Philippe Colson

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

16,159 369 56 119 h-index g-index citations papers 19,989 7.8 7.36 433 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
369	The puzzling mutational landscape of the SARS-2-variant Omicron <i>Journal of Medical Virology</i> , 2022 ,	19.7	14
368	Limited spread of a rare spike E484K-harboring SARS-CoV-2 in Marseille, France <i>Archives of Virology</i> , 2022 , 167, 583	2.6	1
367	Characteristics of the first 1,119 SARS-CoV-2 Omicron variant cases, in Marseille, France, November-December 2021 <i>Journal of Medical Virology</i> , 2022 ,	19.7	12
366	Introduction of the SARS-CoV-2 Beta variant from Comoros into the Marseille geographical area <i>Travel Medicine and Infectious Disease</i> , 2022 , 102277	8.4	O
365	Omicron variant escapes therapeutic mAbs including recently released Evusheld [], contrary to eight prior main VOC <i>Clinical Infectious Diseases</i> , 2022 ,	11.6	3
364	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	2.6	5
363	SARS-CoV-2 reinfection and COVID-19 severity Emerging Microbes and Infections, 2022, 1-41	18.9	5
362	Hepatitis delta treatment with bulevirtide in real life: a case report.: HDV treatment with bulevirtide <i>International Journal of Antimicrobial Agents</i> , 2022 , 106562	14.3	0
361	First cases of infection with the 21L/BA.2 Omicron variant in Marseille, France <i>Journal of Medical Virology</i> , 2022 ,	19.7	6
360	Severity of the first 207 infections with the SARS-CoV-2 Omicron BA.2 variant, in Marseille, France, December, 2021-February, 2022 <i>Journal of Medical Virology</i> , 2022 ,	19.7	1
359	Control of common viral epidemics but not of SARS-CoV-2 through the application of hygiene and distancing measures <i>Journal of Clinical Virology</i> , 2022 , 150-151, 105163	14.5	1
358	Culture and identification of a "Deltamicron" SARS-CoV-2 in a three cases cluster in southern France <i>Journal of Medical Virology</i> , 2022 ,	19.7	3
357	High rate of reinfection with the SARS-CoV-2 Omicron variant Journal of Infection, 2022,	18.9	1
356	SARS-CoV-2 Vaccination and Protection Against Clinical Disease: A Retrospective Study, Bouches-du-Rh\(^{\text{B}}\)e District, Southern France, 2021 <i>Frontiers in Microbiology</i> , 2021 , 12, 796807	5.7	0
355	Epidemiology and genomic characterisation of travel-associated and locally-acquired influenza, Marseille, France <i>Travel Medicine and Infectious Disease</i> , 2021 , 45, 102236	8.4	
354	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	5.7	11
353	Occurrence of a substitution or deletion of SARS-CoV-2 spike amino acid 677 in various lineages in Marseille, France. <i>Virus Genes</i> , 2021 , 1	2.3	2

Drug repurposing against SARS-CoV-1, SARS-CoV-2 and MERS-CoV. Future Microbiology, 2021, 16, 1341-12370 352 Isolation of 4000 SARS-CoV-2 shows that contagiousness is associated with viral load, not vaccine 18.9 351 or symptomatic status. Emerging Microbes and Infections, 2021, 10, 2276-2278 High Individual Heterogeneity of Neutralizing Activities against the Original Strain and Nine 6.2 350 7 Different Variants of SARS-CoV-2. Viruses, 2021, 13, Hepatitis A seroprevalence in homeless persons, in Marseille, France. Clinics and Research in 349 2.4 Hepatology and Gastroenterology, 2021, 45, 101571 Automated Western immunoblotting detection of anti-SARS-CoV-2 serum antibodies. European 8 348 5.3 Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 1309-1317 Introduction into the Marseille geographical area of a mild SARS-CoV-2 variant originating from 347 8.4 21 sub-Saharan Africa: An investigational