

Markus Riegler

List of Publications by Year in descending order

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111
papers

6,685
citations

117571

34
h-index

69214

77
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114
all docs

114
docs citations

114
times ranked

6160
citing authors

#	ARTICLE	IF	CITATIONS
1	Isolation and molecular characterization of five entomopathogenic nematode species and their bacterial symbionts from eastern Australia. <i>BioControl</i> , 2022, 67, 63-74.	0.9	5
2	Substantial rearrangements, single nucleotide frameshift deletion and low diversity in mitogenome of <i>Wolbachia</i> -infected strepsipteran endoparasitoid in comparison to its tephritid hosts. <i>Scientific Reports</i> , 2022, 12, 477.	1.6	1
3	Endosymbionts moderate constrained sex allocation in a haplodiploid thrips species in a temperature-sensitive way. <i>Heredity</i> , 2022, , .	1.2	4
4	Common endosymbionts affect host fitness and sex allocation via egg size provisioning. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2022, 289, 20212582.	1.2	7
5	Virulence, penetration rate and reproductive potential of entomopathogenic nematodes from eastern Australia in Queensland fruit fly, <i>Bactrocera tryoni</i> . <i>Biological Control</i> , 2022, 169, 104871.	1.4	9
6	Two New Phoretic Species of Heterostigmatic Mites (Acari: Prostigmata: Neopygmephoridae) <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50</i>	1.0	1
7	Egg size-mediated sex allocation and mating-regulated reproductive investment in a haplodiploid thrips species. <i>Functional Ecology</i> , 2021, 35, 485-498.	1.7	7
8	Tephritid fruit flies have a large diversity of co-occurring RNA viruses. <i>Journal of Invertebrate Pathology</i> , 2021, 186, 107569.	1.5	15
9	Constrained sex allocation after mating in a haplodiploid thrips species depends on maternal condition. <i>Evolution; International Journal of Organic Evolution</i> , 2021, 75, 1525-1536.	1.1	6
10	Genome analyses of four <i>Wolbachia</i> strains and associated mitochondria of <i>Rhagoletis cerasi</i> expose cumulative modularity of cytoplasmic incompatibility factors and cytoplasmic hitchhiking across host populations. <i>BMC Genomics</i> , 2021, 22, 616.	1.2	4
11	Host-endoparasitoid-endosymbiont relationships: concealed Strepsiptera provide new twist to <i>Wolbachia</i> in Australian tephritid fruit flies. <i>Environmental Microbiology</i> , 2021, 23, 5587-5604.	1.8	7
12	Vulnerability of island insect pollinator communities to pathogens. <i>Journal of Invertebrate Pathology</i> , 2021, 186, 107670.	1.5	2
13	Major biogeographic barriers in eastern Australia have shaped the population structure of widely distributed <i>Eucalyptus moluccana</i> and its putative subspecies. <i>Ecology and Evolution</i> , 2021, 11, 14828-14842.	0.8	4
14	Sheltered life beneath elytra: three new species of <i>Eutarsopolipus</i> (Acari, Heterostigmata,) <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50</i>	0.8	2
15	Temporal changes in the microbiome of stingless bee foragers following colony relocation. <i>FEMS Microbiology Ecology</i> , 2020, 97, .	1.3	16
16	Scientific note on small hive beetle infestation of stingless bee (<i>Tetragonula carbonaria</i>) colony following a heat wave. <i>Apidologie</i> , 2020, 51, 1199-1201.	0.9	8
17	Characterization of the bacterial communities of psyllids associated with Rutaceae in Bhutan by high throughput sequencing. <i>BMC Microbiology</i> , 2020, 20, 215.	1.3	21
18	Tiny hitchhikers and parasites: a review of Australian heterostigmatic mites (Acari: Prostigmata) associated with insects, with description of three new species. <i>Austral Entomology</i> , 2020, 59, 401-421.	0.8	5

