Philippe Colson

List of Publications by Year
in descending order

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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	Incomplete tricarboxylic acid cycle and proton gradient in Pandoravirus massiliensis: is it still a virus?. ISME Journal, 2022, 16, 695-704.	4.4	16
2	Occurrence of a substitution or deletion of SARS-CoV-2 spike amino acid 677 in various lineages in Marseille, France. Virus Genes, 2022, 58, 53-58.	0.7	8
3	The puzzling mutational landscape of the SARSâ€2â€variant Omicron. Journal of Medical Virology, 2022, 94, 2019-2025.	2.5	63
4	Epidemiology and genomic characterisation of travel-associated and locally-acquired influenza, Marseille, France. Travel Medicine and Infectious Disease, 2022, 45, 102236.	1.5	0
5	Limited spread of a rare spike E484K-harboring SARS-CoV-2 in Marseille, France. Archives of Virology, 2022, 167, 583.	0.9	3
6	The emergence, dynamics and significance of SARS-CoV-2 variants. New Microbes and New Infections, 2022, 45, 100962.	0.8	5
7	Characteristics of the first 1119 SARSâ€CoVâ€2 Omicron variant cases, in Marseille, France, Novemberâˆ'December 2021. Journal of Medical Virology, 2022, 94, 2290-2295.	2.5	56
8	Introduction of the SARS-CoV-2 Beta variant from Comoros into the Marseille geographical area. Travel Medicine and Infectious Disease, 2022, 46, 102277.	1.5	3
9	Omicron Variant Escapes Therapeutic Monoclonal Antibodies (mAbs) Including Recently Released EvusheldÂ®, Contrary to 8 Prior Main Variant of Concern (VOC). Clinical Infectious Diseases, 2022, 75, e534-e535.	2.9	25
10	Emergence in southern France of a new SARS-CoV-2 variant harbouring both N501Y and E484K substitutions in the spike protein. Archives of Virology, 2022, 167, 1185-1190.	0.9	39
11	SARS-CoV-2 reinfection and COVID-19 severity. Emerging Microbes and Infections, 2022, 11, 894-901.	3.0	46
12	Hepatitis delta treatment with bulevirtide in real life: a case report. International Journal of Antimicrobial Agents, 2022, 59, 106562.	1.1	5
13	First cases of infection with the 21L/BA.2 Omicron variant in Marseille, France. Journal of Medical Virology, 2022, 94, 3421-3430.	2.5	19
14	The severity of the first 207 infections with the SARSâ€CoVâ€2 Omicron BA.2 variant, in Marseille, France, December 2021â€"FebruaryÂ2022. Journal of Medical Virology, 2022, 94, 3494-3497.	2.5	18
15	Influenza A imported from Comoros to France in the summer months of 2021. Journal of Travel Medicine, 2022, 29, .	1.4	0
16	Emerging SARS-CoV-2 Genotypes Show Different Replication Patterns in Human Pulmonary and Intestinal Epithelial Cells. Viruses, 2022, 14, 23.	1.5	16
17	Control of common viral epidemics but not of SARS-CoV-2 through the application of hygiene and distancing measures. Journal of Clinical Virology, 2022, 150-151, 105163.	1.6	6
18	Culture and identification of a â€œDeltamicronâ€•SARSâ€CoVâ€2 in a three cases cluster in southern France. Journal of Medical Virology, 2022, 94, 3739-3749.	2.5	58

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19	Who Were Hospitalized Deceased Patients from COVID-19 During the First Year of Pandemic? Retrospective Analysis of 1104 Deceased Patients in South of France. Journal of Epidemiology and Global Health, 2022, , .	1.1	0
20	High rate of reinfection with the SARS-CoV-2 Omicron variant. Journal of Infection, 2022, 85, 174-211.	1.7	26
21	Screening and Whole Genome Sequencing of SARS-CoV-2 Circulating During the First Three Waves of the COVID-19 Pandemic in Libreville and the Haut-OgoouÃ© Province in Gabon. Frontiers in Medicine, 2022, 9, .	1.2	7
22	Sequential Appearance and Isolation of a SARS-CoV-2 Recombinant between Two Major SARS-CoV-2 Variants in a Chronically Infected Immunocompromised Patient. Viruses, 2022, 14, 1266.	1.5	13
