List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Molecular Analysis of Vibrio cholerae O1, O139, non-O1, and non-O139 Strains: Clonal Relationships between Clinical and Environmental Isolates. Applied and Environmental Microbiology, 2001, 67, 910-921.	1.4	217
2	Development of a Hexaplex PCR Assay for Rapid Detection of Virulence and Regulatory Genes in Vibrio cholerae and Vibrio mimicus. Journal of Clinical Microbiology, 2002, 40, 4321-4324.	1.8	113
3	Characterization of Listeria monocytogenes isolated from Ganges water, human clinical and milk samples at Varanasi, India. Infection, Genetics and Evolution, 2013, 14, 83-91.	1.0	65
4	Extracellular vesicles: An emerging platform in gram-positive bacteria. Microbial Cell, 2020, 7, 312-322.	1.4	60
5	Virulence and genotypic characterization of Listeria monocytogenes isolated from vegetable and soil samples. BMC Microbiology, 2014, 14, 241.	1.3	58
6	Molecular Characterization of Vibrio cholerae O139 Bengal Isolated from Water and the Aquatic Plant Eichhornia crassipes in the River Ganga, Varanasi, India. Applied and Environmental Microbiology, 2003, 69, 2389-2394.	1.4	53
7	Emergence and dissemination of antibiotic resistance: A global problem. Indian Journal of Medical Microbiology, 2012, 30, 384-390.	0.3	49
8	Attachment of non-culturable toxigenicVibrio cholerae01 and non-01 andAeromonasspp. to the aquatic arthropodGerris spinolaeand plants in the River Ganga, Varanasi. FEMS Immunology and Medical Microbiology, 1995, 12, 113-120.	2.7	48
9	Multiplex PCR for detection of antibiotic resistance genes and the SXT element: application in the characterization of Vibrio cholerae. Journal of Medical Microbiology, 2007, 56, 346-351.	0.7	40
10	<i>>Vibrio cholerae</i> nonâ€O1, nonâ€O139 strains isolated before 1992 from Varanasi, India are multiple drug resistant, contain <i>int</i> SXT, <i>dfr18</i> and <i>aadA5</i> genes. Environmental Microbiology, 2008, 10, 866-873.	1.8	38
11	Biofilm Formation by ica-Negative Ocular Isolates of Staphylococcus haemolyticus. Frontiers in Microbiology, 2018, 9, 2687.	1.5	35
12	Production of haemolysis and its correlation with enterotoxicity in Aeromonas spp Journal of Medical Microbiology, 1992, 37, 262-267.	0.7	30
13	Analysis of 19 Highly Conserved Vibrio cholerae Bacteriophages Isolated from Environmental and Patient Sources Over a Twelve-Year Period. Viruses, 2018, 10, 299.	1.5	29
14	Binding efficiencies of carbohydrate ligands with different genotypes of cholera toxin B: molecular modeling, dynamics and docking simulation studies. Journal of Molecular Modeling, 2012, 18, 1-10.	0.8	28
15	Pregnancy - associated human listeriosis: Virulence and genotypic analysis of Listeria monocytogenes from clinical samples. Journal of Microbiology, 2015, 53, 653-660.	1.3	27
16	Antibiotic Susceptibility, Virulence Pattern, and Typing of Staphylococcus aureus Strains Isolated From Variety of Infections in India. Frontiers in Microbiology, 2019, 10, 2763.	1.5	27
17	Enterotoxicity of clinical and environmental isolates of Aeromonas spp Journal of Medical Microbiology, 1992, 36, 269-272.	0.7	27
18	Septaplex PCR assay for rapid identification ofVibrio choleraeincluding detection of virulence and intSXT genes. FEMS Microbiology Letters, 2006, 265, 208-214.	0.7	24

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19	Comparative structural analysis of two proteins belonging to quorum sensing system in <i>Vibrio cholerae</i> . Journal of Biomolecular Structure and Dynamics, 2012, 30, 574-584.	2.0	24
20	In vitro susceptibility of Vibrio cholerae O1 biotype El Tor strains associated with an outbreak of cholera in Kerala, Southern India. Journal of Antimicrobial Chemotherapy, 2001, 47, 361-362.	1.3	23
