

# Waldan K Kwong

## List of Publications by Year in descending order

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times ranked

2981  
citing authors

#	ARTICLE	IF	CITATIONS
1	Gene Transfer Agents in Bacterial Endosymbionts of Microbial Eukaryotes. <i>Genome Biology and Evolution</i> , 2022, 14, .	2.5	8
2	Taxonomy of the Apicomplexan Symbionts of Coral, including Corallicolida ord. nov., Reassignment of the Genus <i>Gemmocystis</i> , and Description of New Species <i>Corallicola aquarius</i> gen. nov. sp. nov. and <i>Anthozoaphila gnarlus</i> gen. nov. sp. nov.. <i>Journal of Eukaryotic Microbiology</i> , 2021, 68, e12852.	1.7	9
3	Corallicolids: The elusive coral-infecting apicomplexans. <i>PLoS Pathogens</i> , 2021, 17, e1009845.	4.7	5
4	Phylogenomics Identifies a New Major Subgroup of Apicomplexans, Marosporida class nov., with Extreme Apicoplast Genome Reduction. <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	23
5	Highly Reduced Genomes of Protist Endosymbionts Show Evolutionary Convergence. <i>Current Biology</i> , 2020, 30, 925-933.e3.	3.9	41
6	Bee microbiomes go viral. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 11197-11199.	7.1	3
7	Microbiome Evolution: Having the Guts to Be Different. <i>Current Biology</i> , 2020, 30, R766-R768.	3.9	3
8	A letter to Denis Lynn. <i>Aquatic Ecosystem Health and Management</i> , 2020, 23, 17-18.	0.6	0
9	A widespread coral-infecting apicomplexan with chlorophyll biosynthesis genes. <i>Nature</i> , 2019, 568, 103-107.	27.8	102
10	Division of labor in honey bee gut microbiota for plant polysaccharide digestion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 25909-25916.	7.1	191
11	Genome Sequences of <i>Apibacter</i> spp., Gut Symbionts of Asian Honey Bees. <i>Genome Biology and Evolution</i> , 2018, 10, 1174-1179.	2.5	27
12	Microbiome Structure Influences Infection by the Parasite <i>Crithidia bombi</i> in Bumble Bees. <i>Applied and Environmental Microbiology</i> , 2018, 84, .	3.1	86
13	Is Host Filtering the Main Driver of Phyllosymbiosis across the Tree of Life?. <i>MSystems</i> , 2018, 3, .	3.8	119
14	Competitive organelle-specific adaptors recruit Vps13 to membrane contact sites. <i>Journal of Cell Biology</i> , 2018, 217, 3593-3607.	5.2	122
15	Convergent evolution of a modified, acetate-driven TCA cycle in bacteria. <i>Nature Microbiology</i> , 2017, 2, 17067.	13.3	60
16	Immune system stimulation by the native gut microbiota of honey bees. <i>Royal Society Open Science</i> , 2017, 4, 170003.	2.4	276
17	Dynamic microbiome evolution in social bees. <i>Science Advances</i> , 2017, 3, e1600513.	10.3	349
18	Diversification of Type VI Secretion System Toxins Reveals Ancient Antagonism among Bee Gut Microbes. <i>MBio</i> , 2017, 8, .	4.1	94

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19	The Bee Microbiome: Impact on Bee Health and Model for Evolution and Ecology of Host-Microbe Interactions. MBio, 2016, 7, e02164-15.	4.1	215
20	Gut microbial communities of social bees. Nature Reviews Microbiology, 2016, 14, 374-384.	28.6	648
21	Metabolism of Toxic Sugars by Strains of the Bee Gut Symbiont <i>Gilliamella apicola</i> . MBio, 2016, 7, .	4.1	216
22	Genome-wide screen identifies host colonization determinants in a bacterial gut symbiont. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 13887-13892.	7.1	112
23	<i>Apibacter adventoris</i> gen. nov., sp. nov., a member of the phylum Bacteroidetes isolated from honey bees. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 1323-1329.	1.7	39
24	Evolution of host specialization in gut microbes: the bee gut as a model. Gut Microbes, 2015, 6, 214-220.	9.8	86
25	Genome Sequences of <i>Lactobacillus</i> sp. Strains wkb8 and wkb10, Members of the Firm-5 Clade, from Honey Bee Guts. Genome Announcements, 2014, 2, .	0.8	30
26	Genomics and host specialization of honey bee and bumble bee gut symbionts. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 11509-11514.	7.1	305
27	Complex integrons containing qnrB4-ampC (blaDHA-1) in plasmids of multidrug-resistant <i>Citrobacter freundii</i> from wastewater. Canadian Journal of Microbiology, 2013, 59, 110-116.	1.7	21
28	<i>Frischella perrara</i> gen. nov., sp. nov., a gammaproteobacterium isolated from the gut of the honeybee, <i>Apis mellifera</i> . International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 3646-3651.	1.7	96
29	Cultivation and characterization of the gut symbionts of honey bees and bumble bees: description of <i>Snodgrassella alvi</i> gen. nov., sp. nov., a member of the family Neisseriaceae of the Betaproteobacteria, and <i>Gilliamella apicola</i> gen. nov., sp. nov., a member of Orbaceae fam. nov., Orbales ord. nov., a sister taxon to the order "Enterobacteriales. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 2008-2018.	1.7	257
30	Standard methods for research on <i>Apis mellifera</i> gut symbionts. Journal of Apicultural Research, 2013, 52, 1-24.	1.5	98
31	Long-Term Exposure to Antibiotics Has Caused Accumulation of Resistance Determinants in the Gut Microbiota of Honeybees. MBio, 2012, 3, .	4.1	161