23	Outbreak of central nervous system infections among children in Thai Binh, Viet Nam. Emerging Microbes and Infections, 2022, 11, 1683-1692.	3.0	3
24	SARS-CoV-2 Testing of Aircraft Wastewater Shows That Mandatory Tests and Vaccination Pass before Boarding Did Not Prevent Massive Importation of Omicron Variant into Europe. Viruses, 2022, 14, 1511.	1.5	9
25	Evaluation of the Panbio COVID-19 Rapid Antigen Detection Test Device for the Screening of Patients with COVID-19. Journal of Clinical Microbiology, 2021, 59, .	1.8	136
26	Hostâ€“virus interactions and defense mechanisms for giant viruses. Annals of the New York Academy of Sciences, 2021, 1486, 39-57.	1.8	3
27	Different pattern of the second outbreak of COVID-19 in Marseille, France. International Journal of Infectious Diseases, 2021, 102, 17-19.	1.5	12
28	Correlation Between 3790 Quantitative Polymerase Chain Reactionâ€“Positives Samples and Positive Cell Cultures, Including 1941 Severe Acute Respiratory Syndrome Coronavirus 2 Isolates. Clinical Infectious Diseases, 2021, 72, e921-e921.	2.9	158
29	Resistance to human immunodeficiency virus infection: a rare but neglected state. Annals of the New York Academy of Sciences, 2021, 1485, 22-42.	1.8	0
30	Interpretation of SARS-CoV-2 PCR results for the diagnosis of COVID-19. International Journal of Antimicrobial Agents, 2021, 57, 106238.	1.1	0
31	Evaluating the serological status of COVID-19 patients using an indirect immunofluorescent assay, France. European Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 361-371.	1.3	30
32	Hydroxychloroquine and azithromycin as a treatment of COVID-19: results of an open label non-randomized clinical trial revisited. International Journal of Antimicrobial Agents, 2021, 57, 106243.	1.1	33
33	Coinfections with SARSâ€CoVâ€2 and other respiratory viruses in Southeastern France: A matter of sampling time. Journal of Medical Virology, 2021, 93, 1878-1881.	2.5	14
34	Evidence of SARS-CoV-2 re-infection with a different genotype. Journal of Infection, 2021, 82, 84-123.	1.7	47
35	Early combination therapy with hydroxychloroquine and azithromycin reduces mortality in 10,429 COVID-19 outpatients. Reviews in Cardiovascular Medicine, 2021, 22, 1063.	0.5	21
36	Epidemiological serosurvey and molecular characterization of sexually transmitted infections among 1890 sheltered homeless people in Marseille: Cross-sectional one day-surveys (2000â€“2015). Journal of Infection, 2021, 82, 60-66.	1.7	3

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37	Giant virus-related sequences in the 5300-year-old Ã–tzi mummy metagenome. Virus Genes, 2021, 57, 222-227.	0.7	2
38	Development and validation of a multiplex qPCR assay for detection and relative quantification of HPV16 and HPV18 E6 and E7 oncogenes. Scientific Reports, 2021, 11, 4039.	1.6	10
39	No barrier to care, yet disparities in the HIV care continuum in France: a nationwide population study. Journal of Antimicrobial Chemotherapy, 2021, 76, 1573-1579.	1.3	2
40	Automated Western immunoblotting detection of anti-SARS-CoV-2 serum antibodies. European Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 1309-1317.	1.3	23
41	Introduction into the Marseille geographical area of a mild SARS-CoV-2 variant originating from sub-Saharan Africa: An investigational study. Travel Medicine and Infectious Disease, 2021, 40, 101980.	1.5	31
42	Direct next-generation sequencing diagnosis of echovirus 9 meningitis, France. European Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 2037-2039.	1.3	8
43	COVIDâ€19 reâ€infection. European Journal of Clinical Investigation, 2021, 51, e13537.	1.7	51
44	Effect of hydroxychloroquine and azithromycin on the viral clearance of SARS-CoV-2: response to HervÃ© Seligmann. International Journal of Antimicrobial Agents, 2021, 57, 106306.	1.1	1
45	Mink, SARS-CoV-2, and the Human-Animal Interface. Frontiers in Microbiology, 2021, 12, 663815.	1.5	106
