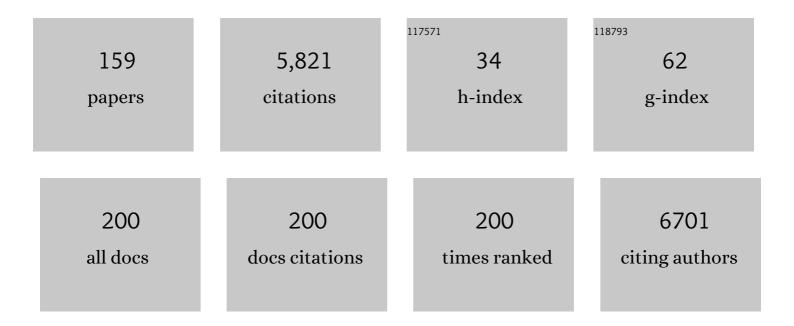
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Neutralization of Variant Under Investigation B.1.617.1 With Sera of BBV152 Vaccinees. Clinical Infectious Diseases, 2022, 74, 366-368.	2.9	200
2	Neutralization Potential of Covishield Vaccinated Individuals Sera Against B.1.617.1. Clinical Infectious Diseases, 2022, 74, 558-559.	2.9	35
3	Predominance of delta variant among the COVID-19 vaccinated and unvaccinated individuals, India, May 2021. Journal of Infection, 2022, 84, 94-118.	1.7	59
4	Community transmission of SARS-CoV-2 with B.1.1.7 lineage in Mumbai, India. Journal of Microbiology, Immunology and Infection, 2022, 55, 1116-1121.	1.5	1
5	Global emergence of SARS-CoV-2 variants: new foresight needed for improved vaccine efficacy. Lancet Infectious Diseases, The, 2022, 22, 298-299.	4.6	8
6	Isolation and characterization of SARS-CoV-2 Beta variant from UAE travelers. Journal of Infection and Public Health, 2022, 15, 182-186.	1.9	22
7	Isolation of SARS-CoV-2 B.1.1.28.2 (P2) variant and pathogenicity comparison with D614G variant in hamster model. Journal of Infection and Public Health, 2022, 15, 164-171.	1.9	7
8	Detection of Zika virus disease in Thiruvananthapuram, Kerala, India 2021 during the second wave of COVIDâ€19 pandemic. Journal of Medical Virology, 2022, 94, 2346-2349.	2.5	7
9	Substantial immune response in Omicron infected breakthrough and unvaccinated individuals against SARS-CoV-2 variants of concern. Journal of Infection, 2022, 84, e80-e81.	1.7	13
10	OUP accepted manuscript. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2022, , .	0.7	1
11	Nipah Virus Outbreak in Kerala State, India Amidst of COVID-19 Pandemic. Frontiers in Public Health, 2022, 10, 818545.	1.3	24
12	Isolation and Genomic Characterization of SARS-CoV-2 Omicron Variant Obtained from Human Clinical Specimens. Viruses, 2022, 14, 461.	1.5	9
13	Elevated neutralization of Omicron with sera of COVID-19 recovered and breakthrough cases vaccinated with Covaxin than two dose naÃīve vaccinees. Journal of Infection, 2022, 84, 834-872.	1.7	13
14	Immune responses against different variants of SARS-CoV-2 including Omicron following 6Âmonths of administration of heterologous prime-boost COVID-19 vaccine. Journal of Travel Medicine, 2022, 29, .	1.4	9
15	Detection and isolation of SARS oVâ€2 EtaÂvariant from the international travelers and local residents of India. Journal of Medical Virology, 2022, 94, 3404-3409.	2.5	6
16	Protective Immunity of the Primary SARS-CoV-2 Infection Reduces Disease Severity Post Re-Infection with Delta Variants in Syrian Hamsters. Viruses, 2022, 14, 596.	1.5	6
17	Antibody responses to Sputnik Vaccination in naÃ⁻ve and COVID 19-recovered vaccine recipients, India. Journal of Travel Medicine, 2022, 29, .	1.4	2
18	Booster dose of the inactivated COVID-19 vaccine BBV152 (Covaxin) enhances the neutralizing antibody response against Alpha, Beta, Delta and Omicron variants of concern. Journal of Travel Medicine, 2022, 29	1.4	20

#	Article	lF	CITATIONS
19	Development of a Reverse Transcription Loop - Mediated Isothermal Amplification [RT-LAMP] as a early rapid detection assay for Crimean Congo Hemorrhagic Fever virus. Acta Tropica, 2022, 231, 106435.	0.9	5
20	SARS-CoV-2 Delta and delta derivatives impact on neutralization of Covishield recipient sera. Journal of Infection, 2022, 84, e36-e38.	1.7	0
