

Cinzia Cantacessi

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125
papers

4,228
citations

36
h-index

60
g-index

131
ext. papers

5,176
ext. citations

5.9
avg, IF

5.21
L-index

#	Paper	IF	Citations
125	Whole-genome sequence of <i>Schistosoma haematobium</i> . <i>Nature Genetics</i> , 2012 , 44, 221-5	36.3	325
124	<i>Ascaris suum</i> draft genome. <i>Nature</i> , 2011 , 479, 529-33	50.4	217
123	A portrait of the "SCP/TAPS" proteins of eukaryotes--developing a framework for fundamental research and biotechnological outcomes. <i>Biotechnology Advances</i> , 2009 , 27, 376-88	17.8	126
122	Unlocking the transcriptomes of two carcinogenic parasites, <i>Clonorchis sinensis</i> and <i>Opisthorchis viverrini</i> . <i>PLoS Neglected Tropical Diseases</i> , 2010 , 4, e719	4.8	122
121	The role of wild canids and felids in spreading parasites to dogs and cats in Europe. Part II: Helminths and arthropods. <i>Veterinary Parasitology</i> , 2015 , 213, 24-37	2.8	114
120	Carcinogenic Liver Fluke Secretes Extracellular Vesicles That Promote Cholangiocytes to Adopt a Tumorigenic Phenotype. <i>Journal of Infectious Diseases</i> , 2015 , 212, 1636-45	7	103
119	Elucidating the transcriptome of <i>Fasciola hepatica</i> - a key to fundamental and biotechnological discoveries for a neglected parasite. <i>Biotechnology Advances</i> , 2010 , 28, 222-31	17.8	102
118	Impact of experimental hookworm infection on the human gut microbiota. <i>Journal of Infectious Diseases</i> , 2014 , 210, 1431-4	7	100
117	<i>Phortica variegata</i> as an intermediate host of <i>Thelazia callipaeda</i> under natural conditions: evidence for pathogen transmission by a male arthropod vector. <i>International Journal for Parasitology</i> , 2006 , 36, 1167-73	4.3	93
116	Human thelaziosis--a neglected parasitic disease of the eye. <i>Journal of Parasitology</i> , 2006 , 92, 872-5	0.9	83
115	Lungworms and gastrointestinal parasites of domestic cats: a European perspective. <i>International Journal for Parasitology</i> , 2017 , 47, 517-528	4.3	78
114	The zoophilic fruitfly <i>Phortica variegata</i> : morphology, ecology and biological niche. <i>Medical and Veterinary Entomology</i> , 2006 , 20, 358-64	2.4	74
113	Efficacy of a combination of 10% imidacloprid/50% permethrin for the prevention of leishmaniasis in kennelled dogs in an endemic area. <i>Veterinary Parasitology</i> , 2007 , 144, 270-8	2.8	73
112	Nematode biology and larval development of <i>Thelazia callipaeda</i> (Spirurida, Thelaziidae) in the drosophilid intermediate host in Europe and China. <i>Parasitology</i> , 2005 , 131, 847-55	2.7	73
111	The role of wild canids and felids in spreading parasites to dogs and cats in Europe. Part I: Protozoa and tick-borne agents. <i>Veterinary Parasitology</i> , 2015 , 213, 12-23	2.8	72
110	Experimental hookworm infection and escalating gluten challenges are associated with increased microbial richness in celiac subjects. <i>Scientific Reports</i> , 2015 , 5, 13797	4.9	70
109	A portrait of the transcriptome of the neglected trematode, <i>Fasciola gigantica</i> --biological and biotechnological implications. <i>PLoS Neglected Tropical Diseases</i> , 2011 , 5, e1004	4.8	70

