List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Reliability of Self-Sampling for Accurate Assessment of Respiratory Virus Viral and Immunologic Kinetics. Journal of Infectious Diseases, 2022, 226, 278-286.	1.9	10
2	Clinical and Infection Prevention Applications of Severe Acute Respiratory Syndrome Coronavirus 2 Genotyping: An Infectious Diseases Society of America/American Society for Microbiology Consensus Review Document. Clinical Infectious Diseases, 2022, 74, 1496-1502.	2.9	20
3	Retrospective Detection of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) in Symptomatic Patients Prior to Widespread Diagnostic Testing in Southern California. Clinical Infectious Diseases, 2022, 74, 271-277.	2.9	4
4	Variants of Concern Are Overrepresented Among Postvaccination Breakthrough Infections of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) in Washington State. Clinical Infectious Diseases, 2022, 74, 1089-1092.	2.9	38
5	Clinical and Infection Prevention Applications of Severe Acute Respiratory Syndrome Coronavirus 2 Genotyping: an Infectious Diseases Society of America/American Society for Microbiology Consensus Review Document. Journal of Clinical Microbiology, 2022, 60, JCM0165921.	1.8	13
6	Predicting infectivity: comparing four PCRâ€based assays to detect culturable SARS oVâ€2 in clinical samples. EMBO Molecular Medicine, 2022, 14, e15290.	3.3	38
7	SARS-CoV-2 breakthrough infections elicit potent, broad, and durable neutralizing antibody responses. Cell, 2022, 185, 872-880.e3.	13.5	165
8	Measuring infectious SARS-CoV-2 in clinical samples reveals a higher viral titer:RNA ratio for Delta and Epsilon vs. Alpha variants. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	3.3	35
9	Mutations in viral nucleocapsid protein and endoRNase are discovered to associate with COVID19 hospitalization risk. Scientific Reports, 2022, 12, 1206.	1.6	12
10	Trajectory of Viral RNA Load Among Persons With Incident SARS-CoV-2 G614 Infection (Wuhan Strain) in Association With COVID-19 Symptom Onset and Severity. JAMA Network Open, 2022, 5, e2142796.	2.8	57
11	Evaluating Antibody Mediated Protection against Alpha, Beta, and Delta SARS-CoV-2 Variants of Concern in K18-hACE2 Transgenic Mice. Journal of Virology, 2022, 96, jvi0218421.	1.5	14
12	Elucidating Pathways Mediating the Relationship Between Male Sex and COVID-19 Severity. Clinical Epidemiology, 2022, Volume 14, 115-125.	1.5	3
13	Self-Assessed Severity as a Determinant of COVID-19 Symptom Specificity: A Longitudinal Cohort Study. Clinical Infectious Diseases, 2022, , .	2.9	0
14	Detection and kinetics of subgenomic SARS-CoV-2 RNA viral load in longitudinal diagnostic RNA positive samples. Journal of Infectious Diseases, 2022, , .	1.9	4
15	Performance of anterior nares and tongue swabs for nucleic acid, Nucleocapsid, and Spike antigen testing for detecting SARS-CoV-2 against nasopharyngeal PCR and viral culture. International Journal of Infectious Diseases, 2022, 117, 287-294.	1.5	7
16	The Impact of Bâ€cell Directed Therapy on SARSâ€CoVâ€2 Vaccine Efficacy in CLL. British Journal of Haematology, 2022, , .	1.2	11
17	Fragment size-based enrichment of viral sequences in plasma cell-free DNA. Journal of Molecular Diagnostics, 2022, 24, 476-484.	1.2	4
18	Modeling Infection and Tropism of Human Parainfluenza Virus Type 3 in Ferrets. MBio, 2022, 13, e0383121.	1.8	5

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19	The SARS-CoV-2 Omicron Variant Does Not Have Higher Nasal Viral Loads Compared to the Delta Variant in Symptomatic and Asymptomatic Individuals. Journal of Clinical Microbiology, 2022, 60, e0013922.	1.8	28