21	Clinical, immunological and genomic analysis of the post vaccinated SARS-CoV-2 infected cases with Delta derivatives from Maharashtra, India, 2021. Journal of Infection, 2022, 85, e26-e29.	1.7	2
22	Previous SARS-CoV-2 Infection Status Among the Current RT-PCR-Positive Individuals Affected During the Second Wave of COVID-19 Infections in Chennai, India. Frontiers in Public Health, 2022, 10, 836454.	1.3	0
23	Development and evaluation of indirect antibody ELISA assay for early diagnosis and surveillance of Crimean-Congo hemorrhagic fever infection in humans. Virus Research, 2022, 313, 198717.	1.1	0
24	Pathogenicity of SARS-CoV-2 Omicron (R346K) variant in Syrian hamsters and its cross-neutralization with different variants of concern. EBioMedicine, 2022, 79, 103997.	2.7	29
25	Serosurvey for Nipah virus in bat population of southern part of India. Comparative Immunology, Microbiology and Infectious Diseases, 2022, 85, 101800.	0.7	3
26	A case of breakthrough infection with SARS-CoV-2 Delta derivative and reinfection with Omicron variant in a fully vaccinated health care professional. Journal of Infection, 2022, 85, e15-e17.	1.7	2
27	Reduced neutralizing antibody response in naÃ ⁻ ve Covishield vaccinees against Omicron emphasizes booster vaccination. Journal of Infection, 2022, 85, 90-122.	1.7	4
28	SARS-CoV-2 Kappa Variant Shows Pathogenicity in a Syrian Hamster Model. Vector-Borne and Zoonotic Diseases, 2022, 22, 289-296.	0.6	2
29	Persistence of immunity and impact of third dose of inactivated COVID-19 vaccine against emerging variants. Scientific Reports, 2022, 12, .	1.6	23
30	Delta variant SARS-CoV-2 infections in pediatric cases during the second wave in India. Journal of Microbiology, Immunology and Infection, 2022, 55, 1060-1068.	1.5	2
31	Effectiveness of BBV152/Covaxin and AZD1222/Covishield vaccines against severe COVID-19 and B.1.617.2/Delta variant in India, 2021: a multi-centric hospital-based case-control study. International Journal of Infectious Diseases, 2022, 122, 693-702.	1.5	25
32	Development and Evaluation of Reverse Transcription Loop-Mediated Isothermal Amplification for Rapid and Real-Time Detection of Kyasanur Forest Disease Virus. International Journal of Infectious Diseases, 2021, 112, 346-351.	1.5	4
33	Molecular epidemiology of a familial cluster of SARS-CoV-2 infection during lockdown period in Sant Kabir Nagar, Uttar Pradesh, India. Epidemiology and Infection, 2021, 149, .	1.0	3
34	Authors' response. Indian Journal of Medical Research, 2021, 153, 703.	0.4	0
35	Electron microscopy imaging of SARS-CoV-2 at different temperatures. Indian Journal of Medical Research, 2021, 153, 692-695.	0.4	2
36	Detection of Nipah virus in Pteropus medius in 2019 outbreak from Ernakulam district, Kerala, India. BMC Infectious Diseases, 2021, 21, 162.	1.3	21

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37	Immunogenicity and protective efficacy of BBV152, whole virion inactivated SARS- CoV-2 vaccine candidates in the Syrian hamster model. IScience, 2021, 24, 102054.	1.9	70
38	Immunogenicity and protective efficacy of inactivated SARS-CoV-2 vaccine candidate, BBV152 in rhesus macaques. Nature Communications, 2021, 12, 1386.	5.8	74
39	Ebola virus outbreak preparedness plan for developing Nations: Lessons learnt from affected countries. Journal of Infection and Public Health, 2021, 14, 293-305.	1.9	12