study. Travel Medicine and Infectious Disease, 2021, 40, 101980 Direct next-generation sequencing diagnosis of echovirus 9 meningitis, France. European Journal of 346 5.3 3 Clinical Microbiology and Infectious Diseases, 2021, 40, 2037-2039 COVID-19 re-infection. European Journal of Clinical Investigation, 2021, 51, e13537 345 4.6 22 Effect of hydroxychloroquine and azithromycin on the viral clearance of SARS-CoV-2: response to 344 14.3 0 Herv'Seligmann. International Journal of Antimicrobial Agents, 2021, 57, 106306 Mink, SARS-CoV-2, and the Human-Animal Interface. Frontiers in Microbiology, 2021, 12, 663815 343 5.7 47 Global Discrepancies between Numbers of Available SARS-CoV-2 Genomes and Human 342 6.2 3 Development Indexes at Country Scales. Viruses, 2021, 13, Direct Diagnosis of Echovirus 12 Meningitis Using Metagenomic Next Generation Sequencing. 341 4.5 Pathogens, **2021**, 10, SARS-CoV-2 Persistent Viral Shedding in the Context of Hydroxychloroguine-Azithromycin 6.2 2 340 Treatment. Viruses, 2021, 13, Rhizomal Reclassification of Living Organisms. International Journal of Molecular Sciences, 2021, 22, 6.3 6 339 Clinical outcomes in COVID-19 patients infected with different SARS-CoV-2 variants in Marseille, 338 9.5 13 France. Clinical Microbiology and Infection, 2021, 27, 1516.e1-1516.e6 Emergence and outcomes of the SARS-CoV-2 'Marseille-4' variant. International Journal of Infectious 10.5 337 Diseases, 2021, 106, 228-236 SARS-CoV-2 Infectivity and Severity of COVID-19 According to SARS-CoV-2 Variants: Current 336 5.1 13 Evidence. Journal of Clinical Medicine, 2021, 10, Implementation of an in-house real-time reverse transcription-PCR assay for the rapid detection of 13 335 the SARS-CoV-2 Marseille-4 variant. Journal of Clinical Virology, 2021, 139, 104814

334	Marseilleviruses: An Update in 2021. Frontiers in Microbiology, 2021, 12, 648731	5.7	2
333	Diverse Trajectories Drive the Expression of a Giant Virus in the Oomycete Plant Pathogen. <i>Frontiers in Microbiology</i> , 2021 , 12, 662762	5.7	3
332	Implementation of an in-house real-time reverse transcription-PCR assay to detect the emerging SARS-CoV-2 N501Y variants. <i>Journal of Clinical Virology</i> , 2021 , 140, 104868	14.5	10
331	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,	9.7	79
330	Different pattern of the second outbreak of COVID-19 in Marseille, France. <i>International Journal of Infectious Diseases</i> , 2021 , 102, 17-19	10.5	8
329	Correlation Between 3790 Quantitative Polymerase Chain Reaction-Positives Samples and Positive Cell Cultures, Including 1941 Severe Acute Respiratory Syndrome Coronavirus 2 Isolates. <i>Clinical Infectious Diseases</i> , 2021 , 72, e921	11.6	101
328	Resistance to human immunodeficiency virus infection: a rare but neglected state. <i>Annals of the New York Academy of Sciences</i> , 2021 , 1485, 22-42	6.5	
327	Interpretation of SARS-CoV-2 PCR results for the diagnosis of COVID-19. <i>International Journal of Antimicrobial Agents</i> , 2021 , 57, 106238	14.3	
326	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	5.3	18
325	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	14.3	14
324	Coinfections with SARS-CoV-2 and other respiratory viruses in Southeastern France: A matter of sampling time. <i>Journal of Medical Virology</i> , 2021 , 93, 1878-1881	19.7	7
323	Evidence of SARS-CoV-2 re-infection with a different genotype. <i>Journal of Infection</i> , 2021 , 82, 84-123	18.9	27
322	Early combination therapy with hydroxychloroquine and azithromycin reduces mortality in 10,429 COVID-19 outpatients. <i>Reviews in Cardiovascular Medicine</i> , 2021 , 22, 1063-1072	3.9	3