20	De novo emergence of a remdesivir resistance mutation during treatment of persistent SARS-CoV-2 infection in an immunocompromised patient: a case report. Nature Communications, 2022, 13, 1547.	5.8	159
21	Response of Human Liver Tissue to Innate Immune Stimuli. Frontiers in Immunology, 2022, 13, 811551.	2.2	1
22	Host–pathogen dynamics in longitudinal clinical specimens from patients with COVID-19. Scientific Reports, 2022, 12, 5856.	1.6	3
23	Associations Between Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Variants and Risk of Coronavirus Disease 2019 (COVID-19) Hospitalization Among Confirmed Cases in Washington State: A Retrospective Cohort Study. Clinical Infectious Diseases, 2022, 75, e536-e544.	2.9	38
24	T cell receptor sequencing identifies prior SARS-CoV-2 infection and correlates with neutralizing antibodies and disease severity. JCI Insight, 2022, 7, .	2.3	26
25	Identification of Omicron-Delta Coinfections Using PCR-Based Genotyping. Microbiology Spectrum, 2022, 10, e0060522.	1.2	6
26	Two Novel Iflaviruses Discovered in Bat Samples in Washington State. Viruses, 2022, 14, 994.	1.5	3
27	Thermodynamically coupled biosensors for detecting neutralizing antibodies against SARS-CoV-2 variants. Nature Biotechnology, 2022, 40, 1336-1340.	9.4	23
28	Molecular Analysis of SARS-CoV-2 Lineages in Armenia. Viruses, 2022, 14, 1074.	1.5	7
29	Case Study: Impact of Diurnal Variations and Stormwater Dilution on SARS-CoV-2 RNA Signal Intensity at Neighborhood Scale Wastewater Pumping Stations. ACS ES&T Water, 2022, 2, 1964-1975.	2.3	4
30	The Clinical and Genomic Epidemiology of Rhinovirus in Homeless Shelters—King County, Washington. Journal of Infectious Diseases, 2022, 226, S304-S314.	1.9	6
31	Narrow transmission bottlenecks and limited within-host viral diversity during a SARS-CoV-2 outbreak on a fishing boat. Virus Evolution, 2022, 8, .	2.2	7
32	Rapid and accurate identification of SARS-CoV-2 Omicron variants using droplet digital PCR (RT-ddPCR). Journal of Clinical Virology, 2022, 154, 105218.	1.6	12
33	A Method for Variant Agnostic Detection of SARS-CoV-2, Rapid Monitoring of Circulating Variants, and Early Detection of Emergent Variants Such as Omicron. Journal of Clinical Microbiology, 2022, 60,	1.8	14
34	Clinical Performance Characteristics of the Swift Normalase Amplicon Panel for Sensitive Recovery of Severe Acute Respiratory Syndrome Coronavirus 2 Genomes. Journal of Molecular Diagnostics, 2022, 24, 963-976.	1.2	7
35	Low Prevalence of Severe Acute Respiratory Syndrome Coronavirus 2 Among Pregnant and Postpartum Patients With Universal Screening in Seattle, Washington. Clinical Infectious Diseases, 2021, 72, 869-872.	2.9	31
36	Occurrence and Timing of Subsequent Severe Acute Respiratory Syndrome Coronavirus 2 Reverse-transcription Polymerase Chain Reaction Positivity Among Initially Negative Patients. Clinical Infectious Diseases, 2021, 72, 323-326.	2.9	78

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37	Dynamics of Neutralizing Antibody Titers in the Months After Severe Acute Respiratory Syndrome Coronavirus 2 Infection. Journal of Infectious Diseases, 2021, 223, 197-205.	1.9	216
38	Hydroxychloroquine as Postexposure Prophylaxis to Prevent Severe Acute Respiratory Syndrome Coronavirus 2 Infection. Annals of Internal Medicine, 2021, 174, 344-352.	2.0	73