40	Inactivated COVID-19 vaccine BBV152/COVAXIN effectively neutralizes recently emerged B.1.1.7 variant of SARS-CoV-2. Journal of Travel Medicine, 2021, 28, .	1.4	97
41	Th1 skewed immune response of whole virion inactivated SARS CoV 2 vaccine and its safety evaluation. IScience, 2021, 24, 102298.	1.9	70
42	An Epidemiological Analysis of SARS-CoV-2 Genomic Sequences from Different Regions of India. Viruses, 2021, 13, 925.	1.5	29
43	Imported SARS-CoV-2 V501Y.V2 variant (B.1.351) detected in travelers from South Africa and Tanzania to India. Travel Medicine and Infectious Disease, 2021, 41, 102023.	1.5	31
44	Neutralization of VUI B.1.1.28 P2 variant with sera of COVID-19 recovered cases and recipients of Covaxin an inactivated COVID-19 vaccine. Journal of Travel Medicine, 2021, 28, .	1.4	38
45	Safety and immunogenicity of an inactivated SARS-CoV-2 vaccine, BBV152: a double-blind, randomised, phase 1 trial. Lancet Infectious Diseases, The, 2021, 21, 637-646.	4.6	326
46	A qualitative IgG ELISA for detection of SARS-CoV-2-specific antibodies in Syrian hamster serum samples. STAR Protocols, 2021, 2, 100573.	0.5	13
47	Neutralization of Beta and Delta variant with sera of COVID-19 recovered cases and vaccinees of inactivated COVID-19 vaccine BBV152/Covaxin. Journal of Travel Medicine, 2021, 28, .	1.4	94
48	Point of care real-time polymerase chain reaction-based diagnostic for Kyasanur forest disease. International Journal of Infectious Diseases, 2021, 108, 226-230.	1.5	4
49	Safety and immunogenicity of an inactivated SARS-CoV-2 vaccine, BBV152: interim results from a double-blind, randomised, multicentre, phase 2 trial, and 3-month follow-up of a double-blind, randomised phase 1 trial. Lancet Infectious Diseases, The, 2021, 21, 950-961.	4.6	271
50	SARS-CoV-2 Spike Mutations, L452R, T478K, E484Q and P681R, in the Second Wave of COVID-19 in Maharashtra, India. Microorganisms, 2021, 9, 1542.	1.6	521
51	Development of double antibody sandwich ELISA as potential diagnostic tool for rapid detection of Crimean-Congo hemorrhagic fever virus. Scientific Reports, 2021, 11, 14699.	1.6	9
52	Neutralization of Delta variant with sera of Covishieldâ"¢ vaccinees and COVID-19-recovered vaccinated individuals. Journal of Travel Medicine, 2021, 28, .	1.4	28
53	Detection of possible Nipah virus infection in Rousettus leschenaultii and Pipistrellus Pipistrellus bats in Maharashtra, India. Journal of Infection and Public Health, 2021, 14, 1010-1012.	1.9	8
54	Performance assessment of seven SARSâ€CoVâ€2 IgG enzymeâ€linked immunosorbent assays. Journal of Medical Virology, 2021, 93, 6696-6702.	2.5	11

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55	Comparison of SARS-CoV-2 Variants of Concern 202012/01 (U.K. Variant) and D614G Variant Transmission by Different Routes in Syrian Hamsters. Vector-Borne and Zoonotic Diseases, 2021, 21, 638-641.	0.6	9
56	Sequential determination of viral load, humoral responses and phylogenetic analysis in fatal and non-fatal cases of Crimean-Congo hemorrhagic fever patients from Gujarat, India, 2019. PLoS Neglected Tropical Diseases, 2021, 15, e0009718.	1.3	4
57	SARS-CoV-2 Delta Variant Pathogenesis and Host Response in Syrian Hamsters. Viruses, 2021, 13, 1773.	1.5	43
58	Longitudinal clinico-serological analysis of anti-nucleocapsid and anti-receptor binding domain of spike protein antibodies against SARS-CoV-2. International Journal of Infectious Diseases, 2021, 112, 103-110.	1.5	11