321	Epidemiological serosurvey and molecular characterization of sexually transmitted infections among 1890 sheltered homeless people in Marseille: Cross-sectional one day-surveys (2000-2015). <i>Journal of Infection</i> , 2021 , 82, 60-66	18.9	2
320	Giant virus-related sequences in the 5300-year-old Ezi mummy metagenome. <i>Virus Genes</i> , 2021 , 57, 222-227	2.3	О
319	Development and validation of a multiplex qPCR assay for detection and relative quantification of HPV16 and HPV18 E6 and E7 oncogenes. <i>Scientific Reports</i> , 2021 , 11, 4039	4.9	3
318	No barrier to care, yet disparities in the HIV care continuum in France: a nationwide population study. <i>Journal of Antimicrobial Chemotherapy</i> , 2021 , 76, 1573-1579	5.1	O
317	A Possible Role of Remdesivir and Plasma Therapy in the Selective Sweep and Emergence of New SARS-CoV-2 Variants. <i>Journal of Clinical Medicine</i> , 2021 , 10,	5.1	8

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316	SARS-CoV-2 variant from India to Marseille: The still active role of ports in the introduction of epidemics. <i>Travel Medicine and Infectious Disease</i> , 2021 , 42, 102085	8.4	8
315	Consequences of the COVID-19 Outbreak Lockdown on Non-Viral Infectious Agents as Reported by a Laboratory-Based Surveillance System at the IHU Mditerrane Infection, Marseille, France. <i>Journal of Clinical Medicine</i> , 2021 , 10,	5.1	1
314	Clinical characteristics of paediatric COVID-19 patients followed for up to 13 months. <i>Acta Paediatrica, International Journal of Paediatrics</i> , 2021 , 110, 3331-3333	3.1	2
313	Monitoring the Circulation of SARS-CoV-2 Variants by Genomic Analysis of Wastewater in Marseille, South-East France. <i>Pathogens</i> , 2021 , 10,	4.5	4
312	Whole Genome Sequencing of SARS-CoV-2 Strains in COVID-19 Patients From Djibouti Shows Novel Mutations and Clades Replacing Over Time. <i>Frontiers in Medicine</i> , 2021 , 8, 737602	4.9	1
311	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	9.5	20
310	Epidemiological and genetic characterization of measles virus circulating strains at Marseille, France during 2017-2019 measles outbreak. <i>Journal of Infection</i> , 2021 , 83, 361-370	18.9	2
309	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		5
308	Incomplete tricarboxylic acid cycle and proton gradient in Pandoravirus massiliensis: is it still a virus?. <i>ISME Journal</i> , 2021 ,	11.9	4
307	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. <i>Infection, Genetics and Evolution</i> , 2021 , 95, 105092	4.5	7
306	Concurrent Nanopore Next-Generation Sequencing of Hepatitis B and Delta Virus Genomes Directly From Patient Plasma. <i>Annals of Laboratory Medicine</i> , 2021 , 41, 608-611	3.1	
305	Microelimination or Not? The Changing Epidemiology of Human Immunodeficiency Virus-Hepatitis C Virus Coinfection in France 2012-2018. <i>Clinical Infectious Diseases</i> , 2021 , 73, e3266-e3274	11.6	3
304	Emerging SARS-CoV-2 Genotypes Show Different Replication Patterns in Human Pulmonary and Intestinal Epithelial Cells <i>Viruses</i> , 2021 , 14,	6.2	3
303	Heterogeneity in susceptibility to hydroxychloroquine of SARS-CoV-2 isolates <i>Frontiers in Bioscience</i> , 2021 , 26, 1493-1502		O
302	Incidence and Outcome of Coinfections with SARS-CoV-2 and Rhinovirus Viruses, 2021, 13,	6.2	4
301	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	8.4	300
300	Analysis of a Marseillevirus Transcriptome Reveals Temporal Gene Expression Profile and Host Transcriptional Shift. <i>Frontiers in Microbiology</i> , 2020 , 11, 651	5.7	10
