

Shah M Faruque

List of Publications by Year in descending order

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117
papers

9,382
citations

34016

52
h-index

42291

92
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118
all docs

118
docs citations

118
times ranked

5643
citing authors

#	ARTICLE	IF	CITATIONS
1	Environmental prevalence of toxigenic <i>Vibrio cholerae</i> O1 in Bangladesh coincides with <i>V. cholerae</i> non-O1 non-O139 genetic variants which overproduce autoinducer-2. <i>PLoS ONE</i> , 2021, 16, e0254068.	1.1	3
2	In silico comparative genomics of SARS-CoV-2 to determine the source and diversity of the pathogen in Bangladesh. <i>PLoS ONE</i> , 2021, 16, e0245584.	1.1	27
3	<i>Vibrio cholerae</i> strains with inactivated <i>cqsS</i> gene overproduce autoinducer-2 which enhances resuscitation of dormant environmental <i>V. cholerae</i> . <i>PLoS ONE</i> , 2019, 14, e0223226.	1.1	4
4	Toxigenic properties and <i>stx</i> phage characterization of <i>Escherichia coli</i> O157 isolated from animal sources in a developing country setting. <i>BMC Microbiology</i> , 2018, 18, 98.	1.3	13
5	Environmental Spread of New Delhi Metallo- β -Lactamase-1-Producing Multidrug-Resistant Bacteria in Dhaka, Bangladesh. <i>Applied and Environmental Microbiology</i> , 2017, 83, .	1.4	67
6	Analysis of the CRISPR-Cas system in bacteriophages active on epidemic strains of <i>Vibrio cholerae</i> in Bangladesh. <i>Scientific Reports</i> , 2017, 7, 14880.	1.6	31
7	Environmental bacteriophages active on biofilms and planktonic forms of toxigenic <i>Vibrio cholerae</i> : Potential relevance in cholera epidemiology. <i>PLoS ONE</i> , 2017, 12, e0180838.	1.1	22
8	Quorum Regulated Resistance of <i>Vibrio cholerae</i> against Environmental Bacteriophages. <i>Scientific Reports</i> , 2016, 6, 37956.	1.6	70
9	Safe distances between groundwater-based water wells and pit latrines at different hydrogeological conditions in the Ganges Atrai floodplains of Bangladesh. <i>Journal of Health, Population and Nutrition</i> , 2016, 35, 26.	0.7	23
10	Faecal contamination of commuters' hands in main vehicle stations in Dhaka city, Bangladesh. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2016, 110, 367-372.	0.7	5
11	Species-wide whole genome sequencing reveals historical global spread and recent local persistence in <i>Shigella flexneri</i> . <i>ELife</i> , 2015, 4, e07335.	2.8	94
12	Intercontinental dissemination of azithromycin-resistant shigellosis through sexual transmission: a cross-sectional study. <i>Lancet Infectious Diseases</i> , The, 2015, 15, 913-921.	4.6	204
13	Suppression of Virulence of Toxigenic <i>Vibrio cholerae</i> by Anethole through the Cyclic AMP (cAMP)-cAMP Receptor Protein Signaling System. <i>PLoS ONE</i> , 2015, 10, e0137529.	1.1	21
14	Fluoroquinolone Resistance Mechanisms of <i>Shigella flexneri</i> Isolated in Bangladesh. <i>PLoS ONE</i> , 2014, 9, e102533.	1.1	46
15	Genetic and phenotypic analysis of <i>Vibrio cholerae</i> non-O1, non-O139 isolated from German and Austrian patients. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2014, 33, 767-778.	1.3	69
16	Phenotypic and genotypic characteristics of <i>Vibrio cholerae</i> O1 isolated from the Sierra Leone cholera outbreak in 2012. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2014, 108, 715-720.	0.7	7
17	Molecular characterization of cytolethal distending toxin gene-positive <i>Escherichia coli</i> from healthy cattle and swine in Nara, Japan. <i>BMC Microbiology</i> , 2014, 14, 97.	1.3	48
18	RS1 Satellite Phage Promotes Diversity of Toxigenic <i>Vibrio cholerae</i> by Driving CTX Prophage Loss and Elimination of Lysogenic Immunity. <i>Infection and Immunity</i> , 2014, 82, 3636-3643.	1.0	14