39	Prolonged persistence of PCR-detectable virus during an outbreak of SARS-CoV-2 in an inpatient geriatric psychiatry unit in King County, Washington. American Journal of Infection Control, 2021, 49, 293-298.	1.1	16
40	Evolutionary History of Endogenous Human Herpesvirus 6 Reflects Human Migration out of Africa. Molecular Biology and Evolution, 2021, 38, 96-107.	3.5	31
41	CrAssphage and its bacterial host in cat feces. Scientific Reports, 2021, 11, 815.	1.6	14
42	NGSocomial Infections: High-Resolution Views of Hospital-Acquired Infections Through Genomic Epidemiology. Journal of the Pediatric Infectious Diseases Society, 2021, 10, S88-S95.	0.6	7
43	Analytical Sensitivity of the Abbott BinaxNOW COVID-19 Ag Card. Journal of Clinical Microbiology, 2021, 59, .	1.8	69
44	In silico detection of SARS-CoV-2 specific B-cell epitopes and validation in ELISA for serological diagnosis of COVID-19. Scientific Reports, 2021, 11, 4290.	1.6	22
45	CRISPR-Cas9 gene editing of hepatitis B virus in chronically infected humanized mice. Molecular Therapy - Methods and Clinical Development, 2021, 20, 258-275.	1.8	62
46	Hydroxychloroquine with or without azithromycin for treatment of early SARS-CoV-2 infection among high-risk outpatient adults: A randomized clinical trial. EClinicalMedicine, 2021, 33, 100773.	3.2	55
47	Endogenously Produced SARS-CoV-2 Specific IgG Antibodies May Have a Limited Impact on Clearing Nasal Shedding of Virus during Primary Infection in Humans. Viruses, 2021, 13, 516.	1.5	5
48	Clinical and Virologic Characteristics and Outcomes of Coronavirus Disease 2019 at a Cancer Center. Open Forum Infectious Diseases, 2021, 8, ofab193.	0.4	4
49	Inactivation of genes in oxidative respiration and iron acquisition pathways in pediatric clinical isolates of Small colony variant Enterobacteriaceae. Scientific Reports, 2021, 11, 7457.	1.6	4
50	SARS-CoV-2 ORF6 Disrupts Bidirectional Nucleocytoplasmic Transport through Interactions with Rae1 and Nup98. MBio, 2021, 12, .	1.8	92
51	A human coronavirus evolves antigenically to escape antibody immunity. PLoS Pathogens, 2021, 17, e1009453.	2.1	183
52	<i>In Vivo</i> Generation of BK and JC Polyomavirus Defective Viral Genomes in Human Urine Samples Associated with Higher Viral Loads. Journal of Virology, 2021, 95, .	1.5	9
53	Viral genomes reveal patterns of the SARS-CoV-2 outbreak in Washington State. Science Translational Medicine, 2021, 13, .	5.8	58
54	Molecular Features of the Measles Virus Viral Fusion Complex That Favor Infection and Spread in the Brain. MBio, 2021, 12, e0079921.	1.8	24

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55	Specific allelic discrimination of N501Y and other SARSâ€CoVâ€2 mutations by ddPCR detects B.1.1.7 lineage in Washington State. Journal of Medical Virology, 2021, 93, 5931-5941.	2.5	31
56	Performance characteristics of the Abbott Alinity m SARS-CoV-2 assay. Journal of Clinical Virology, 2021, 140, 104869.	1.6	16
57	Reducing COVID-19 quarantine with SARS-CoV-2 testing: a simulation study. BMJ Open, 2021, 11, e050473.	0.8	8
58	Estimating the False-Positive Rate of Highly Automated SARS-CoV-2 Nucleic Acid Amplification Testing. Journal of Clinical Microbiology, 2021, 59, e0108021.	1.8	12
59	Genomics and transcriptomics yields a system-level view of the biology of the pathogen Naegleria fowleri. BMC Biology, 2021, 19, 142.	1.7	18
60	Genetic engineering of Treponema pallidum subsp. pallidum, the Syphilis Spirochete. PLoS Pathogens, 2021, 17, e1009612.	2.1	27
61	Test it earlier, result it faster, makes us stronger: how rapid viral diagnostics enable therapeutic success. Current Opinion in Virology, 2021, 49, 111-116.	2.6	7