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19	Parallel Sequencing of Wolbachia wCer2 from Donor and Novel Hosts Reveals Multiple Incompatibility Factors and Genome Stability after Host Transfers. <i>Genome Biology and Evolution</i> , 2020, 12, 720-735.	1.1	14
20	The fate of carbon in a mature forest under carbon dioxide enrichment. <i>Nature</i> , 2020, 580, 227-231.	13.7	218
21	Occurrence of honey bee-associated pathogens in Varroa-free pollinator communities. <i>Journal of Invertebrate Pathology</i> , 2020, 171, 107344.	1.5	19
22	<p><p>A review of the distribution and host plant associations of the platypodine ambrosia beetles (Coleoptera: Curculionidae: Platypodinae) of Australia, with an electronic species identification key</p><p>. <i>Zootaxa</i>, 2020, 4894, 69-80.</p>	0.2	2
23	Symbiotic microbiota may reflect host adaptation by resident to invasive ant species. <i>PLoS Pathogens</i> , 2019, 15, e1007942.	2.1	27
24	Models and Nomenclature for Cytoplasmic Incompatibility: Caution over Premature Conclusions – A Response to Beckmann et al.. <i>Trends in Genetics</i> , 2019, 35, 397-399.	2.9	33
25	Additions of sugar and nitrogenous fertiliser affect plant nitrogen status and soil microbial communities. <i>Applied Soil Ecology</i> , 2019, 139, 47-55.	2.1	3
26	Diet and irradiation effects on the bacterial community composition and structure in the gut of domesticated teneral and mature Queensland fruit fly, <i>Bactrocera tryoni</i> (Diptera: Tephritidae). <i>BMC Microbiology</i> , 2019, 19, 281.	1.3	26
27	Tephritid-microbial interactions to enhance fruit fly performance in sterile insect technique programs. <i>BMC Microbiology</i> , 2019, 19, 287.	1.3	39
28	Impacts of recent climate change on terrestrial flora and fauna: Some emerging Australian examples. <i>Austral Ecology</i> , 2019, 44, 3-27.	0.7	105
29	High nymphal host density and mortality negatively impact parasitoid complex during an insect herbivore outbreak. <i>Insect Science</i> , 2019, 26, 351-365.	1.5	7
30	A temperate pollinator with high thermal tolerance is still susceptible to heat events predicted under future climate change. <i>Ecological Entomology</i> , 2018, 43, 506-512.	1.1	17
31	Insect threats to food security. <i>Science</i> , 2018, 361, 846-846.	6.0	41
32	Near full-length 16S rRNA gene next-generation sequencing revealed <i>Asaia</i> as a common midgut bacterium of wild and domesticated Queensland fruit fly larvae. <i>Microbiome</i> , 2018, 6, 85.	4.9	82
33	An ancient and a recent colonization of islands by an Australian sap-feeding insect. <i>Journal of Biogeography</i> , 2018, 45, 2389-2399.	1.4	1
34	Primary parasitoids of red scale (<i>Aonidiella aurantii</i>) in Australia and a review of their introductions from Asia. <i>Insect Science</i> , 2017, 24, 150-168.	1.5	3
35	Local coexistence and genetic isolation of three pollinator species on the same fig tree species. <i>Heredity</i> , 2017, 118, 486-490.	1.2	15
36	First confirmed report of a bacterial brood disease in stingless bees. <i>Journal of Invertebrate Pathology</i> , 2017, 144, 7-10.	1.5	19

