Mohammed Z Rahman

List of Publications by Year in descending order

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68 papers

1,220 citations

18 h-index 29 g-index

71 all docs

71 docs citations

71 times ranked

1749 citing authors

#	Article	IF	CITATIONS
1	Association of C-Reactive Protein With Bacterial and Respiratory Syncytial Virus–Associated Pneumonia Among Children Aged <5 Years in the PERCH Study. Clinical Infectious Diseases, 2017, 64, S378-S386.	2.9	84
2	COVID-19 reinfections among naturally infected and vaccinated individuals. Scientific Reports, 2022, 12, 1438.	1.6	79
3	Global burden of acute lower respiratory infection associated with human metapneumovirus in children under 5 years in 2018: a systematic review and modelling study. The Lancet Global Health, 2021, 9, e33-e43.	2.9	71
4	Biofilm Acts as a Microenvironment for Planktonâ€Associated <i>Vibrio cholerae</i> in the Aquatic Environment of Bangladesh. Microbiology and Immunology, 2007, 51, 369-379.	0.7	68
5	Avian influenza surveillance in domestic waterfowl and environment of live bird markets in Bangladesh, 2007–2012. Scientific Reports, 2018, 8, 9396.	1.6	54
6	Efficacy of a Russian-backbone live attenuated influenza vaccine among young children in Bangladesh: a randomised, double-blind, placebo-controlled trial. The Lancet Global Health, 2016, 4, e946-e954.	2.9	46
7	Detection of enteric―and nonâ€enteric adenoviruses in gastroenteritis patients, Bangladesh, 2012â€2015. Journal of Medical Virology, 2018, 90, 677-684.	2.5	43
8	Nipah Virus Contamination of Hospital Surfaces during Outbreaks, Bangladesh, 2013–2014. Emerging Infectious Diseases, 2018, 24, 15-21.	2.0	39
9	Isolation and Full-Genome Characterization of Nipah Viruses from Bats, Bangladesh. Emerging Infectious Diseases, 2019, 25, 166-170.	2.0	32
10	Epidemiology and genetic characterization of Peste des petits ruminants virus in Bangladesh. Veterinary Medicine and Science, 2018, 4, 161-171.	0.6	24
11	Epidemiology and genetic diversity of human astrovirus infection among hospitalized patients with acute diarrhea in Bangladesh from 2010 to 2012. Journal of Clinical Virology, 2013, 58, 612-618.	1.6	23
12	Genetic characterization of a rare bovine-like human VP4 mono-reassortant G6P[8] rotavirus strain detected from an infant in Bangladesh. Infection, Genetics and Evolution, 2013, 19, 120-126.	1.0	23
13	The emergence of SARSâ€CoVâ€2 variants in Dhaka city, Bangladesh. Transboundary and Emerging Diseases, 2021, 68, 3000-3001.	1.3	22
14	Safety of Russian-backbone seasonal trivalent, live-attenuated influenza vaccine in a phase II randomized placebo-controlled clinical trial among children in urban Bangladesh. Vaccine, 2015, 33, 3415-3421.	1.7	21
15	Effectiveness of a Behavior Change Intervention with Hand Sanitizer Use and Respiratory Hygiene in Reducing Laboratory-Confirmed Influenza among Schoolchildren in Bangladesh: A Cluster Randomized Controlled Trial. American Journal of Tropical Medicine and Hygiene, 2019, 101, 1446-1455.	0.6	21
16	SARS-CoV-2 and influenza virus coinfection among patients with severe acute respiratory infection during the first wave of COVID-19 pandemic in Bangladesh: a hospital-based descriptive study. BMJ Open, 2021, 11, e053768.	0.8	21
17	Maternal vitamin D supplementation during pregnancy and lactation to prevent acute respiratory infections in infancy in Dhaka, Bangladesh (MDARI trial): protocol for a prospective cohort study nested within a randomized controlled trial. BMC Pregnancy and Childbirth, 2016, 16, 309.	0.9	20
18	The Epidemiology of Melioidosis and Its Association with Diabetes Mellitus: A Systematic Review and Meta-Analysis. Pathogens, 2022, 11, 149.	1.2	20

