

# Philippe Colson

## List of Publications by Year in descending order

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Version: 2024-02-01

390  
papers

21,725  
citations

20759

60  
h-index

12558

132  
g-index

433  
all docs

433  
docs citations

433  
times ranked

25546  
citing authors

#	ARTICLE	IF	CITATIONS
1	Hydroxychloroquine and azithromycin as a treatment of COVID-19: results of an open-label non-randomized clinical trial. <i>International Journal of Antimicrobial Agents</i> , 2020, 56, 105949.	1.1	3,955
2	New insights on the antiviral effects of chloroquine against coronavirus: what to expect for COVID-19?. <i>International Journal of Antimicrobial Agents</i> , 2020, 55, 105938.	1.1	842
3	Viral RNA load as determined by cell culture as a management tool for discharge of SARS-CoV-2 patients from infectious disease wards. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2020, 39, 1059-1061.	1.3	767
4	Chloroquine and hydroxychloroquine as available weapons to fight COVID-19. <i>International Journal of Antimicrobial Agents</i> , 2020, 55, 105932.	1.1	724
5	Clinical and microbiological effect of a combination of hydroxychloroquine and azithromycin in 80 COVID-19 patients with at least a six-day follow up: A pilot observational study. <i>Travel Medicine and Infectious Disease</i> , 2020, 34, 101663.	1.5	605
6	Pig Liver Sausage as a Source of Hepatitis E Virus Transmission to Humans. <i>Journal of Infectious Diseases</i> , 2010, 202, 825-834.	1.9	571
7	rpoB -Based Identification of Nonpigmented and Late-Pigmenting Rapidly Growing Mycobacteria. <i>Journal of Clinical Microbiology</i> , 2003, 41, 5699-5708.	1.8	456
8	Early treatment of COVID-19 patients with hydroxychloroquine and azithromycin: A retrospective analysis of 1061 cases in Marseille, France. <i>Travel Medicine and Infectious Disease</i> , 2020, 35, 101738.	1.5	372
9	Identification of Rare Pathogenic Bacteria in a Clinical Microbiology Laboratory: Impact of Matrix-Assisted Laser Desorption Ionization-Flight Mass Spectrometry. <i>Journal of Clinical Microbiology</i> , 2013, 51, 2182-2194.	1.8	362
10	Chloroquine for the 2019 novel coronavirus SARS-CoV-2. <i>International Journal of Antimicrobial Agents</i> , 2020, 55, 105923.	1.1	354
11	Recycling of chloroquine and its hydroxyl analogue to face bacterial, fungal and viral infections in the 21st century. <i>International Journal of Antimicrobial Agents</i> , 2007, 30, 297-308.	1.1	332
12	Chronic Hepatitis E with Cirrhosis in a Kidney-Transplant Recipient. <i>New England Journal of Medicine</i> , 2008, 358, 859-860.	13.9	326
13	Modern clinical microbiology: new challenges and solutions. <i>Nature Reviews Microbiology</i> , 2013, 11, 574-585.	13.6	264
14	“Megavirales”, a proposed new order for eukaryotic nucleocytoplasmic large DNA viruses. <i>Archives of Virology</i> , 2013, 158, 2517-2521.	0.9	256
15	A comprehensive repertoire of prokaryotic species identified in human beings. <i>Lancet Infectious Diseases</i> , The, 2015, 15, 1211-1219.	4.6	250
16	Tailed giant Tupanvirus possesses the most complete translational apparatus of the known virosphere. <i>Nature Communications</i> , 2018, 9, 749.	5.8	247
17	In vitro testing of combined hydroxychloroquine and azithromycin on SARS-CoV-2 shows synergistic effect. <i>Microbial Pathogenesis</i> , 2020, 145, 104228.	1.3	246
18	Transfusion-associated Hepatitis E, France. <i>Emerging Infectious Diseases</i> , 2007, 13, 648-649.	2.0	226