299	Comparison of mortality associated with respiratory viral infections between December 2019 and March 2020 with that of the previous year in Southeastern France. <i>International Journal of Infectious Diseases</i> , 2020 , 96, 154-156	10.5	9

298	Child with liver transplant recovers from COVID-19 infection. A case report. <i>Archives De Pediatrie</i> , 2020 , 27, 275-276	1.8	25
297	Vermamoeba vermiformis CDC-19 draft genome sequence reveals considerable gene trafficking including with candidate phyla radiation and giant viruses. <i>Scientific Reports</i> , 2020 , 10, 5928	4.9	7
296	A virophage cross-species infection through mutant selection represses giant virus propagation, promoting host cell survival. <i>Communications Biology</i> , 2020 , 3, 248	6.7	5
295	Does spitting in public play a role in transmitting SARS-CoV-2?. <i>Travel Medicine and Infectious Disease</i> , 2020 , 36, 101759	8.4	3
294	Insight into the Lifestyle of Amoeba during Bioreactor Growth Using Transcriptomics and Proteomics. <i>Microorganisms</i> , 2020 , 8,	4.9	2
293	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	4.1	48
292	Testing the repatriated for SARS-Cov2: Should laboratory-based quarantine replace traditional quarantine?. <i>Travel Medicine and Infectious Disease</i> , 2020 , 34, 101624	8.4	31
291	Description of Virulent Factors and Horizontal Gene Transfers of Keratitis-Associated Amoeba Acanthamoeba Triangularis by Genome Analysis. <i>Pathogens</i> , 2020 , 9,	4.5	5
290	SARS-CoV-2: fear versus data. International Journal of Antimicrobial Agents, 2020, 55, 105947	14.3	48
289	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	14.3	663
288	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	8.4	82
287	Yaravirus: A novel 80-nm virus infecting. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 16579-16586	11.5	15
286	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	5.3	494
285	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	8.4	162
284	Core gene-based molecular detection and identification of Acanthamoeba species. <i>Scientific Reports</i> , 2020 , 10, 1583	4.9	6
283	Full-length title: Dramatic HIV DNA degradation associated with spontaneous HIV suppression and disease-free outcome in a young seropositive woman following her infection. <i>Scientific Reports</i> , 2020 , 10, 2548	4.9	1
282	Kaposi sarcoma among people living with HIV in the French DAT'AIDS cohort between 2010 and 2015. <i>Journal of the European Academy of Dermatology and Venereology</i> , 2020 , 34, 1065-1073	4.6	3
281	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	8.4	479

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280	In vitro testing of combined hydroxychloroquine and azithromycin on SARS-CoV-2 shows synergistic effect. <i>Microbial Pathogenesis</i> , 2020 , 145, 104228	3.8	171
279	Letter to the editor: Plenty of coronaviruses but no SARS-CoV-2. <i>Eurosurveillance</i> , 2020 , 25,	19.8	8
278	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	14.3	3043
277	Natural history of COVID-19 and therapeutic options. <i>Expert Review of Clinical Immunology</i> , 2020 , 16, 1159-1184	5.1	41
276	Respiratory Syncytial Virus Infection: Its Propensity for Bacterial Coinfection and Related Mortality in Elderly Adults. <i>Open Forum Infectious Diseases</i> , 2020 , 7, ofaa546	1	7
275	Hepatitis B Virus Genomics Knocking at the Door of Routine Diagnostic Laboratories. <i>Journal of Infectious Diseases</i> , 2020 , 221, 1026-1029	7	1
274	Hepatitis Delta recurrence post-liver transplantation in absence of detectable hepatitis B surface antigen and hepatitis B virus DNA in peripheral blood. <i>Clinics and Research in Hepatology and Gastroenterology</i> , 2020 , 44, e41-e44	2.4	2