46	Global Discrepancies between Numbers of Available SARS-CoV-2 Genomes and Human Development Indexes at Country Scales. Viruses, 2021, 13, 775.	1.5	7
47	Extensive Comparative Genomic Analysis of Enterococcus faecalis and Enterococcus faecium Reveals a Direct Association between the Absence of CRISPRâ€"Cas Systems, the Presence of Anti-Endonuclease (ardA) and the Acquisition of Vancomycin Resistance in E. faecium. Microorganisms, 2021, 9, 1118.	1.6	6
48	Direct Diagnosis of Echovirus 12 Meningitis Using Metagenomic Next Generation Sequencing. Pathogens, 2021, 10, 610.	1.2	5
49	SARS-CoV-2 Persistent Viral Shedding in the Context of Hydroxychloroquine-Azithromycin Treatment. Viruses, 2021, 13, 890.	1.5	7
50	Rhizomal Reclassification of Living Organisms. International Journal of Molecular Sciences, 2021, 22, 5643.	1.8	16
51	Clinical outcomes in COVID-19 patients infected with different SARS-CoV-2 variants in Marseille, France. Clinical Microbiology and Infection, 2021, 27, 1516.e1-1516.e6.	2.8	18
52	Emergence and outcomes of the SARS-CoV-2 â€˜Marseille-4â€™ variant. International Journal of Infectious Diseases, 2021, 106, 228-236.	1.5	44
53	SARS-CoV-2 Infectivity and Severity of COVID-19 According to SARS-CoV-2 Variants: Current Evidence. Journal of Clinical Medicine, 2021, 10, 2635.	1.0	36
54	Implementation of an in-house real-time reverse transcription-PCR assay for the rapid detection of the SARS-CoV-2 Marseille-4 variant. Journal of Clinical Virology, 2021, 139, 104814.	1.6	27

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55	Evaluation of Strategies to Fight COVID-19: The French Paradigm. Journal of Clinical Medicine, 2021, 10, 2942.	1.0	4
56	Marseilleviruses: An Update in 2021. Frontiers in Microbiology, 2021, 12, 648731.	1.5	14
57	Diverse Trajectories Drive the Expression of a Giant Virus in the Oomycete Plant Pathogen Phytophthora parasitica. Frontiers in Microbiology, 2021, 12, 662762.	1.5	12
58	Implementation of an in-house real-time reverse transcription-PCR assay to detect the emerging SARS-CoV-2 N501Y variants. Journal of Clinical Virology, 2021, 140, 104868.	1.6	15
59	A Possible Role of Remdesivir and Plasma Therapy in the Selective Sweep and Emergence of New SARS-CoV-2 Variants. Journal of Clinical Medicine, 2021, 10, 3276.	1.0	18
60	SARS-CoV-2 variant from India to Marseille: The still active role of ports in the introduction of epidemics. Travel Medicine and Infectious Disease, 2021, 42, 102085.	1.5	11
61	Consequences of the COVID-19 Outbreak Lockdown on Non-Viral Infectious Agents as Reported by a Laboratory-Based Surveillance System at the IHU MÃ©diterranÃ©e Infection, Marseille, France. Journal of Clinical Medicine, 2021, 10, 3210.	1.0	3
62	Clinical characteristics of paediatric COVIDâ€19 patients followed for up to 13Âmonths. Acta Paediatrica, International Journal of Paediatrics, 2021, 110, 3331-3333.	0.7	20
63	Monitoring the Circulation of SARS-CoV-2 Variants by Genomic Analysis of Wastewater in Marseille, South-East France. Pathogens, 2021, 10, 1042.	1.2	23
64	Whole Genome Sequencing of SARS-CoV-2 Strains in COVID-19 Patients From Djibouti Shows Novel Mutations and Clades Replacing Over Time. Frontiers in Medicine, 2021, 8, 737602.	1.2	4
65	Spreading of a new SARS-CoV-2 N501Y spike variant in a new lineage. Clinical Microbiology and Infection, 2021, 27, 1352.e1-1352.e5.	2.8	30
66	Epidemiological and genetic characterization of measles virus circulating strains at Marseille, France during 2017â€“2019 measles outbreak. Journal of Infection, 2021, 83, 361-370.	1.7	6
67	Co-infection of SARS-CoV-2 and influenza viruses: A systematic review and meta-analysis. Journal of Clinical Virology Plus, 2021, 1, 100036.	0.4	29
68	Clinical outcomes in patients infected with different SARS-CoV-2 variants at one hospital during three phases of the COVID-19 epidemic in Marseille, France. Infection, Genetics and Evolution, 2021, 95, 105092.	1.0	22