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19	In vitro and in vivo bactericidal activity of <i>Vitex negundo</i> leaf extract against diverse multidrug resistant enteric bacterial pathogens. <i>Asian Pacific Journal of Tropical Medicine</i> , 2013, 6, 352-359.	0.4	17
20	Genetic diversity and antibiotic resistance in <i>Escherichia coli</i> from environmental surface water in Dhaka City, Bangladesh. <i>Diagnostic Microbiology and Infectious Disease</i> , 2013, 76, 222-226.	0.8	26
21	Role of Phages in the Epidemiology of Cholera. <i>Current Topics in Microbiology and Immunology</i> , 2013, 379, 165-180.	0.7	12
22	Coupling mutagenesis and parallel deep sequencing to probe essential residues in a genome or gene. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E848-57.	3.3	36
23	Quorum-sensing autoinducers resuscitate dormant <i>Vibrio cholerae</i> in environmental water samples. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 9926-9931.	3.3	69
24	Changing Trends in the Prevalence of <i>Shigella</i> Species: Emergence of Multi-Drug Resistant <i>Shigella sonnei</i> Biotype g in Bangladesh. <i>PLoS ONE</i> , 2013, 8, e82601.	1.1	79
25	Molecular Characterizations of Cytolethal Distending Toxin Produced by <i>Providencia alcalifaciens</i> Strains Isolated from Patients with Diarrhea. <i>Infection and Immunity</i> , 2012, 80, 1323-1332.	1.0	37
26	Phage-bacterial interactions in the evolution of toxigenic <i>Vibrio cholerae</i> . <i>Virulence</i> , 2012, 3, 556-565.	1.8	153
27	Phase Variable O Antigen Biosynthetic Genes Control Expression of the Major Protective Antigen and Bacteriophage Receptor in <i>Vibrio cholerae</i> O1. <i>PLoS Pathogens</i> , 2012, 8, e1002917.	2.1	138
28	Molecular Epidemiology of Toxigenic <i>Vibrio cholerae</i> . , 2011, , 115-127.		2
29	Phenotypic, Genotypic, and Antibiotic Sensitivity Patterns of Strains Isolated from the Cholera Epidemic in Zimbabwe. <i>Journal of Clinical Microbiology</i> , 2011, 49, 2325-2327.	1.8	33
30	An experimental study of phage mediated bactericidal selection & emergence of the El Tor <i>Vibrio cholerae</i> . <i>Indian Journal of Medical Research</i> , 2011, 133, 218-24.	0.4	3
31	El Tor cholera with severe disease: a new threat to Asia and beyond. <i>Epidemiology and Infection</i> , 2010, 138, 347-352.	1.0	121
32	Development of simple and rapid PCR fingerprinting methods for <i>Vibrio cholerae</i> on the basis of genetic diversity of the superintegron. <i>Journal of Applied Microbiology</i> , 2010, 109, 304-312.	1.4	16
33	Satellite phage ϕ T enables toxigenic conversion by CTX phage through dif site alteration. <i>Nature</i> , 2010, 467, 982-985.	13.7	104
34	The Cyclic AMP (cAMP)-cAMP Receptor Protein Signaling System Mediates Resistance of <i>Vibrio cholerae</i> O1 Strains to Multiple Environmental Bacteriophages. <i>Applied and Environmental Microbiology</i> , 2010, 76, 4233-4240.	1.4	18
35	Quorum-regulated biofilms enhance the development of conditionally viable, environmental <i>Vibrio cholerae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 1588-1593.	3.3	74
36	Functional Analysis of VopF Activity Required for Colonization in <i>Vibrio cholerae</i> . <i>MBio</i> , 2010, 1, .	1.8	45