21	Characterization of the genetic background of Vibrio cholerae O1 biotype El Tor serotype Inaba strains isolated in Trivandrum, southern India. Journal of Medical Microbiology, 2007, 56, 260-265.	0.7	23
22	Determination of relationships among non-toxigenic Vibrio cholerae O1 biotype El Tor strains from housekeeping gene sequences and ribotype patterns. Research in Microbiology, 2009, 160, 57-62.	1.0	23
23	MULTIPLEX PCR ASSAY FOR THE DETECTION OF ANOPHELES FLUVIATILIS SPECIES COMPLEX, HUMAN HOST PREFERENCE, AND PLASMODIUM FALCIPARUM SPOROZOITE PRESENCE, USING A UNIQUE MOSQUITO PROCESSING METHOD. American Journal of Tropical Medicine and Hygiene, 2007, 76, 837-843.	0.6	22
24	Antibiotic Resistance Profile, Outer Membrane Proteins, Virulence Factors and Genome Sequence Analysis Reveal Clinical Isolates of Enterobacter Are Potential Pathogens Compared to Environmental Isolates. Frontiers in Cellular and Infection Microbiology, 2020, 10, 54.	1.8	21
25	Development and evaluation of hexaplex PCR for rapid detection of methicillin, cadmium/zinc and antiseptic-resistant staphylococci, with simultaneous identification of PVL-positive and -negativeStaphylococcus aureusand coagulase negative staphylococci. FEMS Microbiology Letters, 2014, 352, 114-122.	0.7	20
26	Identification of Major Sequence Types among Multidrug-Resistant Staphylococcus epidermidis Strains Isolated from Infected Eyes and Healthy Conjunctiva. Frontiers in Microbiology, 2017, 8, 1430.	1.5	20
27	<i>Vibrio cholerae</i> infection, novel drug targets and phage therapy. Future Microbiology, 2011, 6, 1199-1208.	1.0	19
28	Identification of Novel Sequence Types among Staphylococcus haemolyticus Isolated from Variety of Infections in India. PLoS ONE, 2016, 11, e0166193.	1.1	19
29	Multidrug-resistant Staphylococcus haemolyticus isolates from infected eyes and healthy conjunctivae in India. Journal of Clobal Antimicrobial Resistance, 2016, 6, 154-159.	0.9	15
30	Application of DNA-based methods in typing <i>Vibrio cholerae</i> strains. Future Microbiology, 2008, 3, 87-96.	1.0	13
31	Phenotypic and genetic characteristics of Vibrio cholerae O1 carrying Haitian ctxB and attributes of classical and El Tor biotypes isolated from Silvassa, India. Journal of Medical Microbiology, 2016, 65, 720-728.	0.7	13
32	Characterization of a Toxigenic Vibrio cholerae O139 Strain Belonging to a New Ribotype and Isolated from a Diarrheal Patient. Journal of Clinical Microbiology, 2002, 40, 4779-4781.	1.8	12
33	Characterization of Vibrio cholerae O139 belonging to multiple ribotypes and isolated from diarrhoeal patients in Kerala, southern India. Infection, Genetics and Evolution, 2011, 11, 454-459.	1.0	12
34	Extracellular <scp>DNA</scp> builds and interacts with vibrio polysaccharide in the biofilm matrix formed by <i>Vibrio cholerae</i> . Environmental Microbiology Reports, 2020, 12, 594-606.	1.0	11
35	Hexaplex PCR for rapid detection of virulence factors. Expert Review of Molecular Diagnostics, 2003, 3, 781-784.	1.5	8
36	A unique methodology for detecting the spread of chloroquine-resistant strains of Plasmodium falciparum, in previously unreported areas, by analyzing anophelines of malaria endemic zones of Orissa, India. Infection, Genetics and Evolution, 2009, 9, 462-467.	1.0	8

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37	Sequence analysis of <i>Vibrio cholerae orfU</i> and <i>zot</i> from preâ€CTXΦ and CTXΦ reveals multiple origin of preâ€CTXΦ and CTXΦ reveals multiple.	1.0	8
38	Effect of storage and sodium chloride on excision of CTXФor pre-CTXФand CTXФfrom Vibrio cholerae O139 strains. Infection, Genetics and Evolution, 2010, 10, 925-930.	1.0	8