108	Massively parallel sequencing and analysis of the <i>Necator americanus</i> transcriptome. <i>PLoS Neglected Tropical Diseases</i> , 2010 , 4, e684	4.8	66
107	The past, present, and future of <i>Leishmania</i> genomics and transcriptomics. <i>Trends in Parasitology</i> , 2015 , 31, 100-8	6.4	65
106	Secreted proteomes of different developmental stages of the gastrointestinal nematode <i>Nippostrongylus brasiliensis</i> . <i>Molecular and Cellular Proteomics</i> , 2014 , 13, 2736-51	7.6	63
105	First reports of autochthonous eyeworm infection by <i>Thelazia callipaeda</i> (Spirurida, Thelaziidae) in dogs and cat from France. <i>Veterinary Parasitology</i> , 2007 , 149, 294-7	2.8	63
104	A practical, bioinformatic workflow system for large data sets generated by next-generation sequencing. <i>Nucleic Acids Research</i> , 2010 , 38, e171	20.1	60
103	Morphological and molecular differentiation between <i>Dicrocoelium dendriticum</i> (Rudolphi, 1819) and <i>Dicrocoelium chinensis</i> (Sudarikov and Ryjikov, 1951) Tang and Tang, 1978 (Platyhelminthes: Digenea). <i>Acta Tropica</i> , 2007 , 104, 91-8	3.2	54
102	Infections by human gastrointestinal helminths are associated with changes in faecal microbiota diversity and composition. <i>PLoS ONE</i> , 2017 , 12, e0184719	3.7	52
101	First transcriptomic analysis of the economically important parasitic nematode, <i>Trichostrongylus colubriformis</i> , using a next-generation sequencing approach. <i>Infection, Genetics and Evolution</i> , 2010 , 10, 1199-207	4.5	50
100	This Gut Ain't Big Enough for Both of Us. Or Is It? Helminth-Microbiota Interactions in Veterinary Species. <i>Trends in Parasitology</i> , 2017 , 33, 619-632	6.4	49
99	<i>Schistosoma mansoni</i> infection is associated with quantitative and qualitative modifications of the mammalian intestinal microbiota. <i>Scientific Reports</i> , 2018 , 8, 12072	4.9	47
98	The specific identification of anisakid larvae from fishes from the Yellow Sea, China, using mutation scanning-coupled sequence analysis of nuclear ribosomal DNA. <i>Molecular and Cellular Probes</i> , 2007 , 21, 386-90	3.3	47
97	Prevention of canine leishmaniosis in a hyper-endemic area using a combination of 10% imidacloprid/4.5% flumethrin. <i>PLoS ONE</i> , 2013 , 8, e56374	3.7	46
96	Changes in duodenal tissue-associated microbiota following hookworm infection and consecutive gluten challenges in humans with coeliac disease. <i>Scientific Reports</i> , 2016 , 6, 36797	4.9	44
95	Bovine theileriosis--an emerging problem in south-eastern Australia?. <i>Infection, Genetics and Evolution</i> , 2011 , 11, 2095-7	4.5	44
94	Differences in transcription between free-living and CO ₂ -activated third-stage larvae of <i>Haemonchus contortus</i> . <i>BMC Genomics</i> , 2010 , 11, 266	4.5	44
93	Efficacy of a combination of imidacloprid 10%/permethrin 50% versus fipronil 10%/(S)-methoprene 12%, against ticks in naturally infected dogs. <i>Veterinary Parasitology</i> , 2005 , 130, 293-304	2.8	43
92	Efficacy of an imidacloprid/flumethrin collar against fleas, ticks and tick-borne pathogens in dogs. <i>Parasites and Vectors</i> , 2013 , 6, 245	4	41
91	The transcriptome of <i>Trichuris suis</i> --first molecular insights into a parasite with curative properties for key immune diseases of humans. <i>PLoS ONE</i> , 2011 , 6, e23590	3.7	40