69	Concurrent Nanopore Next-Generation Sequencing of Hepatitis B and Delta Virus Genomes Directly From Patient Plasma. Annals of Laboratory Medicine, 2021, 41, 608-611.	1.2	1
70	Microelimination or Not? The Changing Epidemiology of Human Immunodeficiency Virus-Hepatitis C Virus Coinfection in France 2012â€“2018. Clinical Infectious Diseases, 2021, 73, e3266-e3274.	2.9	8
71	High Individual Heterogeneity of Neutralizing Activities against the Original Strain and Nine Different Variants of SARS-CoV-2. Viruses, 2021, 13, 2177.	1.5	21
72	Drug repurposing against SARS-CoV-1, SARS-CoV-2Âand MERS-CoV. Future Microbiology, 2021, 16, 1341-1370.	1.0	25

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73	Isolation of 4000 SARS-CoV-2 shows that contagiousness is associated with viral load, not vaccine or symptomatic status. Emerging Microbes and Infections, 2021, 10, 2276-2278.	3.0	5
74	SARS-CoV-2 Vaccination and Protection Against Clinical Disease: A Retrospective Study, Bouches-du-RhÃ´ne District, Southern France, 2021. Frontiers in Microbiology, 2021, 12, 796807.	1.5	5
75	Analysis of SARS-CoV-2 Variants From 24,181 Patients Exemplifies the Role of Globalization and Zoonosis in Pandemics. Frontiers in Microbiology, 2021, 12, 786233.	1.5	46
76	Heterogeneity in susceptibility to hydroxychloroquine of SARS-CoV-2 isolates. Frontiers in Bioscience, 2021, 26, 1493-1502.	0.8	2
77	Incidence and Outcome of Coinfections with SARS-CoV-2 and Rhinovirus. Viruses, 2021, 13, 2528.	1.5	20
78	International Journal of Antimicrobial Agents Themed Section Issue: â€œTherapeutics of hepatitis virusesâ€• International Journal of Antimicrobial Agents, 2020, 56, 105402.	1.1	2
79	Chronic hepatitis E in absence of severe immune deficiency. Clinics and Research in Hepatology and Gastroenterology, 2020, 44, e1-e4.	0.7	4
80	Hepatitis B Virus Genomics Knocking at the Door of Routine Diagnostic Laboratories. Journal of Infectious Diseases, 2020, 221, 1026-1029.	1.9	2
81	Hepatitis Delta recurrence post-liver transplantation in absence of detectable hepatitis B surface antigen and hepatitis B virus DNA in peripheral blood. Clinics and Research in Hepatology and Gastroenterology, 2020, 44, e41-e44.	0.7	2
82	Multimorbidity in Elderly Persons According to the Year of Diagnosis of Human Immunodeficiency Virus Infection: A Cross-sectional Datâ€™AIDS Cohort Study. Clinical Infectious Diseases, 2020, 71, 2880-2888.	2.9	10
83	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
84	Temporal and age distributions of SARS-CoV-2 and other coronaviruses, southeastern France. International Journal of Infectious Diseases, 2020, 101, 121-125.	1.5	6
85	Proteomics and Lipidomics Investigations to Decipher the Behavior of Willaertia magna C2c Maky According to Different Culture Modes. Microorganisms, 2020, 8, 1791.	1.6	1
86	Hepatitis A seroprevalence in homeless persons, in Marseille, France. Clinics and Research in Hepatology and Gastroenterology, 2020, 45, 101571.	0.7	1
87	Antibiotic resistance surveillance systems: A review. Journal of Global Antimicrobial Resistance, 2020, 23, 430-438.	0.9	69
88	Ultrarapid diagnosis, microscope imaging, genome sequencing, and culture isolation of SARS-CoV-2. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 1601-1603.	1.3	52
89	Major discrepancy between factual antibiotic resistance and consumption in South of France: analysis of 539,037 bacterial strains. Scientific Reports, 2020, 10, 18262.	1.6	5
90	Contribution of VitaPCR SARS-CoV-2 to the emergency diagnosis of COVID-19. Journal of Clinical Virology, 2020, 133, 104682.	1.6	31

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91	Promiscuous Enzyme Activity as a Driver of Allo and Iso Convergent Evolution, Lessons from the Î²-Lactamases. International Journal of Molecular Sciences, 2020, 21, 6260.	1.8	5