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19	Outcomes of 3,737 COVID-19 patients treated with hydroxychloroquine/azithromycin and other regimens in Marseille, France: A retrospective analysis. <i>Travel Medicine and Infectious Disease</i> , 2020, 36, 101791.	1.5	209
20	Teicoplanin: an alternative drug for the treatment of COVID-19?. <i>International Journal of Antimicrobial Agents</i> , 2020, 55, 105944.	1.1	205
21	Faustovirus, an Asfarvirus-Related New Lineage of Giant Viruses Infecting Amoebae. <i>Journal of Virology</i> , 2015, 89, 6585-6594.	1.5	191
22	Correlation Between 3790 Quantitative Polymerase Chain Reactionâ€“Positives Samples and Positive Cell Cultures, Including 1941 Severe Acute Respiratory Syndrome Coronavirus 2 Isolates. <i>Clinical Infectious Diseases</i> , 2021, 72, e921-e921.	2.9	158
23	Pepper Mild Mottle Virus, a Plant Virus Associated with Specific Immune Responses, Fever, Abdominal Pains, and Pruritus in Humans. <i>PLoS ONE</i> , 2010, 5, e10041.	1.1	153
24	Reclassification of Giant Viruses Composing a Fourth Domain of Life in the New Order &#x2013; Megavirales. <i>Intervirology</i> , 2012, 55, 321-332.	1.2	149
25	Treatment of severe acute hepatitis E by ribavirin. <i>Journal of Clinical Virology</i> , 2011, 52, 60-62.	1.6	137
26	Evaluation of the Panbio COVID-19 Rapid Antigen Detection Test Device for the Screening of Patients with COVID-19. <i>Journal of Clinical Microbiology</i> , 2021, 59, .	1.8	136
27	Mimivirus: leading the way in the discovery of giant viruses of amoebae. <i>Nature Reviews Microbiology</i> , 2017, 15, 243-254.	13.6	132
28	First Isolation of Mimivirus in a Patient With Pneumonia. <i>Clinical Infectious Diseases</i> , 2013, 57, e127-e134.	2.9	131
29	Dolutegravir and weight gain. <i>Aids</i> , 2017, 31, 1499-1500.	1.0	129
30	Related Giant Viruses in Distant Locations and Different Habitats: Acanthamoeba polyphaga mouloumouvirus Represents a Third Lineage of the Mimiviridae That Is Close to the Megavirus Lineage. <i>Genome Biology and Evolution</i> , 2012, 4, 1324-1330.	1.1	118
31	Giant Viruses of Amoebas: An Update. <i>Frontiers in Microbiology</i> , 2016, 7, 349.	1.5	110
32	Rapid viral diagnosis and ambulatory management of suspected COVID-19 cases presenting at the infectious diseases referral hospital in Marseille, France, - January 31st to March 1st, 2020: A respiratory virus snapshot. <i>Travel Medicine and Infectious Disease</i> , 2020, 36, 101632.	1.5	109
33	â€œMarseilleviridaeâ€“, a new family of giant viruses infecting amoebae. <i>Archives of Virology</i> , 2013, 158, 915-920.	0.9	106
34	Mink, SARS-CoV-2, and the Human-Animal Interface. <i>Frontiers in Microbiology</i> , 2021, 12, 663815.	1.5	106
35	Autochthonous Infections with Hepatitis E Virus Genotype 4, France. <i>Emerging Infectious Diseases</i> , 2012, 18, 1361-1364.	2.0	103
36	The Large Marseillevirus Explores Different Entry Pathways by Forming Giant Infectious Vesicles. <i>Journal of Virology</i> , 2016, 90, 5246-5255.	1.5	103