62	A SARS-CoV-2 Nucleocapsid Variant that Affects Antigen Test Performance. Journal of Clinical Virology, 2021, 141, 104900.	1.6	53
63	Immunogenicity of a heterologous COVID-19 vaccine after failed vaccination in a lymphoma patient. Cancer Cell, 2021, 39, 1037-1038.	7.7	20
64	Whole Genome Sequence Analysis of Brucella melitensis Phylogeny and Virulence Factors. Microbiology Research, 2021, 12, 698-710.	0.8	3
65	Anti-SARS-CoV-2 Antibody Levels Measured by the AdviseDx SARS-CoV-2 Assay Are Concordant with Previously Available Serologic Assays but Are Not Fully Predictive of Sterilizing Immunity. Journal of Clinical Microbiology, 2021, 59, e0098921.	1.8	48
66	Fast SARS-CoV-2 Variant Detection Using Snapback Primer High-Resolution Melting. Diagnostics, 2021, 11, 1788.	1.3	8
67	Unbiased Pandemic Pathogen Detection and the Federal Register. Journal of Clinical Microbiology, 2021, 59, e0134621.	1.8	0
68	Phylogenetic estimates of SARS-CoV-2 introductions into Washington State. The Lancet Regional Health Americas, 2021, 1, 100018.	1.5	8
69	Longitudinal TprK profiling of in vivo and in vitro-propagated Treponema pallidum subsp. pallidum reveals accumulation of antigenic variants in absence of immune pressure. PLoS Neglected Tropical Diseases, 2021, 15, e0009753.	1.3	15
70	A Bifluorescent-Based Assay for the Identification of Neutralizing Antibodies against SARS-CoV-2 Variants of Concern <i>In Vitro</i> and <i>In Vivo</i> . Journal of Virology, 2021, 95, e0112621.	1.5	13
71	Analysis of SARS-CoV-2 infection dynamic in vivo using reporter-expressing viruses. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	25
72	Development of the RealTime SARS-CoV-2 quantitative Laboratory Developed Test and correlation with viral culture as a measure of infectivity. Journal of Clinical Virology, 2021, 143, 104945.	1.6	22

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73	Modifying laboratory testing via home brew during the COVID-19 pandemic. Journal of Clinical and Translational Science, 2021, 5, e93.	0.3	3
74	RADx Variant Task Force Program for Assessing the Impact of Variants on SARS-CoV-2 Molecular and Antigen Tests. IEEE Open Journal of Engineering in Medicine and Biology, 2021, 2, 1-1.	1.7	6
75	Rapid adaptation to human protein kinase R by a unique genomic rearrangement in rhesus cytomegalovirus. PLoS Pathogens, 2021, 17, e1009088.	2.1	9
76	Human parainfluenza virus evolution during lung infection of immunocompromised individuals promotes viral persistence. Journal of Clinical Investigation, 2021, 131, .	3.9	12
77	Hamster organotypic modeling of SARS-CoV-2 lung and brainstem infection. Nature Communications, 2021, 12, 5809.	5.8	37
78	Oral prodrug of remdesivir parent GS-441524 is efficacious against SARS-CoV-2 in ferrets. Nature Communications, 2021, 12, 6415.	5.8	74
79	Treponema pallidum genome sequencing from six continents reveals variability in vaccine candidate genes and dominance of Nichols clade strains in Madagascar. PLoS Neglected Tropical Diseases, 2021, 15, e0010063.	1.3	30
80	Implementation of a fully remote randomized clinical trial with cardiac monitoring. Communications Medicine, 2021, 1, .	1.9	4
81	International Spread of Multidrug-Resistant Campylobacter coli in Men Who Have Sex With Men in Washington State and Québec, 2015–2018. Clinical Infectious Diseases, 2020, 71, 1896-1904.	2.9	20
82	Direct RT-qPCR detection of SARS-CoV-2 RNA from patient nasopharyngeal swabs without an RNA extraction step. PLoS Biology, 2020, 18, e3000896.	2.6	119
