List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Detection, Molecular Identification and Transmission of the Intestinal Protozoa Blastocystis sp. in Guinea from a Large-Scale Epidemiological Study Conducted in the Conakry Area. Microorganisms, 2022, 10, 446.	3.6	8
2	Animal, Herd and Feed Characteristics Associated with Blastocystis Prevalence and Molecular Diversity in Dairy Cattle from the North of France. Parasitologia, 2022, 2, 45-53.	1.3	2
3	Blastocystis sp. Prevalence and Subtypes Distribution amongst Syrian Refugee Communities Living in North Lebanon. Microorganisms, 2021, 9, 184.	3.6	16
4	Changes in the Human Gut Microbiota Associated With Colonization by Blastocystis sp. and Entamoeba spp. in Non-Industrialized Populations. Frontiers in Cellular and Infection Microbiology, 2021, 11, 533528.	3.9	26
5	Genomic insights into the host specific adaptation of the Pneumocystis genus. Communications Biology, 2021, 4, 305.	4.4	23
6	Revisiting the Pneumocystis host specificity paradigm and transmission ecology in wild Southeast Asian rodents. Infection, Genetics and Evolution, 2021, 93, 104978.	2.3	6
7	Editorial: Enteric Unicellular Eukaryotic Parasites and Gut Microbiota: Mechanisms and Ecology. Frontiers in Microbiology, 2021, 12, 779412.	3.5	0
8	Persistent Cryptosporidium parvum Infection Leads to the Development of the Tumor Microenvironment in an Experimental Mouse Model: Results of a Microarray Approach. Microorganisms, 2021, 9, 2569.	3.6	6
9	Prevalence and Subtype Distribution of Blastocystis sp. in Senegalese School Children. Microorganisms, 2020, 8, 1408.	3.6	63
10	Cryptosporidium and Colon Cancer: Cause or Consequence?. Microorganisms, 2020, 8, 1665.	3.6	31
11	Genetic basis for virulence differences of various Cryptosporidium parvum carcinogenic isolates. Scientific Reports, 2020, 10, 7316.	3.3	10
12	Respiratory mycobiome and suggestion of inter-kingdom network during acute pulmonary exacerbation in cystic fibrosis. Scientific Reports, 2020, 10, 3589.	3.3	71
13	First Report on the Prevalence and Subtype Distribution of Blastocystis sp. in Edible Marine Fish and Marine Mammals: A Large Scale-Study Conducted in Atlantic Northeast and on the Coasts of Northern France. Microorganisms, 2020, 8, 460.	3.6	21
14	Diversity and Complexity of the Large Surface Protein Family in the Compacted Genomes of Multiple <i>Pneumocystis</i> Species. MBio, 2020, 11, .	4.1	11
15	The Impact of Bioinformatics Pipelines on Microbiota Studies: Does the Analytical "Microscope―Affect the Biological Interpretation?. Microorganisms, 2019, 7, 393.	3.6	17
16	First report on the prevalence and subtype distribution of Blastocystis sp. in dairy cattle in Lebanon and assessment of zoonotic transmission. Acta Tropica, 2019, 194, 23-29.	2.0	45
17	Use of shotgun metagenomics for the identification of protozoa in the gut microbiota of healthy individuals from worldwide populations with various industrialization levels. PLoS ONE, 2019, 14, e0211139.	2.5	44
18	Pneumocystis Species Co-evolution: State-of-the-Art Review. OBM Genetics, 2019, 3, 1-1.	0.4	3

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19	Genetic diversity and evolution of <i>Pneumocystis</i> fungi infecting wild Southeast Asian murid rodents. Parasitology, 2018, 145, 885-900.	1.5	17
20	Prevalence and subtype distribution of Blastocystis sp. isolates from poultry in Lebanon and evidence of zoonotic potential. Parasites and Vectors, 2018, 11, 389.	2.5	70