59	Identification of Phasi Charoen-Like Phasivirus in Field Collected <i>Aedes aegypti</i> from Karnataka State, India. Vector-Borne and Zoonotic Diseases, 2021, 21, 900-909.	0.6	5
60	Clinical Characterization and Genomic Analysis of Samples from COVID-19 Breakthrough Infections during the Second Wave among the Various States of India. Viruses, 2021, 13, 1782.	1.5	70
61	Comparable neutralization of SARS-CoV-2 Delta AY.1 and Delta with individuals sera vaccinated with BBV152. Journal of Travel Medicine, 2021, 28, .	1.4	25
62	Isolation and characterization of the new SARS-CoV-2 variant in travellers from the United Kingdom to India: VUI-202012/01 of the B.1.1.7 lineage. Journal of Travel Medicine, 2021, 28, .	1.4	46
63	Natural Selection Plays an Important Role in Shaping the Codon Usage of Structural Genes of the Viruses Belonging to the Coronaviridae Family. Viruses, 2021, 13, 3.	1.5	12
64	Inactivation of SARS-CoV-2 by gamma irradiation. Indian Journal of Medical Research, 2021, 153, 196.	0.4	12
65	Factors associated with mortality among moderate and severe patients with COVID-19 in India: a secondary analysis of a randomised controlled trial. BMJ Open, 2021, 11, e050571.	0.8	15
66	Immunogenicity and safety of a heterologous prime-boost COVID-19 vaccine schedule: ChAdOx1 vaccine Covishield followed by BBV152 Covaxin. Journal of Travel Medicine, 2021, 28, .	1.4	17
67	SARS-CoV-2 & influenza A virus co-infection in an elderly patient with pneumonia. Indian Journal of Medical Research, 2021, 153, 190-195.	0.4	1
68	Experiences of Indian Council of Medical Research with tick-borne zoonotic infections: Kyasanur Forest disease & Crimean-Congo haemorrhagic fever in India with One Health focus. Indian Journal of Medical Research, 2021, 153, 339-347.	0.4	0
69	A case with SARS-CoV-2 reinfection from India. Indian Journal of Medical Microbiology, 2021, 40, 166-166.	0.3	3
70	Efficacy, safety, and lot-to-lot immunogenicity of an inactivated SARS-CoV-2 vaccine (BBV152): interim results of a randomised, double-blind, controlled, phase 3 trial. Lancet, The, 2021, 398, 2173-2184.	6.3	261
71	Standardization & validation of Truenatâ"¢ point-of-care test for rapid diagnosis of Nipah. Indian Journal of Medical Research, 2021, 154, 645.	0.4	3
72	Genome Sequencing Reveals a Mixed Picture of SARS-CoV-2 Variant of Concern Circulation in Eastern Uttar Pradesh, India. Frontiers in Medicine, 2021, 8, 781287.	1.2	5

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73	Antibody response in symptomatic & asymptomatic Nipah virus cases from Kerala, India. Indian Journal of Medical Research, 2021, 154, 533.	0.4	7
74	ZRC3308 Monoclonal Antibody Cocktail Shows Protective Efficacy in Syrian Hamsters against SARS-CoV-2 Infection. Viruses, 2021, 13, 2424.	1.5	4
75	Possible Role of Accessory Proteins in the Viral Replication for the 20I/501Y.V1 (B.1.1.7) SARS CoV-2 Variant. Pathogens, 2021, 10, 1586.	1.2	0
76	Comparison of neutralizing antibody response in first and second waves of SARS-CoV-2 pandemic in India. Journal of Travel Medicine, 2021, , .	1.4	3
77	Study of Kyasanur forest disease viremia, antibody kinetics, and virus infection in target organs of Macaca radiata. Scientific Reports, 2020, 10, 12561.	1.6	12