83	Pooling of SARS-CoV-2 samples to increase molecular testing throughput. Journal of Clinical Virology, 2020, 131, 104570.	1.6	51
84	Optimization and clinical validation of dual-target RT-LAMP for SARS-CoV-2. Journal of Virological Methods, 2020, 286, 113972.	1.0	36
85	Sensitive Identification of Bacterial DNA in Clinical Specimens by Broad-Range 16S rRNA Gene Enrichment. Journal of Clinical Microbiology, 2020, 58, .	1.8	15
86	Orally efficacious broad-spectrum allosteric inhibitor of paramyxovirus polymerase. Nature Microbiology, 2020, 5, 1232-1246.	5.9	18
87	Clinical evaluation of the BioFire® Respiratory Panel 2.1 and detection of SARS-CoV-2. Journal of Clinical Virology, 2020, 129, 104538.	1.6	60
88	Estimation of Full-Length TprK Diversity in Treponema pallidum subsp. <i>pallidum</i> . MBio, 2020, 11, .	1.8	19
89	Sensitive Recovery of Complete SARS-CoV-2 Genomes from Clinical Samples by Use of Swift Biosciences' SARS-CoV-2 Multiplex Amplicon Sequencing Panel. Journal of Clinical Microbiology, 2020, 59, .	1.8	58
90	The First Quarter of SARS-CoV-2 Testing: the University of Washington Medicine Experience. Journal of Clinical Microbiology, 2020, 58, .	1.8	12

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91	Western Washington State COVID-19 Experience: Keys to Flattening the Curve and Effective Health System Response. Journal of the American College of Surgeons, 2020, 231, 316-324e1.	0.2	26
92	Retrospective clinical evaluation of 4 lateral flow assays for the detection of SARS-CoV-2 lgG. Diagnostic Microbiology and Infectious Disease, 2020, 98, 115161.	0.8	14
93	Inhibition of Coronavirus Entry <i>In Vitro</i> and <i>Ex Vivo</i> by a Lipid-Conjugated Peptide Derived from the SARS-CoV-2 Spike Glycoprotein HRC Domain. MBio, 2020, 11, .	1.8	63
94	Neutralizing Antibodies Correlate with Protection from SARS-CoV-2 in Humans during a Fishery Vessel Outbreak with a High Attack Rate. Journal of Clinical Microbiology, 2020, 58, .	1.8	494
95	Herpes Simplex Virus Mistyping due to HSV-1 × HSV-2 Interspecies Recombination in Viral Gene Encoding Glycoprotein B. Viruses, 2020, 12, 860.	1.5	5
96	Detection of SARS-CoV-2 with SHERLOCK One-Pot Testing. New England Journal of Medicine, 2020, 383, 1492-1494.	13.9	506
97	In vivo antiviral host transcriptional response to SARS-CoV-2 by viral load, sex, and age. PLoS Biology, 2020, 18, e3000849.	2.6	225
98	Cryptic transmission of SARS-CoV-2 in Washington state. Science, 2020, 370, 571-575.	6.0	217
99	Gene editing and elimination of latent herpes simplex virus in vivo. Nature Communications, 2020, 11, 4148.	5.8	46
100	Pathogen or Bystander: Clinical Significance of Detecting Human Herpesvirus 6 in Pediatric Cerebrospinal Fluid. Journal of Clinical Microbiology, 2020, 58, .	1.8	26
101	Stability of SARS-CoV-2 in Phosphate-Buffered Saline for Molecular Detection. Journal of Clinical Microbiology, 2020, 58, .	1.8	36
102	Outbreak Investigation of COVID-19 Among Residents and Staff of an Independent and Assisted Living Community for Older Adults in Seattle, Washington. JAMA Internal Medicine, 2020, 180, 1101.	2.6	101
103	Metagenomic Analysis Reveals Clinical SARS-CoV-2 Infection and Bacterial or Viral Superinfection and Colonization. Clinical Chemistry, 2020, 66, 966-972.	1.5	63
104	Changes in SARS-CoV-2 Positivity Rate in Outpatients in Seattle and Washington State, March 1–April 16, 2020. JAMA - Journal of the American Medical Association, 2020, 323, 2334.	3.8	25
105	Coast-to-Coast Spread of SARS-CoV-2 during the Early Epidemic in the United States. Cell, 2020, 181, 990-996.e5.	13.5	321