21	Diffusion of <i>Pneumocystis jirovecii</i> in the surrounding air of patients with <i>Pneumocystis</i> colonization: frequency and putative risk factors: Table 1 Medical Mycology, 2017, 55, myw113.	0.7	13
22	Pathogenic Mechanisms of Cryptosporidium and Giardia. Trends in Parasitology, 2017, 33, 561-576.	3.3	148
23	Prevalence, transmission, and host specificity of Cryptosporidium spp. in various animal groups from two French zoos. Parasitology Research, 2017, 116, 3419-3422.	1.6	18
24	Gut Protozoa: Friends or Foes of the Human Gut Microbiota?. Trends in Parasitology, 2017, 33, 925-934.	3.3	136
25	Proteogenomic Insights into the Intestinal Parasite <i>Blastocystis</i> sp. Subtype 4 Isolate WR1. Proteomics, 2017, 17, 1700211.	2.2	5
26	Three-dimensional (3D) culture of adult murine colon as an in vitro model of cryptosporidiosis: Proof of concept. Scientific Reports, 2017, 7, 17288.	3.3	28
27	Targeted metagenomic sequencing data of human gut microbiota associated with Blastocystis colonization. Scientific Data, 2017, 4, 170081.	5.3	8
28	High Prevalence of Pneumocystis jirovecii Dihydropteroate Synthase Gene Mutations in Patients with a First Episode of Pneumocystis Pneumonia in Santiago, Chile, and Clinical Response to Trimethoprim-Sulfamethoxazole Therapy. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	41
29	Molecular Epidemiology of Blastocystis sp. in Various Animal Groups from Two French Zoos and Evaluation of Potential Zoonotic Risk. PLoS ONE, 2017, 12, e0169659.	2.5	135
30	High association of Cryptosporidium spp. infection with colon adenocarcinoma in Lebanese patients. PLoS ONE, 2017, 12, e0189422.	2.5	39
31	Colonization with the enteric protozoa Blastocystis is associated with increased diversity of human gut bacterial microbiota. Scientific Reports, 2016, 6, 25255.	3.3	210
32	Relationship Between <i>Pneumocystis carinii</i> Burden and the Degree of Host Immunosuppression in an Airborne Transmission Experimental Model. Journal of Eukaryotic Microbiology, 2016, 63, 309-317.	1.7	4
33	Draft genome sequence of the intestinal parasite Blastocystis subtype 4-isolate WR1. Genomics Data, 2015, 4, 22-23.	1.3	27
34	Cryptosporidium parvum-induced ileo-caecal adenocarcinoma and WNT signaling in a rodent model. DMM Disease Models and Mechanisms, 2014, 7, 693-700.	2.4	34
35	Aerially transmitted human fungal pathogens: What can we learn from metagenomics and comparative genomics?. Revista Iberoamericana De Micologia, 2014, 31, 54-61.	0.9	7
36	An improved single-round PCR leads to rapid and highly sensitive detection of Pneumocystis spp Medical Mycology, 2014, 52, 841-846.	0.7	6

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37	Complementation of a manganese-dependent superoxide dismutase-deficient yeast strain with Pneumocystis carinii sod2 gene. Fungal Biology, 2014, 118, 885-895.	2.5	0
38	MOLECULAR DETECTION OF <i>HISTOPLASMA CAPSULATUM</i> IN THE LUNG OF A FREE-RANGING COMMON NOCTULE (<i>NYC-TALUS NOCTULA</i>) FROM FRANCE USING THE <i>Hcp100</i> GENE. Journal of Zoo and Wildlife Medicine, 2013, 44, 15-20.	0.6	10
39	Near-Universal Prevalence of Pneumocystis and Associated Increase in Mucus in the Lungs of Infants With Sudden Unexpected Death. Clinical Infectious Diseases, 2013, 56, 171-179.	5.8	58
40	Growth and Airborne Transmission of Cell-Sorted Life Cycle Stages of Pneumocystis carinii. PLoS ONE, 2013, 8, e79958.	2.5	33