78	Clinico-epidemiological and genomic profile of first Zika Virus outbreak in India at Jaipur city of Rajasthan state. Journal of Infection and Public Health, 2020, 13, 1920-1926.	1.9	8
79	Preparedness of public health-care system for Zika virus outbreak: An Indian perspective. Journal of Infection and Public Health, 2020, 13, 949-955.	1.9	9
80	Detection of African genotype in Hyalomma tick pools during Crimean Congo hemorrhagic fever outbreak, Rajasthan, India, 2019. Virus Research, 2020, 286, 198046.	1.1	8
81	Phylogeography of Kyasanur Forest Disease virus in India (1957–2017) reveals evolution and spread in the Western Ghats region. Scientific Reports, 2020, 10, 1966.	1.6	27
82	Experiential learnings from the Nipah virus outbreaks in Kerala towards containment of infectious public health emergencies in India. Epidemiology and Infection, 2020, 148, e90.	1.0	17
83	Clinical profile and outcome of patients with Crimean Congo haemorrhagic fever: a hospital based observational study from Rajasthan, India. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2020, 114, 650-656.	0.7	7
84	First isolation of SARS-CoV-2 from clinical samples in India. Indian Journal of Medical Research, 2020, 151, 244.	0.4	98
85	Genomic analysis of SARS-CoV-2 strains among Indians returning from Italy, Iran & China, & Italian tourists in India. Indian Journal of Medical Research, 2020, 151, 255.	0.4	44
86	Development of indigenous IgG ELISA for the detection of anti-SARS-CoV-2 IgG. Indian Journal of Medical Research, 2020, 151, 444.	0.4	57
87	Evaluation of the susceptibility of mice & hamsters to SARS-CoV-2 infection. Indian Journal of Medical Research, 2020, 151, 479.	0.4	25
88	Quasispecies analysis of the SARS-CoV-2 from representative clinical samples: A preliminary analysis. Indian Journal of Medical Research, 2020, 152, 105.	0.4	5
89	Transcriptome & viral growth analysis of SARS-CoV-2-infected Vero CCL-81 cells. Indian Journal of Medical Research, 2020, 152, 70.	0.4	9
90	Neutralizing antibody responses to SARS-CoV-2 in COVID-19 patients. Indian Journal of Medical Research, 2020, 152, 82.	0.4	64

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91	Identification of SARS-CoV-2 clusters from symptomatic cases in India. Indian Journal of Medical Research, 2020, 152, 111.	0.4	1
92	Prevalence of SARS-CoV-2 infection in India: Findings from the national serosurvey, May-June 2020. Indian Journal of Medical Research, 2020, 152, 48.	0.4	168
93	Full-genome sequences of the first two SARS-CoV-2 viruses from India. Indian Journal of Medical Research, 2020, 151, 200.	0.4	130
94	Biorisk assessment for infrastructure & biosafety requirements for the laboratories providing coronavirus SARS-CoV-2/(COVID-19) diagnosis. Indian Journal of Medical Research, 2020, 151, 172-176.	0.4	23
95	Detection of coronaviruses in Pteropus & Rousettus species of bats from different States of India. Indian Journal of Medical Research, 2020, 151, 226.	0.4	21
96	Proactive preparedness for Cat Que virus: An Orthobunyavirus existing in India. Indian Journal of Medical Research, 2020, 151, 571.	0.4	4
97	Zika virus outbreak in Rajasthan, India in 2018 was caused by a virus endemic to Asia. Infection, Genetics and Evolution, 2019, 69, 199-202.	1.0	45
98	Distribution and prevalence of ticks on livestock population in endemic area of Kyasanur forest disease in Western Ghats of Kerala, South India. Journal of Parasitic Diseases, 2019, 43, 256-262.	0.4	16