273	Multimorbidity in Elderly Persons According to the Year of Diagnosis of Human Immunodeficiency Virus Infection: A Cross-sectional Dat'AIDS Cohort Study. <i>Clinical Infectious Diseases</i> , 2020 , 71, 2880-288	8 ^{11.6}	4
272	Ribavirin for Hepatitis E Virus Infection After Organ Transplantation: A Large European Retrospective Multicenter Study. <i>Clinical Infectious Diseases</i> , 2020 , 71, 1204-1211	11.6	36
271	Temporal and age distributions of SARS-CoV-2 and other coronaviruses, southeastern France. <i>International Journal of Infectious Diseases</i> , 2020 , 101, 121-125	10.5	3
270	Antibiotic resistance surveillance systems: A review. <i>Journal of Global Antimicrobial Resistance</i> , 2020 , 23, 430-438	3.4	18
269	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	5.3	27
268	Major discrepancy between factual antibiotic resistance and consumption in South of France: analysis of 539,037 bacterial strains. <i>Scientific Reports</i> , 2020 , 10, 18262	4.9	1
267	Contribution of VitaPCR SARS-CoV-2 to the emergency diagnosis of COVID-19. <i>Journal of Clinical Virology</i> , 2020 , 133, 104682	14.5	16
266	Promiscuous Enzyme Activity as a Driver of Allo and Iso Convergent Evolution, Lessons from the ELactamases. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	4
265	Hepatitis A outbreak in HIV-infected patients in Southeastern France: questions and responses?. <i>Epidemiology and Infection</i> , 2020 , 148, e79	4.3	O
264	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	5.3	17
263	Incidence of diabetes in HIV-infected patients treated with first-line integrase strand transfer inhibitors: a French multicentre retrospective study. <i>Journal of Antimicrobial Chemotherapy</i> , 2020 , 75, 3344-3348	5.1	8

262	pidmiologie des infections ^coronavirus HCoV-HKU1 [^] Marseille, France. <i>Mdecine Et Maladies Infectieuses</i> , 2020 , 50, S195	4	78
261	A protein of the metallo-hydrolase/oxidoreductase superfamily with both beta-lactamase and ribonuclease activity is linked with translation in giant viruses. <i>Scientific Reports</i> , 2020 , 10, 21685	4.9	4
260	Chronic hepatitis E in absence of severe immune deficiency. <i>Clinics and Research in Hepatology and Gastroenterology</i> , 2020 , 44, e1-e4	2.4	0
259	Host-virus interactions and defense mechanisms for giant viruses. <i>Annals of the New York Academy of Sciences</i> , 2020 , 1486, 39	6.5	1
258	Human papillomavirus type 7-associated anal condyloma after renal transplantation in a child. <i>Pediatric Transplantation</i> , 2019 , 23, e13470	1.8	O
257	Discovery and Further Studies on Giant Viruses at the IHU Mediterranee Infection That Modified the Perception of the Virosphere. <i>Viruses</i> , 2019 , 11,	6.2	14
256	Virophages of Giant Viruses: An Update at Eleven. Viruses, 2019, 11,	6.2	17
255	Human metallo-Elactamase enzymes degrade penicillin. Scientific Reports, 2019 , 9, 12173	4.9	18
254	The Methodology Behind Network Thinking: Graphs to Analyze Microbial Complexity and Evolution. <i>Methods in Molecular Biology</i> , 2019 , 1910, 271-308	1.4	2
253	Measles, the need for a paradigm shift. European Journal of Epidemiology, 2019, 34, 897-915	12.1	11
252	Multidisciplinary evaluation of Clostridium butyricum clonality isolated from preterm neonates with necrotizing enterocolitis in South France between 2009 and 2017. <i>Scientific Reports</i> , 2019 , 9, 2077	4.9	5
251	Evidence of a Cellulosic Layer in Tegument and the Mystery of the Genetic Support of Its Biosynthesis. <i>Frontiers in Microbiology</i> , 2019 , 10, 2932	5.7	8