90	Heterochronic faecal transplantation boosts gut germinal centres in aged mice. <i>Nature Communications</i> , 2019 , 10, 2443	17.4	39
89	Genetic characterization of three unique operational taxonomic units of <i>Eimeria</i> from chickens in Australia based on nuclear spacer ribosomal DNA. <i>Veterinary Parasitology</i> , 2008 , 152, 226-34	2.8	35
88	Application of 10% imidacloprid/50% permethrin to prevent <i>Ehrlichia canis</i> exposure in dogs under natural conditions. <i>Veterinary Parasitology</i> , 2008 , 153, 320-8	2.8	35
87	Key strongylid nematodes of animals - Impact of next-generation transcriptomics on systems biology and biotechnology. <i>Biotechnology Advances</i> , 2012 , 30, 469-88	17.8	34
86	Suppression of inflammation by helminths: a role for the gut microbiota?. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2015 , 370,	5.8	32
85	Probing of a human proteome microarray with a recombinant pathogen protein reveals a novel mechanism by which hookworms suppress B-cell receptor signaling. <i>Journal of Infectious Diseases</i> , 2015 , 211, 416-25	7	31
84	Feline lungworms unlock a novel mode of parasite transmission. <i>Scientific Reports</i> , 2015 , 5, 13105	4.9	31
83	Molecular mechanisms of hookworm disease: stealth, virulence, and vaccines. <i>Journal of Allergy and Clinical Immunology</i> , 2012 , 130, 13-21	11.5	29
82	A comprehensive analysis of the faecal microbiome and metabolome of <i>Strongyloides stercoralis</i> infected volunteers from a non-endemic area. <i>Scientific Reports</i> , 2018 , 8, 15651	4.9	29
81	SCP/TAPS proteins in helminths--where to from now?. <i>Molecular and Cellular Probes</i> , 2012 , 26, 54-9	3.3	28
80	DNA technological progress toward advanced diagnostic tools to support human hookworm control. <i>Biotechnology Advances</i> , 2008 , 26, 35-45	17.8	28
79	First report of <i>Thelazia callipaeda</i> (Spirurida, Thelaziidae) in wolves in Italy. <i>Journal of Wildlife Diseases</i> , 2007 , 43, 508-11	1.3	28
78	Occurrence of strongyloidiasis in privately owned and sheltered dogs: clinical presentation and treatment outcome. <i>Parasites and Vectors</i> , 2017 , 10, 345	4	27
77	Comparative analyses of mitochondrial and nuclear genetic markers for the molecular identification of <i>Rhipicephalus</i> spp. <i>Infection, Genetics and Evolution</i> , 2013 , 20, 422-7	4.5	27
76	Deep insights into <i>Dictyocaulus viviparus</i> transcriptomes provides unique prospects for new drug targets and disease intervention. <i>Biotechnology Advances</i> , 2011 , 29, 261-71	17.8	27
75	The relationships between faecal egg counts and gut microbial composition in UK Thoroughbreds infected by cyathostomins. <i>International Journal for Parasitology</i> , 2018 , 48, 403-412	4.3	26
74	Gastropod-Borne Helminths: A Look at the Snail-Parasite Interplay. <i>Trends in Parasitology</i> , 2016 , 32, 255-264	11.4	26
73	A deep exploration of the transcriptome and "excretory/secretory" proteome of adult <i>Fascioloides magna</i> . <i>Molecular and Cellular Proteomics</i> , 2012 , 11, 1340-53	7.6	26

72	Genomic-bioinformatic analysis of transcripts enriched in the third-stage larva of the parasitic nematode <i>Ascaris suum</i> . <i>PLoS Neglected Tropical Diseases</i> , 2008 , 2, e246	4.8	26
71	Atypical (RIO) protein kinases from <i>Haemonchus contortus</i> --promise as new targets for nematocidal drugs. <i>Biotechnology Advances</i> , 2011 , 29, 338-50	17.8	25
70	Tissue-specific transcriptomes of <i>Anisakis simplex</i> (sensu stricto) and <i>Anisakis pegreffii</i> reveal potential molecular mechanisms involved in pathogenicity. <i>Parasites and Vectors</i> , 2018 , 11, 31	4	24
69	Helminth infections and gut microbiota - a feline perspective. <i>Parasites and Vectors</i> , 2016 , 9, 625	4	24
68	The <i>Anisakis</i> Transcriptome Provides a Resource for Fundamental and Applied Studies on Allergy-Causing Parasites. <i>PLoS Neglected Tropical Diseases</i> , 2016 , 10, e0004845	4.8	23
67	Getting the most out of parasitic helminth transcriptomes using HelmDB: implications for biology and biotechnology. <i>Biotechnology Advances</i> , 2013 , 31, 1109-19	17.8	22
66	A genome-wide analysis of annexins from parasitic organisms and their vectors. <i>Scientific Reports</i> , 2013 , 3, 2893	4.9	22
65	Progress on the transcriptomics of carcinogenic liver flukes of humans--unique biological and biotechnological prospects. <i>Biotechnology Advances</i> , 2010 , 28, 859-70	17.8	22
64	Risk factors for canine neosporosis in farm and kennel dogs in southern Italy. <i>Veterinary Parasitology</i> , 2007 , 145, 240-4	2.8	22
63	Insights into SCP/TAPS proteins of liver flukes based on large-scale bioinformatic analyses of sequence datasets. <i>PLoS ONE</i> , 2012 , 7, e31164	3.7	22
62	Helminth Microbiomes - A Hidden Treasure Trove?. <i>Trends in Parasitology</i> , 2019 , 35, 13-22	6.4	22
61	<i>Angiostrongylus chabaudi</i> in felids: New findings and a review of the literature. <i>Veterinary Parasitology</i> , 2016 , 228, 188-192	2.8	21
60	Molecular characterization of selected dermatophytes and their identification by electrophoretic mutation scanning. <i>Electrophoresis</i> , 2009 , 30, 3555-64	3.6	21
59	Helminth-microbiota cross-talk - A journey through the vertebrate digestive system. <i>Molecular and Biochemical Parasitology</i> , 2019 , 233, 111222	1.9	20
58	Hookworm Treatment for Relapsing Multiple Sclerosis: A Randomized Double-Blinded Placebo-Controlled Trial. <i>JAMA Neurology</i> , 2020 , 77, 1089-1098	17.2	20
57	Chronic polyarthritis associated to <i>Cercopithifilaria baina</i> infection in a dog. <i>Veterinary Parasitology</i> , 2014 , 205, 401-4	2.8	20
56	Tick vectors of <i>Cercopithifilaria baina</i> in dogs: <i>Rhipicephalus sanguineus sensu lato</i> versus <i>Ixodes ricinus</i> . <i>Parasitology Research</i> , 2013 , 112, 3013-7	2.4	20
55	Evaluation of blood and bone marrow in selected canine vector-borne diseases. <i>Parasites and Vectors</i> , 2014 , 7, 534	4	19