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37	Gut symbiont enhances insecticide resistance in a significant pest, the oriental fruit fly <i>Bactrocera dorsalis</i> (Hendel). <i>Microbiome</i> , 2017, 5, 13.	4.9	318
38	Unpacking boxes: Integration of molecular, morphological and ecological approaches reveals extensive patterns of reticulate evolution in box eucalypts. <i>Molecular Phylogenetics and Evolution</i> , 2017, 108, 70-87.	1.2	20
39	Independent cytoplasmic incompatibility induced by <i>Cardinium</i> and <i>Wolbachia</i> maintains endosymbiont coinfections in haplodiploid thrips populations. <i>Evolution; International Journal of Organic Evolution</i> , 2017, 71, 995-1008.	1.1	45
40	Relative Abundance and Strain Diversity in the Bacterial Endosymbiont Community of a Sap-Feeding Insect Across Its Native and Introduced Geographic Range. <i>Microbial Ecology</i> , 2017, 74, 722-734.	1.4	13
41	Feminizing <i>Wolbachia</i> endosymbiont disrupts maternal sex chromosome inheritance in a butterfly species. <i>Evolution Letters</i> , 2017, 1, 232-244.	1.6	33
42	Symbionts in waiting: the dynamics of incipient endosymbiont complementation and replacement in minimal bacterial communities of psyllids. <i>Microbiome</i> , 2017, 5, 58.	4.9	47
43	Unravelling mummies: cryptic diversity, host specificity, trophic and coevolutionary interactions in psyllid parasitoid food webs. <i>BMC Evolutionary Biology</i> , 2017, 17, 127.	3.2	14
44	Climate and atmospheric change impacts on sap-feeding herbivores: a mechanistic explanation based on functional groups of primary metabolites. <i>Functional Ecology</i> , 2017, 31, 161-171.	1.7	21
45	Diversity and specificity of sap-feeding herbivores and their parasitoids on Australian fig trees. <i>Insect Conservation and Diversity</i> , 2017, 10, 107-119.	1.4	4
46	Belowground Ecology of Scarabs Feeding on Grass Roots: Current Knowledge and Future Directions for Management in Australasia. <i>Frontiers in Plant Science</i> , 2016, 7, 321.	1.7	45
47	Above-Belowground Herbivore Interactions in Mixed Plant Communities Are Influenced by Altered Precipitation Patterns. <i>Frontiers in Plant Science</i> , 2016, 7, 345.	1.7	33
48	Codivergence of the primary bacterial endosymbiont of psyllids versus host switches and replacement of their secondary bacterial endosymbionts. <i>Environmental Microbiology</i> , 2016, 18, 2591-2603.	1.8	50
49	Boom and bust: rapid feedback responses between insect outbreak dynamics and canopy leaf area impacted by rainfall and CO_2 . <i>Global Change Biology</i> , 2016, 22, 3632-3641.	4.2	28
50	The hitchhiker's guide to Europe: the infection dynamics of an ongoing <i>Wolbachia</i> invasion and mitochondrial selective sweep in <i>Rhagoletis cerasi</i> . <i>Molecular Ecology</i> , 2016, 25, 1595-1609.	2.0	68
51	Phylogeographic analyses of bacterial endosymbionts in fig homotomids (Hemiptera: Psylloidea) reveal codiversification of both primary and secondary endosymbionts. <i>FEMS Microbiology Ecology</i> , 2016, 92, fiw205.	1.3	19
52	<i>Wolbachia</i> in Parasitoids Attacking Native European and Introduced Eastern Cherry Fruit Flies in Europe. <i>Environmental Entomology</i> , 2016, 45, 1424-1431.	0.7	13
53	Insect herbivory in a mature Eucalyptus woodland canopy depends on leaf phenology but not CO_2 enrichment. <i>BMC Ecology</i> , 2016, 16, 47.	3.0	19
54	One step ahead: a parasitoid disperses farther and forms a wider geographic population than its fig wasp host. <i>Molecular Ecology</i> , 2016, 25, 882-894.	2.0	27