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37	Zamilon, a Novel Virophage with Mimiviridae Host Specificity. PLoS ONE, 2014, 9, e94923.	1.1	101
38	Natural history of COVID-19 and therapeutic options. Expert Review of Clinical Immunology, 2020, 16, 1159-1184.	1.3	101
39	Evidence of the megavirome in humans. Journal of Clinical Virology, 2013, 57, 191-200.	1.6	100
40	Pacmanvirus, a New Giant Icosahedral Virus at the Crossroads between Asfarviridae and Faustoviruses. Journal of Virology, 2017, 91, .	1.5	99
41	Cedratvirus, a Double-Cork Structured Giant Virus, is a Distant Relative of Pithoviruses. Viruses, 2016, 8, 300.	1.5	98
42	Mimiviridae: clusters of orthologous genes, reconstruction of gene repertoire evolution and proposed expansion of the giant virus family. Virology Journal, 2013, 10, 106.	1.4	97
43	Hepatitis E virus as an emerging cause of chronic liver disease in organ transplant recipients. Journal of Hepatology, 2009, 50, 622-624.	1.8	91
44	Safety of TNF-blocking agents in rheumatic patients with serology suggesting past hepatitis B state: results from a cohort of 21 patients. Arthritis Research and Therapy, 2009, 11, R179.	1.6	91
45	A Decade of Improvements in Mimiviridae and Marseilleviridae Isolation from Amoeba. Intervirology, 2013, 56, 354-363.	1.2	88
46	Clinical and virological factors associated with hepatitis B virus reactivation in HBsAg-negative and anti-HBc antibodies-positive patients undergoing chemotherapy and/or autologous stem cell transplantation for cancer. Journal of Viral Hepatitis, 2010, 17, 807-815.	1.0	85
47	Structure of faustovirus, a large dsDNA virus. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 6206-6211.	3.3	84
48	Viruses with More Than 1,000 Genes: Mamavirus, a New Acanthamoeba polyphagomimivirus Strain, and Reannotation of Mimivirus Genes. Genome Biology and Evolution, 2011, 3, 737-742.	1.1	83
49	Hepatitis E in an HIV-infected patient. Journal of Clinical Virology, 2009, 45, 269-271.	1.6	78
50	Hepatitis E virus infection in patients infected with the human immunodeficiency virus. Journal of Medical Virology, 2011, 83, 1704-1716.	2.5	78
51	Clinical phenotype and outcome of hepatitis E virus-associated neuralgic amyotrophy. Neurology, 2017, 89, 909-917.	1.5	75
52	Shan Virus: A New Mimivirus Isolated from the Stool of a Tunisian Patient with Pneumonia. Intervirology, 2013, 56, 424-429.	1.2	74
53	A Brazilian Marseillevirus Is the Founding Member of a Lineage in Family Marseilleviridae. Viruses, 2016, 8, 76.	1.5	74
54	Ribavirin for Hepatitis E Virus Infection After Organ Transplantation: A Large European Retrospective Multicenter Study. Clinical Infectious Diseases, 2020, 71, 1204-1211.	2.9	74

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55	Emergence of Autochthonous Infections with Hepatitis E Virus of Genotype 4 in Europe. <i>Intervirolgy</i> , 2014, 57, 43-48.	1.2	73
56	Traditional and syndromic surveillance of infectious diseases and pathogens. <i>International Journal of Infectious Diseases</i> , 2016, 48, 22-28.	1.5	71
57	Repertoire of human gut microbes. <i>Microbial Pathogenesis</i> , 2017, 106, 103-112.	1.3	70
58	Complete genome sequence of Tunisivirus, a new member of the proposed family <i>Marseilleviridae</i> . <i>Archives of Virology</i> , 2014, 159, 2349-2358.	0.9	69
59	Antibiotic resistance surveillance systems: A review. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 23, 430-438.	0.9	69
60	Can Plant Viruses Cross the Kingdom Border and Be Pathogenic to Humans?. <i>Viruses</i> , 2015, 7, 2074-2098.	1.5	68
61	High-throughput isolation of giant viruses of the <i>Mimiviridae</i> and <i>Marseilleviridae</i> families in the Tunisian environment. <i>Environmental Microbiology</i> , 2013, 15, 2000-2007.	1.8	67
62	Prevalence of drug resistant mutants and virological response to combination therapy in patients with primary HIV-1 infection. , 2000, 61, 181-186.		64
63	Clinical and virological significance of the co-existence of HBsAg and anti-HBs antibodies in hepatitis B chronic carriers. <i>Virology</i> , 2007, 367, 30-40.	1.1	63
64	The puzzling mutational landscape of the SARS-CoV-2 variant Omicron. <i>Journal of Medical Virology</i> , 2022, 94, 2019-2025.	2.5	63
65	Fighting viruses with antibiotics: an overlooked path. <i>International Journal of Antimicrobial Agents</i> , 2016, 48, 349-352.	1.1	61
66	Giant Viruses of Amoebae: A Journey Through Innovative Research and Paradigm Changes. <i>Annual Review of Virology</i> , 2017, 4, 61-85.	3.0	61
67	Clinical efficacy of chloroquine derivatives in COVID-19 infection: comparative meta-analysis between the big data and the real world. <i>New Microbes and New Infections</i> , 2020, 38, 100709.	0.8	61
68	SARS-CoV-2: fear versus data. <i>International Journal of Antimicrobial Agents</i> , 2020, 55, 105947.	1.1	61
69	Multidrug Resistance Genotypes (Insertions in the P23 <sup>4</sup> Finger Subdomain and MDR Mutations) of HIV-1 Reverse Transcriptase from Extensively Treated Patients: Incidence and Association with Other Resistance Mutations. <i>Virology</i> , 2000, 270, 310-316.	1.1	58
70	Polymorphism and Drug-Selected Mutations in the Protease Gene of Human Immunodeficiency Virus Type 2 from Patients Living in Southern France. <i>Journal of Clinical Microbiology</i> , 2004, 42, 570-577.	1.8	58
71	Culture and identification of a Δomicron-SARS-CoV-2 in a three cases cluster in southern France. <i>Journal of Medical Virology</i> , 2022, 94, 3739-3749.	2.5	58
72	Infection with hepatitis E virus in kidney transplant recipients in southeastern France. <i>Journal of Medical Virology</i> , 2013, 85, 462-471.	2.5	57