92	Hepatitis A outbreak in HIV-infected patients in Southeastern France: questions and responses?. Epidemiology and Infection, 2020, 148, e79.	1.0	4
93	Children account for a small proportion of diagnoses of SARS-CoV-2 infection and do not exhibit greater viral loads than adults. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 1983-1987.	1.3	40
94	Incidence of diabetes in HIV-infected patients treated with first-line integrase strand transfer inhibitors: a French multicentre retrospective study. Journal of Antimicrobial Chemotherapy, 2020, 75, 3344-3348.	1.3	22
95	A protein of the metallo-hydrolase/oxidoreductase superfamily with both beta-lactamase and ribonuclease activity is linked with translation in giant viruses. Scientific Reports, 2020, 10, 21685.	1.6	9
96	Early treatment of COVID-19 patients with hydroxychloroquine and azithromycin: A retrospective analysis of 1061 cases in Marseille, France. Travel Medicine and Infectious Disease, 2020, 35, 101738.	1.5	372
97	Analysis of a Marseillevirus Transcriptome Reveals Temporal Gene Expression Profile and Host Transcriptional Shift. Frontiers in Microbiology, 2020, 11, 651.	1.5	20
98	Comparison of mortality associated with respiratory viral infections between December 2019 and March 2020 with that of the previous year in Southeastern France. International Journal of Infectious Diseases, 2020, 96, 154-156.	1.5	16
99	Child with liver transplant recovers from COVID-19 infection. A case report. Archives De Pediatrie, 2020, 27, 275-276.	0.4	31
100	Vermamoeba vermiformis CDC-19 draft genome sequence reveals considerable gene trafficking including with candidate phyla radiation and giant viruses. Scientific Reports, 2020, 10, 5928.	1.6	20
101	A virophage cross-species infection through mutant selection represses giant virus propagation, promoting host cell survival. Communications Biology, 2020, 3, 248.	2.0	15
102	Does spitting in public play a role in transmitting SARS-CoV-2?. Travel Medicine and Infectious Disease, 2020, 36, 101759.	1.5	5
103	Insight into the Lifestyle of Amoeba Willaertia magna during Bioreactor Growth Using Transcriptomics and Proteomics. Microorganisms, 2020, 8, 771.	1.6	5
104	Clinical efficacy of chloroquine derivatives in COVID-19 infection: comparative meta-analysis between the big data and the real world. New Microbes and New Infections, 2020, 38, 100709.	0.8	61
105	Testing the repatriated for SARS-Cov2: Should laboratory-based quarantine replace traditional quarantine?. Travel Medicine and Infectious Disease, 2020, 34, 101624.	1.5	39
106	Description of Virulent Factors and Horizontal Gene Transfers of Keratitis-Associated Amoeba Acanthamoeba Triangularis by Genome Analysis. Pathogens, 2020, 9, 217.	1.2	14
107	SARS-CoV-2: fear versus data. International Journal of Antimicrobial Agents, 2020, 55, 105947.	1.1	61
108	New insights on the antiviral effects of chloroquine against coronavirus: what to expect for COVID-19?. International Journal of Antimicrobial Agents, 2020, 55, 105938.	1.1	842

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109	Teicoplanin: an alternative drug for the treatment of COVID-19?. International Journal of Antimicrobial Agents, 2020, 55, 105944.	1.1	205
110	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. Travel Medicine and Infectious Disease, 2020, 36, 101632.	1.5	109
111	Chloroquine and hydroxychloroquine as available weapons to fight COVID-19. International Journal of Antimicrobial Agents, 2020, 55, 105932.	1.1	724
112	Yaravirus: A novel 80-nm virus infecting Acanthamoeba castellanii. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 16579-16586.	3.3	41
113	Viral RNA load as determined by cell culture as a management tool for discharge of SARS-CoV-2 patients from infectious disease wards. European Journal of Clinical Microbiology and Infectious Diseases, 2020, 39, 1059-1061.	1.3	767