99	Nipah Virus Sequences from Humans and Bats during Nipah Outbreak, Kerala, India, 2018. Emerging Infectious Diseases, 2019, 25, 1003-1006.	2.0	59
100	A case report of the enterovirus-D68 associated severe acute respiratory illness in a pediatric case from India. Journal of Infection and Public Health, 2019, 12, 900-903.	1.9	1
101	Growth Kinetics of Kyasanur Forest Disease Virus in Mammalian Cell Lines and Development of Plaque Reduction Neutralization Test. Vector-Borne and Zoonotic Diseases, 2019, 19, 630-636.	0.6	0
102	Persistence of IgG antibodies in survivors of Crimean Congo hemorrhagic fever virus infection, India. Journal of Infection and Public Health, 2019, 12, 598-599.	1.9	6
103	Characterization of a strain of quaranfil virus isolated from soft ticks in India. Is quaranfil virus an unrecognized cause of disease in human and animals?― Heliyon, 2019, 5, e01368.	1.4	8
104	Characterization of Novel Reoviruses Wad Medani Virus (Orbivirus) and Kundal Virus (Coltivirus) Collected from <i>Hyalomma anatolicum</i> Ticks in India during Surveillance for Crimean Congo Hemorrhagic Fever. Journal of Virology, 2019, 93, .	1.5	20
105	Crimean Congo hemorrhagic fever serosurvey in humans for identifying high-risk populations and high-risk areas in the endemic state of Gujarat, India. BMC Infectious Diseases, 2019, 19, 104.	1.3	24
106	Laboratory-Confirmed Avian Influenza A(H9N2) Virus Infection, India, 2019. Emerging Infectious Diseases, 2019, 25, 2328-2330.	2.0	20
107	Spatial Association Between a Nipah Virus Outbreak in India and Nipah Virus Infection in Pteropus Bats. Clinical Infectious Diseases, 2019, 69, 378-379.	2.9	4
108	Outbreak Investigation of Nipah Virus Disease in Kerala, India, 2018. Journal of Infectious Diseases, 2019. 219. 1867-1878.	1.9	173

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109	Identification and phylogenetic analysis of herpes simplex virus-1 from clinical isolates in India. Access Microbiology, 2019, 1, e000047.	0.2	4
110	Emerging/re-emerging viral diseases & new viruses on the Indian horizon. Indian Journal of Medical Research, 2019, 149, 447.	0.4	93
111	Kinetics of viral RNA, immunoglobulin-M & G antibodies in Kyasanur forest disease. Indian Journal of Medical Research, 2019, 150, 186.	0.4	9
112	Kyasanur Forest Disease Prevalence in Western Ghats Proven and Confirmed by Recent Outbreak in Maharashtra, India, 2016. Vector-Borne and Zoonotic Diseases, 2018, 18, 164-172.	0.6	30
113	Development of single step RT-PCR for detection of Kyasanur forest disease virus from clinical samples. Heliyon, 2018, 4, e00549.	1.4	6
114	First laboratory confirmation on the existence of Zika virus disease in India. Journal of Infection, 2018, 76, 314-317.	1.7	22
115	Equine Encephalosis Virus in India, 2008. Emerging Infectious Diseases, 2018, 24, 898-901.	2.0	25
116	Acute Encephalitis with Atypical Presentation of Rubella in Family Cluster, India. Emerging Infectious Diseases, 2018, 24, 1923-1925.	2.0	3
117	Positivity of dengue and chikungunya among Crimean–Congo hemorrhagic fever-negative cases in India: 2013–2016. Journal of Infection and Public Health, 2018, 11, 900-901.	1.9	2
118	Molecular diversity of Coxsackievirus A10 circulating in the southern and northern region of India [2009–17]. Infection, Genetics and Evolution, 2018, 66, 101-110.	1.0	18