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73	Discrepancy between anti-hepatitis E virus immunoglobulin G prevalence assessed by two assays in kidney and liver transplant recipients. <i>Journal of Clinical Virology</i> , 2013, 56, 62-64.	1.6	57
74	Influence of the HCV subtype on the virological response to pegylated interferon and ribavirin therapy. <i>Journal of Medical Virology</i> , 2009, 81, 2029-2035.	2.5	56
75	The Giant Cafeteria roenbergensis Virus That Infects a Widespread Marine Phagocytic Protist Is a New Member of the Fourth Domain of Life. <i>PLoS ONE</i> , 2011, 6, e18935.	1.1	56
76	Characteristics of the first 1119 SARS-CoV-2 Omicron variant cases, in Marseille, France, November~December 2021. <i>Journal of Medical Virology</i> , 2022, 94, 2290-2295.	2.5	56
77	First Isolation of a Marseillevirus in the Diptera Syrphidae <i>Eristalis tenax</i> . <i>Intervirology</i> , 2013, 56, 386-394.	1.2	55
78	Virological and Pharmacological Parameters Predicting the Response to Lopinavir-Ritonavir in Heavily Protease Inhibitor-Experienced Patients. <i>Antimicrobial Agents and Chemotherapy</i> , 2005, 49, 1720-1726.	1.4	52
79	Detection of hepatitis E virus in wild boar ( <i>Sus scrofa</i> ) livers. <i>Veterinary Journal</i> , 2010, 186, 259-261.	0.6	52
80	Neuropsychiatric adverse effects on dolutegravir. <i>Aids</i> , 2017, 31, 1201-1203.	1.0	52
81	Ultrarapid diagnosis, microscope imaging, genome sequencing, and culture isolation of SARS-CoV-2. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2020, 39, 1601-1603.	1.3	52
82	<i>Acanthamoeba polyphaga</i> mimivirus and other giant viruses: an open field to outstanding discoveries. <i>Virology Journal</i> , 2014, 11, 120.	1.4	51
83	COVID-19 re-infection. <i>European Journal of Clinical Investigation</i> , 2021, 51, e13537.	1.7	51
84	Frequent transmission of hepatitis E virus among piglets in farms in Southern France. <i>Journal of Medical Virology</i> , 2009, 81, 1750-1759.	2.5	50
85	NS3 Protease Polymorphism and Natural Resistance to Protease Inhibitors in French Patients Infected with HCV Genotypes 1-5. <i>Antiviral Therapy</i> , 2011, 16, 1093-1102.	0.6	50
86	Neurologic Disorders and Hepatitis E, France, 2010. <i>Emerging Infectious Diseases</i> , 2011, 17, 1510-2.	2.0	50
87	Mimivirus inaugurated in the 21st century the beginning of a reclassification of viruses. <i>Current Opinion in Microbiology</i> , 2016, 31, 16-24.	2.3	50
88	A Large Open Pangenome and a Small Core Genome for Giant Pandoraviruses. <i>Frontiers in Microbiology</i> , 2018, 9, 1486.	1.5	48
89	Evidence of SARS-CoV-2 re-infection with a different genotype. <i>Journal of Infection</i> , 2021, 82, 84-123.	1.7	47
90	Analysis of SARS-CoV-2 Variants From 24,181 Patients Exemplifies the Role of Globalization and Zoonosis in Pandemics. <i>Frontiers in Microbiology</i> , 2021, 12, 786233.	1.5	46