114	Outcomes of 3,737 COVID-19 patients treated with hydroxychloroquine/azithromycin and other regimens in Marseille, France: A retrospective analysis. Travel Medicine and Infectious Disease, 2020, 36, 101791.	1.5	209
115	Core gene-based molecular detection and identification of Acanthamoeba species. Scientific Reports, 2020, 10, 1583.	1.6	12
116	Dramatic HIV DNA degradation associated with spontaneous HIV suppression and disease-free outcome in a young seropositive woman following her infection. Scientific Reports, 2020, 10, 2548.	1.6	3
117	Chloroquine for the 2019 novel coronavirus SARS-CoV-2. International Journal of Antimicrobial Agents, 2020, 55, 105923.	1.1	354
118	Kaposi sarcoma among people living with <scp>HIV</scp> in the French <scp>DAT</scp>â€™<scp>AIDS</scp> cohort between 2010 and 2015. Journal of the European Academy of Dermatology and Venereology, 2020, 34, 1065-1073.	1.3	4
119	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. Travel Medicine and Infectious Disease, 2020, 34, 101663.	1.5	605
120	In vitro testing of combined hydroxychloroquine and azithromycin on SARS-CoV-2 shows synergistic effect. Microbial Pathogenesis, 2020, 145, 104228.	1.3	246
121	Hydroxychloroquine and azithromycin as a treatment of COVID-19: results of an open-label non-randomized clinical trial. International Journal of Antimicrobial Agents, 2020, 56, 105949.	1.1	3,955
122	Natural history of COVID-19 and therapeutic options. Expert Review of Clinical Immunology, 2020, 16, 1159-1184.	1.3	101
123	Respiratory Syncytial Virus Infection: Its Propensity for Bacterial Coinfection and Related Mortality in Elderly Adults. Open Forum Infectious Diseases, 2020, 7, ofaa546.	0.4	16
124	Letter to the editor: Plenty of coronaviruses but no SARS-CoV-2. Eurosurveillance, 2020, 25, .	3.9	8
125	Virophages of Giant Viruses: An Update at Eleven. Viruses, 2019, 11, 733.	1.5	28
126	Human metallo-Î²-lactamase enzymes degrade penicillin. Scientific Reports, 2019, 9, 12173.	1.6	34

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127	The Methodology Behind Network Thinking: Graphs to Analyze Microbial Complexity and Evolution. Methods in Molecular Biology, 2019, 1910, 271-308.	0.4	4
128	Measles, the need for a paradigm shift. European Journal of Epidemiology, 2019, 34, 897-915.	2.5	19
129	Human papillomavirus type 7â€associated anal condyloma after renal transplantation in a child. Pediatric Transplantation, 2019, 23, e13470.	0.5	1
130	Discovery and Further Studies on Giant Viruses at the IHU Mediterranee Infection That Modified the Perception of the Virosphere. Viruses, 2019, 11, 312.	1.5	23
131	Multidisciplinary evaluation of Clostridium butyricum clonality isolated from preterm neonates with necrotizing enterocolitis in South France between 2009 and 2017. Scientific Reports, 2019, 9, 2077.	1.6	10
132	The PACASurvE laboratory network for real-time infection surveillance and alert at a regional scale. European Journal of Public Health, 2019, 29, .	0.1	0
133	Sequence Similarities between Viroids and Human MicroRNAs. Intervirology, 2019, 62, 227-234.	1.2	5
134	Evidence of a Cellulosic Layer in Pandoravirus massiliensis Tegument and the Mystery of the Genetic Support of Its Biosynthesis. Frontiers in Microbiology, 2019, 10, 2932.	1.5	14
135	Investigation of potential pathogenicity of Willaertia magna by investigating the transfer of bacteria pathogenicity genes into its genome. Scientific Reports, 2019, 9, 18318.	1.6	12
136	Recent data on hepatitis E. Current Opinion in Infectious Diseases, 2019, 32, 475-481.	1.3	10
137	Hepatitis E virus as an agent of hepatocellular carcinoma. International Journal of Infectious Diseases, 2019, 80, 62-63.	1.5	8