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55	Climate change, nutrition and immunity: Effects of elevated CO ₂ and temperature on the immune function of an insect herbivore. <i>Journal of Insect Physiology</i> , 2016, 85, 57-64.	0.9	24
56	Precipitation, not CO ₂ enrichment, drives insect herbivore frass deposition and subsequent nutrient dynamics in a mature Eucalyptus woodland. <i>Plant and Soil</i> , 2016, 399, 29-39.	1.8	15
57	Delimiting the species within the genus <i>Austroplebeia</i> , an Australian stingless bee, using multiple methodologies. <i>Apidologie</i> , 2016, 47, 76-89.	0.9	15
58	Loss of <i>Wolbachia</i> but not <i>Cardinium</i> in the invasive range of the Australian thrips species, <i>Pezothrips kellyanus</i> . <i>Biological Invasions</i> , 2016, 18, 197-214.	1.2	31
59	Characterisation of 14 microsatellite markers for the Australian fig psyllid, <i>Mycopsylla fici</i> . <i>Australian Journal of Zoology</i> , 2015, 63, 233.	0.6	2
60	Genetic variation amongst biotypes of <i>Dactylopius tomentosus</i> . <i>Insect Science</i> , 2015, 22, 360-374.	1.5	3
61	<i>Wolbachia</i> pseudogenes and low prevalence infections in tropical but not temperate Australian tephritid fruit flies: manifestations of lateral gene transfer and endosymbiont spillover?. <i>BMC Evolutionary Biology</i> , 2015, 15, 202.	3.2	37
62	Amino acid-mediated impacts of elevated carbon dioxide and simulated root herbivory on aphids are neutralized by increased air temperatures. <i>Journal of Experimental Botany</i> , 2015, 66, 613-623.	2.4	28
63	Responses of leaf beetle larvae to elevated [CO ₂] and temperature depend on Eucalyptus species. <i>Oecologia</i> , 2015, 177, 607-617.	0.9	24
64	Phylogenetic placement of Australian carrion beetles (Coleoptera: Silphidae). <i>Austral Entomology</i> , 2015, 54, 366-375.	0.8	9
65	Anatomy of an outbreak: the biology and population dynamics of a <i>Cardiaspina</i> psyllid species in an endangered woodland ecosystem. <i>Agricultural and Forest Entomology</i> , 2015, 17, 292-301.	0.7	32
66	Characterisation of microsatellite markers for fig-pollinating wasps in the <i>Pleistodontes imperialis</i> species complex. <i>Australian Journal of Zoology</i> , 2015, 63, 122.	0.6	2
67	The Microbiome of Field-Caught and Laboratory-Adapted Australian Tephritid Fruit Fly Species with Different Host Plant Use and Specialisation. <i>Microbial Ecology</i> , 2015, 70, 498-508.	1.4	125
68	Polyploidy versus endosymbionts in obligately thelytokous thrips. <i>BMC Evolutionary Biology</i> , 2015, 15, 23.	3.2	20
69	Double trouble: combined action of meiotic drive and <i>Wolbachia</i> feminization in <i>Eurema</i> butterflies. <i>Biology Letters</i> , 2015, 11, 20150095.	1.0	39
70	<i>Wolbachia</i> Influences the Production of Octopamine and Affects <i>Drosophila</i> Male Aggression. <i>Applied and Environmental Microbiology</i> , 2015, 81, 4573-4580.	1.4	46
71	Nutritional enhancement of leaves by a psyllid through senescence-like processes: insect manipulation or plant defence?. <i>Oecologia</i> , 2014, 176, 1061-1074.	0.9	35
72	Expression patterns of sex-determination genes in single male and female embryos of two <i>Bactrocera</i> fruit fly species during early development. <i>Insect Molecular Biology</i> , 2014, 23, 754-767.	1.0	29