39	Vibrio cholerae O1 biotype El Tor strains isolated in 1992 from Varanasi, India harboured El Tor CTXΦ and classical ctxB on the chromosome-I and classical CTXΦ and classical ctxB on the chromosome-II. Environmental Microbiology Reports, 2011, 3, 783-790.	1.0	8
40	Synthesis, Photophysical Studies on Some Anthracene–based Ionic Liquids and their Application as Biofilm Formation Inhibitor. ChemistrySelect, 2017, 2, 2426-2432.	0.7	8
41	Antibiotic resistance in clinical and environmental isolates of Aeromonas spp. Journal of Antimicrobial Chemotherapy, 1994, 33, 368-369.	1.3	7
42	Influence of animal passage on haemolysin and enterotoxin production in Vibrio cholerae O1 biotype El Tor strains. Journal of Medical Microbiology, 1994, 40, 246-251.	0.7	7
43	Homology modelling of a sensor histidine kinase from Aeromonas hydrophila. Journal of Molecular Modeling, 2010, 16, 1003-1009.	0.8	7
44	Genomic profile of antibiotic resistant, classical ctxB positive Vibrio cholerae O1 biotype El Tor isolated in 2003 and 2005 from Puri, India: A retrospective study. Indian Journal of Medical Microbiology, 2016, 34, 462-470.	0.3	7
45	Haemagglutinating activity, serum sensitivity and enterotoxigenicity of Aeromonas spp Journal of Medical Microbiology, 1993, 38, 49-53.	0.7	6
46	A putative heat-labile enterotoxin expressed by strains of Aeromonas media. Journal of Medical Microbiology, 2000, 49, 685-689.	0.7	6
47	Haemolysin produced by Vibrio cholerae non-O1 is not enterotoxic. Journal of Medical Microbiology, 1996, 45, 35-39.	0.7	4
48	Production of the new cholera toxin by environmental isolates of Vibrio cholerae non-O1. Journal of Medical Microbiology, 1996, 45, 31-34.	0.7	4
49	Characterization of cytotoxin-producing Aeromonas caviae (strain HT10) isolated from a sulfur spring in Orissa, India. Letters in Applied Microbiology, 2007, 44, 338-341.	1.0	4
50	Detection and molecular characterization of Vibrio cholerae O1 Inaba biotype El Tor strain in Kerala, S. India. World Journal of Microbiology and Biotechnology, 2008, 24, 433-434.	1.7	4
51	Characteristics of Vibrio cholerae O1 isolated from water of the River Ganga, Varanasi, India. Indian Journal of Medical Microbiology, 2015, 33, 507-515.	0.3	4
52	Analysis of clonally related environmental <i>Vibrio cholerae</i> O1 El Tor isolated before 1992 from Varanasi, India reveals origin of SXTâ€ŀCEs belonging to O139 and O1 serogroups. Environmental Microbiology Reports, 2010, 2, 50-57.	1.0	3
53	Whole-Genome Sequence of Vibrio alginolyticus Isolated from the Mucus of the Coral <i>Fungia danai</i> in the Andaman Sea, India. Genome Announcements, 2016, 4, .	0.8	3
54	Whole-Genome Sequences of <i>Staphylococcus haemolyticus</i> Isolated from Infected Eyes and Healthy Conjunctiva in Bhubaneswar, India. Genome Announcements, 2016, 4, .	0.8	3

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55	Purification and Characterization of Cytotoxin Produced by a Clinical Isolate of Vibrio cholerae O54 TV113. Applied Biochemistry and Biotechnology, 2012, 167, 809-823.	1.4	2
56	Virulence Pattern and Genomic Diversity of Vibrio cholerae O1 and O139 Strains Isolated From Clinical and Environmental Sources in India. Frontiers in Microbiology, 2020, 11, 1838.	1.5	2
57	Mechanism of Antibiotic Resistance and Pathogenicity of Vibrio cholerae. , 2020, , 273-299.		2
58	Enteropathogenicity of and. FEMS Immunology and Medical Microbiology, 1997, 17, 243-250.	2.7	1
59	Characterization of a Cholera Toxin Gene-Negative Clinical Strain of Vibrio cholerae O139 Bengal. Journal of Clinical Microbiology, 2004, 42, 1381-1381.	1.8	1