level phylogeny and evolutionary history of Pentatomomorpha (Hemiptera: Heteroptera) inferred from mitochondrial genome sequences. <i>Systematic Entomology</i> , 2019 , 44, 810-819	3.4	35
50	Assessing the value of DNA barcodes and other priority gene regions for molecular phylogenetics of Lepidoptera. <i>PLoS ONE</i> , 2010 , 5, e10525	3.7	34
49	Building a DNA barcode reference library for the true butterflies (Lepidoptera) of Peninsula Malaysia: what about the subspecies?. <i>PLoS ONE</i> , 2013 , 8, e79969	3.7	32
48	Field calibration of blowfly-derived DNA against traditional methods for assessing mammal diversity in tropical forests. <i>Genome</i> , 2016 , 59, 1008-1022	2.4	26
47	Impact of urbanisation and agriculture on the diet of fruit bats. <i>Urban Ecosystems</i> , 2018 , 21, 61-70	2.8	22
46	Reading Mammal Diversity from Flies: The Persistence Period of Amplifiable Mammal mtDNA in Blowfly Guts (Chrysomya megacephala) and a New DNA Mini-Barcode Target. <i>PLoS ONE</i> , 2015 , 10, e012	23871	22
45	Identity of the ailanthus webworm moth (Lepidoptera, Yponomeutidae), a complex of two species: evidence from DNA barcoding, morphology and ecology. <i>ZooKeys</i> , 2010 , 46, 41-60	1.2	22
44	Mercury accumulation in bats near hydroelectric reservoirs in Peninsular Malaysia. <i>Ecotoxicology</i> , 2014 , 23, 1164-71	2.9	20
43	DNA barcoding for biosecurity: case studies from the UK plant protection program. <i>Genome</i> , 2016 , 59, 1033-1048	2.4	19

Urban parks: refuges for tropical butterflies in Southeast Asia?. Urban Ecosystems, 2016, 19, 1131-1147 2.8 18 42 DNA Barcodes and Insect Biodiversity417-431 18 41 Citizen Science: The First Peninsular Malaysia Butterfly Count. Biodiversity Data Journal, 2015, e7159 1.8 40 17 Pollination implications of the diverse diet of tropical nectar-feeding bats roosting in an urban 3.1 39 17 cave. PeerJ, 2018, 6, e4572 Application of DNA barcodes in wildlife conservation in Tropical East Asia. Conservation Biology, 6 38 17 2016. 30. 982-9 Utility of DNA barcoding for rapid and accurate assessment of bat diversity in Malaysia in the 16 1.2 37 absence of formally described species. Genetics and Molecular Research, 2014, 13, 920-5 Can butterflies cope with city life? Butterfly diversity in a young megacity in southern China. 36 15 2.4 Genome, 2016, 59, 751-61 Diversity and human perceptions of bees (Hymenoptera: Apoidea) in Southeast Asian megacities. 35 2.4 14 Genome, 2016, 59, 827-839 High-throughput terrestrial biodiversity assessments: mitochondrial metabarcoding, metagenomics or metatranscriptomics?. Mitochondrial DNA Part A: DNA Mapping, Sequencing, and 1.3 34 14 Analysis, 2019, 30, 60-67 Comparison of Butterflies, Bats and Beetles as Bioindicators Based on Four Key Criteria and DNA 33 1.4 13 Barcodes. Tropical Conservation Science, 2015, 8, 138-149 Assessing the value of DNA barcodes for molecular phylogenetics: effect of increased taxon 32 13 3.7 sampling in lepidoptera. PLoS ONE, 2011, 6, e24769 Trends in DNA barcoding and metabarcoding. Genome, 2019, 62, v-viii 2.4 A checklist of the bats of Peninsular Malaysia and progress towards a DNA barcode reference 30 3.7 11 library. PLoS ONE, 2017, 12, e0179555 DNA Barcodes and Insect Biodiversity 2017, 575-592 29 11 Mitochondrial genome of Phalantus geniculatus (Hemiptera: Reduviidae): trnT duplication and 28 7.9 10 phylogenetic implications. International Journal of Biological Macromolecules, 2019, 129, 110-115 Public Perceptions and Knowledge of, and Responses to, Bats in Urban Areas in Peninsular 27 10 2.4 Malaysia. Anthrozoos, 2019, 32, 825-834 First report of brown widow spider sightings in Peninsular Malaysia and notes on its global 26 2.2 10 distribution. Journal of Venomous Animals and Toxins Including Tropical Diseases, 2015, 21, 11 Ectoparasites of murids in peninsular Malaysia and their associated diseases. Parasites and Vectors, 8 4 **2015**, 8, 254