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37	Prevalence and characteristics of cytolethal distending toxin-producing <i>Escherichia coli</i> from children with diarrhea in Japan. <i>Microbiology and Immunology</i> , 2009, 53, 206-215.	0.7	44
38	Effect of Phage on the Infectivity of <i>Vibrio cholerae</i> and Emergence of Genetic Variants. <i>Infection and Immunity</i> , 2008, 76, 5266-5273.	1.0	60
39	Acquisition of classical CTX prophage from <i>Vibrio cholerae</i> O141 by El Tor strains aided by lytic phages and chitin-induced competence. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 11951-11956.	3.3	79
40	Distribution of Genes for Virulence and Ecological Fitness among Diverse <i>Vibrio cholerae</i> Population in a Cholera Endemic Area: Tracking the Evolution of Pathogenic Strains. <i>DNA and Cell Biology</i> , 2008, 27, 347-355.	0.9	85
41	Molecular Epidemiological Studies of <i>Vibrio cholerae</i> in Bengal Region. <i>Biocontrol Science</i> , 2008, 13, 1-8.	0.2	8
42	Seasonal Cholera from Multiple Small Outbreaks, Rural Bangladesh. <i>Emerging Infectious Diseases</i> , 2008, 14, 831-833.	2.0	73
43	An inducible lambdaoid prophage encoding cytolethal distending toxin (Cdt-I) and a type III effector protein in enteropathogenic <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 14483-14488.	3.3	57
44	Genomic analysis of the Mozambique strain of <i>Vibrio cholerae</i> O1 reveals the origin of El Tor strains carrying classical CTX prophage. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 5151-5156.	3.3	101
45	Peru-15, a live attenuated oral cholera vaccine, is safe and immunogenic in Bangladeshi toddlers and infants. <i>Vaccine</i> , 2007, 25, 231-238.	1.7	97
46	CYTOLETHAL DISTENDING TOXIN (CDT): GENETIC DIVERSITY, STRUCTURE AND ROLE IN DIARRHEAL DISEASE. <i>Toxin Reviews</i> , 2006, 25, 61-88.	1.5	51
47	Cholera stool bacteria repress chemotaxis to increase infectivity. <i>Molecular Microbiology</i> , 2006, 60, 417-426.	1.2	75
48	Transmissibility of cholera: In vivo-formed biofilms and their relationship to infectivity and persistence in the environment. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 6350-6355.	3.3	301
49	An Improved Technique for Isolation of Environmental <i>Vibrio cholerae</i> with Epidemic Potential: Monitoring the Emergence of a Multiple-Antibiotic-Resistant Epidemic Strain in Bangladesh. <i>Journal of Infectious Diseases</i> , 2006, 193, 1029-1036.	1.9	52
50	Cholera Due to Altered El Tor Strains of <i>Vibrio cholerae</i> O1 in Bangladesh. <i>Journal of Clinical Microbiology</i> , 2006, 44, 4211-4213.	1.8	222
51	Phenotypic and Molecular Characteristics of <i>Escherichia coli</i> Isolated from Aquatic Environment of Bangladesh. <i>Microbiology and Immunology</i> , 2006, 50, 359-370.	0.7	17
52	Modeling the role of bacteriophage in the control of cholera outbreaks. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 4652-4657.	3.3	173
53	<i>Vibrios.</i> , 2006, , 332-372.		5
54	Randomized, Controlled Study of the Safety and Immunogenicity of Peru-15, a Live Attenuated Oral Vaccine Candidate for Cholera, in Adult Volunteers in Bangladesh. <i>Journal of Infectious Diseases</i> , 2005, 192, 573-579.	1.9	78