41	Evidence of Airborne Excretion of Pneumocystis carinii during Infection in Immunocompetent Rats. Lung Involvement and Antibody Response. PLoS ONE, 2013, 8, e62155.	2.5	12
42	Characterizing Pneumocystis in the Lungs of Bats: Understanding Pneumocystis Evolution and the Spread of Pneumocystis Organisms in Mammal Populations. Applied and Environmental Microbiology, 2012, 78, 8122-8136.	3.1	29
43	Pneumocystis Molecular Phylogeny: A Way to Understand Both Pneumocystosis Natural History and Host Taxonomy. , 2012, , 149-178.		4
44	Mixed human intra- and inter-subtype infections with the parasite Blastocystis sp Parasitology International, 2012, 61, 719-722.	1.3	24
45	The Airway Microbiota in Cystic Fibrosis: A Complex Fungal and Bacterial Community—Implications for Therapeutic Management. PLoS ONE, 2012, 7, e36313.	2.5	312
46	Microplanktonic Community Structure in a Coastal System Relative to a Phaeocystis Bloom Inferred from Morphological and Tag Pyrosequencing Methods. PLoS ONE, 2012, 7, e39924.	2.5	68
47	Cryptosporidium parvum Infection in SCID Mice Infected with Only One Oocyst: qPCR Assessment of Parasite Replication in Tissues and Development of Digestive Cancer. PLoS ONE, 2012, 7, e51232.	2.5	53
48	Exploring and quantifying fungal diversity in freshwater lake ecosystems using rDNA cloning/sequencing and SSU tag pyrosequencing. Environmental Microbiology, 2011, 13, 1433-1453.	3.8	161
49	Pneumocystis: from a doubtful unique entity to a group of highly diversified fungal species. FEMS Yeast Research, 2011, 11, 2-17.	2.3	60
50	Molecular subtyping of Blastocystis sp. isolates from symptomatic patients in Italy. Parasitology Research, 2011, 109, 613-619.	1.6	76
51	<i>Pneumocystis carinii</i> and <i>Pneumocystis wakefieldiae</i> in Wild <i>Rattus norvegicus</i> Trapped in Thailand. Journal of Eukaryotic Microbiology, 2010, 57, 213-217.	1.7	20
52	Vertical Transmission of <i>Pneumocystis jirovecii</i> in Humans. Emerging Infectious Diseases, 2009, 15, 125-127.	4.3	37
53	Transmission de Pneumocystis. Journal De Mycologie Medicale, 2009, 19, 276-284.	1.5	7
54	Pneumocystis species, co-evolution and pathogenic power. Infection, Genetics and Evolution, 2008, 8, 708-726.	2.3	103

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55	Nosocomial <i>Pneumocystis jirovecii</i> infections. Parasite, 2008, 15, 359-365.	2.0	41
56	Exploring transplacental transmission of <i>Pneumocystisoryctolagi</i> in first-time pregnant and multiparous rabbit does. Medical Mycology, 2007, 45, 701-707.	0.7	15
57	Pneumocystis oryctolagisp. nov., an uncultured fungus causing pneumonia in rabbits at weaning: review of current knowledge, and description of a new taxon on genotypic, phylogenetic and phenotypic bases. FEMS Microbiology Reviews, 2006, 30, 853-871.	8.6	82
58	Molecular Identification of Tritrichomonas foetus-Like Organisms as Coinfecting Agents of Human Pneumocystis Pneumonia. Journal of Clinical Microbiology, 2006, 44, 1165-1168.	3.9	56
59	Molecular diagnosis ofPneumocystispneumonia. FEMS Immunology and Medical Microbiology, 2005, 45, 405-410.	2.7	73
60	Molecular typing of Pneumocystis jirovecii found in formalin-fixed paraffin-embedded lung tissue sections from sudden infant death victims. Microbiology (United Kingdom), 2004, 150, 1167-1172.	1.8	25
61	Long-Term Colonization with Pneumocystis jirovecii in Hospital Staffs: A Challenge to Prevent Nosocomial Pneumocystosis. Journal of Eukaryotic Microbiology, 2003, 50, 614-615.	1.7	29