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73	Tropical tephritid fruit fly community with high incidence of shared <i>Wolbachia</i> strains as platform for horizontal transmission of endosymbionts. <i>Environmental Microbiology</i> , 2014, 16, 3622-3637.	1.8	58
74	Susceptibility of Queensland fruit fly, <i>Bactrocera tryoni</i> (Froggatt) (Diptera: Tephritidae), to entomopathogenic nematodes. <i>Biological Control</i> , 2014, 69, 34-39.	1.4	58
75	Comprehensive transcriptome analysis of early male and female <i>Bactrocera jarvisi</i> embryos. <i>BMC Genetics</i> , 2014, 15, S7.	2.7	11
76	Australian endemic pest tephritids: genetic, molecular and microbial tools for improved Sterile Insect Technique. <i>BMC Genetics</i> , 2014, 15, S9.	2.7	23
77	Do eucalypt plantation management practices create understory reservoirs of scarab beetle pests in the soil?. <i>Forest Ecology and Management</i> , 2013, 306, 275-280.	1.4	18
78	Interactive effects of pre-industrial, current and future [CO ₂] and temperature on an insect herbivore of <i>Eucalyptus</i> . <i>Oecologia</i> , 2013, 171, 1025-1035.	0.9	19
79	Interactive direct and plant-mediated effects of elevated atmospheric [CO ₂] and temperature on a eucalypt-feeding insect herbivore. <i>Global Change Biology</i> , 2013, 19, 1407-1416.	4.2	69
80	Genomic Evolution of the Pathogenic <i>Wolbachia</i> Strain, wMelPop. <i>Genome Biology and Evolution</i> , 2013, 5, 2189-2204.	1.1	96
81	Biology and trophic interactions of lucerne aphids. <i>Agricultural and Forest Entomology</i> , 2013, 15, 335-350.	0.7	28
82	Evidence for a recent horizontal transmission and spatial spread of <i>Wolbachia</i> from endemic <i>Rhagoletis cerasi</i> (Diptera: Tephritidae) to invasive <i>Rhagoletis cingulata</i> in Europe. <i>Molecular Ecology</i> , 2013, 22, 4101-4111.	2.0	70
83	Effects of elevated temperature and CO ₂ on aboveground-belowground systems: a case study with plants, their mutualistic bacteria and root/shoot herbivores. <i>Frontiers in Plant Science</i> , 2013, 4, 445.	1.7	32
84	Root Damage by Insects Reverses the Effects of Elevated Atmospheric CO ₂ on Eucalypt Seedlings. <i>PLoS ONE</i> , 2013, 8, e79479.	1.1	15
85	Uncovering <i>Wolbachia</i> Diversity upon Artificial Host Transfer. <i>PLoS ONE</i> , 2013, 8, e82402.	1.1	24
86	Tandem repeat markers as novel diagnostic tools for high resolution fingerprinting of <i>Wolbachia</i> . <i>BMC Microbiology</i> , 2012, 12, S12.	1.3	48
87	Allele Intersection Analysis: A Novel Tool for Multi Locus Sequence Assignment in Multiply Infected Hosts. <i>PLoS ONE</i> , 2011, 6, e22198.	1.1	22
88	Multiple <i>Wolbachia</i> infections in <i>Rhagoletis pomonella</i> . <i>Entomologia Experimentalis Et Applicata</i> , 2011, 139, 138-144.	0.7	27
89	Mitochondrial phylogenies in the light of pseudogenes and <i>Wolbachia</i> : re-assessment of a bark beetle dataset. <i>ZooKeys</i> , 2010, 56, 269-280.	0.5	24
90	Evidence for Metabolic Provisioning by a Common Invertebrate Endosymbiont, <i>Wolbachia pipientis</i> , during Periods of Nutritional Stress. <i>PLoS Pathogens</i> , 2009, 5, e1000368.	2.1	306