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55	Self-limiting nature of seasonal cholera epidemics: Role of host-mediated amplification of phage. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 6119-6124.	3.3	213
56	Transcriptional Profiling of <i>Vibrio cholerae</i> Recovered Directly from Patient Specimens during Early and Late Stages of Human Infection. Infection and Immunity, 2005, 73, 4488-4493.	1.0	103
57	Genomic Sequence and Receptor for the <i>Vibrio cholerae</i> Phage KSF-1 ϕ : Evolutionary Divergence among Filamentous Vibriophages Mediating Lateral Gene Transfer. Journal of Bacteriology, 2005, 187, 4095-4103.	1.0	41
58	Reduction in Capsular Content and Enhanced Bacterial Susceptibility to Serum Killing of <i>Vibrio cholerae</i> O139 Associated with the 2002 Cholera Epidemic in Bangladesh. Infection and Immunity, 2005, 73, 6577-6583.	1.0	22
59	Genomic characterization of non-O1, non-O139 <i>Vibrio cholerae</i> reveals genes for a type III secretion system. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 3465-3470.	3.3	217
60	Seasonal epidemics of cholera inversely correlate with the prevalence of environmental cholera phages. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 1702-1707.	3.3	317
61	Free-Living to Freewheeling: The Evolution of <i>Vibrio cholerae</i> from Innocence to Infamy. , 2004, , 198-221.		0
62	Diverse CTX Phages among Toxigenic <i>Vibrio cholerae</i> O1 and O139 Strains Isolated between 1994 and 2002 in an Area Where Cholera is Endemic in Bangladesh. Journal of Clinical Microbiology, 2004, 42, 5854-5856.	1.8	68
63	Genetic diversity and virulence potential of environmental <i>Vibrio cholerae</i> population in a cholera-endemic area. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 2123-2128.	3.3	182
64	Molecular characterisation of rough strains of <i>Vibrio cholerae</i> isolated from diarrhoeal cases in India and their comparison to smooth strains. FEMS Microbiology Letters, 2004, 232, 23-30.	0.7	12
65	Genetics of Stress Adaptation and Virulence in Toxigenic <i>Vibrio cholerae</i> . DNA and Cell Biology, 2004, 23, 723-741.	0.9	28
66	Pathogenicity islands and phages in <i>Vibrio cholerae</i> evolution. Trends in Microbiology, 2003, 11, 505-510.	3.5	179
67	<i>Shigella dysenteriae</i> Type 1-Specific Bacteriophage from Environmental Waters in Bangladesh. Applied and Environmental Microbiology, 2003, 69, 7028-7031.	1.4	31
68	A 4 $\frac{1}{2}$ -Year Study of the Epidemiology of <i>Vibrio cholerae</i> in Four Rural Areas of Bangladesh. Journal of Infectious Diseases, 2003, 187, 96-101.	1.9	189
69	CTX ϕ -independent production of the RS1 satellite phage by <i>Vibrio cholerae</i> . Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 1280-1285.	3.3	34
70	ToxR regulon of <i>Vibrio cholerae</i> and its expression in vibrios shed by cholera patients. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 2801-2806.	3.3	177
71	Examination of Diverse Toxin-Coregulated Pilus-Positive <i>Vibrio cholerae</i> Strains Fails To Demonstrate Evidence for <i>Vibrio</i> Pathogenicity Island Phage. Infection and Immunity, 2003, 71, 2993-2999.	1.0	51
72	Emergence and evolution of <i>Vibrio cholerae</i> O139. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 1304-1309.	3.3	166

