

# Christopher C M Baker

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3676462/publications.pdf>

Version: 2024-02-01

12  
papers

258  
citations

1162889

8  
h-index

1281743

11  
g-index

15  
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15  
docs citations

15  
times ranked

480  
citing authors

#	ARTICLE	IF	CITATIONS
1	Measuring protected-area effectiveness using vertebrate distributions from leech iDNA. <i>Nature Communications</i> , 2022, 13, 1555.	5.8	8
2	Large-herbivore nemabiomes: patterns of parasite diversity and sharing. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2022, 289, 20212702.	1.2	6
3	Resource availability and heterogeneity shape the self-organisation of regular spatial patterning. <i>Ecology Letters</i> , 2021, 24, 1880-1891.	3.0	5
4	Spatial patterning of soil microbial communities created by fungus-farming termites. <i>Molecular Ecology</i> , 2020, 29, 4487-4501.	2.0	15
5	Combining stable isotope analysis with DNA metabarcoding improves inferences of trophic ecology. <i>PLoS ONE</i> , 2019, 14, e0219070.	1.1	15
6	Discovery of Four Novel Circular Single-Stranded DNA Viruses in Fungus-Farming Termites. <i>Genome Announcements</i> , 2018, 6, .	0.8	11
7	Virus discovery in all three major lineages of terrestrial arthropods highlights the diversity of single-stranded DNA viruses associated with invertebrates. <i>PeerJ</i> , 2018, 6, e5761.	0.9	51
8	Distinctive fungal communities in an obligate African ant-plant mutualism. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20162501.	1.2	19
9	Dissecting host-associated communities with DNA barcodes. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2016, 371, 20150328.	1.8	23
10	Metabarcoding as a tool for investigating arthropod diversity in <i>Nepenthes</i> pitcher plants. <i>Austral Ecology</i> , 2016, 41, 120-132.	0.7	24
11	Kin Selection and the Evolution of Social Information Use in Animal Conflict. <i>PLoS ONE</i> , 2012, 7, e31664.	1.1	0
12	Bacterial Diversity across Individual Lichens. <i>Applied and Environmental Microbiology</i> , 2011, 77, 4249-4252.	1.4	74