138	Giant mimiviruses escape many canonical criteria of the virus definition. Clinical Microbiology and Infection, 2019, 25, 147-154.	2.8	17
139	Hepatitis E virus infection in heart transplant recipients, Southeastern France. Clinics and Research in Hepatology and Gastroenterology, 2019, 43, 108-111.	0.7	5
140	â€œTupanvirusâ€, a new genus in the family Mimiviridae. Archives of Virology, 2019, 164, 325-331.	0.9	23
141	Tailed giant Tupanvirus possesses the most complete translational apparatus of the known virosphere. Nature Communications, 2018, 9, 749.	5.8	247
142	Mimivirus in the urine of a kidney-transplant recipient. Clinical Microbiology and Infection, 2018, 24, 561-563.	2.8	8
143	A foretold hepatitis A outbreak in France. Liver International, 2018, 38, 955-956.	1.9	4
144	Deciphering the genomes of 16 Acanthamoeba species does not provide evidence of integration of known giant virus-associated mobile genetic elements. Virus Research, 2018, 251, 14-16.	1.1	4

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145	Nucleotide and amino acid diversity of hepatitisÂB virus surface antigen in acute infections, southeastern France. Clinics and Research in Hepatology and Gastroenterology, 2018, 42, e32-e34.	0.7	1
146	Hepatocellular carcinoma complicating hepatitis E virusâ€"related cirrhosis. Hepatology, 2018, 67, 446-448.	3.6	27
147	Faustovirus E12 Transcriptome Analysis Reveals Complex Splicing in Capsid Gene. Frontiers in Microbiology, 2018, 9, 2534.	1.5	31
148	Ancestrality and Mosaicism of Giant Viruses Supporting the Definition of the Fourth TRUC of Microbes. Frontiers in Microbiology, 2018, 9, 2668.	1.5	44
149	Ocular syphilis, an old adversary is back in the old world too!. Aids, 2018, 32, 2433-2434.	1.0	2
150	Investigations by the Institut Hospitalo-Universitaire MÃ©diterranÃ©e Infection of food and food-borne infections in the Mediterranean Basin and in sub-Saharan Africa. New Microbes and New Infections, 2018, 26, S37-S42.	0.8	4
151	A Phylogenomic Study of Acanthamoeba polyphaga Draft Genome Sequences Suggests Genetic Exchanges With Giant Viruses. Frontiers in Microbiology, 2018, 9, 2098.	1.5	27
152	Emergence of uncommon HIVâ€1 nonâ€B subtypes and circulating recombinant forms and trends in transmission of antiretroviral drug resistance in patients with primary infection during the 2013â€2015 period in Marseille, Southeastern France. Journal of Medical Virology, 2018, 90, 1559-1567.	2.5	5
153	Higher education at the Institut Hospitalo-Universitaire MÃ©diterranÃ©e Infection. New Microbes and New Infections, 2018, 26, S109-S113.	0.8	0
154	A Large Open Pangenome and a Small Core Genome for Giant Pandoraviruses. Frontiers in Microbiology, 2018, 9, 1486.	1.5	48
155	Experimental Inoculation in Rats and Mice by the Giant Marseillevirus Leads to Long-Term Detection of Virus. Frontiers in Microbiology, 2018, 9, 463.	1.5	5
156	Protozoal Giant Viruses. , 2018, , .		0
157	Repertoire of human gut microbes. Microbial Pathogenesis, 2017, 106, 103-112.	1.3	70
158	Mimiviruses and the Human Interferon System: Viral Evasion of Classical Antiviral Activities, But Inhibition By a Novel Interferon-Î² Regulated Immunomodulatory Pathway. Journal of Interferon and Cytokine Research, 2017, 37, 1-8.	0.5	11
159	Detection of rare hepatitis C viruses of subtype 4r in Southeastern France. Clinical Microbiology and Infection, 2017, 23, 493-495.	2.8	0
160	Mimivirus: leading the way in the discovery of giant viruses of amoebae. Nature Reviews Microbiology, 2017, 15, 243-254.	13.6	132
161	Toward chronic hepatitis C eradication in HIV-positive patients, including those cirrhotic and infected with genotype 3 viruses. Gut, 2017, 66, 1868-1869.	6.1	1
162	Daclatasvir plasma concentration assessment in HIVâ€"HCV-coinfected real-life patients. Antiviral Therapy, 2017, 22, 731-733.	0.6	1

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