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73	Reemergence of Epidemic <i>Vibrio cholerae</i> O139, Bangladesh. <i>Emerging Infectious Diseases</i> , 2003, 9, 1116-1122.	2.0	86
74	Pathogenic Potential of Environmental <i>Vibrio cholerae</i> Strains Carrying Genetic Variants of the Toxin-Coregulated Pilus Pathogenicity Island. <i>Infection and Immunity</i> , 2003, 71, 1020-1025.	1.0	50
75	Prevalence of the Pandemic Genotype of <i>Vibrio parahaemolyticus</i> in Dhaka, Bangladesh, and Significance of Its Distribution across Different Serotypes. <i>Journal of Clinical Microbiology</i> , 2002, 40, 284-286.	1.8	99
76	Cloning and Characterization of Genes Encoding Homologues of the B Subunit of Cholera Toxin and the <i>Escherichia coli</i> Heat-Labile Enterotoxin from Clinical Isolates of <i>Citrobacter freundii</i> and <i>E. coli</i> . <i>Infection and Immunity</i> , 2002, 70, 7153-7155.	1.0	25
77	New Variants of <i>Vibrio cholerae</i> O1 Biotype El Tor with Attributes of the Classical Biotype from Hospitalized Patients with Acute Diarrhea in Bangladesh. <i>Journal of Clinical Microbiology</i> , 2002, 40, 3296-3299.	1.8	258
78	Isolation of <i>Shigella dysenteriae</i> Type 1 and <i>S. flexneri</i> Strains from Surface Waters in Bangladesh: Comparative Molecular Analysis of Environmental <i>Shigella</i> Isolates versus Clinical Strains. <i>Applied and Environmental Microbiology</i> , 2002, 68, 3908-3913.	1.4	75
79	RS1 Element of <i>Vibrio cholerae</i> Can Propagate Horizontally as a Filamentous Phage Exploiting the Morphogenesis Genes of CTX ϕ . <i>Infection and Immunity</i> , 2002, 70, 163-170.	1.0	62
80	Molecular Ecology of Toxigenic <i>Vibrio cholerae</i> . <i>Microbiology and Immunology</i> , 2002, 46, 59-66.	0.7	86
81	Molecular Analysis of Antibiotic Resistance Gene Clusters in <i>Vibrio cholerae</i> O139 and O1 SXT Constins. <i>Antimicrobial Agents and Chemotherapy</i> , 2001, 45, 2991-3000.	1.4	300
82	Diminished Diarrheal Response to <i>Vibrio cholerae</i> Strains Carrying the Replicative Form of the CTX ϕ Genome instead of CTX ϕ Lysogens in Adult Rabbits. <i>Infection and Immunity</i> , 2001, 69, 6084-6090.	1.0	22
83	The efficacy of bismuth subsalicylate in the treatment of acute diarrhoea and the prevention of persistent diarrhoea. <i>Acta Paediatrica, International Journal of Paediatrics</i> , 2001, 90, 605-610.	0.7	12
84	Immune Response of Bangladeshi Children With Acute Diarrhea Who Subsequently Have Persistent Diarrhea. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2000, 31, 528-535.	0.9	8
85	Genomic diversity among <i>Vibrio cholerae</i> O139 strains isolated in Bangladesh and India between 1992 and 1998. <i>FEMS Microbiology Letters</i> , 2000, 184, 279-284.	0.7	42
86	Virulence Genes in Environmental Strains of <i>Vibrio cholerae</i> . <i>Applied and Environmental Microbiology</i> , 2000, 66, 4022-4028.	1.4	146
87	Sunlight-Induced Propagation of the Lysogenic Phage Encoding Cholera Toxin. <i>Infection and Immunity</i> , 2000, 68, 4795-4801.	1.0	58
88	Lysogenic Conversion of Environmental <i>Vibrio mimicus</i> Strains by CTX ϕ . <i>Infection and Immunity</i> , 1999, 67, 5723-5729.	1.0	62
89	Case-Control Study of Enteropathogens Associated with Childhood Diarrhea in Dhaka, Bangladesh. <i>Journal of Clinical Microbiology</i> , 1999, 37, 3458-3464.	1.8	268
90	Molecular Characterization of a New Ribotype of <i>Vibrio cholerae</i> O139 Bengal Associated with an Outbreak of Cholera in Bangladesh. <i>Journal of Clinical Microbiology</i> , 1999, 37, 1313-1318.	1.8	44