24	First Case Report of Canthariasis in an Infant Caused by the Larvae of Lasioderma serricorne (Coleoptera: Anobiidae). <i>Journal of Medical Entomology</i> , 2016 , 53, 1234-1237	2.2	8
23	Intestinal Myiasis in a Malaysian Patient Caused by Larvae of Clogmia albipunctatus (Diptera: Psychodidae). <i>Journal of Medical Entomology</i> , 2016 , 53, 957-960	2.2	8
22	Hookworm infections among migrant workers in Malaysia: Molecular identification of Necator americanus and Ancylostoma duodenale. <i>Acta Tropica</i> , 2017 , 173, 109-115	3.2	7
21	Towards monitoring the sandflies (Diptera: Psychodidae) of Thailand: DNA barcoding the sandflies of Wihan Cave, Uttaradit. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2016 , 27, 3795-801	1.3	7
20	DNA barcoding implicates 23 species and four orders as potential pollinators of Chinese knotweed (Persicaria chinensis) in Peninsular Malaysia. <i>Bulletin of Entomological Research</i> , 2015 , 105, 515-20	1.7	7
19	Evolution of tRNA gene rearrangement in the mitochondrial genome of ichneumonoid wasps (Hymenoptera: Ichneumonoidea). <i>International Journal of Biological Macromolecules</i> , 2020 , 164, 540-547	, 7.9	7
18	DNA barcodes and citizen science provoke a diversity reappraisal for the "ring" butterflies of Peninsular Malaysia (Ypthima: Satyrinae: Nymphalidae: Lepidoptera). <i>Genome</i> , 2016 , 59, 879-888	2.4	7
17	A two-step DNA barcoding approach for delimiting moth species: moths of Dongling Mountain (Beijing, China) as a case study. <i>Scientific Reports</i> , 2018 , 8, 14256	4.9	7
16	DNA barcodes for dragonflies and damselflies (Odonata) of Mindanao, Philippines. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2018 , 29, 206-211	1.3	6
15	Tracking the southern river terrapin (Batagur affinis) through environmental DNA: prospects and challenges. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2018 , 29, 862-866	1.3	6
14	Unexpected diversity of sandflies (Diptera: Psychodidae) in tourist caves in Northern Thailand. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2017 , 28, 949-955	1.3	6
13	Conserved gene arrangement in the mitochondrial genomes of barklouse families Stenopsocidae and Psocidae. <i>Frontiers of Agricultural Science and Engineering</i> , 2017 , 4, 358	1.7	6
12	Mitochondrial phylogeny and comparative mitogenomics of closely related pine moth pests (Lepidoptera:). <i>PeerJ</i> , 2019 , 7, e7317	3.1	6
11	Ring roads and urban biodiversity: distribution of butterflies in urban parks in Beijing city and correlations with other indicator species. <i>Scientific Reports</i> , 2019 , 9, 7653	4.9	5
10	Temporal changes in arthropod activity in tropical anthropogenic forests. <i>Bulletin of Entomological Research</i> , 2018 , 108, 792-799	1.7	5
9	Feeding behavior of Mimomyia (Etorleptiomyia) luzonensis (Ludlow, 1905) (Diptera, Culicidae) in Peninsular Malaysia. <i>Acta Tropica</i> , 2017 , 171, 138-140	3.2	4
8	Plant-herbivorous insect networks: who is eating what revealed by long barcodes using high-throughput sequencing and Trinity assembly. <i>Insect Science</i> , 2021 , 28, 127-143	3.6	3
7	DNA Barcoding: Bioinformatics Workflows for Beginners 2019 , 985-995		2

LIST OF PUBLICATIONS

6	Genetic Diversity of Pediculus humanus capitis (Phthiraptera: Pediculidae) in Peninsular Malaysia and Molecular Detection of Its Potential Associated Pathogens. <i>Journal of Medical Entomology</i> , 2020 , 57, 915-926	2.2	2
5	Using full-length metabarcoding and DNA barcoding to infer community assembly for speciose taxonomic groups: a case study. <i>Evolutionary Ecology</i> , 2020 , 34, 1063-1088	1.8	2
4	Analysis of Gene Expression in an Inbred Line of Soft-Shell Clams () Displaying Growth Heterosis: Regulation of Structural Genes and the NOD2 Pathway. <i>International Journal of Genomics</i> , 2016 , 2016, 6720947	2.5	2
3	Complete mitochondrial genome of the soft-shell clam Mya arenaria. <i>Mitochondrial DNA Part A:</i> DNA Mapping, Sequencing, and Analysis, 2016 , 27, 3553-4	1.3	1
3		0.1	1