119	Characterization of Unknown Orthobunya-Like Viruses from India. Viruses, 2018, 10, 451.	1.5	9
120	Identification and characterization of novel mosquito-borne (Kammavanpettai virus) and tick-borne (Wad Medani) reoviruses isolated in India. Journal of General Virology, 2018, 99, 991-1000.	1.3	9
121	Experimental Zika virus infection in Aedes aegypti: Susceptibility, transmission & co-infection with dengue & chikungunya viruses. Indian Journal of Medical Research, 2018, 147, 88.	0.4	28
122	Detection of Kyasanur forest disease in newer areas of Sindhudurg district of Maharashtra State. Indian Journal of Medical Research, 2018, 148, 453.	0.4	5
123	Circulation of Nipah virus in Pteropus giganteus bats in northeast region of India, 2015. Indian Journal of Medical Research, 2018, 147, 318.	0.4	19
124	Clinical & epidemiological significance of Kyasanur forest disease. Indian Journal of Medical Research, 2018, 148, 145.	0.4	40
125	<i>Zika virus</i> Pathogenesis in Infant Mice after Natural Transmission by the Bite of Infected Mosquitoes. Intervirology, 2017, 60, 227-234.	1.2	2
126	Crimean-Congo Hemorrhagic Fever in Migrant Worker Returning from Oman to India, 2016. Emerging Infectious Diseases, 2017, 23, 1005-1008.	2.0	21

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127	A mini-review of Bunyaviruses recorded in India. Indian Journal of Medical Research, 2017, 145, 601-610.	0.4	6
128	Nosocomial infection of CCHF among health care workers in Rajasthan, India. BMC Infectious Diseases, 2016, 16, 624.	1.3	43
129	Isolation and characterization of Oya virus a member of Simbu serogroup, family Bunyaviridae, isolated from Karnataka, India. Infection, Genetics and Evolution, 2016, 44, 122-126.	1.0	8
130	Isolation of Tioman virus from Pteropus giganteus bat in North-East region of India. Infection, Genetics and Evolution, 2016, 45, 224-229.	1.0	6
131	Recent Scenario of Emergence of Kyasanur Forest Disease in India and Public Health Importance. Current Tropical Medicine Reports, 2016, 3, 7-13.	1.6	34
132	An Early Passage Human Isolate of Kyasanur Forest Disease Virus Shows Acute Neuropathology in Experimentally Infected CD-1 Mice. Vector-Borne and Zoonotic Diseases, 2016, 16, 496-498.	0.6	10
133	Zika virus: Indian perspectives. Indian Journal of Medical Research, 2016, 143, 553.	0.4	20
134	Use of hydrogen peroxide vapour & plasma irradiation in combination for quick decontamination of closed chambers. Indian Journal of Medical Research, 2016, 144, 245.	0.4	6
135	Difference in vector ticks dropping rhythm governs the epidemiology of Crimean-Congo haemorrhagic fever & Kyasanur forest disease in India. Indian Journal of Medical Research, 2016, 144, 633-635.	0.4	3
136	On the transmission pattern of Kyasanur Forest disease (KFD) in India. Infectious Diseases of Poverty, 2015, 4, 37.	1.5	55
137	Cross-sectional Serosurvey of Crimean-Congo Hemorrhagic Fever Virus IgG in Livestock, India, 2013–2014. Emerging Infectious Diseases, 2015, 21, 1837-1839.	2.0	37
138	New focus of Kyasanur Forest disease virus activity in a tribal area in Kerala, India, 2014. Infectious Diseases of Poverty, 2015, 4, 12.	1.5	37
139	First confirmed case of Crimean-Congo haemorrhagic fever from Sirohi district in Rajasthan State, India. Indian Journal of Medical Research, 2015, 142, 489.	0.4	16
140	Serosurvey of Crimean-Congo Hemorrhagic Fever Virus in Domestic Animals, Gujarat, India, 2013. Vector-Borne and Zoonotic Diseases, 2014, 14, 690-692.	0.6	21