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91	SARS-CoV-2 reinfection and COVID-19 severity. <i>Emerging Microbes and Infections</i> , 2022, 11, 894-901.	3.0	46
92	Genomic and phylogenetic analysis of hepatitis C virus isolates: A survey of 535 strains circulating in southern France. <i>Journal of Medical Virology</i> , 2003, 71, 391-398.	2.5	44
93	Severe Thrombocytopenia Associated with Acute Hepatitis E Virus Infection. <i>Journal of Clinical Microbiology</i> , 2008, 46, 2450-2452.	1.8	44
94	Complete genome sequence of Cannes 8 virus, a new member of the proposed family "Marseilleviridae". <i>Virus Genes</i> , 2013, 47, 550-555.	0.7	44
95	Marseillevirus prevalence in multitransfused patients suggests blood transmission. <i>Journal of Clinical Virology</i> , 2013, 58, 722-725.	1.6	44
96	Ancestrality and Mosaicism of Giant Viruses Supporting the Definition of the Fourth TRUC of Microbes. <i>Frontiers in Microbiology</i> , 2018, 9, 2668.	1.5	44
97	Emergence and outcomes of the SARS-CoV-2 "Marseille-4"™ variant. <i>International Journal of Infectious Diseases</i> , 2021, 106, 228-236.	1.5	44
98	Group A streptococcal infections during the seasonal influenza outbreak 2010/11 in South East England. <i>Eurosurveillance</i> , 2011, 16, .	3.9	43
99	Acute and chronic hepatitis E in patients infected with human immunodeficiency virus. <i>Journal of Viral Hepatitis</i> , 2011, 18, 227-228.	1.0	42
100	Chronic Hepatitis E Virus Infection Is Specifically Associated With an Interferon-Related Transcriptional Program. <i>Journal of Infectious Diseases</i> , 2013, 207, 125-132.	1.9	42
101	Gene Repertoire of Amoeba-Associated Giant Viruses. <i>Intervirology</i> , 2010, 53, 330-343.	1.2	41
102	Hepatitis E in three immunocompromized children in southeastern France. <i>Journal of Clinical Virology</i> , 2012, 53, 162-166.	1.6	41
103	DNA-Dependent RNA Polymerase Detects Hidden Giant Viruses in Published Databanks. <i>Genome Biology and Evolution</i> , 2014, 6, 1603-1610.	1.1	41
104	Severe Pulmonary Arterial Hypertension in Patients Treated for Hepatitis C With Sofosbuvir. <i>Chest</i> , 2016, 149, e69-e73.	0.4	41
105	Yaravirus: A novel 80-nm virus infecting <i>Acanthamoeba castellanii</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 16579-16586.	3.3	41
106	Children account for a small proportion of diagnoses of SARS-CoV-2 infection and do not exhibit greater viral loads than adults. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2020, 39, 1983-1987.	1.3	40
107	Hepatitis E virus as a newly identified cause of acute viral hepatitis during human immunodeficiency virus infection. <i>Clinical Microbiology and Infection</i> , 2008, 14, 1176-1180.	2.8	39
108	The Expanding Family of Virophages. <i>Viruses</i> , 2016, 8, 317.	1.5	39