106	Performance Characteristics of the Abbott Architect SARS-CoV-2 IgG Assay and Seroprevalence in Boise, Idaho. Journal of Clinical Microbiology, 2020, 58, .	1.8	496
107	Expedited SARS-CoV-2 screening of donors and recipients supports continued solid organ transplantation. American Journal of Transplantation, 2020, 20, 3106-3112.	2.6	13
108	Validation of SARS-CoV-2 detection across multiple specimen types. Journal of Clinical Virology, 2020, 128, 104438.	1.6	66

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109	Validation and verification of the Abbott RealTime SARS-CoV-2 assay analytical and clinical performance. Journal of Clinical Virology, 2020, 129, 104474.	1.6	58
110	Multiplexing primer/probe sets for detection of SARS-CoV-2 by qRT-PCR. Journal of Clinical Virology, 2020, 129, 104499.	1.6	35
111	Preprocedural Surveillance Testing for SARS-CoV-2 in an Asymptomatic Population in the Seattle Region Shows Low Rates of Positivity. Journal of Clinical Microbiology, 2020, 58, .	1.8	6
112	Prevalence of Coronavirus Disease 2019 Infection and Outcomes Among Symptomatic Healthcare Workers in Seattle, Washington. Clinical Infectious Diseases, 2020, 71, 2702-2707.	2.9	61
113	Detection of SARS-CoV-2 by bronchoscopy after negative nasopharyngeal testing: Stay vigilant for COVID-19. Respiratory Medicine Case Reports, 2020, 30, 101120.	0.2	24
114	When To Retest: an Examination of Repeat COVID-19 PCR Patterns in an Ambulatory Population. Journal of Clinical Microbiology, 2020, 58, .	1.8	3
115	The Laboratory Diagnosis of Coronavirus Disease 2019— Frequently Asked Questions. Clinical Infectious Diseases, 2020, 71, 2996-3001.	2.9	52
116	Genomic surveillance reveals multiple introductions of SARS-CoV-2 into Northern California. Science, 2020, 369, 582-587.	6.0	253
117	Covid-19 in Critically III Patients in the Seattle Region — Case Series. New England Journal of Medicine, 2020, 382, 2012-2022.	13.9	2,120
118	Identification of multiple large deletions in ORF7a resulting in in-frame gene fusions in clinical SARS-CoV-2 isolates. Journal of Clinical Virology, 2020, 129, 104523.	1.6	71
119	High-resolution profiling of human cytomegalovirus cell-free DNA in human plasma highlights its exceptionally fragmented nature. Scientific Reports, 2020, 10, 3734.	1.6	7
120	Identification of Mycobacterium porcinum in patients with cystic Fibrosis: Pathogen or contaminant?. Journal of Cystic Fibrosis, 2020, 19, 580-586.	0.3	7
121	Comparison of Commercially Available and Laboratory-Developed Assays for <i>In Vitro</i> Detection of SARS-CoV-2 in Clinical Laboratories. Journal of Clinical Microbiology, 2020, 58, .	1.8	215
122	Comparative Performance of SARS-CoV-2 Detection Assays Using Seven Different Primer-Probe Sets and One Assay Kit. Journal of Clinical Microbiology, 2020, 58, .	1.8	401
123	Cell free DNA from respiratory pathogens is detectable in the blood plasma of Cystic Fibrosis patients. Scientific Reports, 2020, 10, 6903.	1.6	15
124	Prevalent and Diverse Intratumoral Oncoprotein-Specific CD8+ T Cells within Polyomavirus-Driven Merkel Cell Carcinomas. Cancer Immunology Research, 2020, 8, 648-659.	1.6	28
125	Comparative genomics and full-length Tprk profiling of Treponema pallidum subsp. pallidum reinfection. PLoS Neglected Tropical Diseases, 2020, 14, e0007921.	1.3	18
126	Evaluation of Genotypic Antiviral Resistance Testing as an Alternative to Phenotypic Testing in a Patient with DOCK8 Deficiency and Severe HSV-1 Disease. Journal of Infectious Diseases, 2020, 221, 2035-2042.	1.9	9

