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List of Publications by Year in descending order

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Version: 2024-02-01

933447 940533 21 318 10 16 citations g-index h-index papers 30 30 30 376 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	The genome of Eimeria falciformisv - reduction and specialization in a single host apicomplexan parasite. BMC Genomics, 2014, 15, 696.	2.8	44
2	Population expansion and individual age affect endoparasite richness and diversity in a recolonising large carnivore population. Scientific Reports, 2017, 7, 41730.	3.3	35
3	The Intestinal Eukaryotic and Bacterial Biome of Spotted Hyenas: The Impact of Social Status and Age on Diversity and Composition. Frontiers in Cellular and Infection Microbiology, 2017, 7, 262.	3.9	34
4	Eimeria falciformis infection of the mouse caecum identifies opposing roles of IFN \hat{I}^3 -regulated host pathways for the parasite development. Mucosal Immunology, 2014, 7, 969-982.	6.0	21
5	Dual RNA-seq reveals no plastic transcriptional response of the coccidian parasite Eimeria falciformis to host immune defenses. BMC Genomics, 2017, 18, 686.	2.8	20
6	Massive encapsulation of larval Anguillicoloides crassus in the intestinal wall of Japanese eels. Parasites and Vectors, 2009, 2, 48.	2.5	19
7	Detection and quantification of house mouse Eimeria at the species level – Challenges and solutions for the assessment of coccidia in wildlife. International Journal for Parasitology: Parasites and Wildlife, 2019, 10, 29-40.	1.5	18
8	The P-glycoprotein repertoire of the equine parasitic nematode Parascaris univalens. Scientific Reports, 2020, 10, 13586.	3.3	16
9	Recolonizing gray wolves increase parasite infection risk in their prey. Ecology and Evolution, 2018, 8, 2160-2170.	1.9	13
10	Generalist <i>Eimeria</i> species in rodents: Multilocus analyses indicate inadequate resolution of established markers. Ecology and Evolution, 2020, 10, 1378-1389.	1.9	12
11	Intensity of infection with intracellular <i>Eimeria</i> spp. and pinworms is reduced in hybrid mice compared to parental subspecies. Journal of Evolutionary Biology, 2020, 33, 435-448.	1.7	11
12	Bat Community Responses to Structural Habitat Complexity Resulting from Management Practices Within Different Land Use Types — A Case Study from North-Eastern Germany. Acta Chiropterologica, 2019, 20, 387.	0.6	10
13	The transcriptome of the invasive eel swimbladder nematode parasite Anguillicola crassus. BMC Genomics, 2013, 14, 87.	2.8	8
14	A Novel Non-invasive Method to Detect RELM Beta Transcript in Gut Barrier Related Changes During a Gastrointestinal Nematode Infection. Frontiers in Immunology, 2019, 10, 445.	4.8	7
15	Coupling between tolerance and resistance for two related <i>Eimeria</i> parasite species. Ecology and Evolution, 2020, 10, 13938-13948.	1.9	7
16	Transcriptome analyses of <i>Anguillicola crassus </i> from native and novel hosts. PeerJ, 2014, 2, e684.	2.0	7
17	Shifting focus from resistance to disease tolerance: A review on hybrid house mice. Ecology and Evolution, 2022, 12, e8889.	1.9	6
18	Dual RNA Sequencing Meta-analysis in Plasmodium Infection Identifies Host-Parasite Interactions. MSystems, $2021, 6, .$	3.8	5

#	Article	IF	CITATIONS
19	Whip- and pinworm infections elicit contrasting effector and distinct regulatory responses in wild house mice. International Journal for Parasitology, 2022, 52, 519-524.	3.1	4
20	Toxoplasma and Eimeria co-opt the host cFos expression for intracellular development in mammalian cells. Computational and Structural Biotechnology Journal, 2021, 19, 719-731.	4.1	3
21	DNA-based quantification and counting of transmission stages provides different but complementary parasite load estimates: an example from rodent coccidia (Eimeria). Parasites and Vectors, 2022, 15, 45.	2.5	1