250	Investigation of potential pathogenicity of Willaertia magna by investigating the transfer of bacteria pathogenicity genes into its genome. <i>Scientific Reports</i> , 2019 , 9, 18318	4.9	8
249	Recent data on hepatitis E. Current Opinion in Infectious Diseases, 2019, 32, 475-481	5.4	5
248	Giant mimiviruses escape many canonical criteria of the virus definition. <i>Clinical Microbiology and Infection</i> , 2019 , 25, 147-154	9.5	8
247	Hepatitis E virus infection in heart transplant recipients, Southeastern France. <i>Clinics and Research in Hepatology and Gastroenterology</i> , 2019 , 43, 108-111	2.4	4
246	"Tupanvirus", a new genus in the family Mimiviridae. Archives of Virology, 2019, 164, 325-331	2.6	11
245	Tailed giant Tupanvirus possesses the most complete translational apparatus of the known virosphere. <i>Nature Communications</i> , 2018 , 9, 749	17.4	136

244	Mimivirus in the urine of a kidney-transplant recipient. Clinical Microbiology and Infection, 2018, 24, 561	-563	5
243	A foretold hepatitis A outbreak in France. Liver International, 2018, 38, 955-956	7.9	3
242	Deciphering the genomes of 16 Acanthamoeba species does not provide evidence of integration of known giant virus-associated mobile genetic elements. <i>Virus Research</i> , 2018 , 251, 14-16	6.4	2
241	Nucleotide and amino acid diversity of hepatitis B virus surface antigen in acute infections, southeastern France. <i>Clinics and Research in Hepatology and Gastroenterology</i> , 2018 , 42, e32-e34	2.4	1
240	Hepatocellular carcinoma complicating hepatitis E virus-related cirrhosis. <i>Hepatology</i> , 2018 , 67, 446-448	8 11.2	17
239	A Large Open Pangenome and a Small Core Genome for Giant Pandoraviruses. <i>Frontiers in Microbiology</i> , 2018 , 9, 1486	5.7	23
238	Experimental Inoculation in Rats and Mice by the Giant Marseillevirus Leads to Long-Term Detection of Virus. <i>Frontiers in Microbiology</i> , 2018 , 9, 463	5.7	4
237	Faustovirus E12 Transcriptome Analysis Reveals Complex Splicing in Capsid Gene. <i>Frontiers in Microbiology</i> , 2018 , 9, 2534	5.7	6
236	Ancestrality and Mosaicism of Giant Viruses Supporting the Definition of the Fourth TRUC of Microbes. <i>Frontiers in Microbiology</i> , 2018 , 9, 2668	5.7	30
235	Ocular syphilis, an old adversary is back in the old world too!. <i>Aids</i> , 2018 , 32, 2433-2434	3.5	2
235	Ocular syphilis, an old adversary is back in the old world too!. <i>Aids</i> , 2018 , 32, 2433-2434 Investigations by the Institut Hospitalo-Universitaire Mditerrane Infection of food and food-borne infections in the Mediterranean Basin and in sub-Saharan Africa. <i>New Microbes and New Infections</i> , 2018 , 26, S37-S42	3.5	2
	Investigations by the Institut Hospitalo-Universitaire Mditerrane Infection of food and food-borne infections in the Mediterranean Basin and in sub-Saharan Africa. <i>New Microbes and New</i>		
234	Investigations by the Institut Hospitalo-Universitaire Mditerrane Infection of food and food-borne infections in the Mediterranean Basin and in sub-Saharan Africa. <i>New Microbes and New Infections</i> , 2018 , 26, S37-S42 A Phylogenomic Study of Draft Genome Sequences Suggests Genetic Exchanges With Giant	4.1	21
234	Investigations by the Institut Hospitalo-Universitaire Mditerrane 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 A Phylogenomic Study of Draft Genome Sequences Suggests Genetic Exchanges With Giant Viruses. Frontiers in Microbiology, 2018, 9, 2098 Emergence of uncommon HIV-1 non-B subtypes and circulating recombinant forms and trends in transmission of antiretroviral drug resistance in patients with primary infection during the	4.1 5·7	21
