

Waldan K Kwong

List of Publications by Year in descending order

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Version: 2024-02-01

31
papers

3,819
citations

304743

22
h-index

434195

31
g-index

34
all docs

34
docs citations

34
times ranked

2981
citing authors

#	ARTICLE	IF	CITATIONS
1	Gut microbial communities of social bees. <i>Nature Reviews Microbiology</i> , 2016, 14, 374-384.	28.6	648
2	Dynamic microbiome evolution in social bees. <i>Science Advances</i> , 2017, 3, e1600513.	10.3	349
3	Genomics and host specialization of honey bee and bumble bee gut symbionts. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 11509-11514.	7.1	305
4	Immune system stimulation by the native gut microbiota of honey bees. <i>Royal Society Open Science</i> , 2017, 4, 170003.	2.4	276
5	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 α -Enterobacteriales. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 2008-2018.	1.7	257
6	Metabolism of Toxic Sugars by Strains of the Bee Gut Symbiont <i>Gilliamella apicola</i> . <i>MBio</i> , 2016, 7, .	4.1	216
7	The Bee Microbiome: Impact on Bee Health and Model for Evolution and Ecology of Host-Microbe Interactions. <i>MBio</i> , 2016, 7, e02164-15.	4.1	215
8	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
9	Long-Term Exposure to Antibiotics Has Caused Accumulation of Resistance Determinants in the Gut Microbiota of Honeybees. <i>MBio</i> , 2012, 3, .	4.1	161
10	Competitive organelle-specific adaptors recruit Vps13 to membrane contact sites. <i>Journal of Cell Biology</i> , 2018, 217, 3593-3607.	5.2	122
11	Is Host Filtering the Main Driver of Phyllosymbiosis across the Tree of Life?. <i>MSystems</i> , 2018, 3, .	3.8	119
12	Genome-wide screen identifies host colonization determinants in a bacterial gut symbiont. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 13887-13892.	7.1	112
13	A widespread coral-infecting apicomplexan with chlorophyll biosynthesis genes. <i>Nature</i> , 2019, 568, 103-107.	27.8	102
14	Standard methods for research on <i>Apis mellifera</i> gut symbionts. <i>Journal of Apicultural Research</i> , 2013, 52, 1-24.	1.5	98
15	<i>Frischella perrara</i> gen. nov., sp. nov., a gammaproteobacterium isolated from the gut of the honeybee, <i>Apis mellifera</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 3646-3651.	1.7	96
16	Diversification of Type VI Secretion System Toxins Reveals Ancient Antagonism among Bee Gut Microbes. <i>MBio</i> , 2017, 8, .	4.1	94
17	Evolution of host specialization in gut microbes: the bee gut as a model. <i>Gut Microbes</i> , 2015, 6, 214-220.	9.8	86
18	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

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19	Convergent evolution of a modified, acetate-driven TCA cycle in bacteria. <i>Nature Microbiology</i> , 2017, 2, 17067.	13.3	60
20	Highly Reduced Genomes of Protist Endosymbionts Show Evolutionary Convergence. <i>Current Biology</i> , 2020, 30, 925-933.e3.	3.9	41
21	<i>Apibacter adventoris</i> gen. nov., sp. nov., a member of the phylum Bacteroidetes isolated from honey bees. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2016, 66, 1323-1329.	1.7	39
22	Genome Sequences of <i>Lactobacillus</i> sp. Strains wkB8 and wkB10, Members of the Firm-5 Clade, from Honey Bee Guts. <i>Genome Announcements</i> , 2014, 2, .	0.8	30
23	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
24	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
25	Complex integrons containing qnrB4-ampC (blaDHA-1) in plasmids of multidrug-resistant <i>Citrobacter freundii</i> from wastewater. <i>Canadian Journal of Microbiology</i> , 2013, 59, 110-116.	1.7	21
26	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
27	Gene Transfer Agents in Bacterial Endosymbionts of Microbial Eukaryotes. <i>Genome Biology and Evolution</i> , 2022, 14, .	2.5	8
28	Corallicolids: The elusive coral-infecting apicomplexans. <i>PLoS Pathogens</i> , 2021, 17, e1009845.	4.7	5
29	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
30	Microbiome Evolution: Having the Guts to Be Different. <i>Current Biology</i> , 2020, 30, R766-R768.	3.9	3
31	A letter to Denis Lynn. <i>Aquatic Ecosystem Health and Management</i> , 2020, 23, 17-18.	0.6	0