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19	Middle East Respiratory Syndrome Coronavirus Antibodies in Dromedary Camels, Bangladesh, 2015. Emerging Infectious Diseases, 2018, 24, 926-928.	2.0	19
20	The Pattern of Highly Pathogenic Avian Influenza H5N1 Outbreaks in South Asia. Tropical Medicine and Infectious Disease, 2019, 4, 138.	0.9	19
21	Challenges to Evaluating Respiratory Syncytial Virus Mortality in Bangladesh, 2004–2008. PLoS ONE, 2013, 8, e53857.	1.1	19
22	Immunogenicity and Viral Shedding of Russian-Backbone, Seasonal, Trivalent, Live, Attenuated Influenza Vaccine in a Phase II, Randomized, Placebo-Controlled Trial Among Preschool-Aged Children in Urban Bangladesh. Clinical Infectious Diseases, 2019, 69, 777-785.	2.9	18
23	The epidemiology and estimated etiology of pathogens detected from the upper respiratory tract of adults with severe acute respiratory infections in multiple countries, 2014–2015. PLoS ONE, 2020, 15, e0240309.	1.1	18
24	Association of Biosecurity and Hygiene Practices with Environmental Contamination with Influenza A Viruses in Live Bird Markets, Bangladesh. Emerging Infectious Diseases, 2020, 26, 2087-2096.	2.0	18
25	Inference of Nipah virus evolution, 1999–2015. Virus Evolution, 2021, 7, veaa062.	2.2	18
26	Genome Sequence of a SARS-CoV-2 Strain from Bangladesh That Is Nearly Identical to United Kingdom SARS-CoV-2 Variant B.1.1.7. Microbiology Resource Announcements, 2021, 10, .	0.3	18
27	Variation of Toxigenic <i>Vibrio cholerae </i> O1 in the Aquatic Environment of Bangladesh and Its Correlation with the Clinical Strains. Microbiology and Immunology, 2004, 48, 773-777.	0.7	17
28	Novel intergenotype human norovirus recombinant GII.16/GII.3 in Bangladesh. Infection, Genetics and Evolution, 2013, 20, 325-329.	1.0	17
29	Epidemiologic and Genotypic Distribution of Noroviruses Among Children With Acute Diarrhea and Healthy Controls in a Low-income Rural Setting. Clinical Infectious Diseases, 2019, 69, 505-513.	2.9	17
30	Molecular characterization of group A rotavirus from rhesus macaques (<i>Macaca mulatta ⟨i⟩) at human–wildlife interfaces in Bangladesh. Transboundary and Emerging Diseases, 2020, 67, 956-966.</i>	1.3	17
31	High incidence of reassortant G9P[4] rotavirus strain in Bangladesh: Fully heterotypic from vaccine strains. Journal of Clinical Virology, 2013, 58, 755-756.	1.6	16
32	Genetic diversity of Nipah virus in Bangladesh. International Journal of Infectious Diseases, 2021, 102, 144-151.	1.5	15
33	Antigenic heterogeneity of capsid protein VP1 in foot-and-mouth disease virus (FMDV) serotype Asia1. Advances and Applications in Bioinformatics and Chemistry, 2013, 6, 37.	1.6	14
34	Efficacy of trivalent influenza vaccine against laboratory-confirmed influenza among young children in a randomized trial in Bangladesh. Vaccine, 2017, 35, 6967-6976.	1.7	14
35	Detection of highly pathogenic avian influenza A(H5N6) viruses in waterfowl in Bangladesh. Virology, 2019, 534, 36-44.	1.1	13
36	Major zoonotic diseases of public health importance in Bangladesh. Veterinary Medicine and Science, 2021, 7, 1199-1210.	0.6	13

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37	Genomics, social media and mobile phone data enable mapping of SARS-CoV-2 lineages to inform health policy in Bangladesh. Nature Microbiology, 2021, 6, 1271-1278.	5.9	13
38	Serological Cross-Reaction Between O-Antigens of Shigella dysenteriae Type 4 and an Environmental Escherichia albertii Isolate. Current Microbiology, 2013, 67, 590-595.	1.0	12
39	Evaluation of recombinase-based isothermal amplification assays for point-of-need detection of SARS-CoV-2 in resource-limited settings. International Journal of Infectious Diseases, 2022, 114, 105-111.	1.5	12
40	Recovery and Characterization of Environmental Variants of Shigella flexneri from Surface Water in Bangladesh. Current Microbiology, 2011, 63, 372-6.	1.0	11
41	Serological Evidence of Coxiella burnetii Infection in Cattle and Goats in Bangladesh. EcoHealth, 2015, 12, 354-358.	0.9	11
42	Serological Cross-Reactivity of Environmental Isolates of Enterobacter, Escherichia, Stenotrophomonas, and Aerococcus with Shigella sppSpecific Antisera. Current Microbiology, 2007, 54, 63-67.	1.0	10
43	An outbreak of classical swine fever in pigs in Bangladesh, 2015. Veterinary Medicine and Science, 2018, 4, 45-52.	0.6	10
44	Identification of Novel Mutations in the N Gene of SARS-CoV-2 That Adversely Affect the Detection of the Virus by Reverse Transcription-Quantitative PCR. Microbiology Spectrum, 2021, 9, e0054521.	1.2	10
45	Organization of the CTX Prophage in Environmental Isolates of <i>Vibrio mimicus</i> . Microbiology and Immunology, 2005, 49, 779-784.	0.7	9
46	Epidemiology and Molecular Characterization of Rotavirus A in Fruit Bats in Bangladesh. EcoHealth, 2020, 17, 398-405.	0.9	9
47	A Case Series Describing the Recurrence of COVID-19 in Patients Who Recovered from Initial Illness in Bangladesh. Tropical Medicine and Infectious Disease, 2021, 6, 41.	0.9	9
48	Epidemiology of the Rhinovirus (RV) in African and Southeast Asian Children: A Case-Control Pneumonia Etiology Study. Viruses, 2021, 13, 1249.	1.5	9
49	Evidence of interspecies <scp>O</scp> antigen gene cluster transfer between <i><scp>S</scp>higella boydii</i> 15 and <i><scp>E</scp>scherichia fergusonii</i> Apmis, 2012, 120, 959-966.	0.9	8
50	The Etiology of Childhood Pneumonia in Bangladesh. Pediatric Infectious Disease Journal, 2021, 40, S79-S90.	1.1	8
51	Wastewater surveillance of SARS-CoV-2 in Bangladesh: Opportunities and challenges. Current Opinion in Environmental Science and Health, 2022, 27, 100334.	2.1	8
52	Genetic characterization of human metapneumovirus identified through community and facilityâ€based surveillance of infants in Dhaka, Bangladesh. Journal of Medical Virology, 2019, 91, 549-554.	2.5	7
53	Clinical evaluation of SARS-CoV-2 antigen-based rapid diagnostic test kit for detection of COVID-19 cases in Bangladesh. Heliyon, 2021, 7, e08455.	1.4	7
54	Prevalence of a Novel Division-Level Bacterial Lineage in Lake Dhanmondi, Dhaka, Bangladesh, as Revealed by Deep Sequencing of 16S rRNA Gene Amplicons. Current Microbiology, 2012, 65, 356-360.	1.0	5