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91	Hidden <i>Wolbachia</i> diversity in field populations of the European cherry fruit fly, <i>Rhagoletis cerasi</i> (Diptera, Tephritidae). <i>Molecular Ecology</i> , 2009, 18, 3816-3830.	2.0	96
92	Absence of the symbiont <i>Candidatus</i> <i>Midichloria mitochondrii</i> in the mitochondria of the tick <i>Ixodes holocyclus</i> . <i>FEMS Microbiology Letters</i> , 2009, 299, 241-247.	0.7	28
93	Evidence for low-titre infections in insect symbiosis: <i>Wolbachia</i> in the bark beetle <i>Pityogenes chalcographus</i> (Coleoptera, Scolytinae). <i>Environmental Microbiology</i> , 2009, 11, 1923-1933.	1.8	64
94	A <i>Wolbachia</i> Symbiont in <i>Aedes aegypti</i> Limits Infection with Dengue, Chikungunya, and Plasmodium. <i>Cell</i> , 2009, 139, 1268-1278.	13.5	1,384
95	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 May 2009–31 July 2009. <i>Molecular Ecology Resources</i> , 2009, 9, 1460-1466.	2.2	128
96	<i>Wolbachia</i> infections and superinfections in cytoplasmically incompatible populations of the European cherry fruit fly <i>Rhagoletis cerasi</i> (Diptera, Tephritidae). <i>Molecular Ecology</i> , 2008, 11, 2425-2434.	2.0	109
97	Male Development Time Influences the Strength of <i>Wolbachia</i> -Induced Cytoplasmic Incompatibility Expression in <i>Drosophila melanogaster</i> . <i>Genetics</i> , 2007, 177, 801-808.	1.2	96
98	Evolutionary dynamics of insect symbiont associations. <i>Trends in Ecology and Evolution</i> , 2007, 22, 625-627.	4.2	19
99	The Genus <i>Wolbachia</i> . , 2006, , 547-561.		5
100	Evolutionary Dynamics of wAu-Like <i>Wolbachia</i> Variants in Neotropical <i>Drosophila</i> spp. <i>Applied and Environmental Microbiology</i> , 2006, 72, 826-835.	1.4	50
101	Evidence for a Global <i>Wolbachia</i> Replacement in <i>Drosophila melanogaster</i> . <i>Current Biology</i> , 2005, 15, 1428-1433.	1.8	216
102	Distribution, Expression, and Motif Variability of Ankyrin Domain Genes in <i>Wolbachia pipientis</i> . <i>Journal of Bacteriology</i> , 2005, 187, 5136-5145.	1.0	126
103	New names for old strains? <i>Wolbachia</i> wSim is actually wRi. <i>Genome Biology</i> , 2005, 6, 401.	13.9	11
104	Phylogenomics of the Reproductive Parasite <i>Wolbachia pipientis</i> wMel: A Streamlined Genome Overrun by Mobile Genetic Elements. <i>PLoS Biology</i> , 2004, 2, e69.	2.6	713
105	<i>Wolbachia</i> Transfer from <i>Rhagoletis cerasi</i> to <i>Drosophila simulans</i> : Investigating the Outcomes of Host-Symbiont Coevolution. <i>Applied and Environmental Microbiology</i> , 2004, 70, 273-279.	1.4	78
106	INCIPIENT EVOLUTION OF WOLBACHIA COMPATIBILITY TYPES. <i>Evolution; International Journal of Organic Evolution</i> , 2004, 58, 1901.	1.1	5
107	INCIPIENT EVOLUTION OF WOLBACHIA COMPATIBILITY TYPES. <i>Evolution; International Journal of Organic Evolution</i> , 2004, 58, 1901-1908.	1.1	17
108	<i>Wolbachia</i> -induced cytoplasmic incompatibility as a means for insect pest population control. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 15042-15045.	3.3	345

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109	Development of a Physical and Genetic Map of the Virulent Wolbachia Strain w MelPop. Journal of Bacteriology, 2003, 185, 7077-7084.	1.0	30
110	Genetic evidence for natural hybridization between the Dutch elm disease pathogens <i>Ophiostoma novo-ulmi</i> sp. nov. and <i>O. novo-ulmi</i> sp. americana. Plant Pathology, 2002, 51, 78-84.	1.2	43
111	Morphological and genetic identification of the three pine pests of the genus <i>Tomicus</i> (Coleoptera, Tj ETQq1 1 0.784314 rgBT /Overl	0.7	41