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109	Testing the repatriated for SARS-Cov2: Should laboratory-based quarantine replace traditional quarantine?. <i>Travel Medicine and Infectious Disease</i> , 2020, 34, 101624.	1.5	39
110	Emergence in southern France of a new SARS-CoV-2 variant harbouring both N501Y and E484K substitutions in the spike protein. <i>Archives of Virology</i> , 2022, 167, 1185-1190.	0.9	39
111	Real-Time Microbiology Laboratory Surveillance System to Detect Abnormal Events and Emerging Infections, Marseille, France. <i>Emerging Infectious Diseases</i> , 2015, 21, 1302-1310.	2.0	38
112	The Investigation of Promoter Sequences of Marseilleviruses Highlights a Remarkable Abundance of the AAATATTT Motif in Intergenic Regions. <i>Journal of Virology</i> , 2017, 91, .	1.5	37
113	Usefulness of rpoB Gene Sequencing for Identification of Afipia and Bosea Species, Including a Strategy for Choosing Discriminative Partial Sequences. <i>Applied and Environmental Microbiology</i> , 2003, 69, 6740-6749.	1.4	36
114	Marseillevirus in lymphoma: a giant in the lymph node. <i>Lancet Infectious Diseases</i> , The, 2016, 16, e225-e234.	4.6	36
115	The analysis of translation-related gene set boosts debates around origin and evolution of mimiviruses. <i>PLoS Genetics</i> , 2017, 13, e1006532.	1.5	36
116	SARS-CoV-2 Infectivity and Severity of COVID-19 According to SARS-CoV-2 Variants: Current Evidence. <i>Journal of Clinical Medicine</i> , 2021, 10, 2635.	1.0	36
117	Investigation of <i>Acinetobacter baumannii</i> resistance to carbapenems in Marseille hospitals, south of France: a transition from an epidemic to an endemic situation. <i>Apmis</i> , 2013, 121, 64-71.	0.9	34
118	HIV infection en route to endogenization: two cases. <i>Clinical Microbiology and Infection</i> , 2014, 20, 1280-1288.	2.8	34
119	MG-Digger: An Automated Pipeline to Search for Giant Virus-Related Sequences in Metagenomes. <i>Frontiers in Microbiology</i> , 2016, 7, 428.	1.5	34
120	Human metallo- $\beta$ -lactamase enzymes degrade penicillin. <i>Scientific Reports</i> , 2019, 9, 12173.	1.6	34
121	Prevalence of resistance mutations in antiretroviral-naïve chronically HIV-infected patients in 1998: a French nationwide study. <i>Aids</i> , 2001, 15, 1777-1782.	1.0	33
122	Giant Viruses of Amoebae as Potential Human Pathogens. <i>Intervirology</i> , 2013, 56, 376-385.	1.2	33
123	Hydroxychloroquine and azithromycin as a treatment of COVID-19: results of an open label non-randomized clinical trial revisited. <i>International Journal of Antimicrobial Agents</i> , 2021, 57, 106243.	1.1	33
124	Natural presence of substitution R155K within hepatitis C virus NS3 protease from a treatment-naïve chronically infected patient. <i>Hepatology</i> , 2008, 47, 766-767.	3.6	31
125	Factors influencing peripheral blood mononuclear cell-associated HIV-1 DNA level after long-term suppressive antiretroviral therapy in 236 patients. <i>Aids</i> , 2009, 23, 2165-2171.	1.0	31
126	Faustovirus E12 Transcriptome Analysis Reveals Complex Splicing in Capsid Gene. <i>Frontiers in Microbiology</i> , 2018, 9, 2534.	1.5	31



