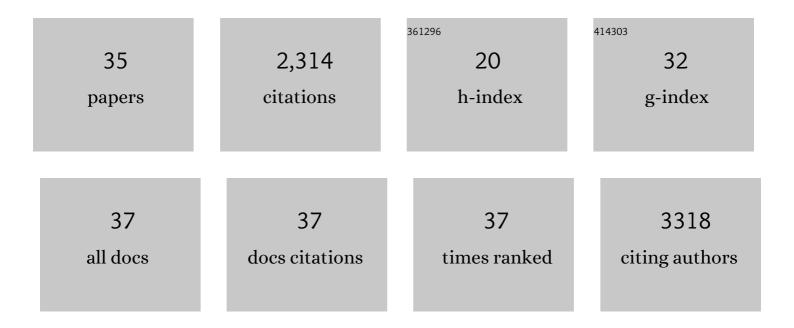
## Sarah C L Knowles

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3529572/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Marked seasonal variation in the wild mouse gut microbiota. ISME Journal, 2015, 9, 2423-2434.	4.4	282
2	Chronic malaria infections increase family inequalities and reduce parental fitness: experimental evidence from a wild bird population. Journal of Evolutionary Biology, 2010, 23, 557-569.	0.8	204
3	Withinâ€population variation in prevalence and lineage distribution of avian malaria in blue tits, <i>Cyanistes caeruleus</i> . Molecular Ecology, 2007, 16, 3263-3273.	2.0	194
4	Elevated reproductive effort increases blood parasitaemia and decreases immune function in birds: a metaâ€regression approach. Functional Ecology, 2009, 23, 405-415.	1.7	173
5	Parasite-Microbiota Interactions With the Vertebrate Gut: Synthesis Through an Ecological Lens. Frontiers in Microbiology, 2018, 9, 843.	1.5	146
6	Fitness effects of endemic malaria infections in a wild bird population: the importance of ecological structure. Journal of Animal Ecology, 2011, 80, 1196-1206.	1.3	136
7	Stability of within-host–parasite communities in a wild mammal system. Proceedings of the Royal Society B: Biological Sciences, 2013, 280, 20130598.	1.2	121
8	Molecular epidemiology of malaria prevalence and parasitaemia in a wild bird population. Molecular Ecology, 2011, 20, 1062-1076.	2.0	118
9	One health – an ecological and evolutionary framework for tackling Neglected Zoonotic Diseases. Evolutionary Applications, 2016, 9, 313-333.	1.5	112
10	Phenotypic correlates of <i>Clock</i> gene variation in a wild blue tit population: evidence for a role in seasonal timing of reproduction. Molecular Ecology, 2009, 18, 2444-2456.	2.0	97
11	Siteâ€occupancy modelling as a novel framework for assessing test sensitivity and estimating wildlife disease prevalence from imperfect diagnostic tests. Methods in Ecology and Evolution, 2012, 3, 339-348.	2.2	93
12	Infection dynamics of endemic malaria in a wild bird population: parasite species-dependent drivers of spatial and temporal variation in transmission rates. Journal of Animal Ecology, 2011, 80, 1207-1216.	1.3	87
13	The reliability of observational approaches for detecting interspecific parasite interactions: comparison with experimental results. International Journal for Parasitology, 2014, 44, 437-445.	1.3	76
14	The effect of helminth co-infection on malaria in mice: A meta-analysis. International Journal for Parasitology, 2011, 41, 1041-1051.	1.3	66
15	Social networks strongly predict the gut microbiota of wild mice. ISME Journal, 2021, 15, 2601-2613.	4.4	64
16	Identifying Microbiome-Mediated Behaviour in Wild Vertebrates. Trends in Ecology and Evolution, 2020, 35, 972-980.	4.2	53
17	The evolution of ecological facilitation within mixed-species biofilms in the mouse gastrointestinal tract. ISME Journal, 2018, 12, 2770-2784.	4.4	34
18	Spatial determinants of infection risk in a multiâ€species avian malaria system. Ecography, 2013, 36, 587-598	2.1	30

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19	Context-dependent effects of parental effort on malaria infection in a wild bird population, and their role in reproductive trade-offs. Oecologia, 2010, 164, 87-97.	0.9	29
20	NO EVIDENCE FOR AVIAN MALARIA INFECTION DURING THE NESTLING PHASE IN A PASSERINE BIRD. Journal of Parasitology, 2006, 92, 1302-1304.	0.3	25
21	Epidemiology and fitness effects of wood mouse herpesvirus in a natural host population. Journal of General Virology, 2012, 93, 2447-2456.	1.3	23
22	The impact of an 8-year mass drug administration programme on prevalence, intensity and co-infections of soil-transmitted helminthiases in Burundi. Parasites and Vectors, 2016, 9, 513.	1.0	21
23	Effects of laboratory domestication on the rodent gut microbiome. ISME Communications, 2021, 1, .	1.7	21
24	Optimising cluster survey design for planning schistosomiasis preventive chemotherapy. PLoS Neglected Tropical Diseases, 2017, 11, e0005599.	1.3	19
25	Dispersal in a patchy landscape reveals contrasting determinants of infection in a wild avian malaria system. Journal of Animal Ecology, 2014, 83, 429-439.	1.3	17
26	Juvenile Female Aggression in Cooperatively Breeding Pied Babblers: Causes and Contexts. Ethology, 2008, 114, 452-458.	0.5	14
27	A 16S rRNA Gene and Draft Genome Database for the Murine Oral Bacterial Community. MSystems, 2021, 6, .	1.7	14
28	Epidemiological Interactions between Urogenital and Intestinal Human Schistosomiasis in the Context of Praziquantel Treatment across Three West African Countries. PLoS Neglected Tropical Diseases, 2015, 9, e0004019.	1.3	14
29	Synchronous Seasonality in the Gut Microbiota of Wild Mouse Populations. Frontiers in Microbiology, 2022, 13, 809735.	1.5	14
30	How should we store avian faecal samples for microbiota analyses? Comparing efficacy and cost-effectiveness. Journal of Microbiological Methods, 2019, 165, 105689.	0.7	5
31	Evolutionary Biology: Parasite, Know Thyself. Current Biology, 2008, 18, R655-R657.	1.8	2
32	The impact of albendazole treatment on the incidence of viral- and bacterial-induced diarrhea in school children in southern Vietnam: study protocol for a randomized controlled trial. Trials, 2016, 17, 279.	0.7	2
33	Sex Ratios: Human Twins and Fraternal Effects. Current Biology, 2007, 17, R801-R804.	1.8	Ο
34	Response to Nguyen et al. â€~Laboratory-Inspired Manipulations Hold Value for Wild Microbiome-Behaviour Research'. Trends in Ecology and Evolution, 2021, 36, 278-280.	4.2	0
35	<i>Bifidobacterium castoris</i> strains isolated from wild mice show evidence of frequent host switching and diverse carbohydrate metabolism potential. ISME Communications, 2022, 2, .	1.7	0