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91	Aetiological, Clinical and Epidemiological Characteristics of a Seasonal Peak of Diarrhoea in Dhaka, Bangladesh. <i>Scandinavian Journal of Infectious Diseases</i> , 1998, 30, 393-396.	1.5	20
92	Epidemiology, Genetics, and Ecology of Toxigenic <i>Vibrio cholerae</i> . <i>Microbiology and Molecular Biology Reviews</i> , 1998, 62, 1301-1314.	2.9	808
93	Small Bowel and Fecal Microbiology in Children Suffering from Persistent Diarrhea in Bangladesh. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 1998, 26, 9-15.	0.9	23
94	Analysis of Clinical and Environmental Strains of Nontoxigenic <i>Vibrio cholerae</i> for Susceptibility to CTX ϕ : Molecular Basis for Origination of New Strains with Epidemic Potential. <i>Infection and Immunity</i> , 1998, 66, 5819-5825.	1.0	97
95	Induction of the Lysogenic Phage Encoding Cholera Toxin in Naturally Occurring Strains of Toxigenic <i>Vibrio cholerae</i> O1 and O139. <i>Infection and Immunity</i> , 1998, 66, 3752-3757.	1.0	82
96	Phenotypic and genotypic changes in <i>Vibrio cholerae</i> O139 Bengal. <i>Journal of Clinical Microbiology</i> , 1997, 35, 2588-2592.	1.8	34
97	Emergence of a new clone of toxigenic <i>Vibrio cholerae</i> O1 biotype El Tor displacing <i>V. cholerae</i> O139 Bengal in Bangladesh. <i>Journal of Clinical Microbiology</i> , 1997, 35, 624-630.	1.8	70
98	Molecular analysis of toxigenic <i>Vibrio cholerae</i> O139 Bengal strains isolated in Bangladesh between 1993 and 1996: evidence for emergence of a new clone of the Bengal vibrios. <i>Journal of Clinical Microbiology</i> , 1997, 35, 2299-2306.	1.8	62
99	Controlled study of cytolethal distending toxin-producing <i>Escherichia coli</i> infections in Bangladeshi children. <i>Journal of Clinical Microbiology</i> , 1996, 34, 717-719.	1.8	62
100	Demonstration of a lack of synergistic effect of rotavirus with other diarrheal pathogens on severity of diarrhea in children. <i>Journal of Clinical Microbiology</i> , 1996, 34, 1340-1342.	1.8	15
101	Hypotonic oral rehydration solution in acute diarrhoea: a controlled clinical trial. <i>Acta Paediatrica, International Journal of Paediatrics</i> , 1995, 84, 289-293.	0.7	29
102	Molecular epidemiology of toxigenic <i>Vibrio cholerae</i> in Bangladesh studied by numerical analysis of rRNA gene restriction patterns. <i>Journal of Clinical Microbiology</i> , 1995, 33, 2833-2838.	1.8	54
103	Controlled study of <i>Escherichia coli</i> diarrheal infections in Bangladeshi children. <i>Journal of Clinical Microbiology</i> , 1995, 33, 973-977.	1.8	153
104	Hemagglutinating properties of enteroaggregative <i>Escherichia coli</i> . <i>Journal of Clinical Microbiology</i> , 1994, 32, 510-514.	1.8	28
105	Molecular analysis of rRNA and cholera toxin genes carried by the new epidemic strain of toxigenic <i>Vibrio cholerae</i> O139 synonym Bengal. <i>Journal of Clinical Microbiology</i> , 1994, 32, 1050-1053.	1.8	63
106	A monoclonal antibody to <i>Shigella dysenteriae</i> serotype 13 cross-reacting with Shiga toxin. <i>FEMS Microbiology Letters</i> , 1993, 107, 343-347.	0.7	6
107	Large outbreak of clinical cholera due to <i>Vibrio cholerae</i> non-O1 in Bangladesh. <i>Lancet</i> , The, 1993, 341, 704.	6.3	365
108	Characterisation of <i>Plesiomonas shigelloides</i> strains that share type-specific antigen with <i>Shigella flexneri</i> 6 and common group 1 antigen with <i>Shigella flexneri</i> spp. and <i>Shigella dysenteriae</i> 1. <i>Journal of Medical Microbiology</i> , 1993, 39, 211-217.	0.7	17

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109	Clonal relationships among classical <i>Vibrio cholerae</i> O1 strains isolated between 1961 and 1992 in Bangladesh. <i>Journal of Clinical Microbiology</i> , 1993, 31, 2513-2516.	1.8	50
110	DNA probe analysis of diarrhoeagenic <i>Escherichia coli</i> : detection of EAF-positive isolates of traditional enteropathogenic <i>E. coli</i> serotypes among Bangladeshi paediatric diarrhoea patients. <i>Molecular and Cellular Probes</i> , 1992, 6, 93-99.	0.9	25
111	A comparative study of specific gene probes and standard bioassays to identify diarrhoeagenic <i>Escherichia coli</i> in paediatric patients with diarrhoea in Bangladesh. <i>Journal of Medical Microbiology</i> , 1992, 36, 37-40.	0.7	34
112	Sharing of virulence-associated properties at the phenotypic and genetic levels between enteropathogenic <i>Escherichia coli</i> and <i>Hafnia alvei</i> . <i>Journal of Medical Microbiology</i> , 1992, 37, 310-314.	0.7	146
113	Differentiation of <i>Shigella flexneri</i> strains by rRNA gene restriction patterns. <i>Journal of Clinical Microbiology</i> , 1992, 30, 2996-2999.	1.8	25
114	Comparison of a modified adherence assay with existing assay methods for identification of enteroaggregative <i>Escherichia coli</i> . <i>Journal of Clinical Microbiology</i> , 1992, 30, 1614-1616.	1.8	16
115	An ELISA for the Detection of Localized Adherent Classic Enteropathogenic <i>Escherichia coli</i> Serogroups. <i>Journal of Infectious Diseases</i> , 1991, 164, 986-989.	1.9	16
116	Outbreak of keratoconjunctivitis due to <i>Salmonella weltevreden</i> in a guinea pig colony. <i>Journal of Clinical Microbiology</i> , 1991, 29, 2002-2006.	1.8	6
117	Evolution of <i>Vibrio cholerae</i> and Cholera Epidemics. , 0, , 361-371.		0