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127	Contribution of VitaPCR SARS-CoV-2 to the emergency diagnosis of COVID-19. <i>Journal of Clinical Virology</i> , 2020, 133, 104682.	1.6	31
128	Child with liver transplant recovers from COVID-19 infection. A case report. <i>Archives De Pediatrie</i> , 2020, 27, 275-276.	0.4	31
129	Introduction into the Marseille geographical area of a mild SARS-CoV-2 variant originating from sub-Saharan Africa: An investigational study. <i>Travel Medicine and Infectious Disease</i> , 2021, 40, 101980.	1.5	31
130	Polymorphism and drug-selected mutations in the reverse transcriptase gene of HIV-2 from patients living in southeastern France. <i>Journal of Medical Virology</i> , 2005, 75, 381-390.	2.5	30
131	The expanding family <i>Marseilleviridae</i> . <i>Virology</i> , 2014, 466-467, 27-37.	1.1	30
132	Evaluating the serological status of COVID-19 patients using an indirect immunofluorescent assay, France. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021, 40, 361-371.	1.3	30
133	Spreading of a new SARS-CoV-2 N501Y spike variant in a new lineage. <i>Clinical Microbiology and Infection</i> , 2021, 27, 1352.e1-1352.e5.	2.8	30
134	Entecavir as a first-line treatment for HBV reactivation following polychemotherapy for lymphoma. <i>British Journal of Haematology</i> , 2008, 143, 148-150.	1.2	29
135	First Isolation of a Giant Virus from Wild <i>Hirudo medicinalis</i> Leech: <i>Mimiviridae</i> isolation in <i>Hirudo medicinalis</i> . <i>Viruses</i> , 2013, 5, 2920-2930.	1.5	29
136	Co-infection of SARS-CoV-2 and influenza viruses: A systematic review and meta-analysis. <i>Journal of Clinical Virology Plus</i> , 2021, 1, 100036.	0.4	29
137	Tobacco mosaic virus in cigarettes and saliva of smokers. <i>Journal of Clinical Virology</i> , 2012, 55, 374-376.	1.6	28
138	Virophages of Giant Viruses: An Update at Eleven. <i>Viruses</i> , 2019, 11, 733.	1.5	28
139	Acute polyarthritis revealing hepatitis E. <i>Clinical Rheumatology</i> , 2007, 26, 1973-1975.	1.0	27
140	Detection of hepatitis E virus of genotype 3 in a farm pig in Kinshasa (Democratic Republic of the Congo). <i>Journal of Clinical Microbiology</i> , 2010, 148, 1010-1012.	1.0	27
141	First Human Case of Co-Infection with Two Different Subtypes of Hepatitis E Virus. <i>Intervirology</i> , 2012, 55, 484-487.	1.2	27
142	Chronic hepatitis E resolution in a human immunodeficiency virus (HIV)-infected patient treated with ribavirin. <i>International Journal of Antimicrobial Agents</i> , 2013, 41, 595-597.	1.1	27
143	Hepatocellular carcinoma complicating hepatitis E virus-related cirrhosis. <i>Hepatology</i> , 2018, 67, 446-448.	3.6	27
144	A Phylogenomic Study of <i>Acanthamoeba polyphaga</i> Draft Genome Sequences Suggests Genetic Exchanges With Giant Viruses. <i>Frontiers in Microbiology</i> , 2018, 9, 2098.	1.5	27

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145	Implementation of an in-house real-time reverse transcription-PCR assay for the rapid detection of the SARS-CoV-2 Marseille-4 variant. <i>Journal of Clinical Virology</i> , 2021, 139, 104814.	1.6	27
146	SVARAP and aSVARAP: simple tools for quantitative analysis of nucleotide and amino acid variability and primer selection for clinical microbiology. <i>BMC Microbiology</i> , 2006, 6, 21.	1.3	26
147	Hepatitis E associated with surgical training on pigs. <i>Lancet, The</i> , 2007, 370, 935.	6.3	26
148	Extensive Gene Remodeling in the Viral World: New Evidence for Nongradual Evolution in the Mobilome Network. <i>Genome Biology and Evolution</i> , 2014, 6, 2195-2205.	1.1	26
149	Pan-Genome Analysis of Brazilian Lineage A Amoebal Mimiviruses. <i>Viruses</i> , 2015, 7, 3483-3499.	1.5	26
150	High rate of reinfection with the SARS-CoV-2 Omicron variant. <i>Journal of Infection</i> , 2022, 85, 174-211.	1.7	26
151	Drug repurposing against SARS-CoV-1, SARS-CoV-2 and MERS-CoV. <i>Future Microbiology</i> , 2021, 16, 1341-1370.	1.0	25
152	Omicron Variant Escapes Therapeutic Monoclonal Antibodies (mAbs) Including Recently Released Evusheld <sup>®</sup> , Contrary to 8 Prior Main Variant of Concern (VOC). <i>Clinical Infectious Diseases</i> , 2022, 75, e534-e535.	2.9	25
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