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55	Outbreak of diarrhoea in piglets caused by novel rotavirus genotype G4P[49] in northâ€western district of Bangladesh, February 2014. Transboundary and Emerging Diseases, 2020, 67, 442-449.	1.3	5
56	Upper Respiratory Tract Co-detection of Human Endemic Coronaviruses and High-density Pneumococcus Associated With Increased Severity Among HIV-Uninfected Children Under 5 Years Old in the PERCH Study. Pediatric Infectious Disease Journal, 2021, 40, 503-512.	1.1	5
57	Genome Sequences of SARS-CoV-2 Sublineage B.1.617.2 Strains from 12 Children in Chattogram, Bangladesh. Microbiology Resource Announcements, 2021, 10, e0091221.	0.3	5
58	Factors influencing the performance of rapid SARSâ€CoVâ€2 antigen tests under field condition. Journal of Clinical Laboratory Analysis, 2022, 36, e24203.	0.9	5
59	Effects of Maternal Vitamin D Supplementation During Pregnancy and Lactation on Infant Acute Respiratory Infections: Follow-up of a Randomized Trial in Bangladesh. Journal of the Pediatric Infectious Diseases Society, 2021, 10, 901-909.	0.6	4
60	Nipah Virus Detection at Bat Roosts after Spillover Events, Bangladesh, 2012–2019. Emerging Infectious Diseases, 2022, 28, 1384-1392.	2.0	3
61	An Environmental Escherichia albertii Strain, DM104, Induces Protective Immunity to Shigella dysenteriae in Guinea Pig Eye Model. Current Microbiology, 2014, 68, 642-647.	1.0	2
62	Protection against shigellosis caused by Shigella dysenteriae serotype 4 in guinea pigs using Escherichia albertii DM104 as a live vaccine candidate strain. Acta Microbiologica Et Immunologica Hungarica, 2017, 64, 151-164.	0.4	2
63	Viral etiology of acute gastroenteritis among Forcibly Displaced Myanmar Nationals and adjacent host population in Bangladesh. Journal of Infectious Diseases, 2021, , .	1.9	2
64	A Case Report: Genetically Distinct Severe Acute Respiratory Syndrome Coronavirus-2 Variant Causing Reinfection. Frontiers in Microbiology, 2021, 12, 792514.	1.5	2
65	Detection and Molecular Characterization of Canine Alphacoronavirus in Free-Roaming Dogs, Bangladesh. Viruses, 2022, 14, 67.	1.5	1
66	Coding-Complete Sequence of a SARS-CoV-2 B.1.1.25 Lineage Obtained from an 8-Day-Old Deceased Neonate. Microbiology Resource Announcements, 2021, 10, e0075621.	0.3	0
67	Effect of maternal vitamin D supplementation on nasal pneumococcal acquisition, carriage dynamics and carriage density in infants in Dhaka, Bangladesh. BMC Infectious Diseases, 2022, 22, 52.	1.3	0
68	Epidemiology of Group A rotavirus in rodents and shrews in Bangladesh. Veterinary Research Communications, 2022, , $1.$	0.6	0