54	Dollfustrema durum n. sp. and Heterobucephalopsis perardua n. sp. (Digenea: Bucephalidae) from the giant moray eel, Gymnothorax javanicus (Bleeker) (Anguilliformes: Muraenidae), and proposal of the Heterobucephalopsinae n. subfam. <i>Parasitology International</i> , 2015 , 64, 559-70	2.1	18
53	Bioinformatics meets parasitology. <i>Parasite Immunology</i> , 2012 , 34, 265-75	2.2	18
52	Cryptic parasite revealed improved prospects for treatment and control of human cryptosporidiosis through advanced technologies. <i>Advances in Parasitology</i> , 2011 , 77, 141-73	3.2	18
51	Vertical transmission of Anaplasma platys and Leishmania infantum in dogs during the first half of gestation. <i>Parasites and Vectors</i> , 2016 , 9, 269	4	18
50	Helminths and microbes within the vertebrate gut - not all studies are created equal. <i>Parasitology</i> , 2019 , 146, 1371-1378	2.7	17
49	Improved molecular diagnostic tools for human hookworms. <i>Expert Review of Molecular Diagnostics</i> , 2009 , 9, 17-21	3.8	16
48	Bioinformatic analysis of abundant, gender-enriched transcripts of adult Ascaris suum (Nematoda) using a semi-automated workflow platform. <i>Molecular and Cellular Probes</i> , 2009 , 23, 205-17	3.3	16
47	Molecular characterization and phylogenesis of Steganinae (Diptera, Drosophilidae) inferred by the mitochondrial cytochrome c oxidase subunit 1. <i>Medical and Veterinary Entomology</i> , 2008 , 22, 37-47	2.4	16
46	Dysbiosis associated with acute helminth infections in herbivorous youngstock - observations and implications. <i>Scientific Reports</i> , 2019 , 9, 11121	4.9	15
45	TIMPs of parasitic helminths - a large-scale analysis of high-throughput sequence datasets. <i>Parasites and Vectors</i> , 2013 , 6, 156	4	15
44	Exclusive dependence of IL-10R signalling on intestinal microbiota homeostasis and control of whipworm infection. <i>PLoS Pathogens</i> , 2019 , 15, e1007265	7.6	15
43	A preliminary investigation of serological tools for the detection of Onchocerca lupi infection in dogs. <i>Parasitology Research</i> , 2014 , 113, 1989-91	2.4	14
42	Cardicola beveridgei n. sp. (Digenea: Aporocotylidae) from the mangrove jack, Lutjanus argentimaculatus (Perciformes: Lutjanidae), and C. bullardi n. sp. from the Australian spotted mackerel, Scomberomorus munroi (Perciformes: Scombridae), from the northern Great Barrier Reef. <i>Parasitology International</i> , 2014 , 63, 735-45	2.1	14
41	Proteomic profile of Bithynia siamensis goniomphalos snails upon infection with the carcinogenic liver fluke Opisthorchis viverrini. <i>Journal of Proteomics</i> , 2015 , 113, 281-91	3.9	13
40	Comparative Transcriptomics Reveals Clues for Differences in Pathogenicity between , sensu stricto and. <i>Genes</i> , 2020 , 11,	4.2	13
39	MICHELINdb: a web-based tool for mining of helminth-microbiota interaction datasets, and a meta-analysis of current research. <i>Microbiome</i> , 2020 , 8, 10	16.6	13
38	Coming out of the shell: building the molecular infrastructure for research on parasite-harboring snails. <i>PLoS Neglected Tropical Diseases</i> , 2013 , 7, e2284	4.8	13
37	Investigation of the regulation of transcriptional changes in Ancylostoma caninum larvae following serum activation, with a focus on the insulin-like signalling pathway. <i>Veterinary Parasitology</i> , 2009 , 159, 139-48	2.8	13