141	Emergence of Crimean-Congo hemorrhagic fever in Amreli District of Gujarat State, India, June to July 2013. International Journal of Infectious Diseases, 2014, 18, 97-100.	1.5	24
142	Crimean-Congo Hemorrhagic Fever: Current Scenario in India. Proceedings of the National Academy of Sciences India Section B - Biological Sciences, 2014, 84, 9-18.	0.4	10
143	Outbreak of Kyasanur Forest disease in Thirthahalli, Karnataka, India, 2014. International Journal of Infectious Diseases, 2014, 26, 132-134.	1.5	35
144	Expediency of dengue illness classification: the Sri Lankan perspective Highly infectious tick-borne viral diseases: Kyasanur forest disease and Crimean-Congo haemorrhagic fever in India. WHO South-East Asia Journal of Public Health, 2014, 3, 8.	1.7	45

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145	Establishment of Biosafety Level-3 (BSL-3) laboratory: important criteria to consider while designing, constructing, commissioning & operating the facility in Indian setting. Indian Journal of Medical Research, 2014, 140, 171-83.	0.4	14
146	Genetic characterization and molecular clock analyses of the Crimean-Congo hemorrhagic fever virus from human and ticks in India, 2010–2011. Infection, Genetics and Evolution, 2013, 14, 223-231.	1.0	35
147	Spread of Kyasanur Forest Disease, Bandipur Tiger Reserve, India, 2012–2013. Emerging Infectious Diseases, 2013, 19, 1540-1.	2.0	44
148	Kyasanur Forest Disease, India, 2011–2012. Emerging Infectious Diseases, 2013, 19, 278-281.	2.0	48
149	Establishment of cell line from embryonic tissue of Pipistrellus ceylonicus bat species from India & its susceptibility to different viruses. Indian Journal of Medical Research, 2013, 138, 224-31.	0.4	9
150	Detection, Isolation and Confirmation of Crimean-Congo Hemorrhagic Fever Virus in Human, Ticks and Animals in Ahmadabad, India, 2010–2011. PLoS Neglected Tropical Diseases, 2012, 6, e1653.	1.3	95
151	Detection of Nipah Virus RNA in Fruit Bat (Pteropus giganteus) from India. American Journal of Tropical Medicine and Hygiene, 2012, 87, 576-578.	0.6	80
152	Diagnosis of Kyasanur forest disease by nested RT-PCR, real-time RT-PCR and IgM capture ELISA. Journal of Virological Methods, 2012, 186, 49-54.	1.0	64
153	Buffalopox outbreak in humans and animals in Western Maharashtra, India. Preventive Veterinary Medicine, 2011, 100, 242-247.	0.7	44
154	Genomic analysis reveals Nairobi sheep disease virus to be highly diverse and present in both Africa, and in India in the form of the Ganjam virus variant. Infection, Genetics and Evolution, 2011, 11, 1111-1120.	1.0	27
155	Recent Ancestry of Kyasanur Forest Disease Virus. Emerging Infectious Diseases, 2009, 15, 1431-1437.	2.0	78
156	Molecular characterization of Umbre virus (Bunyaviridae). Virology Journal, 2008, 5, 115.	1.4	8
157	Evaluation of Safety and Immunogenicity of an Adjuvanted, TH-1 Skewed, Whole Virion Inactivated SARS-CoV-2 Vaccine - BBV152. SSRN Electronic Journal, 0, , .	0.4	4
158	Effectiveness of BBV152/Covaxin and AZD1222/Covishield Vaccines Against Severe COVID-19 and B.1.617.2/Delta Variant in India, 2021: A Multi-Centric Hospital-Based Case-Control Study. SSRN Electronic Journal, 0, , .	0.4	3
159	Zika a Vector Borne Disease Detected in Newer States of India Amidst the COVID-19 Pandemic. Frontiers in Microbiology, 0, 13, .	1.5	10