234233232	Investigations by the Institut Hospitalo-Universitaire Mälterrane Infection of food and food-borne infections in the Mediterranean Basin and in sub-Saharan Africa. <i>New Microbes and New Infections</i> , 2018 , 26, S37-S42 A Phylogenomic Study of Draft Genome Sequences Suggests Genetic Exchanges With Giant Viruses. <i>Frontiers in Microbiology</i> , 2018 , 9, 2098 Emergence of uncommon HIV-1 non-B subtypes and circulating recombinant forms and trends in transmission of antiretroviral drug resistance in patients with primary infection during the 2013-2015 period in Marseille, Southeastern France. <i>Journal of Medical Virology</i> , 2018 , 90, 1559-1567 Higher education at the Institut Hospitalo-Universitaire Mälterrane Infection. <i>New Microbes and</i>	4.1 5·7 19.7	21
234233232231	Investigations by the Institut Hospitalo-Universitaire Mäiterrane Infection of food and food-borne infections in the Mediterranean Basin and in sub-Saharan Africa. <i>New Microbes and New Infections</i> , 2018 , 26, S37-S42 A Phylogenomic Study of Draft Genome Sequences Suggests Genetic Exchanges With Giant Viruses. <i>Frontiers in Microbiology</i> , 2018 , 9, 2098 Emergence of uncommon HIV-1 non-B subtypes and circulating recombinant forms and trends in transmission of antiretroviral drug resistance in patients with primary infection during the 2013-2015 period in Marseille, Southeastern France. <i>Journal of Medical Virology</i> , 2018 , 90, 1559-1567 Higher education at the Institut Hospitalo-Universitaire Mäiterrane Infection. <i>New Microbes and New Infections</i> , 2018 , 26, S109-S113	4.1 5·7 19.7	4 21 3
234 233 232 231 230	Investigations by the Institut Hospitalo-Universitaire Mälterrane Infection of food and food-borne infections in the Mediterranean Basin and in sub-Saharan Africa. <i>New Microbes and New Infections</i> , 2018 , 26, S37-S42 A Phylogenomic Study of Draft Genome Sequences Suggests Genetic Exchanges With Giant Viruses. <i>Frontiers in Microbiology</i> , 2018 , 9, 2098 Emergence of uncommon HIV-1 non-B subtypes and circulating recombinant forms and trends in transmission of antiretroviral drug resistance in patients with primary infection during the 2013-2015 period in Marseille, Southeastern France. <i>Journal of Medical Virology</i> , 2018 , 90, 1559-1567 Higher education at the Institut Hospitalo-Universitaire Mälterrane Infection. <i>New Microbes and New Infections</i> , 2018 , 26, S109-S113 Repertoire of human gut microbes. <i>Microbial Pathogenesis</i> , 2017 , 106, 103-112	4.1 5·7 19.7 4.1 3.8	4 21 3

226	Toward chronic hepatitis C eradication in HIV-positive patients, including those cirrhotic and infected with genotype 3 viruses. <i>Gut</i> , 2017 , 66, 1868-1869	19.2	1
225	Daclatasvir plasma concentration assessment in HIV-HCV-coinfected real-life patients. <i>Antiviral Therapy</i> , 2017 , 22, 731-733	1.6	1
224	Spontaneous clearance of chronic hepatitis C virus infection in HIV-positive patients, southeastern France. <i>Clinics and Research in Hepatology and Gastroenterology</i> , 2017 , 41, e87-e89	2.4	1
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11	Emergence in Southern France of a new SARS-CoV-2 variant of probably Cameroonian origin harbouring both substitutions N501Y and E484K in the spike protein		13

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8	Hydroxychloroquine and azithromycin as a treatment of COVID-19: results of an open-label non-randomized clinical trial	59
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