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127	SARS-CoV-2 Viral Load on Admission Is Associated With 30-Day Mortality. Open Forum Infectious Diseases, 2020, 7, ofaa535.	0.4	31
128	Detection of SARS-CoV-2 Among Residents and Staff Members of an Independent and Assisted Living Community for Older Adults — Seattle, Washington, 2020. Morbidity and Mortality Weekly Report, 2020, 69, 416-418.	9.0	108
129	One future of clinical metagenomic sequencing for infectious diseases. Expert Review of Molecular Diagnostics, 2019, 19, 849-851.	1.5	6
130	Characterization of orally efficacious influenza drug with high resistance barrier in ferrets and human airway epithelia. Science Translational Medicine, 2019, 11, .	5.8	253
131	VAPiD: a lightweight cross-platform viral annotation pipeline and identification tool to facilitate virus genome submissions to NCBI GenBank. BMC Bioinformatics, 2019, 20, 48.	1.2	47
132	Genome-Wide Approach to the CD4 T-Cell Response to Human Herpesvirus 6B. Journal of Virology, 2019, 93, .	1.5	6
133	Large, Stable, Contemporary Interspecies Recombination Events in Circulating Human Herpes Simplex Viruses. Journal of Infectious Diseases, 2019, 221, 1271-1279.	1.9	21
134	Whole-genome analysis of extraintestinal pathogenic Escherichia coli (ExPEC) MDR ST73 and ST127 isolated from endangered southern resident killer whales (Orcinus orca). Journal of Antimicrobial Chemotherapy, 2019, 74, 2176-2180.	1.3	13
135	A Novel, Widespread <i>qacA</i> Allele Results in Reduced Chlorhexidine Susceptibility in <i>Staphylococcus epidermidis</i> . Antimicrobial Agents and Chemotherapy, 2019, 63, .	1.4	12
136	Societal Implications of the Internet of Pathogens. Journal of Clinical Microbiology, 2019, 57, .	1.8	7
137	Inherited Chromosomally Integrated Human Herpesvirus 6 Demonstrates Tissue-Specific RNA Expression <i>In Vivo</i> That Correlates with an Increased Antibody Immune Response. Journal of Virology, 2019, 94, .	1.5	27
138	The Brief Case: Inherited Chromosomally Integrated Human Herpesvirus 6 (HHV-6) in the Age of Multiplex HHV-6 Testing. Journal of Clinical Microbiology, 2019, 57, .	1.8	4
139	Closing the Brief Case: Inherited Chromosomally Integrated Human Herpesvirus 6 (HHV-6) in the Age of Multiplex HHV-6 Testing. Journal of Clinical Microbiology, 2019, 57, .	1.8	3
140	Comparison of Three Adenovirus Quantitative PCR Assays with ATCC Reference Strains and Clinical Samples. Journal of Clinical Microbiology, 2019, 57, .	1.8	3
141	Trillions and Trillions: Herpes Simplex Virus–1 Hepatitis in an Immunocompetent Adult. Open Forum Infectious Diseases, 2019, 6, ofz465.	0.4	4
142	Prospective, Real-time Metagenomic Sequencing During Norovirus Outbreak Reveals Discrete Transmission Clusters. Clinical Infectious Diseases, 2019, 69, 941-948.	2.9	21
143	<i>Mycobacterium talmoniae</i> , a Potential Pulmonary Pathogen Isolated from Multiple Patients with Bronchiectasis in the United States, Including the First Case of Clinical Disease in a Patient with Cystic Fibrosis. Journal of Clinical Microbiology, 2019, 57, .	1.8	3
144	Metagenomics to Assist in the Diagnosis of Bloodstream Infection. journal of applied laboratory medicine, The, 2019, 3, 643-653.	0.6	49

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145	Model-based estimation of superinfection prevalence from limited datasets. Journal of the Royal Society Interface, 2018, 15, 20170968.	1.5	1