36	Classic Models for New Perspectives: Delving into Helminth-Microbiota-Immune System Interactions. <i>Trends in Parasitology</i> , 2018 , 34, 640-654	6.4	12
35	RNA-Seq reveals infection-induced gene expression changes in the snail intermediate host of the carcinogenic liver fluke, <i>Opisthorchis viverrini</i> . <i>PLoS Neglected Tropical Diseases</i> , 2014 , 8, e2765	4.8	12
34	Novel inhibitor cystine knot peptides from <i>Momordica charantia</i> . <i>PLoS ONE</i> , 2013 , 8, e75334	3.7	12
33	Infection with the sheep gastrointestinal nematode <i>Teladorsagia circumcincta</i> increases luminal pathobionts. <i>Microbiome</i> , 2020 , 8, 60	16.6	11
32	Cyathostomine egg reappearance period following ivermectin treatment in a cohort of UK Thoroughbreds. <i>Parasites and Vectors</i> , 2018 , 11, 61	4	10
31	Exploring transcriptional conservation between <i>Ancylostoma caninum</i> and <i>Haemonchus contortus</i> by oligonucleotide microarray and bioinformatic analyses. <i>Molecular and Cellular Probes</i> , 2009 , 23, 1-9	3.3	10
30	ROR γ Treg to Th17 ratios correlate with susceptibility to <i>Giardia</i> infection. <i>Scientific Reports</i> , 2019 , 9, 20328	4.9	10
29	Helminths, hosts, and their microbiota: new avenues for managing gastrointestinal helminthiasis in ruminants. <i>Expert Review of Anti-Infective Therapy</i> , 2020 , 18, 977-985	5.5	9
28	High-intensity cardiac infections of <i>Phthinomita heinigeriae</i> n. sp. (Digenea: Aporocotylidae) in the orangeline cardinalfish, <i>Taeniamia fucata</i> (Cantor), off Heron Island on the Great Barrier Reef. <i>Parasitology International</i> , 2016 , 65, 371-7	2.1	9
27	A new PCR assay for the detection and differentiation of <i>Babesia canis</i> and <i>Babesia vogeli</i> . <i>Ticks and Tick-borne Diseases</i> , 2017 , 8, 862-865	3.6	9
26	Oestrus ovis causing human ocular myiasis: from countryside to town centre. <i>Clinical and Experimental Ophthalmology</i> , 2009 , 37, 327-8	2.4	9
25	Impact of next-generation technologies on exploring socioeconomically important parasites and developing new interventions. <i>Methods in Molecular Biology</i> , 2015 , 1247, 437-74	1.4	9
24	<i>Filaria martis</i> Gmelin 1790 (Spirurida, Filariidae) affecting beech marten (<i>Martes foina</i>): morphological description and molecular characterisation of the cytochrome oxidase c subunit I. <i>Parasitology Research</i> , 2007 , 101, 877-83	2.4	8
23	Filarial infection caused by <i>Onchocerca boehmi</i> (Supperer, 1953) in a horse from Italy. <i>Parasitology Research</i> , 2017 , 116, 191-198	2.4	7
22	Molecular characterization and phylogenetic inferences of <i>Dermanyssus gallinae</i> isolates in Italy within an European framework. <i>Medical and Veterinary Entomology</i> , 2014 , 28, 447-52	2.4	7
21	Differential Protein Expression in the Hemolymph of <i>Bithynia siamensis</i> goniomphalos Infected with <i>Opisthorchis viverrini</i> . <i>PLoS Neglected Tropical Diseases</i> , 2016 , 10, e0005104	4.8	7
20	The best type of inoculum for testing the antifungal drug susceptibility of <i>Microsporum canis</i> : In vivo and in vitro results. <i>Mycoses</i> , 2020 , 63, 711-716	5.2	7
19	Parasite transmission by insects: a female affair?. <i>Trends in Parasitology</i> , 2008 , 24, 116-20	6.4	6

