

# Asis Khan

## List of Publications by Year in descending order

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Version: 2024-02-01

31  
papers

2,715  
citations

279798

23  
h-index

434195

31  
g-index

33  
all docs

33  
docs citations

33  
times ranked

2047  
citing authors

#	ARTICLE	IF	CITATIONS
1	Life Cycle and Transmission of <i>Cyclospora cayetanensis</i> : Knowns and Unknowns. <i>Microorganisms</i> , 2022, 10, 118.	3.6	7
2	Disruption of <i>Toxoplasma gondii</i> -Induced Host Cell DNA Replication Is Dependent on Contact Inhibition and Host Cell Type. <i>MSphere</i> , 2022, 7, e0016022.	2.9	3
3	Neosporosis: An Overview of Its Molecular Epidemiology and Pathogenesis. <i>Engineering</i> , 2020, 6, 10-19.	6.7	23
4	Molecular epidemiology and population structure of <i>Toxoplasma gondii</i> . , 2020, , 63-116.		9
5	Molecular evidence of hybridization between pig and human <i>Ascaris</i> indicates an interbred species complex infecting humans. <i>ELife</i> , 2020, 9, .	6.0	42
6	Global selective sweep of a highly inbred genome of the cattle parasite <i>Neospora caninum</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 22764-22773.	7.1	20
7	<i>Toxoplasma gondii</i> : Laboratory Maintenance and Growth. <i>Current Protocols in Microbiology</i> , 2017, 44, 20C.1.1-20C.1.17.	6.5	37
8	PopNet: A Markov Clustering Approach to Study Population Genetic Structure. <i>Molecular Biology and Evolution</i> , 2017, 34, 1799-1811.	8.9	5
9	Local admixture of amplified and diversified secreted pathogenesis determinants shapes mosaic <i>Toxoplasma gondii</i> genomes. <i>Nature Communications</i> , 2016, 7, 10147.	12.8	243
10	Genetic Mapping Reveals that Sinefungin Resistance in <i>Toxoplasma gondii</i> Is Controlled by a Putative Amino Acid Transporter Locus That Can Be Used as a Negative Selectable Marker. <i>Eukaryotic Cell</i> , 2015, 14, 140-148.	3.4	29
11	REDHORSE-REcombination and Double crossover detection in Haploid Organisms using next-generation Sequencing data. <i>BMC Genomics</i> , 2015, 16, 133.	2.8	5
12	Rhoptry Proteins ROP5 and ROP18 Are Major Murine Virulence Factors in Genetically Divergent South American Strains of <i>Toxoplasma gondii</i> . <i>PLoS Genetics</i> , 2015, 11, e1005434.	3.5	99
13	Geographic Separation of Domestic and Wild Strains of <i>Toxoplasma gondii</i> in French Guiana Correlates with a Monomorphic Version of Chromosome 1a. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e3182.	3.0	39
14	NextGen sequencing reveals short double crossovers contribute disproportionately to genetic diversity in <i>Toxoplasma gondii</i> . <i>BMC Genomics</i> , 2014, 15, 1168.	2.8	17
15	Globally diverse <i>Toxoplasma gondii</i> isolates comprise six major clades originating from a small number of distinct ancestral lineages. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 5844-5849.	7.1	349
16	Genetic analyses of atypical <i>Toxoplasma gondii</i> strains reveal a fourth clonal lineage in North America. <i>International Journal for Parasitology</i> , 2011, 41, 645-655.	3.1	263
17	Virulence differences in <i>Toxoplasma</i> mediated by amplification of a family of polymorphic pseudokinases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 9631-9636.	7.1	230
18	A Monomorphic Haplotype of Chromosome 1a Is Associated with Widespread Success in Clonal and Nonclonal Populations of <i>Toxoplasma gondii</i> . <i>MBio</i> , 2011, 2, e00228-11.	4.1	45

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19	Genetic diversity of <i>Toxoplasma gondii</i> in animals and humans. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2009, 364, 2749-2761.	4.0	185
20	Selection at a Single Locus Leads to Widespread Expansion of <i>Toxoplasma gondii</i> Lineages That Are Virulent in Mice. <i>PLoS Genetics</i> , 2009, 5, e1000404.	3.5	133
21	Forward Genetics in <i>Toxoplasma gondii</i> Reveals a Family of Rhoptry Kinases That Mediates Pathogenesis. <i>Eukaryotic Cell</i> , 2009, 8, 1085-1093.	3.4	50
22	Phenotypic and Gene Expression Changes among Clonal Type I Strains of <i>Toxoplasma gondii</i> . <i>Eukaryotic Cell</i> , 2009, 8, 1828-1836.	3.4	76
23	Development of a Simple Latex Agglutination Assay for Detection of Shiga Toxin-Producing <i>Escherichia coli</i> (STEC) by Using Polyclonal Antibody against STEC. <i>Vaccine Journal</i> , 2007, 14, 600-604.	3.1	16
24	<i>Toxoplasma gondii</i> Strains Defective in Oral Transmission Are Also Defective in Developmental Stage Differentiation. <i>Infection and Immunity</i> , 2007, 75, 2580-2590.	2.2	73
25	Genetic Divergence of <i>Toxoplasma gondii</i> Strains Associated with Ocular Toxoplasmosis, Brazil. <i>Emerging Infectious Diseases</i> , 2006, 12, 942-949.	4.3	248
26	A human origin type II strain of <i>Toxoplasma gondii</i> causing severe encephalitis in mice. <i>Microbes and Infection</i> , 2006, 8, 2206-2212.	1.9	34
27	Common inheritance of chromosome Ia associated with clonal expansion of <i>Toxoplasma gondii</i> . <i>Genome Research</i> , 2006, 16, 1119-1125.	5.5	51
28	Composite genome map and recombination parameters derived from three archetypal lineages of <i>Toxoplasma gondii</i> . <i>Nucleic Acids Research</i> , 2005, 33, 2980-2992.	14.5	147
29	Association of Cytolethal Distending Toxin Locus <i>cdtB</i> with Enteropathogenic <i>Escherichia coli</i> Isolated from Patients with Acute Diarrhea in Calcutta, India. <i>Journal of Clinical Microbiology</i> , 2003, 41, 5277-5281.	3.9	43
30	Antibiotic Resistance, Virulence Gene, and Molecular Profiles of Shiga Toxin-Producing <i>Escherichia coli</i> Isolates from Diverse Sources in Calcutta, India. <i>Journal of Clinical Microbiology</i> , 2002, 40, 2009-2015.	3.9	90
31	Prevalence and genetic profiling of virulence determinants of non-O157 Shiga toxin-producing <i>Escherichia coli</i> isolated from cattle, beef, and humans, Calcutta, India. <i>Emerging Infectious Diseases</i> , 2002, 8, 54-62.	4.3	30