146	Copy Number Heterogeneity, Large Origin Tandem Repeats, and Interspecies Recombination in Human Herpesvirus 6A (HHV-6A) and HHV-6B Reference Strains. Journal of Virology, 2018, 92, .	1.5	21
147	The Unstructured Paramyxovirus Nucleocapsid Protein Tail Domain Modulates Viral Pathogenesis through Regulation of Transcriptase Activity. Journal of Virology, 2018, 92, .	1.5	23
148	The human clone ST22 SCCmec IV methicillin-resistant Staphylococcus aureus isolated from swine herds and wild primates in Nepal: is man the common source?. FEMS Microbiology Ecology, 2018, 94, .	1.3	26
149	Heterogeneous Antimicrobial Susceptibility Characteristics in Pseudomonas aeruginosa Isolates from Cystic Fibrosis Patients. MSphere, 2018, 3, .	1.3	17
150	A decade of RNA virus metagenomics is (not) enough. Virus Research, 2018, 244, 218-229.	1.1	129
151	Viral Entry Properties Required for Fitness in Humans Are Lost through Rapid Genomic Change during Viral Isolation. MBio, 2018, 9, .	1.8	27
152	Cooperating H3N2 Influenza Virus Variants Are Not Detectable in Primary Clinical Samples. MSphere, 2018, 3, .	1.3	26
153	Limited Marginal Utility of Deep Sequencing for HIV Drug Resistance Testing in the Age of Integrase Inhibitors. Journal of Clinical Microbiology, 2018, 56, .	1.8	18
154	Epidemiological and genomic characterization of community-acquired Clostridium difficile infections. BMC Infectious Diseases, 2018, 18, 443.	1.3	21
155	Private collection: high correlation of sample collection and patient admission date in clinical microbiological testing complicates sharing of phylodynamic metadata. Virus Evolution, 2018, 4, vey005.	2.2	8
156	An optimized methodology for whole genome sequencing of RNA respiratory viruses from nasopharyngeal aspirates. PLoS ONE, 2018, 13, e0199714.	1.1	30
157	Distribution of Staphylococcus species in dairy cows, workers and shared farm environments. FEMS Microbiology Letters, 2018, 365, .	0.7	11
158	Ultrasensitive Capture of Human Herpes Simplex Virus Genomes Directly from Clinical Samples Reveals Extraordinarily Limited Evolution in Cell Culture. MSphere, 2018, 3, .	1.3	49
159	Comparative genomic, transcriptomic, and proteomic reannotation of human herpesvirus 6. BMC Genomics, 2018, 19, 204.	1.2	45
160	Orally Efficacious Broad-Spectrum Ribonucleoside Analog Inhibitor of Influenza and Respiratory Syncytial Viruses. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	162
161	The challenge of diagnostic metagenomics. Expert Review of Molecular Diagnostics, 2018, 18, 605-615.	1.5	130
162	Quantification of BK Virus Standards by Quantitative Real-Time PCR and Droplet Digital PCR Is Confounded by Multiple Virus Populations in the WHO BKV International Standard. Clinical Chemistry, 2017, 63, 761-769.	1.5	53

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163	Genome Sequences of Three Novel Bunyaviruses, Two Novel Rhabdoviruses, and One Novel Nyamivirus from Washington State Moths. Genome Announcements, 2017, 5, .	0.8	11
164	<i>Clostridium perfringens</i> sepsis masquerading as a hemolytic transfusion reaction. Transfusion, 2017, 57, 1112-1112.	0.8	5
165	First Complete Genome Sequence of Corynebacterium riegelii. Genome Announcements, 2017, 5, .	0.8	3
166	Rule-Out Outbreak: 24-Hour Metagenomic Next-Generation Sequencing for Characterizing Respiratory Virus Source for Infection Prevention. Journal of the Pediatric Infectious Diseases Society, 2017, 6, 168-172.	0.6	38
167	Copy Number Heterogeneity of JC Virus Standards. Journal of Clinical Microbiology, 2017, 55, 824-831.	1.8	19
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