18	Molecular identification of <i>Phortica variegata</i> and <i>Phortica semivirgo</i> (Drosophilidae, Steganinae) by PCR-RFLP of the mitochondrial cytochrome oxidase c subunit I gene. <i>Parasitology Research</i> , 2008 , 103, 727-30	2.4	5
17	Baseline Gut Microbiota Composition Is Associated With Infection Burden in Rodent Models. <i>Frontiers in Immunology</i> , 2020 , 11, 593838	8.4	5
16	Experimental infection with the hookworm, <i>Necator americanus</i> , is associated with stable gut microbial diversity in human volunteers with relapsing multiple sclerosis. <i>BMC Biology</i> , 2021 , 19, 74	7.3	5
15	Data set from the proteomic analysis of <i>Bithynia siamensis</i> goniomphalos snails upon infection with the carcinogenic liver fluke <i>Opisthorchis viverrini</i> . <i>Data in Brief</i> , 2015 , 2, 16-20	1.2	4
14	First laboratory culture of <i>Phortica variegata</i> (Diptera, Steganinae), a vector of <i>Thelazia callipaeda</i> . <i>Journal of Vector Ecology</i> , 2012 , 37, 458-61	1.5	4
13	A bug's life: Delving into the challenges of helminth microbiome studies. <i>PLoS Neglected Tropical Diseases</i> , 2020 , 14, e0008446	4.8	4
12	Heartworm genomics: unprecedented opportunities for fundamental molecular insights and new intervention strategies. <i>Topics in Companion Animal Medicine</i> , 2011 , 26, 193-9	1.1	3
11	Major prospects for exploring canine vector borne diseases and novel intervention methods using genomic technologies. <i>Parasites and Vectors</i> , 2011 , 4, 53	4	2
10	MixS-SA: a MixS extension defining the minimum information standard for sequence data from symbiont-associated micro-organisms. <i>ISME Communications</i> , 2022 , 2,		1
9	Vaccination against the brown stomach worm, <i>Teladorsagia circumcincta</i> , followed by parasite challenge, induces inconsistent modifications in gut microbiota composition of lambs. <i>Parasites and Vectors</i> , 2021 , 14, 189	4	1
8	Serum amyloid A levels and alpha 2 and gamma globulins on serum protein electrophoresis in cats exposed to and infected with <i>Leishmania infantum</i> . <i>Parasites and Vectors</i> , 2021 , 14, 217	4	1
7	Virulence and in vitro antifungal susceptibility of <i>Candida albicans</i> and <i>Candida catenulata</i> from laying hens. <i>International Microbiology</i> , 2021 , 24, 57-63	3	1
6	Gut-microbiota-derived extracellular vesicles: Overlooked mediators in host-helminth interactions?. <i>Trends in Parasitology</i> , 2021 , 37, 690-693	6.4	1
5	Recent Progress in Transcriptomics of Key Gastrointestinal Nematodes of Animals [Fundamental Research Toward New Intervention Strategies]61-72		1
4	Harnessing Genomic Technologies to Explore the Molecular Biology of Liver Flukes-Major Implications for Fundamental and Applied Research73-87		1
3	Worms and bugs of the gut: the search for diagnostic signatures using barcoding, and metagenomics-metabolomics.. <i>Parasites and Vectors</i> , 2022 , 15, 118	4	1
2	Helminth Microbiota Profiling Using Bacterial 16S rRNA Gene Amplicon Sequencing: From Sampling to Sequence Data Mining. <i>Methods in Molecular Biology</i> , 2021 , 2369, 263-298	1.4	0
1	Investigation of Host-Microbe-Parasite Interactions in an In Vitro 3D Model of the Vertebrate Gut. <i>Advanced Biology</i> ,2200015		0

